

Search Details

Job Title/ Assay Name : tropicalis(ex)	Amplicon Sequence: GCCTGTTTGAGCGTCATTTCTCCCTCAAACCCCCGGGTTTGGTGTT GAGCAATACGCTAGGTTTGTGTTGAAAGAATTTAACGTGGAAACTTA TTTTAAGCGACTTAGGTTTATCCAAAAACGCTTATTTTGCTAGTGGC CACCACAATTTATTTTCATAACTTTGACCTCCAATCAGGTAGGACTAC CCGCTGAACTTAAGCATATCAATAAGCGGAGG
Effective Date : 2025-10-28	Forward: GCCTGTTTGAGCGTCRTTTC
Database : core_nt	Reverse: CCTCCGCTTATTGATATGCTTA
Alignment algorithm : blastN	Probe: TGCTAGTGGCCACCACAATTTATTTC
Exclusive Organism: Candida tropicalis (5482)	Inclusive Organism:
RID: G12K4WEJ014	Total Result: 5000
App Version: 1.0	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
Candida albicans/5476 Analyzed Accessions: 1172				
<u>2</u>	PP851176 KT876204	NA	No Mutations	✓✓✓ Detected
<u>3</u>	PP851204 PP851185 PP851222	Reverse ✓✓✓✓	+7C	10.00 ✓✓✓ Detected
<u>1</u>	PP851227	Probe ✓✓✓✓	T4A	9.34 ✓✓✓ Detected
		Reverse ✓✓✓✓	+7C	
<u>2</u>	KP675554 PP851203	Reverse ✓✓✓✓	C1-, C2-, T3-, A10-	10.00 ✓✓✓ Detected
<u>3</u>	PP851220 PP851191 PP851254	Reverse ✓✓✓✓	T8C, +8C, A10T	10.00 ✓✓✓ Detected
<u>2</u>	PP851161 PP851206	Reverse ✓✓✓✓	+5C G, A10-	10.00 ✓✓✓ Detected
<u>1</u>	PP851235	Reverse ✓✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, A10T	10.00 ✓✓✓ Detected
<u>1</u>	PP851261	Reverse ✓✓✓✓	T8C, A10T	10.00 ✓✓✓ Detected
<u>2</u>	EF065163 PP851238	Reverse ✓✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-	10.00 ✓✓✓ Detected
<u>1</u>	PP851209	Reverse ✓✓✓✓	+7C, A10T	10.00 ✓✓✓ Detected

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	PP851189	Reverse   	C1-, G6C, A10T	10.00    Detected
1	PP851166	Reverse   	G6C, T8G, A10T	10.00    Detected
1	PP851257	Reverse   	G6C, T8C, A10T	10.00    Detected
2	PP851263 PP851264	Probe   	T4A	9.34    Detected
		Reverse   	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-	
1	PP851245	Reverse   	+6G, T8C, A10T	10.00    Detected
1	PP851182	Reverse   	C1-, C2-, T3-, T8C, A10T	10.00    Detected
1	PP851266	Reverse   	C1-, C2-, T3-, G6C, A10T	10.00    Detected
1	PP851171	Forward   	G11A	9.34    Detected
		Reverse   	T3C, G6C, A10-	
1	PP851178	Reverse 	T3C, G6C, A10T, +12A	6.00  Not Detected
1	PP851258	Reverse 	T3C, G6C, A10T, T12A	6.00  Not Detected
1	KY209906	Probe  	G9C, A18T	6.67  Not Detected
		Reverse   	C1-, C2-, T3-, C4-, C5-, G6-	
1	PP851193	Forward   	G11A, A16C	0.00    Not Detected
		Reverse   	T3C, G6C, A10-, G18T, T20C	
7	OK267646 PQ895596 PP625293 FJ662398 PQ895568 KP674711 OK267636	Forward   	A16G	0.34    Not Detected
		Probe   	T1C, C3G, T7C, G9T, A17G, A18-, T20A, +23C	
443	443	Forward   	A16G	0.34    Not Detected
		Probe   	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
1	KY101880	Forward   	A16G	0.34    Not Detected
		Probe   	T1C, C3G, G6A, T7C, G9T, A17G, A18T, T19A, +23C	
1	KY101865	Forward   	A16G	0.34    Not Detected
		Probe   	T1C, C3G, G6A, T7C, G9T, A17G, A18G, T20A, +23C	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
2	KC905066 OR058918	Forward XXX	A16G	0.34 XXX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, T24C	
38	38	Forward XXX	A16G	0.34 XXX Not Detected
		Probe XXX	T1C, C3G, T7C, G9T, C11T, A17G, A18-, T20A, +23C	
1	PP755001	Forward XXX	A16G	0.34 XXX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9A, A17G, A18-, T20A, +23C	
1	OR429586	Forward XXX	C12A, A16G	0.34 XXX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
31	31	Forward XXX	A16G	0.34 XXX Not Detected
		Probe XXX	T1G, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +1CGGC, +23C	
1	KY101835	Forward XXX	A16G	0.34 XXX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, C11T, A17G, A18G, T20A, +23C	
1	PQ518567	Forward XXX	A16G	0.34 XXX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A18-, T20A, +23C	
4	PP838742 MZ770753 EU266569 MW132095	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	C1-	
1	PV761643	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, T7C, G9T, C11T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	C1-	
1	ON142620	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1G, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +1CGGC, +23C	
		Reverse ✓✓✓	C1-	
1	FJ662389	Forward XXX	A16G	0.34 XXX Not Detected
		Probe XXX	T7C, G9T, C11T, A17G, A18-, T20A, +3GGCGG, +23C	
1	PP214445	Forward XXX	A16G	0.34 XXX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18T, T19A, T21A, A22T	
4	MW081141 KP674966 MG707642 MZ674385	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	C1-, C2-	
1	AY560100	Forward XXX	A16G	0.34 XXX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, T24C, C26T, A27T	
307	307	Forward XXX	A16G	0.34 XXX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, C11T, A17G, A18-, T20A, +23C	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
<u>4</u>	MW307710 OK267674 GU584190 KP675201	Forward XXX	A16G	0.34 XXX Not Detected
		Probe XXX	T1C, C3G, T4C, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
<u>5</u>	KP675564 PV790048 MZ255119 PV130371 LC601969	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	T8C	
<u>3</u>	OR058910 MG599244 OR058932	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	T11A	
<u>2</u>	FJ697168 KP675384	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	C4T	
<u>2</u>	PQ415053 PQ415054	Forward XXX	A16G	0.34 XXX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, C11A, A17G, A18-, T20A, +23C	
<u>1</u>	KU987844	Forward XXX	T14C, A16G	0.34 XXX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
<u>2</u>	KP675529 KP675528	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, T7C, G9T, C11T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	A10T	
<u>1</u>	ON845628	Forward XXX	A16G	0.34 XXX Not Detected
		Probe XXX	T1C, C3G, G6A, T7G, G9T, C11T, A17G, A18-, T20A, +23C	
<u>1</u>	KF309067	Forward XXX	A16G, C20T	0.34 XXX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
<u>1</u>	KP675530	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	A10T	
<u>2</u>	PQ469764 KP675470	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	G6C	
<u>1</u>	KP675496	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	T9C	
<u>1</u>	OK267803	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	C7A	
<u>1</u>	KY101879	Forward XXX	A16G	0.34 XXX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18G, +19A, +23C	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
31	31	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	C1-, C2-, T3-	
1	KP675482	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1G, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +1CGGC, +23C	
		Reverse ✓✓✓	G6C	
17	17	Forward XXX	A16G	0.34 XXX Not Detected
		Probe XXX	T1G, C3T, T4C, A5T, G6C, C11G, A12T, A15C, A18T, +13G, +24CT	
29	29	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	+7C	
20	20	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	+6G	
1	AY560103	Forward XXX	A16G	0.34 XXX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +11T, +23C	
1	MZ254966	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, T7C, G9T, C11T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	+7C	
5	KP675438 PP851217 KP675557 KT876203 KP674655	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	+9T	
4	PQ468274 OP293046 MZ674383 KP675598	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, C11T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	C1-	
2	KP674531 KP675282	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	+6A	
1	AY560104	Forward XXX	A16G	0.34 XXX Not Detected
		Probe XXX	T1C, C3G, T7A, G8C, C10T, A17G, A18-, T20A, +11T, +23C	
4	KP675360 KP675278 OK267701 LC601970	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	+6T	
1	KP675339	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	C1-, G6C	
1	KP674885	Forward XXX	A16G	3.56 XX

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
		Probe XXX	T1G, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +1CGGC, +23C	Not Detected
		Reverse ✓✓✓	C1-, C2-, T3-	
1	MG250508	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	+6N	
1	PP851364	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, T7C, G9T, C11T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	+6G	
1	KY101841	Forward XXX	A16G	0.34 XXX Not Detected
		Probe XXX	T1C, C3G, T4C, A5G, G8A, C11G, A17G, A18-, T20A, +11TT, +23C	
7	PV563060 PQ407901 MW132128 MW139347 MG599195 PQ738309 MG599194	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-	
1	MN027924	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1G, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +1CGGC, +23C	
		Reverse ✓✓✓	+9T	
3	KP674785 LC601967 KP674988	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1G, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +1CGGC, +23C	
		Reverse ✓✓✓	+7C	
2	OK267775 KP674471	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1G, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +1CGGC, +23C	
		Reverse ✓✓✓	+6G	
1	LC601958	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	T9-	
1	PP218386	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	+6G C	
3	KY575033 KY575031 KY575029	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	T12-	
2	MG020714 OK267779	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	+5C G	
2	MW181231 MH016352	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, C11T, A17G, A18-, T20A, +23C	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
		Reverse   	C1-, C2-	
1	AY560102	Forward   	A16G	0.34   
		Probe   	T1C, G6A, T7C, G9T, A17G, A18-, T20A, +2GCGG, +11T, +23C	Not Detected
51	51	Forward   	A16G	0.34   
		Probe   	T1C, C3G, T4C, G6A, T7C, G9T, C11T, A17G, A18-, T20A, +23C	Not Detected
21	21	Forward   	A16G	0.34   
		Probe   	T1C, C3G, G6A, T7C, G9T, C11T, C14T, A17G, A18-, T20A, +23C	Not Detected
1	KY101840	Forward   	A16G	0.34   
		Probe   	T1C, C3G, G6A, T7C, G9T, C11T, A17G, A18G, +11T, +19A, +23C	Not Detected
1	MK026226	Forward   	G1-, C2-, C3-, T4-, G5-, A16G	0.34   
		Probe   	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	Not Detected
3	LC537155 KP674802 KP765021	Forward   	G11A, A16G	0.34   
		Probe   	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	Not Detected
1	KP675320	Forward   	G11A, A16C	0.34   
		Probe   	T1G, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +1CGGC, +23C	Not Detected
1	ON845641	Forward   	A16G	0.34   
		Probe   	T4G, A5-, T7C, C11A, C14G, A17C, T19C, T20G, T24A, A27T, +14TCT	Not Detected
1	MK026352	Forward   	G1-, C2-, C3-, A16G	0.34   
		Probe   	T1C, C3G, G6A, T7C, G9T, C11T, A17G, A18-, T20A, +23C	Not Detected
1	KF746423	Forward   	C12-, G13-, A16G	0.34   
		Probe   	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	Not Detected
1	PQ895585	Forward   	A16G	0.34   
		Probe   	T1C, C3G, T4C, A5G, G9T, C11T, A17G, A18-, T20A, +7CAC, +23C	Not Detected
1	JN159659	Probe   	T1A, C3G, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, A15T, C16T, A18T, T19G, T20C, A22T	5.33 
		Reverse   	C1-, C2-, T3-, C4-, C5-, G6-, C7-	Not Detected
1	ON365597	Forward   	G11A, A16G	0.34   
		Probe   	T4G, G9T, A17G, A18-, T20A, +5CG, +7AAC, +23C	Not Detected
1	GQ411404	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	0.67   
1	OM033483	Forward   	G1-, C2-, C3-, T4-, G5-, A16G	0.34   
		Probe   	T1C, C3G, T4C, G6A, T7C, G9T, C11T, A17G, A18-, T20A, +23C	Not Detected
1	ON398746	Forward   	T8-, A16G	0.34   
		Probe   	T1C, C3G, G6A, T7C, G9T, C11T, A17G, A18-, T20A, +23C	Not Detected
1	KP674909	Forward   	A16T	3.11  

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
		Probe XXX	T1C, C3G, G6C, T7C, G9T, C11T, A17G, A18-, T20A, +23C	Not Detected
		Reverse ✓✓✓	C1-, C7G, C19G	
<u>1</u>	LC612888	Probe XXX	T1C, C3G, G6A, T7C, G9T, C11T, A17G, A18-, T20A, T24C, A27C	0.67 XXX Not Detected
<u>1</u>	KP675235	Forward XXX	A16T	3.11 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, C11T, A17-, A18G, T20A, +23C	
		Reverse ✓✓✓	C1-, C2-, T3-, T12-, C19G	
<u>2</u>	PV363914 PP851262	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X Not Detected
		Reverse ✓✓✓	T8C, A10T	
<u>1</u>	KP674910	Forward XXX	G11C, A16T	2.45 XXX Not Detected
		Probe XXX	T1C, C3G, G6C, T7C, G9T, C11T, A17-, A18G, T20A, +23C	
		Reverse ✓	C1-, +6G, T12-, C19G	
<u>1</u>	PP851195	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX Not Detected
		Reverse X	G6C, T8C, A10T	
<u>1</u>	KP675381	Forward XXX	T4G, G11C, A16T	3.11 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	C1-, C4T, C7G, C19G	
<u>1</u>	KP674795	Forward XXX	G11C, A16T	3.11 XX Not Detected
		Probe XXX	T1C, C3G, G6C, T7C, G9T, C11T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	C1-, C4T, C7G, C19G	
<u>1</u>	PP851181	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX Not Detected
		Reverse X	+6G, T8C, A10T	
<u>1</u>	PP851224	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX Not Detected
		Reverse X	T8C, +8C, A10T	
<u>1</u>	EF065165	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX Not Detected
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-	
<u>1</u>	PP851260	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX Not Detected
		Reverse X	+5C G, T8C, A10T	
<u>2</u>	PQ139370 PP851197	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX Not Detected
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-	
<u>1</u>	OQ134708	Forward XXX	G11C, A16T	0.22 XXX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, C11T, A17-, A18G, T20A, +23C	
		Reverse XXX	C1-, C2-, C5G, T12-, C19G	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	KP674964	Forward XXX	G11A, A16C	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	C1-, C2-, C7T	
1	KP675402	Forward XXX	G11A, A16C	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	C1-, C2-	
1	PP851244	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX Not Detected
		Reverse X	T3C, G6C, A10-, G13A	
1	KP675596	Forward XXX	T4G, G11C, A16T	3.11 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, C11T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	C1-, C4T, C7G, C19G	
1	MH061327	Forward XXX	G11C, A16T, T19C	0.22 XXX Not Detected
		Probe XXX	T1C, C3G, G6C, T7C, G9T, C11T, A17G, A18-, T20A, +23C	
		Reverse XXX	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, C19G	
1	PP851241	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33 X Not Detected
		Reverse ✓✓✓	T8C, +8C, A10T	
1	EF065168	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	0.34 XXX Not Detected
		Reverse XXX	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, T21-	
1	KP674963	Forward XXX	T6G, G11A, A16C	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, T24C, C26T, A27C	
		Reverse ✓✓✓	C1-, C2-, C7G, A10T	
1	OR786727	Forward XXX	T8A, A10T, A16G	0.22 XXX Not Detected
		Probe XXX	T1G, T4A, G6T, T7C, G9T, C11T, A17G, A18-, T20A, +23C	
		Reverse XXX	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T20A	
3	OK054546 OK054543 PQ415055	Forward XXX	G1-, C2-, C3-, T7C, C15T, A16G, T18G	0.34 XXX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, C11T, A17G, A18-, T20A, +23C	
1	EF065166	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	0.67 XXX Not Detected
		Reverse	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, +16N	
1	KP674527	Forward XXX	T6G, G11A, A16C	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, C11T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	C1-, C2-, C7G, A10T	
1	KP674991	Forward XXX	T4G, G11C, A16T	0.22 XXX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9C, A17-, A18G, T20A, +23C	
		Reverse XXX	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, C19G	
1	KP674643	Forward XXX	T4G, G11C, A16T	0.22 XXX

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
		Probe XXXX	T1C, C3G, G6A, T7C, G9T, C11T, A17G, A18-, T20A, +23C	Not Detected
		Reverse XXXX	C1-, C2-, G6C, A10-, C19G	
1	KP674965	Forward XXXX	T6G, G11A, A16C	0.34 XXXX Not Detected
		Probe XXXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, T24C, C26T, A27C	
		Reverse	C1-, C2-, C7T, T15A, C19T	
1	KP674900	Forward XXXX	T6G, G11A, A16C	3.56 XX Not Detected
		Probe XXXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, T24C, C26T, A27C	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, A10T	
1	KP675459	Forward XXXX	T4G, G11C, A16G	0.22 XXXX Not Detected
		Probe XXXX	T1C, C3G, G6A, T7C, G9T, C11A, A17G, A18-, T20A, +23C	
		Reverse XXXX	C4T, A10-, T17A, C19G	
1	PQ651724	Forward XXXX	G11C, G13A, T14C, A16T, T19C	3.22 XX Not Detected
		Probe XXXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-	
1	PP851177	Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-, A14-	6.00 X limited coverage (excluded)
1	PP851265	Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-	6.00 X limited coverage (excluded)
2	ON010016 PP851167	Reverse XXXX	Reverse Missing	0.00 XXXX limited coverage (excluded)
2	EF065160 EF065161	Probe XX	C3G, C10G	1.67 XXXX limited coverage (excluded)
		Reverse XXXX	Reverse Missing	
1	PP851247	Forward ✓✓✓	G1-, G11A	8.67 ✓✓✓ Further Analysis
		Reverse	T3C, G6C, T12-, G18T	
1	EF065164	Probe ✓✓✓	T1G, G6T	4.33 XX limited coverage (excluded)
		Reverse XXXX	Reverse Missing	
1	PP851168	Forward ✓✓	G1-, T14C	4.00 XX limited coverage (excluded)
		Reverse XXXX	Reverse Missing	
1	KP760068	Forward XXXX	G1-, C2-, C3-, T4-, G5-, T6-, T7-, T8-, G9-, A10-, G11-, C12-, G13-, T14-, C15-, A16-	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
1	MK547208	Forward XXXX	G1-, C2-, C3-, T4-, G5-, T6-, T7-, T8-, G9-, A10-, G11-, C12-, G13-, T14-, C15-, A16-, T17-, T18-	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
<u>6</u>	MW895255 MW895254 MW895253 MF767825 MF767817 MG599152	Probe XXXX	Probe Missing	0.34 XXXX limited coverage (excluded)
		Reverse XXXX	Reverse Missing	
<u>1</u>	MH665532	Forward XXXX	A16G	2.22 XXXX limited coverage (excluded)
		Probe XXXX	T1C, C3G, T4C, A5G, G9T, C11T, A17-, A18G, T20A, A27-, +7AC, +23C	
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-, A14-	
<u>1</u>	PP851252	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX limited coverage (excluded)
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-	
<u>1</u>	PX056820	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX limited coverage (excluded)
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-, A14-, T15-	
<u>4</u>	MN121171 MN121170 KY106281 KY106278	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
<u>1</u>	OQ363152	Forward XXXX	G11C, A16T	0.22 XXXX limited coverage (excluded)
		Probe XXXX	T1C, C3G, G6C, T7C, G9T, C11T, A17-, A18G, T20A, +23C	
		Reverse XXXX	Reverse Missing	
<u>1</u>	OK305934	Forward XXXX	T4G, G9A, G11C, A16T, T19C	1.33 XXXX limited coverage (excluded)
		Probe XXXX	T1C, C3G, T7C, G9C, C11T, A17G, A18-, T20A, T24C, A27C	
		Reverse XX	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, C19G	
<u>1</u>	MN419368	Forward XXXX	T4G, G9A, G11C, A16T, T19C	1.33 XXXX limited coverage (excluded)
		Probe XXXX	T1C, C3G, G6A, T7C, G9C, C11T, A17G, A18-, T20A, T24C, A27C	
		Reverse XX	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, C19G	
<u>1</u>	PV259176	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1C, C3G, G6A, T7C, G9T, C11T, A17G, A18-, T20A, +23C	
<u>2</u>	FR772354 MT079126	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, A17G, A18-, T20A, +23C	
<u>2</u>	OK054545 PQ415052	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, T23-	
[Candida] sp. (uncertain placement)/1853550 Analyzed Accessions: 93				
<u>9</u>	MK026354 ON113916 MN327091 ON864501 OR005392	NA	No Mutations	✓✓✓ Detected

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
	ON864514 ON864798 ON864878 PQ552918			
1	OP804126	Reverse   	C1-, +6G	10.00    Detected
1	OP804125	Probe   	T7G	9.34    Detected
		Reverse   	C1-, +6N	
1	KX944465	Reverse   	+5C G, A10-	10.00    Detected
1	KY209907	Probe   	G9C	9.34    Detected
		Reverse   	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-	
1	PV953157	Reverse   	C1-, C2-, T3-, C4-, C5-, G6-, C7-	10.00    Detected
1	PQ608380	Forward  	C15T	9.00    Detected
		Reverse   	T3C, G6C, +11A	
1	KY209904	Reverse   	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T20G	0.00    Not Detected
1	OQ376565	Probe   	T4A, T7G, C14A	0.67    Not Detected
1	KY209900	Reverse 	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, A14C	6.00  Not Detected
1	MN327101	Reverse 	+6G, T11A, G13T	6.00  Not Detected
1	MH368485	Probe   	T7G, A12C, A15C	0.67    Not Detected
1	MN327093	Probe   	T4C, G6C, T7C	3.67   Not Detected
		Reverse 	C1-, C2-, T3-, C4-, C5-, G13A	
1	KY582845	Forward   	C20T	1.67    Not Detected
		Reverse  	C1-, C2-, T3-, T12-	
1	MW894665	Probe   	T7G, A18T, T21A	0.34    Not Detected
		Reverse   	C1-, C2-, T3-, C4-, C7G, T11A, T17A, C19G	
1	KY489858	Probe   	C14-, A15-, C16A	0.34    Not Detected
		Reverse   	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, G13-, T20G	
1	OM349078	Probe   	G2-, C3-, C14A, T19C, A22T, C26T	0.67    Not Detected

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	PQ093787	Forward   	+11G	6.89  Not Detected
		Probe  	+3TT, +18TT, +25A	
		Reverse   	C1-, C2-, T3-, G6C	
1	MZ950606	Probe   	T4-, G9C, A18G, T19C, T21-, A22-, A27-, +7CG	0.67    Not Detected
1	KY209905	Probe   	T1C, A5T, T7G, G9C, C16A, A18T	0.34    Not Detected
		Reverse   	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, C19T, +19G	
5	MN908681 MN908680 MN908679 MN908682 MT013027	Probe   	T4A, A5C, T7C, A18G, T19C, A22T, C26T, A27C, +1CTGG, +8GGGTGT	0.67    Not Detected
28	28	Forward   	A16G	0.34    Not Detected
		Probe   	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
2	KY996540 KY996541	Forward   	A16G	0.34    Not Detected
		Probe   	T1C, C3G, T7C, G9T, C11T, A17G, A18-, T20A, +23C	
9	ON907719 KY996553 ON907754 MW408206 KY996543 KY996538 KY996536 ON907705 ON907706	Forward   	A16G	0.34    Not Detected
		Probe   	T1C, C3G, G6A, T7C, G9T, C11T, A17G, A18-, T20A, +23C	
2	ON907677 ON907674	Forward   	A16G	3.56   Not Detected
		Probe   	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse   	+7C	
3	ON907669 ON907700 ON907704	Forward   	A16G	0.34    Not Detected
		Probe   	T1C, C3G, G6A, T7C, G9T, C11T, C14T, A17G, A18-, T20A, +23C	
2	ON907666 ON907709	Forward   	A16G	0.34    Not Detected
		Probe   	T1C, C3G, T4C, G6A, T7C, G9T, C11T, A17G, A18-, T20A, +23C	
1	KY996547	Forward   	A16G	0.34    Not Detected
		Probe   	T1G, C3T, T4C, A5T, G6C, C11G, A12T, A15C, A18T, +13G, +24CT	
6	MZ568152 PP931188 MZ568153 OP045258 KY458748 OQ376564	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	0.67    Not Detected
1	ON907727	Forward   	T6G, C12G, T14G, A16G, T17G	0.34    Not Detected
		Probe   	T1C, C3G, G6A, T7C, C11T, C14T, A17G, A18-, T20A, +23C	
1	PQ309277	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	0.67    Not Detected

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	MZ568154	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-	Not Detected
1	MG649458	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-	Not Detected
1	ON907710	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX
		Reverse X	+6G, T8C, A10T	Not Detected
1	MG817596	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	0.34 XXX
		Reverse XXX	C7G, C19G	Not Detected
1	ON907684	Forward XXX	G11C, A16T	0.34 XXX
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	Not Detected
1	LC875769	Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-	6.00 X limited coverage (excluded)
3	MK019122 ON907683 ON907761	Reverse XXX	Reverse Missing	0.00 XXX limited coverage (excluded)
1	LC752261	Forward ✓✓✓	T4A	6.22
		Probe ✓✓✓	C26T, A27C	✓ limited coverage (excluded)
		Reverse XXX	Reverse Missing	
7	OQ677531 PP982893 OQ381287 MF940120 MF940118 MF940124 KX757236	Probe XXX	Probe Missing	0.34 XXX
		Reverse XXX	Reverse Missing	limited coverage (excluded)
1	PQ242608	Forward XXX	G9T, C12G, A16C, C20T	0.22
		Probe XXX	Probe Missing	XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
1	MN327090	Forward ✓✓✓	T4C	3.56
		Probe XXX	Probe Missing	XX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
Candida dubliniensis/42374 Analyzed Accessions: 66				
6	MK027122 KT715474 MT132376 MW709487 MH591468 OM033485	NA	No Mutations	✓✓✓ Detected

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	MZ045916	Probe   	T7G	8.67    Detected
		Reverse   	T12-	
1	MZ227294	Reverse   	C1-	10.00    Detected
1	PP475380	Forward   	A16G	0.34    Not Detected
		Probe   	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
55	55	Forward   	A16G	0.34    Not Detected
		Probe   	T1G, C3T, T4C, A5T, G6C, C11G, A12T, A15C, A18T, +13G, +24CT	
1	KY514057	Forward   	A16G	0.34    Not Detected
		Probe   	T1G, A5C, T7G, C10T, A12G, A15-, A18T, +2T, +5T, +24CT	
1	MZ536249	Forward   	A16G	3.56   Not Detected
		Probe   	T1G, A5G, T7C, C10T, A12G, A15-, A18T, +2TCT, +24CT	
		Reverse   	C1-	
1	MN629347	Probe   	C26A	4.33  limited coverage (excluded)
		Reverse   	Reverse Missing	
1	MK752670	Reverse   	Reverse Missing	0.00    limited coverage (excluded)
1	MW357730	Probe   	Probe Missing	0.34    limited coverage (excluded)
		Reverse   	Reverse Missing	
1	KF959835	Forward   	Forward Missing	0.34    limited coverage (excluded)
		Probe   	T1G, C3T, T4C, A5T, G6C, C11G, A12T, A15C, A18T, +13G, +24CT	
1	KF959837	Forward   	Forward Missing	0.34    limited coverage (excluded)
		Probe   	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, A18T, +24CT	
[Candida] sp. VVT-2012/1214789 Analyzed Accessions: 12				
2	JX406284 JX406283	NA	No Mutations	   Detected
1	JX406282	Probe   	C3A, T4G, T7G, G9C, A12C, C16A, A18T, A22T	0.34    Not Detected
		Reverse   	C1-, C2-, C5T, T8G, A10T, C19T, T20G	
1	JX406300	Forward   	A16G	0.34    Not Detected
		Probe   	T1C, C3G, T7C, G9T, A17G, A18-, T20A, +23C	
1	JX406292	Forward   	A16G	0.34    Not Detected
		Probe   	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
1	JX406298	Forward   	A16G	0.34   

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
		Probe XXXX	T1G, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +1CGGC, +23C	Not Detected
1	JX406293	Forward XXXX	A16G	3.56 XX Not Detected
		Probe XXXX	T1C, C3G, G6A, T7C, G9T, C11T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	C1-	
1	JX406290	Forward XXXX	A16G	3.56 XX Not Detected
		Probe XXXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-	
1	JX406302	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X Not Detected
		Reverse ✓✓✓	T11C	
2	JX406303 JX406285	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	0.67 XXXX Not Detected
1	JX406306	Forward XXXX	G1-, C2-, T17C	0.22 XXXX Not Detected
		Probe XXXX	G2-, A5C, G6A, T7A, C13T, A15C, C16T, T19A, C26T	
		Reverse XXXX	G6C, C7T, T8G, T15C, G18C, T21C	
1	JX406308	Forward XXXX	Forward Missing	0.22 XXXX limited coverage (excluded)
		Probe XXXX	G6A, T7C, G9C, A15T, T21G	
		Reverse XXXX	C1-, C2-, T20C	
Cryptococcus neoformans/5207 Analyzed Accessions: 2				
1	KJ175191	NA	No Mutations	✓✓✓ Detected
1	KJ175193	Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-	10.00 ✓✓✓ Detected
1	MZ306032	Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-, A14-	6.00 X limited coverage (excluded)
1	MZ306029	Forward XXXX	A16G	3.00 XX limited coverage (excluded)
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-, A14-	
1	MG995749	Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-	10.00 ✓✓✓ limited coverage (excluded)
1	KJ175192	Probe ✓✓✓	C26-, A27-	4.33 XX limited coverage (excluded)
		Reverse XXXX	Reverse Missing	
Aureobasidium pullulans/5580 Analyzed Accessions: 2				
2	MZ227295 MF380668	NA	No Mutations	✓✓✓ Detected
1	MT549153	Reverse XXXX	Reverse Missing	0.00 XXXX

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
				limited coverage (excluded)
Pichia kudriavzevii/4909 Analyzed Accessions: 2				
<u>2</u>	OR975919 MN795590	NA	No Mutations	✓✓✓✓ Detected
<u>1</u>	ON142638	Probe XXXX	Probe Missing	0.34 XXXX
		Reverse XXXX	Reverse Missing	limited coverage (excluded)
Candida/5475 Analyzed Accessions: 2				
<u>1</u>	KU870989	Reverse ✓	C1-, C2-, T3-, C4-, C5-, T12-	6.67 ✓ Detected
<u>1</u>	KP960518	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	0.67 XXXX Not Detected
<u>1</u>	KT356206	Forward XXXX	T6G, C12G, A16C, C20T	0.00 XXXX
		Reverse XXXX	Reverse Missing	limited coverage (excluded)
<u>1</u>	KR351310	Probe XXXX	C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, T23-, T24-, T25-, C26-, A27-	0.34 XXXX
		Reverse XXXX	Reverse Missing	limited coverage (excluded)
<u>13</u>	LT795047 KR185773 LT795050 LT795043 LT719074 LT795049 LT719073 KU218498 KR185772 KP659258 KP659257 KP659256 KP659259	Probe XXXX	Probe Missing	0.34 XXXX
		Reverse XXXX	Reverse Missing	limited coverage (excluded)
<u>1</u>	LT719069	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-	limited coverage (excluded)
Malassezia restricta/76775 Analyzed Accessions: 1				
<u>1</u>	OR544594	Probe ✓✓✓✓	G6A	9.34 ✓✓✓✓
		Reverse ✓✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-	Detected
<u>1</u>	OR544595	Forward XXXX	G1-, C20T	0.00 XXXX
		Reverse XXXX	Reverse Missing	limited coverage (excluded)
<u>1</u>	PP976978	Forward XXXX	G1-, C2-, C3-, T4-, G5-, T6-, T7-, T8-, G9-, A10-, G11-, C12-, G13-, T14-, C15-, A16-, C20T	0.22 XXXX
		Probe XXXX	A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, T23-, T24-, T25-, C26-, A27-	limited coverage (excluded)
		Reverse XXXX	Reverse Missing	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
Wickerhamomyces anomalus/4927 Analyzed Accessions: 1				
<u>1</u>	MK027125	Reverse   	C1-, C2-, T3-, T9-	10.00    Detected
[Candida] sp. N17/378043 Analyzed Accessions: 1				
<u>1</u>	DQ640769	Reverse   	T3C, T9A	10.00    Detected
Spathaspora/412764 Analyzed Accessions: 1				
<u>1</u>	MH591472	NA	No Mutations	   Detected
[Candida] sp. BIBHUCPB 16/1091136 Analyzed Accessions: 1				
<u>1</u>	JF909923	NA	No Mutations	   Detected
Candida parapsilosis/5480 Analyzed Accessions: 1210				
<u>552</u>	552	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	0.67    Not Detected
<u>2</u>	MN826506 MW080686	Forward   	A16G	0.34    Not Detected
		Probe   	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
<u>1</u>	LC643454	Probe   	G6A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3T, +7AA	0.67    Not Detected
<u>2</u>	PX107604 LC389814	Probe   	G2A, T4A, G6-, T7-, G8-, G9-, C10A, A12T, C16A, T19C, A22C, T25C, A27-	0.67    Not Detected
<u>1</u>	LC389747	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, T23C, A27-	0.67    Not Detected
<u>2</u>	PX096482 JX463249	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C	0.67    Not Detected
<u>1</u>	EF543645	Probe   	C3G, G6-, T7C, G8A, G9A, C10T, A12T, C16A, T19C, A22C, T25C, A27-	0.67    Not Detected
<u>8</u>	PX096533 PX096499 PX096539 PX096531 PX096544 PX096527 PX096534 PX096487	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22-, T23C, A27C	0.67    Not Detected
<u>2</u>	OM756733 KJ451697	Probe   	C3G, G6C, T7A, G8A, G9A, C11T, A12C, C14A, C16A, T19C, A22C, T25C, A27-	0.67    Not Detected
<u>1</u>	LC390086	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22T, T23C, A27-	0.67    Not Detected

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
4	PX096540 PX096526 PX096545 PX096548	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19A, T20C, A22T, T23C, A27C	0.67 XXX Not Detected
6	KU950722 KY657567 OK303414 MN747310 KT191105 MN747311	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X Not Detected
		Reverse ✓✓✓	C1-	
1	MW703702	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22-, A27C	5.33 X Not Detected
		Reverse ✓✓✓	+9T	
35	35	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X Not Detected
		Reverse ✓✓✓	C1-, C2-	
1	KM115143	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25-, C26-, A27-	0.67 XXX Not Detected
2	PP494242 PX096524	Probe XXX	C3G, G6C, T7A, G8A, G9A, C11T, A12C, C16A, T19C, A22-, T23C, A27C	0.67 XXX Not Detected
14	OL687495 EU871509 LC389122 OW985494 OW986219 LC389810 KP783504 LC389139 OW983289 OK303405 OK317693 MW709969 LC389787 OW984120	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X Not Detected
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-	
5	JN989535 EU564204 MZ020644 JN989522 KC777378	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X Not Detected
		Reverse ✓✓✓	T11A	
8	LC389996 LC389120 LC389985 MT814290 OM523857 OM523858 LC390067 LC389984	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X Not Detected
		Reverse ✓✓✓	T8C	
3	AB109223 PP733852 KJ706418	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X Not Detected
		Reverse ✓✓✓	T3C	
1	PP844419	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X Not Detected
		Reverse ✓✓✓	C7A	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	FJ662415	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, T20A, A22C, T25C, A27-	0.67 XXX Not Detected
3	ON181974 MZ769005 MN747315	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	C7T	Not Detected
3	KM113867 KU052789 PP977577	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	G6C	Not Detected
2	MW819639 OM523851	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	C7G	Not Detected
2	MW703694 OR287171	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	A10T	Not Detected
26	26	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-	Not Detected
1	OM523863	Probe XXX	C3G, G6C, T7A, G8A, G9A, C11T, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	C7G, T8C	Not Detected
26	26	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	+7C	Not Detected
25	25	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	+6G	Not Detected
1	LC389146	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-	Not Detected
1	EU484061	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	+6N	Not Detected
4	OP895150 KX652405 MK267727 GQ280288	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	+6T	Not Detected
5	KP131782 KP131764 KP674489 JN989536 KJ734198	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	+9T	Not Detected
2	AB109224 AB109231	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	+5N	Not Detected
3	JN989524 JN989516 JN989532	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
				Not Detected

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
		Reverse   	+10A	
4	LC390139 LC389118 KP674743 LC390144	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 
		Reverse   	+6A	Not Detected
15	LC389788 KP131755 LC389153 LC389151 LC389784 LC389143 LC389132 LC389125 LC389802 LC389743 LC389792 LC389778 LC389756 LC389152 OR784798	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 
		Reverse   	C1-, C2-, T3-, C4-	Not Detected
1	FR774537	Forward   	G1-, C2-, C3-, T4-	4.83
		Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	 Not Detected
1	LC390156	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, T21-, A22-, T23C, A27-	0.67    Not Detected
5	LC390062 LC389954 LC389972 KM113926 OM523853	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 
		Reverse   	T9-	Not Detected
3	KM113884 KM113880 KM113895	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 
		Reverse   	G6-	Not Detected
3	KT067766 OQ184042 PV130509	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 
		Reverse   	+6G C	Not Detected
1	MF767641	Forward   	A16G	0.34
		Probe   	T1-, C3G, G6C, T7A, G8A, G9A, C11T, A12-, C16A, T19C, A22C, T25-	   Not Detected
1	MG599213	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 
		Reverse   	C1-, +9T	Not Detected
1	GQ280301	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 
		Reverse   	+5C G	Not Detected
14	KM113855 KM113854 MH782231 PQ533515 PQ579417 OL639156 MF767651	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	0.67    Not Detected

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
	MF767646 MF767645 PQ530944 MF767668 MW895352 MF767657 MN807052			
1	OR091288	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	4.67 XX
		Reverse ✓✓✓	T12-	Not Detected
1	MT731467	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T24-, T25-, C26-, A27C	0.67 XXX Not Detected
1	KP675355	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	A10-	Not Detected
1	MT729884	Forward ✓✓	T14C	4.33 XX
		Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	Not Detected
9	OL774548 MH299862 MK392005 MW819637 KP674744 KM113857 KM113990 KM113890 AB109235	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	C7G, T8C	Not Detected
1	JN989526	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	0.34 XXX
		Reverse XXXX	T11A, A22T	Not Detected
1	OM959548	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	A10T, T11A	Not Detected
1	LC390097	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	C7A, T8C	Not Detected
1	KJ706736	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C	5.33 X
		Reverse ✓✓✓	A10T	Not Detected
3	PV363933 OM237275 PV363932	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	T8C, A10T	Not Detected
2	LC390068 KM113862	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	C7T, T8C	Not Detected
1	LC389739	Probe XXXX	C3G, G6C, T7A, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-	Not Detected

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	LC389765	Probe XXXX	C3G, G6-, T7C, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 ✗
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-	Not Detected
4	MT815460 JQ993383 LC389986 AB109288	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 ✗
		Reverse ✓✓✓	C1-, C2-, T3-, T8C	Not Detected
1	MN450875	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 ✗
		Reverse ✓✓✓	C1-, T9-	Not Detected
3	OM959620 OM959541 OM959655	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 ✗
		Reverse ✓✓✓	C1-, C2-, T3-, A10T	Not Detected
16	16	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 ✗
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-	Not Detected
1	KP131759	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 ✗
		Reverse ✓✓✓	C1-, C2-, T3-, G6C	Not Detected
1	MN450869	Forward ✓✓✓	T8-	6.89 ✓ Not Detected
		Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	
		Reverse ✓✓✓	C1-	
1	MG599241	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 ✗
		Reverse ✓✓✓	C1-, C2-, T3-, T11A	Not Detected
1	KY075668	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 ✗
		Reverse ✓✓✓	C1-, C2-, +7C	Not Detected
1	KY075670	Probe XXXX	T1G, C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 ✗
		Reverse ✓✓✓	+6G	Not Detected
1	PQ248795	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 ✗
		Reverse ✓✓✓	+6N, T8C	Not Detected
1	KP131758	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	4.67 ✗✗
		Reverse ✓✓✓	A10T, +12T	Not Detected
1	OM523859	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 ✗
		Reverse ✓✓✓	C7G, +7G	Not Detected
1	OM523862	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 ✗
		Reverse ✓✓✓	C7G, +7A	Not Detected

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	LC389119	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 ×
		Reverse ✓✓✓	G6A, T9-	Not Detected
37	37	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 ×
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-	Not Detected
1	KC182053	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C	3.33 × ×
		Reverse ×	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-	Not Detected
2	AB109286 AB109282	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 ×
		Reverse ✓✓✓	T3C, T9-	Not Detected
1	PV363916	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 × ×
		Reverse ×	C1-, C2-, T8C, A10T	Not Detected
1	AB109292	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 ×
		Reverse ✓✓✓	T3C, C7G, T8C	Not Detected
7	AB109233 AB109290 AB109229 AB109291 AB109230 AB109234 KM113868	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 ×
		Reverse ✓✓✓	C1-, C2-, T3-, T9-	Not Detected
1	KP674827	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 ×
		Reverse ✓✓✓	G6C, C7G, A10T	Not Detected
5	PQ206277 OP618190 PQ206278 KP674736 KP674519	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 ×
		Reverse ✓✓✓	C1-, C2-, T3-, A10-	Not Detected
3	KP674908 MH166866 KP674969	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 × ×
		Reverse ×	C7G, T8C, A10T	Not Detected
1	LC389738	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22T, T23C, A27-	5.33 ×
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-	Not Detected
1	KU961981	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	0.34 × × ×
		Reverse XXXX	C1-, C2-, T3-, C4-, C5-, C19G	Not Detected
1	KP674652	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 × ×
		Reverse ×	C7A, T8C, A10T	Not Detected
1	EF195357	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.67 × ×
		Reverse ✓	C1-, C2-, T3-, T12-	Not Detected

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
43	43	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-	Not Detected
3	MF462176 MK503153 KP675363	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, T8C, A10T	Not Detected
5	OP618179 OP618188 OP618187 OP618193 MT153735	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	4.67 XX
		Reverse ✓✓✓	+6G, T12-	Not Detected
1	PV074065	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C7G, A10T	Not Detected
1	OM681576	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX
		Reverse X	+5S, T8C, A10T	Not Detected
1	JX463245	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	0.34 XXX
		Reverse XXXX	T20C	Not Detected
20	20	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-	Not Detected
1	OM681577	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, A10T	Not Detected
3	MW709971 KP675635 KP675616	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX
		Reverse X	+6G, T8C, A10T	Not Detected
1	MN078122	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	0.67 XXX Not Detected
10	EF193066 KP674607 MT729857 MF380666 EF198017 LC790897 MT729856 OP618194 MK392002 MT729855	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	+6G G, A10-	Not Detected
2	KP674651 KP675285	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	+5C G, A10-	Not Detected
2	PV870148 PP195927	Probe XXXX	C3G, G6C, T7A, G8A, G9A, C11T, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-	Not Detected

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	OP618189	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	4.67 XX
		Reverse ✓✓✓	+5C G, T12-	Not Detected
1	AB109284	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX
		Reverse X	T3C, A10-, T11-	Not Detected
8	JN989527 JN989519 JN989525 JN989517 JN989537 KX652403 MH445556 JN989533	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T11A	Not Detected
1	PP033709	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX
		Reverse X	T3C, T8C, A10-	Not Detected
1	PV363915	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, A10T	Not Detected
1	EF192217	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX
		Reverse X	C1-, T8C, A10C, +12T	Not Detected
1	OQ876043	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.67 XX
		Reverse ✓	C1-, C2-, T3-, C4-, C5-, T12-	Not Detected
6	KC422431 OR507588 KP675663 KP204942 KU961982 KP674958	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, A10-	Not Detected
1	EF191046	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	T3C, G6C, A10-	Not Detected
1	KP131779	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX
		Reverse X	+6G, A10T, +11A	Not Detected
1	PX096507	Probe XXXX	C3G, G6C, T7A, G8A, G9A, C11T, A12C, C16A, T19C, A22C, T25C, A27-	0.67 XXX Not Detected
1	KJ451700	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX
		Reverse X	C1-, C2-, T9C, T11C, T12C	Not Detected
66	66	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-	Not Detected

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	MT729838	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.67 XX
		Reverse ✓	C1-, C2-, T3-, G6C, T12-	Not Detected
1	LC389147	Probe XXXX	G2A, T4A, G6-, T7-, G8-, G9-, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-	Not Detected
5	OM959604 PQ796639 EF187252 OM959551 PV363934	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.67 XX
		Reverse ✓	C1-, C2-, T3-, C4-, C5-, G6-, T12-	Not Detected
1	KC254004	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33 X
		Reverse ✓✓✓	+6G	Not Detected
1	OP603449	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, G6C, A10T, T11A	Not Detected
58	58	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-	Not Detected
1	MW710254	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-	Not Detected
1	OM523860	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	G6T	Not Detected
1	OQ713711	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-	Not Detected
1	KY075672	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22-, T23C, T25C	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-	Not Detected
1	MF510442	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	0.34 XXX
		Reverse XXXX	G13C, T15G, T17G, T21G	Not Detected
1	KY102311	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, T21-, A22-, T24C, A27C	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-	Not Detected
2	MT876148 MT876137	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, G13T	Not Detected
1	MT729885	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	0.34 XXX
		Reverse XXXX	C4G, C7G, T11C, T17A, C19G	Not Detected

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
<u>1</u>	OK090870	Probe XXXX	T1-, G2-, C3G, T4G, G6A, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, G6C, A10T, A14C	Not Detected
<u>1</u>	KC182136	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	0.34 XXX
		Reverse XXXX	C1-, +6G, C19G, T21A	Not Detected
<u>1</u>	MW132093	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-	Not Detected
<u>1</u>	AB109281	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16-, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, T9-	Not Detected
<u>1</u>	KY560331	Forward XXXX	A16T	3.56 XX Not Detected
		Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-	
<u>1</u>	OR784800	Probe XXXX	C3G, G6C, T7A, G8A, G9A, C11T, A12C, C14A, C16A, T19C, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-	Not Detected
<u>1</u>	KP131768	Probe XXXX	C3G, G6-, T7-, G8-, G9-, C11A, C13A, A15T, A17C, T19A, T20A, T21A, A22C, T24C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-	Not Detected
<u>1</u>	MW703799	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22-, T23C, T25C	0.34 XXX
		Reverse XXXX	T12G, A16C, T17C, T20G	Not Detected
<u>1</u>	OR243746	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	0.34 XXX
		Reverse XXXX	C1-, C2-, T3-, C4-, C5-, G6-, C7-, C19G, T20G	Not Detected
<u>1</u>	AB109277	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-	Not Detected
<u>1</u>	KY102314	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16-, T19C, T20C, A22C, T23C, A27C	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-	Not Detected
<u>1</u>	EU871507	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-	Not Detected
<u>2</u>	MT322428 MT322429	Forward ✓✓✓	T8C	3.56 XX Not Detected
		Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	
		Reverse XXXX	A10G, T11A, T21G	
<u>1</u>	KY075671	Probe XXXX	G2A, T4A, G6-, T7-, G8-, G9-, C10-, C11-, C14T, T19A, T20A, T21C, A22T, T24C, A27-	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-	Not Detected

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
2	KM065472 KY685084	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-	Not Detected
1	MH748620	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-	Not Detected
7	PP800201 PP800203 MT322430 PP800202 PP800204 PP800206 PP800205	Forward ✓✓✓	T8C	3.56 XX Not Detected
		Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	
		Reverse XXX	C1-, C2-, T8G, A10G, T11A, T21G	
2	PP195928 PP204164	Probe XXX	C3G, G6C, T7A, G8A, G9A, C11T, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-	limited coverage (excluded)
56	56	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-	limited coverage (excluded)
11	OL334739 PV739597 PP858936 PP858935 MH782053 PV384124 PQ570470 OR199906 PP218380 MT378269 MW940754	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-	limited coverage (excluded)
1	PP204162	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-	limited coverage (excluded)
22	22	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-, A14-	limited coverage (excluded)
19	19	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	0.67 XXX
		Reverse	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-, A14-, T15-	limited coverage (excluded)
1	KC556814	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	0.34 XXX
		Reverse XXX	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-, A22G	limited coverage (excluded)

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
7	PV416740 PV053617 KP131791 KT191118 PP229797 KU987850 KY560336	Probe XXXX Reverse XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27- C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-, A14-, T15-, A16-	0.67 XXXX limited coverage (excluded)
1	KP131770	Probe XXXX Reverse X	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, T21-, A22-, T23C, A27- C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-	3.33 XX limited coverage (excluded)
38	38	Probe XXXX Reverse XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27- Reverse Missing	0.67 XXXX limited coverage (excluded)
1	MW181235	Probe XXXX Reverse X	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27- C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-	3.33 XX limited coverage (excluded)
2	PQ345548 MW197694	Probe XXXX Reverse XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27- Reverse Missing	0.67 XXXX limited coverage (excluded)
1	KC858280	Probe XXXX Reverse X	G2A, T4A, G6-, T7-, G8-, G9-, C10A, A12T, C16A, T19C, A22C, T25C, A27- C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-, A14-	3.33 XX limited coverage (excluded)
407	407	Probe XXXX Reverse XXXX	Probe Missing Reverse Missing	0.34 XXXX limited coverage (excluded)
1	FJ545243	Probe XXXX Reverse X	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27- C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-	3.33 XX limited coverage (excluded)
2	KP131765 KP131776	Probe XXXX Reverse XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, T21-, A22-, T23C, A27- Reverse Missing	0.34 XXXX limited coverage (excluded)
1	OP618204	Probe XXXX Reverse	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C14A, C16A, T19C, A22C, T25C, A27- C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-, A14-, T15-	0.67 XXXX limited coverage (excluded)
2	PV870146 MN888780	Probe XXXX Reverse XXXX	T1-, C3G, G6C, T7A, G8A, G9A, C11T, A12-, C16A, T19C, A22C, T25C Reverse Missing	0.34 XXXX limited coverage (excluded)
1	PX107548	Probe XXXX Reverse X	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27- C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-, A14-	3.33 XX limited coverage (excluded)

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	MN749538	Forward   	G1-	3.56  limited coverage (excluded)
		Probe   	Probe Missing	
		Reverse   	Reverse Missing	
1	MN365727	Forward   	G9T	3.56  limited coverage (excluded)
		Probe   	Probe Missing	
		Reverse   	Reverse Missing	
1	KY611781	Forward   	+16A	0.22    limited coverage (excluded)
		Probe   	Probe Missing	
		Reverse   	Reverse Missing	
Candida orthopsilosis/273371 Analyzed Accessions: 233				
1	MK480614	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	0.67    Not Detected
1	EU484058	Probe   	C3G, G6C, T7A, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	0.67    Not Detected
1	OP970298	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27C	0.67    Not Detected
2	PQ576388 MW936651	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33  Not Detected
		Reverse   	C1-, C2-	
115	115	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	0.67    Not Detected
1	PV444326	Probe   	G2A, T4A, G6A, G8-, G9-, C10-, A12T, C16A, T21-, A22-, T23C, A27-	0.67    Not Detected
4	OR024668 MF380959 PQ605669 FJ746067	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33  Not Detected
		Reverse   	C1-	
8	MH166853 LC389307 OP895156 OK305956 OQ024911 LC389303 LC389308 LC389310	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33  Not Detected
		Reverse   	C1-, C2-, T3-	
9	PP906217 PP906219 OR734352 PP906213 PP906220 MZ254979 PP906214 MW358890 MZ254977	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33  Not Detected
		Reverse   	+6G	
2	MW710102 PQ050711	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33 

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
		Reverse   	+7C	Not Detected
1	FJ515199	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	0.67    Not Detected
3	OM943448 LC389314 FJ746066	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33  Not Detected
		Reverse   	+6A	
3	FM178399 LC389313 LC389306	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33  Not Detected
		Reverse   	C1-, C2-, T3-, C4-	
1	MH793868	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33  Not Detected
		Reverse   	C1-, C2-, T3-, C4-, C5-	
2	JN797502 MK394126	Probe   	T1-, G2-, C3-, A5G, G8A, G9C, C10A, C11A, C14T, A15C, T19A, T20A, A22C, T25C	0.67    Not Detected
1	PP218385	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33  Not Detected
		Reverse   	C7G, A10T	
1	PP906215	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33  Not Detected
		Reverse   	C7G, +9T	
2	KU987859 JN980098	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33  Not Detected
		Reverse   	+6G, T11A	
1	MZ045860	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33  Not Detected
		Reverse   	T8C, A10T	
4	OK310783 OK310782 OP763777 LC389312	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33  Not Detected
		Reverse   	C1-, C2-, T3-, C4-, C5-, G6-	
1	MW894663	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	0.34    Not Detected
		Reverse   	C7G, C19G	
1	KY963104	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33  Not Detected
		Reverse   	A10T, +12T	
1	MG599243	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33  Not Detected
		Reverse   	C1-, C2-, T3-, +9T	
9	OK310781 MG009517 OK310780 FM178397 LC389305 MK156300 LC389304	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33  Not Detected
		Reverse   	C1-, C2-, T3-, C4-, C5-, G6-, C7-	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
	OW985499 LC389309			
3	PP906216 KX376255 ON844319	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33 ✗
		Reverse ✓✓✓	C7G, T8C, A10T	Not Detected
2	MN747309 OR734359	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33 ✗
		Reverse ✓✓✓	C1-, C2-, T3-, A10-	Not Detected
1	KJ706562	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	0.34 XXX
		Reverse XXX	T21A	Not Detected
3	MG241511 MZ255125 LC389315	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33 ✗
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-	Not Detected
1	KC544482	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33 ✗
		Reverse ✓✓✓	C1-, C2-, T3-, T8C, A10T	Not Detected
1	EU247754	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	3.33 XX
		Reverse ✗	T8C, A10T, +11A	Not Detected
1	PP906218	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33 ✗
		Reverse ✓✓✓	+5C G, T12-	Not Detected
2	OQ999001 MW181232	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33 ✗
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-	Not Detected
2	MH793831 MN699483	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33 ✗
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, T12-	Not Detected
5	JN989501 JN989504 JN980100 JN989502 JN989500	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33 ✗
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T11A	Not Detected
13	KY102262 OP763778 KY102266 KY102270 OQ860781 KY102271 MZ956741 KP068749 KY102272 KY102261 KY102263 KP131749 MN747849	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	3.33 XX
		Reverse ✗	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-	Not Detected
1	OR734360	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33 ✗
		Reverse ✓✓✓	C1-, C2-, T3-, G6A, T12-	Not Detected

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	KF307728	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	3.33 XX
		Reverse X	T3C, G6C, A10T, T12C	Not Detected
16	16	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-	Not Detected
1	KJ706403	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33 X
		Reverse ✓✓✓	T3C	Not Detected
1	EU247755	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, A10T, G13T	Not Detected
1	KJ706721	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16-, T19A, T20A, A22C, T25C	5.33 X
		Reverse ✓✓✓	A10T	Not Detected
1	PV733663	Probe XXXX	C3G, T4A, G6-, T7-, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T11A	Not Detected
1	KU865179	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, C7G, T12-, G13-	Not Detected
1	MT443950	Probe XXXX	T1-, G2-, C3G, T4G, G6A, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-	Not Detected
6	MF510449 LC713268 MF510448 OW986154 MT875261 MT482655	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-	limited coverage (excluded)
7	MN784461 MH779839 JN989497 JN980099 JN989511 JN989510 JN989507	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-	limited coverage (excluded)
3	MT680660 PV384123 MT150685	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-, A14-	limited coverage (excluded)
1	PP204051	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27C	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-, A14-	limited coverage (excluded)

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
3	MT443926 MF940123 MT443919	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	3.33 XX limited coverage (excluded)
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-, A14-, T15-	
3	JN989509 KY102274 KY560329	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	0.67 XXX limited coverage (excluded)
		Reverse	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-, A14-, T15-, A16-	
6	JN989508 JN944022 OK135353 EF568026 OK135355 KU052049	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	0.67 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
126	126	Probe XXX	Probe Missing	0.34 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
Candida metapsilosis/273372 Analyzed Accessions: 90				
43	43	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	0.67 XXX Not Detected
4	MT815911 LC389044 LC389041 KM982979	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	5.33 X Not Detected
		Reverse ✓✓✓	+7C	
1	JX463240	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	4.33 XX Not Detected
		Reverse ✓✓	G18C	
1	KU200445	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, A22C, T25C, A27-	0.67 XXX Not Detected
3	PQ721032 OQ726380 OR731586	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	5.33 X Not Detected
		Reverse ✓✓✓	C1-	
1	KY102208	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	5.33 X Not Detected
		Reverse ✓✓✓	C1-, C2-	
3	PQ626156 LC389042 KP895602	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	5.33 X Not Detected
		Reverse ✓✓✓	C1-, C2-, T3-	
3	MW358888 PQ605665 MW358889	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	5.33 X Not Detected
		Reverse ✓✓✓	+6G	
1	KX394415	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	5.33 X Not Detected
		Reverse ✓✓✓	+5N	
1	LC317507	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	5.33 X

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
		Reverse   	C1-, C2-, T3-, C4-, C5-, G6-	Not Detected
1	PV945930	Probe   	T1C, G2A, G6T, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-, +4C, +9AACAA	5.33 
		Reverse   	+7C	Not Detected
5	OW984115 MN699482 OW984118 LC389043 MH715400	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	5.33 
		Reverse   	C1-, C2-, T3-, C4-, C5-, G6-, C7-	Not Detected
4	MZ422942 LC317505 MG009520 LC389046	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	5.33 
		Reverse   	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-	Not Detected
1	MT443953	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, T20A, A22T, T23C, A27-	4.33  
		Reverse  	T17G	Not Detected
1	OL873259	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	5.33 
		Reverse   	+5C G, A10-	Not Detected
1	OR264359	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	3.67  
		Reverse 	T3C, G6C, T12-	Not Detected
1	PQ605666	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	5.33 
		Reverse   	C1-, C2-, T3-, C4-, C5-, A10-	Not Detected
8	KY102205 EF190228 KY102207 KY102206 EF193072 GQ280290 KY102204 OL873267	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	3.33  
		Reverse 	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-	Not Detected
1	KP674707	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	3.33  
		Reverse 	C7G, T8C, A10T	Not Detected
1	MH425629	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	3.33  
		Reverse 	T3C, G6C, A10T, T11A	Not Detected
1	MT814298	Probe   	T1C, G2A, G6T, C10A, A12T, C16A, A18C, T19A, A22C, A27-, +4C, +9AACAA	0.34   
		Reverse   	+6G, T8C, T20C	Not Detected
1	OW984116	Probe   	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	3.33  
		Reverse 	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-	Not Detected
1	MT815464	Probe   	T1G, C3-, T4-, G6A, T7C, G8A, G9A, C10A, A12T, C16A, A18C, A22C, T25C, A27-	0.34   
		Reverse   	C1-, T8C, T9C, A10T, T17C, G18-	Not Detected

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	MT814217	Probe XXXX	T1C, G2A, A5C, G6A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-, +9GACAA	5.33 XX
		Reverse ✓✓✓	C1-, C2-, T8C	Not Detected
1	OQ729888	Probe XXXX	T1C, G2A, G6T, C11A, C13A, A15T, A17C, T19A, T20A, T21C, A27C, +4C, +9TA, +23C	5.33 XX
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-	Not Detected
1	MN733071	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-	limited coverage (excluded)
1	MN733072	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-, A14-	limited coverage (excluded)
5	PV478237 MF510445 MF510444 PP326263 MF510446	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-	limited coverage (excluded)
2	OW984114 JN989514	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-	limited coverage (excluded)
1	JN989513	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-, A14-	limited coverage (excluded)
63	63	Probe XXXX	Probe Missing	0.34 XXXX
		Reverse XXXX	Reverse Missing	limited coverage (excluded)
1	MT443918	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	0.67 XXXX
		Reverse	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-, A14-, T15-	limited coverage (excluded)
1	MF940132	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	0.67 XXXX
		Reverse XXXX	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-, A14-, T15-, A16-	limited coverage (excluded)
1	EU484056	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	0.67 XXXX
		Reverse XXXX	Reverse Missing	limited coverage (excluded)
Lodderomyces elongisporus/36914 Analyzed Accessions: 60				
17	17	Probe XXXX	G6A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3T, +7AA	0.67 XXXX Not Detected
4	AY391848 FR819714 ON209375 MK394128	Probe XXXX	G6A, T7A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3TTAA	0.67 XXXX Not Detected

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	PP758275	Probe XXXX	G6A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3T, +7AA	5.33 X
		Reverse ✓✓✓	C1-, C2-	Not Detected
1	EF394941	Probe XXXX	G6A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3T, +7AA	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-	Not Detected
2	KF728813 FJ515179	Probe XXXX	G6A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3T, +7AA	5.33 X
		Reverse ✓✓✓	+6G	Not Detected
3	KP764944 KP764992 MT443894	Probe XXXX	G6A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3T, +7AA	5.33 X
		Reverse ✓✓✓	+7C	Not Detected
1	KY076619	Probe XXXX	G6A, T7A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3TTAA	5.33 X
		Reverse ✓✓✓	T8C	Not Detected
1	ON142344	Probe XXXX	G6A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3T, +7AA	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, A10T	Not Detected
3	OW988504 PP858958 OW986919	Probe XXXX	G6A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3T, +7AA	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-	Not Detected
1	MT443931	Probe XXXX	G6A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3T, +7AA	5.33 X
		Reverse ✓✓✓	C7G, T8C, A10T	Not Detected
4	KP341544 PP758278 LC195009 KY104073	Probe XXXX	G6A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3T, +7AA	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-	Not Detected
1	KP765025	Probe XXXX	G6A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3T, +7AA	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, T8C, A10T	Not Detected
1	NR_111593	Probe XXXX	G6A, T7A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3TTAA	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-	Not Detected
1	KM361506	Probe XXXX	G6A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3T, +7AA	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-	Not Detected
1	ON242338	Probe XXXX	G6A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3T, +7AA	5.33 X
		Reverse ✓✓✓	C1-, C2-, T8C, A10T, T12A	Not Detected
7	KY104069 LC473130 KY104074 KY104070 KY104077 KY104076 KY104071	Probe XXXX	G6A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3T, +7AA	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-	Not Detected

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
2	GU319980 MH059652	Probe XXXX	G6A, T7A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3TTAA	5.33 XX
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-	Not Detected
1	KY985266	Probe XXXX	G6A, T7A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3TTAA	5.33 XX
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, T12-	Not Detected
1	PV478232	Probe XXXX	G6A, T7A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3TTAA	5.33 XX
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-	Not Detected
2	PP858960 GQ280291	Probe XXXX	G6A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3T, +7AA	5.33 XX
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-	Not Detected
3	LC698657 LC698656 LC698660	Probe XXXX	G6A, T7A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3TTAA	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-	Not Detected
1	LC698659	Probe XXXX	G6A, T7A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3TTAA	0.67 XXX
		Reverse	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, C19T	Not Detected
1	KY104075	Probe XXXX	G6A, G8A, G9A, C10A, C11A, C14T, A15C, A17C, T20C, +3T, +22TT	5.33 XX
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-	Not Detected
2	MT819998 PP858963	Probe XXXX	G6A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3T, +7AA	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-	limited coverage (excluded)
1	PP858961	Probe XXXX	G6A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3T, +7AA	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-	limited coverage (excluded)
2	PP229532 MT443903	Probe XXXX	G6A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3T, +7AA	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-, A14-, T15-	limited coverage (excluded)
7	KU052050 PX067037 EU871516 ON262898 OM401653 MW362775 PP844534	Probe XXXX	G6A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3T, +7AA	0.67 XXX
		Reverse XXXX	Reverse Missing	limited coverage (excluded)
17	17	Probe XXXX	Probe Missing	0.34 XXX
		Reverse XXXX	Reverse Missing	limited coverage (excluded)
Candida viswanathii/5486 Analyzed Accessions: 17				
8	MT659652 AY139792 MT659405 KJ651202 DQ249200	Probe XXXX	T4-, G9C, A18G, T19C, T21-, A22-, A27-, +7CG	0.67 XXX Not Detected

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
	LC605168 NR_138345 MK394122			
1	MT659650	Probe XXXX	T4-, G9C, A18G, T19C, T21-, A22-, A27-, +7CG	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-	Not Detected
1	MT652637	Probe XXXX	T4-, G9C, A18G, T19C, T21-, A22-, A27-, +7CG	5.33 X
		Reverse ✓✓✓	+9T	Not Detected
1	KY102515	Probe XXXX	T4-, G9C, A18G, T19C, T21-, A22-, A27-, +7CG	5.33 X
		Reverse ✓✓✓	C1-, C2-, C7T	Not Detected
1	KU729067	Probe XXXX	T4-, G9C, A18G, T19C, T21-, A22-, A27-, +7CG	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-	Not Detected
1	MT797385	Probe XXXX	T4-, G9C, A18G, T19C, T21-, A22-, A27-, +7CG	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, T8C	Not Detected
2	MT797293 KY102511	Probe XXXX	T4-, G9C, A18G, T19C, T21-, A22-, A27-, +7CG	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-	Not Detected
1	JN714127	Probe XXXX	T4-, G9C, A18G, T19C, T21-, A22-, A27-, +7CG	4.33 XX
		Reverse ✓✓	T20A	Not Detected
1	DQ249201	Forward ✓✓✓	T7G	3.56 XX
		Probe XXXX	C3G, A5C, T7-, G9C, A15C, A18-, T19-, T20-, T21C, A22C, C26T	Not Detected
		Reverse XXXX	C1-, C2-, T3-, T11A, T12-, C19A	
1	MZ841639	Probe XXXX	T1G, C3G, T4C, G9C, A18G, T19C, T21-, A22-, A27-, +7CG	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-, A14-	limited coverage (excluded)
13	FJ542757 KC608220 KY102514 KT191121 KM507318 KY102512 KJ540109 MT659651 KU729074 KJ651201 KT191106 MW559567 MW559577	Probe XXXX	T4-, G9C, A18G, T19C, T21-, A22-, A27-, +7CG	0.67 XXX
		Reverse XXXX	Reverse Missing	limited coverage (excluded)
1	JN714128	Forward X	G1-, C2-, C3-, T4-, G5-, T6-, T7-, T8-, G9-, A10-, G11-	3.33 XX
		Probe XXXX	T4-, A12C, A18G, T19C, T21-, A22-, A27-, +7C, +11CA	limited coverage (excluded)
2	KY588577 KY588576	Forward XXXX	G1-, C2-, C3-, T4-, G5-, T6-, T7-, T8-, G9-, A10-, G11-, C12-, G13-, T14-, C15-, A16-	0.45 XXX
		Probe XXXX	T4-, G9C, A18G, T19C, T21-, A22-, A27-, +7CG	limited coverage (excluded)
		Reverse XXXX	Reverse Missing	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
<u>2</u>	KY102516 KY102513	Probe XXX	T4-, G9C, A18-, T19-, T20-, T21G, A22C, A27-, +7CG	0.34 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
<u>9</u>	OQ913732 MW559573 MW559568 KT191107 KT191127 MW559570 MW559582 KT191110 KT191111	Probe XXX	Probe Missing	0.34 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
Candida sojae/52253 Analyzed Accessions: 8				
<u>4</u>	MK267467 MW895281 MK394120 MK267466	Probe XXX	G2T, C3A, T19C, A22-, C26T	0.67 XXX Not Detected
<u>1</u>	PQ159260	Probe XXX	G2T, C3A, T19C, A22-, C26T	5.33 X Not Detected
		Reverse ✓✓✓	+4T	
<u>1</u>	OR507584	Probe XXX	G2-, C3-, C14A, A22T, C26T, +19A	0.67 XXX Not Detected
<u>1</u>	OW988212	Probe XXX	G2T, C3A, T19C, A22-, C26T	5.33 X Not Detected
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-	
<u>1</u>	FN424104	Probe XXX	G2T, C3A, T19C, A22T, T25-, C26-, A27-	5.33 X Not Detected
		Reverse ✓✓✓	+6G	
<u>9</u>	NR_137087 MG589904 PV740047 PV740040 PV740043 PV740044 PV740041 PV740042 PV740039	Probe XXX	G2T, C3A, T19C, A22-, C26T	0.34 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
<u>2</u>	PV740048 PV740096	Probe XXX	G2T, C3A, T19C, A22T, C26-, A27-	0.34 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
<u>1</u>	KJ722419	Probe XXX	G2T, C3A, T19C, A22-, T23-, T24-, T25-, C26-, A27T	0.34 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
<u>1</u>	PV029126	Probe XXX	G2T, C3A, T19C, A22-, T23-, T24-, T25-, C26-, A27-	0.34 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
<u>1</u>	KY102399	Probe XXX	G2T, C3A, T19-, T20-, T21-, A22-, T23-, T24-, T25-, C26-, A27-	0.34 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	PX107596	Probe XXXX	Probe Missing	0.34 XXXX limited coverage (excluded)
		Reverse XXXX	Reverse Missing	
Lodderomyces/36913 Analyzed Accessions: 8				
1	MG757710	Probe XXXX	G6A, T7A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3TTAA	0.67 XXXX Not Detected
1	MG757711	Probe XXXX	G6A, T7A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3TTAA	5.33 X Not Detected
		Reverse ✓✓✓	+7C	
1	MG757708	Probe XXXX	G6A, T7A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3TTAA	5.33 X Not Detected
		Reverse ✓✓✓	C7A, T8C	
4	JF781477 JF781413 JF781414 JF781408	Probe XXXX	G6A, T7A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3TTAA	5.33 X Not Detected
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-	
1	JF781457	Probe XXXX	G6A, T7A, G8A, G9A, C10G, A12T, C16A, A17T, A18C, T20A, +3TTAA	3.33 XX Not Detected
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-	
1	JN831666	Reverse XXXX	Reverse Missing	0.00 XXXX limited coverage (excluded)
Malassezia/55193 Analyzed Accessions: 6				
2	OK599032 OK598975	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X Not Detected
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-	
1	OK599031	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33 X Not Detected
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-	
1	OK598976	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX Not Detected
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-	
1	OK598974	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	5.33 X Not Detected
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-	
1	OK599033	Probe XXXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	3.33 XX Not Detected
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-	
Candida maltosa/5479 Analyzed Accessions: 4				
2	KY102187 MK394118	Probe XXXX	T1G, C3G, G6C, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, A18T, T19G, T20C, A22T	0.67 XXXX Not Detected

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
2	NR_138346 KJ722418	Probe XXX	T1A, C3T, T4G, G8-, G9-, C10-, C11-, A12-, C13A, A15T, A18T, T19G, T20C, A22T	0.67 XXX Not Detected
1	MG367311	Forward XXX	Forward Missing	0.34 XXX
		Probe XXX	C3G, A5G, G9C, C10T, C11T, C13T, A15-, C16-, A17-, A18-, A22T	limited coverage (excluded)
1	KJ722417	Probe XXX	C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, T23-, T24-, T25-, C26-, A27-	0.34 XXX
		Reverse XXX	Reverse Missing	limited coverage (excluded)
1	KY102188	Probe XXX	Probe Missing	0.34 XXX
		Reverse XXX	Reverse Missing	limited coverage (excluded)
fungal sp./1709941 Analyzed Accessions: 3				
1	OL998448	Probe XXX	T4-, G9C, A18G, T19C, T21-, A22-, A27-, +7CG	0.67 XXX Not Detected
1	MK040500	Probe XXX	T4-, G9C, A18G, T19C, T21-, A22-, A27-, +7CG	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-	Not Detected
1	MF435925	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, T8A, A10T, T11A	Not Detected
5	OP518345 MN957137 OP518346 OQ804481 MK101318	Reverse XXX	Reverse Missing	0.67 XXX limited coverage (excluded)
1	MK040499	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	0.67 XXX
		Reverse XXX	Reverse Missing	limited coverage (excluded)
1	MT669261	Probe XXX	Probe Missing	0.34 XXX
		Reverse XXX	Reverse Missing	limited coverage (excluded)
2	MH812223 MZ016062	Forward XXX	Forward Missing	0.34 XXX
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, C26G	limited coverage (excluded)
3	MZ017132 MZ017134 MZ017057	Forward XXX	Forward Missing	0.34 XXX
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, C26G	limited coverage (excluded)
1	MT548314	Forward XXX	Forward Missing	0.34 XXX
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22G, C26G, +23C	limited coverage (excluded)

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	MZ017190	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-	
[Candida] aquae-textoris/693064 Analyzed Accessions: 3				
1	KJ651205	Probe XXX	T4-, G9C, A18G, T19C, T21-, A22-, A27-, +7CG	0.67 XXX Not Detected
1	HM068913	Probe XXX	T4-, G9C, A18G, T19C, T21-, A22-, A27-, +7CG	5.33 X Not Detected
		Reverse ✓✓✓	C1-, C2-, T3-, C4-	
1	KC290944	Forward XXX	C20T	2.22 XXX Not Detected
		Probe XXX	T4-, G9C, A18G, T19C, T21-, A22-, A27-, +7CG	
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-	
3	MW559574 KU752209 KU752208	Probe XXX	T4-, G9C, A18G, T19C, T21-, A22-, A27-, +7CG	0.67 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
3	KU752210 MW559575 MW559584	Probe XXX	Probe Missing	0.34 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
Candida africana/241526 Analyzed Accessions: 3				
1	MK026279	Forward XXX	A16G	0.34 XXX Not Detected
		Probe XXX	T1G, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +1CGGC, +23C	
1	OL602114	Forward XXX	A16G	0.34 XXX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, C11T, A17G, A18-, T20A, +23C	
1	ON811599	Forward XXX	A16G	3.56 XX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
		Reverse ✓✓✓	+7C	
Candida sanyaensis/1176388 Analyzed Accessions: 2				
1	OR507579	Probe XXX	G2-, C3-, C14A, A22T, C26T, +19A	5.33 X Not Detected
		Reverse ✓✓✓	C4T	
1	PQ834795	Probe XXX	G2-, C3-, C14A, T19C, A22T, C26T	5.33 X Not Detected
		Reverse ✓✓✓	T3C, G6C, A10-	
1	MZ507576	Probe XXX	G2-, C3-, C14A, A22T, C26T, +19A	0.67 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
2	NR_138302 JQ901890	Probe XXX	G2-, C3-, C14A, T19C, A22T, C26T	0.34 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	OK050703	Probe XXX	G2-, C3-, C14A, T19-, T20-, T21-, A22-, T23-, T24-, T25-, C26-, A27-	0.34 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
fungal endophyte/410823 Analyzed Accessions: 2				
1	KR014967	Forward XXX	A16G	0.34 XXX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
1	KR016272	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	0.67 XXX Not Detected
1	KR080843	Reverse XXX	Reverse Missing	0.00 XXX limited coverage (excluded)
1	KR080851	Forward ✓✓✓	G1-, G11A	4.33 XX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
1	KR080862	Forward XXX	T4C, T6G, A10G, C12G, T14G, A16C, C20T	0.00 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
3	KR015017 KR015016 KR016890	Probe XXX	Probe Missing	0.34 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
Candida labiduridarum/434042 Analyzed Accessions: 2				
1	NR_111410	Probe XXX	G2T, C3A, T7A, G8A, C14T, C16T, T19C, T21-, A22-, C26A	0.67 XXX Not Detected
1	KY102176	Probe XXX	G2T, C3A, T7A, G8A, C14T, C16T, T19C, T21-, A22-, C26A	5.33 X Not Detected
		Reverse ✓✓✓	C1-, C2-	
2	EF658664 EF658665	Probe XXX	G2T, C3A, T7A, G8A, C14T, C16T, T19C, T21-, A22-, C26A	0.34 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
[Candida] sp. YS12A/399110 Analyzed Accessions: 2				
2	KF680225 DQ857760	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	0.67 XXX Not Detected
Candida hyderabadensis/454356 Analyzed Accessions: 2				
1	NR_159741	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	0.67 XXX Not Detected
1	KF680226	Forward ✓✓✓	G13C	4.67 XX Not Detected
		Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	
Candida cetoniae/1286810 Analyzed Accessions: 2				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	KC315799	Probe XXX	T1G, G2A, T4A, G6-, T7-, G8-, G9-, C10-, A12T, C16A, T21-, A22-, A27-	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-	Not Detected
1	NR_132875	Probe XXX	T1G, G2A, T4A, G6-, T7-, G8-, G9-, C10-, A12T, C16A, T21-, A22-, A27-	3.33 XX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-	Not Detected
Candida kantuleensis/2315425 Analyzed Accessions: 2				
2	NR_164020 LC317103	Probe XXX	G2-, C3-, G9A, T19C, A22T, T25-, C26-, A27-	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-	Not Detected
Candida neerlandica/148634 Analyzed Accessions: 1				
1	MK394121	Probe XXX	G2A, T4G, G8T, C14A, C16T, A18C, T21-, A22-, T25C, C26A, +7A	0.67 XXX Not Detected
1	NR_136988	Probe XXX	G2A, T4G, G8T, C14A, C16T, A18C, T21-, A22-, T25C, C26A, +7A	0.34 XXX
		Reverse XXX	Reverse Missing	limited coverage (excluded)
1	KY102226	Probe XXX	G2A, T4G, G8-, G9-, C10A, C11T, A12G, T20A, T21C, A22T	0.34 XXX
		Reverse XXX	Reverse Missing	limited coverage (excluded)
Candida saraburiensis/694444 Analyzed Accessions: 1				
1	NR_151809	Probe XXX	T4C, A5C, T7A, G8C, T19-, T20-, T21-, A22C, T25C, +3GG	5.33 X
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-	Not Detected
5	ON786580 ON786585 ON786583 ON786587 ON786582	Probe XXX	T4C, A5C, T7A, G8C, T19-, T20-, T21-, A22C, T25C, +3GG	0.34 XXX
		Reverse XXX	Reverse Missing	limited coverage (excluded)
1	ON786584	Probe XXX	Probe Missing	0.34 XXX
		Reverse XXX	Reverse Missing	limited coverage (excluded)
Candida margitis/1775924 Analyzed Accessions: 1				
1	PQ521377	Probe XXX	T1-, G2-, C3-, A5G, G9A, C10A, A12T, C16G, T19A, A22C, T25C, A27-, +14A	0.67 XXX Not Detected
1	NR_159750	Probe XXX	T1-, G2-, C3-, A5G, G9A, C10A, A12T, C16G, T19A, A22C, T25C, A27-, +14A	0.67 XXX
		Reverse XXX	Reverse Missing	limited coverage (excluded)
Colletotrichum/5455 Analyzed Accessions: 1				
1	ON154855	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	0.67 XXX Not Detected
Lichtheimia corymbifera/42458 Analyzed Accessions: 1				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	MT590584	Forward XXX	A16G	0.34
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	XXX Not Detected
[Candida] multigemmis/45569 Analyzed Accessions: 1				
1	EU343847	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-	XX Not Detected
Fusarium dimerum/57145 Analyzed Accessions: 1				
1	KT876500	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	5.33
		Reverse ✓✓✓	T3C	X Not Detected
Candida giganteus/271359 Analyzed Accessions: 1				
1	NR_151799	Forward XXX	A16G	2.22
		Probe XXX	G2A, T4A, G6-, T7-, G8-, G9-, T19A, T21A, A22C, +23C	XXX
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-	Not Detected
fungal endophyte sp. CEY 01/306095 Analyzed Accessions: 1				
1	AY700120	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	0.67 XXX Not Detected
fungal endophyte sp. CEY 08/306102 Analyzed Accessions: 1				
1	AY700127	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	0.67 XXX Not Detected
fungal endophyte sp. CEY 15/306109 Analyzed Accessions: 1				
1	AY700134	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	0.67 XXX Not Detected
Meyerozyma carpophila/374178 Analyzed Accessions: 1				
1	MH777073	Forward XXX	A16G	0.34
		Probe XXX	T1C, C3G, G6A, T7C, G9T, C11T, A17G, A18-, T20A, +23C	XXX Not Detected
[Candida] sp. KER38/420830 Analyzed Accessions: 1				
1	EF197971	Probe XXX	T1-, C3G, G6C, T7A, G8A, G9A, C11T, A12-, C16A, T19C, A22C, T25C, A27T	0.34
		Reverse XXX	T11C, T17G, C19A, T20A	XXX Not Detected
Candida tetragidarum/434046 Analyzed Accessions: 1				
1	NR_111411	Probe XXX	T1A, C3G, T4A, G8-, G9-, C14-, A15-, C16T, A17G, T19A, T21A	0.67 XXX Not Detected
Candida cf. neerlandica ATCC MYA-4367/594597 Analyzed Accessions: 1				
1	FJ623628	Probe XXX	G2A, T4G, G8T, C14A, C16T, A18C, T21-, A22-, T25C, C26A, +7A	0.67 XXX Not Detected
[Candida] sp. MCCF-101/595535 Analyzed Accessions: 1				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	FJ652052	Probe XXX	T4-, G9C, C16A, A18G, T19C, T21-, A22-, A27-, +7CG	0.67 XXX Not Detected
[Candida] sp. S27/600621 Analyzed Accessions: 1				
1	FJ652102	Forward XXX	C20T	0.34 XXX Not Detected
		Probe XXX	T4-, G9C, A18G, T19C, T21-, A22-, A27-, +7CG	
[Candida] sp. SD 302/687784 Analyzed Accessions: 1				
1	GQ387379	Forward ✓✓✓	G1-	6.22 ✓ Not Detected
		Probe XXX	A12C, A15C	
		Reverse ✓✓	T17G	
Saccharomycetes sp. 5B12/747485 Analyzed Accessions: 1				
1	GU931325	Forward XXX	A16G	0.34 XXX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
[Candida] sp. 4 TMS-2011/942762 Analyzed Accessions: 1				
1	HQ631066	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, A18C, T19A, A22C, T25C, A27-	5.33 × Not Detected
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-	
Zymoseptoria tritici/1047171 Analyzed Accessions: 1				
1	PP550783	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33 XX Not Detected
		Reverse ×	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-	
Saccharomycetales sp. Y5/1249556 Analyzed Accessions: 1				
1	KC121052	Probe ✓✓✓	T7G	4.33 XX Not Detected
		Reverse XXX	C4T, C7G, C19G	
fungal sp. SNB-CN36A/1476097 Analyzed Accessions: 1				
1	KJ023713	Probe XXX	T1G, T7G, A12C, A22T	0.67 XXX Not Detected
		Reverse XXX	C1-, C4T, T11A, A16T, T17A	
[Candida] sp. MM 2_9_d/1480165 Analyzed Accessions: 1				
1	KJ361482	Forward XXX	A16G	0.34 XXX Not Detected
		Probe XXX	T1C, C3G, G6A, T7C, G9A, C11T, A17G, A18-, T20A, +23C	
[Candida] sp. MM C_K/1480166 Analyzed Accessions: 1				
1	KJ361494	Probe XXX	C3G, A12C, A27C	0.34 XXX Not Detected
		Reverse XXX	C1-, C2-, G6C, T9A, C19T	
[Candida] sp. MM C_K_n/1480167 Analyzed Accessions: 1				
1	KJ361497	Probe XXX	C3G, A12C, A15C, A17C	0.34 XXX Not Detected
		Reverse XXX	C1-, C2-, T17G, C19T, T21A	
[Candida] sp. Y4/1483466 Analyzed Accessions: 1				
1	KJ183075	Probe XXX	T1G, C3G, T4C, G9C, A18G, T19C, T21-, A22-, A27-, +7CG	0.67 XXX

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
				Not Detected
Saccharomycetales sp. LY12/1491796 Analyzed Accessions: 1				
1	KJ734196	Probe XXX	T1G, T4A, T7G, G9C, A18T, A27T	0.34
		Reverse XXX	C4T, C7G, T11A, A16T, C19G	XXX Not Detected
[Candida] sp. Y3 CTNM-2014/1506481 Analyzed Accessions: 1				
1	KJ608356	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	0.67 XXX Not Detected
[Candida] sp. JS3/1639001 Analyzed Accessions: 1				
1	KP243194	Probe XXX	T4-, G9C, A18G, T19C, T21-, A22-, A27-, +7CG	5.33
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-	X Not Detected
[Candida] sp. Y574/1778432 Analyzed Accessions: 1				
1	KT596908	Probe XXX	T4-, G9C, A18G, T19C, T21-, A22-, A27-, +7CG	5.33
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-	X Not Detected
Saccharomycetales sp./1891166 Analyzed Accessions: 1				
1	MG748563	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	0.67 XXX Not Detected
Rhodotorula/5533 Analyzed Accessions: 0				
1	LC018747	Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-, A14-	6.00 X limited coverage (excluded)
6	LC018770 LC018741 LC018746 LC018752 LC018765 LC018776	Reverse XXX	Reverse Missing	0.00 XXX limited coverage (excluded)
1	LC018742	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	3.33
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-, A14-	XX limited coverage (excluded)
3	LC018785 LC018767 LC018800	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	0.34
		Reverse XXX	Reverse Missing	XXX limited coverage (excluded)
Saccharomyces cerevisiae/4932 Analyzed Accessions: 0				
1	MN809217	Reverse XXX	Reverse Missing	0.00 XXX limited coverage (excluded)
1	KY105066	Probe XXX	G2A, T4G, G8T, C14A, C16T, A18C, T21-, A22-, T25C, C26A, +7A	0.34
		Reverse XXX	Reverse Missing	XXX limited coverage (excluded)
1	KC182625	Probe XXX	Probe Missing	0.34 XXX

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
		Reverse XXX	Reverse Missing	limited coverage (excluded)
Kluyveromyces marxianus/4911 Analyzed Accessions: 0				
1	PP249687	Probe XXX	T4C, T7G, A12C, A15C	0.34 XXX
		Reverse XXX	Reverse Missing	limited coverage (excluded)
1	KF851350	Forward XXX	Forward Missing	2.89
		Probe ✓✓✓	C3G, G6C	XX
		Reverse XXX	C7G, C19G	limited coverage (excluded)
Diutina rugosa/5481 Analyzed Accessions: 0				
1	OR526619	Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-	6.00 X limited coverage (excluded)
[Candida] sp. CLIB 1309/1056988 Analyzed Accessions: 0				
1	HE574643	Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-	6.00 X limited coverage (excluded)
Candida pseudoviswanathii (nom. inval.)/2861751 Analyzed Accessions: 0				
1	KM586736	Probe XXX	T4-, G9C, A18G, T19C, T21-, A22-, A27-, +7CG	0.67 XXX
		Reverse XXX	Reverse Missing	limited coverage (excluded)
Candolleomyces candolleanus/71730 Analyzed Accessions: 0				
33	33	Forward XXX	Forward Missing	0.34 XXX
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, C26G	limited coverage (excluded)
1	OQ789378	Forward XXX	Forward Missing	0.34 XXX
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T19C, C26G	limited coverage (excluded)
1	OL764902	Forward XXX	Forward Missing	0.34 XXX
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, C26G, +22C	limited coverage (excluded)
Cortinarius/34451 Analyzed Accessions: 0				
2	HQ115590 MW031158	Forward XXX	Forward Missing	0.34 XXX
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, C26T	limited coverage (excluded)
1	OR149211	Forward XXX	Forward Missing	0.34 XXX
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24A	limited coverage (excluded)
2	KT339239 KT339229	Forward XXX	Forward Missing	0.34 XXX
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, C26A, A27T	limited coverage (excluded)

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
3	EU655675 OQ449443 KJ421167	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, C26T	
1	OR434469	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22-	
4	KM226607 PX307074 MT345240 PV469624	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, T24A	
1	KP012837	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22C, T23A	
1	OR434421	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T19A, A22T	
3	KM226598 KM226490 KM226586	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22G, C26T	
2	MF599252 MF599239	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A22G	
1	LC483900	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, C26A, +21T	
1	EU569249	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-	
1	KJ635231	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C16A, C26A, A27T	
1	KJ635220	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22T, +19A	
2	MG019346 MG597367	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22-, T23-	
Rhizophagus/1129544 Analyzed Accessions: 0				
23	23	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-	
Inocybe/71953 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
2	PQ562356 OR052575	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T19G, A22G, C26G	
3	PV200024 PV200023 PQ231207	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, T20A, C26G	
2	PV051443 PV690329	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, C26G	
2	KT800171 KT800279	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, T24A, C26T	
2	OR419897 OR419901	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, C26G	
1	MH038110	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, C26G	
2	PV469605 PV684538	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, C26G	
1	PQ231195	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, T20A, A22C, C26G	
5	PP301784 PP301788 PP301791 PP430123 PQ144499	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, C26G	
1	PX289255	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, A15-, T20C, T21-, C26G	
1	PV200025	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, T23-, C26G	
Entoloma clandestinum/566074 Analyzed Accessions: 0				
22	22	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, C26G	
Entoloma/71937 Analyzed Accessions: 0				
1	OL338308	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, T23-, C26G	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
10	AB691983 MZ611678 AB691974 AB691984 AB691982 AB691979 AB691977 AB691990 AB691975 OL338046	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, C26G	
2	MW448568 MW448567	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, C26G	
2	KY774206 KY774205	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, T25G, C26A	
1	PV650203	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, T19C, T21C, A22T, C26G	
2	OL338408 PX273840	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	Probe Missing	
1	PV520971	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22-, C26G	
Cyberlindnera fabianii/36022 Analyzed Accessions: 0				
17	17	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22C, T25-	
Psathyrella/71729 Analyzed Accessions: 0				
15	MZ996976 KT188613 MK226170 KR155036 KP012810 PP852482 MN688087 MK226164 MW551150 MN017262 MK226166 MW947484 KR155010 MZ996971 MK226168	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, C26G	
2	MN688086 MN688096	Forward XXXX	Forward Missing	0.34 XXXX limited coverage (excluded)
		Probe XXXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, C26G, +22C	
Entoloma cetratum/566072 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
13	OL338338 OL338103 PQ653217 PQ652525 KC898450 PV659586 PQ652479 MK169354 PQ653392 PQ652765 PQ652590 OL338132 MZ317361	Forward XXX Probe XXX	Forward Missing T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, C26G	0.34 XXX limited coverage (excluded)
Fuscoporia/175855 Analyzed Accessions: 0				
5	OR771950 OR771948 OR771952 OR771947 OR771945	Forward XXX Probe XXX	Forward Missing T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, C26A	0.34 XXX limited coverage (excluded)
7	ON496501 ON496526 ON496536 ON496487 ON496529 ON496488 ON496509	Forward XXX Probe XXX	Forward Missing T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, C26G	0.34 XXX limited coverage (excluded)
Cortinarius anomalovelatus/1469658 Analyzed Accessions: 0				
12	FJ039651 FJ717604 FJ039653 FJ157008 NR_131809 FJ717605 FJ039699 FJ039652 FJ157006 FJ157011 FJ157012 PX307060	Forward XXX Probe XXX	Forward Missing T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22T, C26T	0.34 XXX limited coverage (excluded)
Entoloma ortonii/2891369 Analyzed Accessions: 0				
12	PQ639191 OL338209 PQ639056 PQ652448 PQ639198 OL338098 OL338136 OL338069 OL338019 OL338141 OL338222 OL338071	Forward XXX Probe XXX	Forward Missing T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, C26G	0.34 XXX limited coverage (excluded)
Tricholoma equestre/76322 Analyzed Accessions: 0				
11	OM506542 OM506540 HM590873 MW597284 MG597430 HM590875	Forward XXX Probe XXX	Forward Missing T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24A, C26G	0.34 XXX limited coverage (excluded)

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
	HM590870 HM590874 MW597235 HM590871 HM590872			
Gymnopus dryophilus/206318 Analyzed Accessions: 0				
1	LC149609	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, C26G	
9	MH589964 MT745588 MT408920 MG920317 MT408923 MH589961 PQ652879 PV388809 OP375147	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T25G, C26A, A27C	
Inocybe flocculosa/225431 Analyzed Accessions: 0				
7	PP982795 OM964556 OK057165 PQ653193 PQ653241 OP821367 AY380375	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, C26G	
2	KT800366 HQ604516	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, T24A, C26T	
1	PQ652639	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, C26G	
Pluteus chrysophaeus/1962868 Analyzed Accessions: 0				
10	OR242693 KX216313 MH010875 MH010881 KX216316 LT838192 MH010879 MH010873 MH010880 KX216314	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T21C, C26G	
Tricholoma/40144 Analyzed Accessions: 0				
8	LC335842 LC335841 LC335840 LC260648 LC335837 LC335838 LC335836 LC335839	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24A, C26G	
1	PV455430	Forward XXX	Forward Missing	0.34 XXX

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22T, C26G	limited coverage (excluded)
Fuscoporia torulosa/167372 Analyzed Accessions: 0				
9	PV203594 PV211417 EF068135 EF068137 AB811863 EF068139 EF068136 MG231517 EF068138	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, C26G	
Tricholoma cf. frondosae/2919555 Analyzed Accessions: 0				
9	LC684252 LC684276 LC684275 LC684271 LC684274 LC684270 LC684255 LC684273 LC684272	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24A, C26G	
Tricholoma flavovirens/80606 Analyzed Accessions: 0				
8	AF458452 LC684250 LC684251 LC684248 JF899574 LC684249 AF458449 AB036895	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24A, C26G	
Aureonarius callisteus/165391 Analyzed Accessions: 0				
8	FJ157137 FJ157014 KU236695 FJ039666 AY669594 KU236697 PQ653120 FJ157015	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-	
Calonarius aureofulvus/304751 Analyzed Accessions: 0				
8	GU363496 FJ157131 FJ717512 FJ039644 EU057061 EU057051 FJ039643 AY669571	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, C26T	
Gymnopus/93827 Analyzed Accessions: 0				
7	OQ597063 PV791548 OQ597062 MW408704 MZ781539 MH589969 MN930611	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T25G, C26A, A27C	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
Tricholoma auratum/419585 Analyzed Accessions: 0				
1	HM590867	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24A, C26G	
6	AB289662 AB289659 AB289661 AB289663 AB289660 LC684266	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, T24A, C26G	
Fuscoporia marquesiana/2204184 Analyzed Accessions: 0				
2	OR771939 OR771962	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-	
3	OR771941 OR771940 OR771942	Forward XXX	Forward Missing	3.22 XX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-	
		Reverse ✓✓✓	C1-	
1	OR771953	Forward XXX	Forward Missing	3.22 XX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-	
		Reverse ✓✓✓	C1-, C2-	
1	OR771966	Forward XXX	Forward Missing	3.22 XX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-	
		Reverse ✓✓✓	T9G	
Inocybe bellidiana/2881236 Analyzed Accessions: 0				
7	MW845863 MW845861 MW845860 PQ639214 MW845864 NR_174854 MW845862	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, C26G, +19G	
Microstroma/5032 Analyzed Accessions: 0				
1	OK053689	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-	
5	OK053673 OK053599 OK053578 QN866278 QN865297	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-	
Inocybe stellatospora/165594 Analyzed Accessions: 0				
6	AY038328 OP821373 OP821370 OP821369	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, C26T, A27G	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
	PQ652771 OP821371			
Entoloma hirtipes/181160 Analyzed Accessions: 0				
6	OL338150 OL338142 OL338140 OL338221 OL338160 OL338224	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
Pluteus chrysophlebius/181528 Analyzed Accessions: 0				
6	OR994065 KX216318 KX216315 KX216317 KX216319 PX094071	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T21C, C26G	
Calonarius pseudoglaucopus/304752 Analyzed Accessions: 0				
6	EU056950 PQ652571 JF273633 PQ652573 PQ653292 AY669573	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, C26T	
Inocybe tigrina/1169566 Analyzed Accessions: 0				
6	MW845933 MW924020 NR_174869 MW924019 OQ678147 OQ678100	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, C26G	
Armillaria/47424 Analyzed Accessions: 0				
5	KY389167 KY389158 KY389168 KY389160 KY389181	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T25G, C26A, A27C	
Pseudomicrostroma phylloplanum/105771 Analyzed Accessions: 0				
4	OQ066808 OQ066734 OQ066735 OQ066807	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-	
1	NR_165979	Forward XXX	Forward Missing	3.22 XX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-	
		Reverse ✓✓✓	C1-, C2-	
Hygrophorus cossus/234945 Analyzed Accessions: 0				
5	PQ653382 PQ652716 AY242852 PQ653204 AY548964	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, T21C, A22C, A27T	
Peniophorella odontiformis/391082 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
4	OQ996170 OQ996167 OQ996169 OQ996171	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-	
1	OQ996168	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-	
Tricholoma frondosae/511179 Analyzed Accessions: 0				
5	LC684245 LC684246 PQ069040 LC684247 PQ653143	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24A, C26G	
Marasmius brunneospermus/700780 Analyzed Accessions: 0				
5	FJ904966 FJ904968 FJ904967 FJ904969 FJ904965	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, C26G	
Peniophorella rude/1611331 Analyzed Accessions: 0				
1	KJ668497	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-	
1	MN062106	Forward XXX	Forward Missing	3.22 XX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-	
		Reverse ✓✓✓	C1-	
3	MN062109 MN062110 MN062107	Forward XXX	Forward Missing	3.22 XX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-	
		Reverse ✓✓✓	C1-, C2-	
Aureonarius infucatus/1895592 Analyzed Accessions: 0				
5	KU236706 KU236703 KU236700 KU236704 KU236702	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-	
Entoloma incanosquamulosum/2488080 Analyzed Accessions: 0				
5	PV388660 PV388654 OL338320 MK169374 PQ652788	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, C26G	
Callistosporium luteo-olivaceum/2545906 Analyzed Accessions: 0				
5	MF161184 OK626227 OK626228 OK626226 OL764904	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T20A, A22T, T24C, C26G	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
Inocybe mycenoides/2864439 Analyzed Accessions: 0				
<u>5</u>	OK057204 OK057156 MZ667617 OK057151 PQ069036	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, C26G	
Inocybe sabuligena/3118401 Analyzed Accessions: 0				
<u>5</u>	PP262650 PP262647 NR_198721 PP262648 PP262646	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, T23A, C26G	
Calonarius odorifer/36062 Analyzed Accessions: 0				
<u>4</u>	EU057031 PQ652548 EU057052 AY174817	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, C26T	
Lepiota/37314 Analyzed Accessions: 0				
<u>1</u>	MT451923	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C13A, C16A, T19A, C26G	
<u>2</u>	PP594592 PP594630	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-	
<u>1</u>	KP012774	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, C26G	
Amanita/41955 Analyzed Accessions: 0				
<u>1</u>	EU071961	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A22G, T24C	
<u>1</u>	KJ786698	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, A22T, C26G	
<u>2</u>	KT724064 KT724065	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, T23-, T24-, T25-	
Inocybe lacera/165587 Analyzed Accessions: 0				
<u>2</u>	HQ604445 HQ604446	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, C26G	
<u>1</u>	HQ604178	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, C26G	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	HQ604089	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, C26G	
mycorrhizal basidiomycete of <i>Epipogium roseum</i> /270613 Analyzed Accessions: 0				
4	AB176572 AB176575 AB176574 AB176579	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, C26G	
Phellinus punctatus/271459 Analyzed Accessions: 0				
4	EF429242 EF429243 EF429241 EF429245	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, C26G	
Wickerhamomyces/599737 Analyzed Accessions: 0				
4	PQ736443 PQ736444 PQ736447 PQ736442	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, T23-, T24-, T25-	
Inocybe pseudodestructa/674016 Analyzed Accessions: 0				
4	OK057132 OK057134 PQ652813 MT437069	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, T24A	
Inocybe cf. flocculosa DL-2015/1738947 Analyzed Accessions: 0				
4	KT800267 KT800349 KT800198 KT800213	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, T24A, C26T	
Naganishia/1851509 Analyzed Accessions: 0				
4	MK942555 MK942573 MK942574 MK942556	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, T23C	
Inocybe paleoveneta/2086435 Analyzed Accessions: 0				
4	ON946183 PV487632 PV487631 OL583808	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, C26G	
Marasmius chrysocephalus/2697436 Analyzed Accessions: 0				
4	MN930524 MN930584 MN930587 MN930583	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, T23-, C26G	
Inocybe galactica/2874931 Analyzed Accessions: 0				
4	NR_176173 OK057178 OK057196 OK057179	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, C26G	
Entoloma aromaticellum/2891363 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
4	OL338398 OL338402 OL338387 OL338459	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, C26G	
Inocybe cygnea/2949339 Analyzed Accessions: 0				
4	NR_184515 ON003447 ON003449 ON003448	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A17C, T21-, C26G	
Inocybe helobia/2970792 Analyzed Accessions: 0				
4	AM882815 OP164024 PQ658735 QQ133561	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, C26G	
Inocybe undatolacera/3232286 Analyzed Accessions: 0				
4	PP301785 PQ639057 PQ652793 PP301786	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, C26G	
Ogataea minuta/36026 Analyzed Accessions: 0				
3	LC580395 KY106579 LC580396	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T19A, A22T, T24A	
Armillaria borealis/47425 Analyzed Accessions: 0				
3	MW418558 MW418537 MW418548	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T25G, C26A, A27C	
Clitocybe/50966 Analyzed Accessions: 0				
3	KJ680983 PX273847 KJ680982	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24G, T25A	
Callistosporium/71868 Analyzed Accessions: 0				
2	PV248403 PV248402	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T25C, +26TT	
1	OP985109	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T24A, C26G	
Inocybe geophylla/71954 Analyzed Accessions: 0				
2	PX273821 MW845949	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, T21C, C26G	
1	HQ604285	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C14A, C16A, T19A, T21C, C26G	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
Cortinarius anomalus/86079 Analyzed Accessions: 0				
3	MZ145077 MZ145076 MZ145075	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A18G, A22T	
Gymnopilus/86085 Analyzed Accessions: 0				
2	GQ159919 FJ039688	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, C26G	
1	PV742653	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T25G, C26A, A27C	
Cortinarius anserinus/222021 Analyzed Accessions: 0				
3	AY174806 AY174807 AY174805	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22-	
Hygrophorus quercetorum/234948 Analyzed Accessions: 0				
3	AY463489 AY463488 AY463487	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, T21C, A22C, A27T	
Calonarius suaveolens/304753 Analyzed Accessions: 0				
3	DQ083816 AY669574 EU056991	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, C26T	
Thaxterogaster cretax/304787 Analyzed Accessions: 0				
3	NR_121304 EU660953 AY669622	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22-	
Cortinarius lacteus/309020 Analyzed Accessions: 0				
3	EU660951 AY669642 EU660952	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C16A, C26A, A27T	
Inocybe splendens/331641 Analyzed Accessions: 0				
3	FN550911 MG367267 OR100705	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, C10G, C11A, C13A, A22-, T23-, T24-, C26G	
Calonarius praetermissus/548933 Analyzed Accessions: 0				
3	EU684534 KJ421146 EU684535	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, C26T	
Tricholoma subsejunctum/596481 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
<u>3</u>	OQ076921 OQ076924 PP387631	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24A, C26G, +18C	
Calonarius xanthodryophilus/930113 Analyzed Accessions: 0				
<u>3</u>	PV749283 GQ159771 NR_119918	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, C26T	
Thaxterogaster variegatus/937735 Analyzed Accessions: 0				
<u>3</u>	KJ421003 FJ039663 PQ652453	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22-	
Amanita crenulata/1004481 Analyzed Accessions: 0				
<u>3</u>	OQ612499 MF161244 MK204468	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22G, T24C, C26G	
Cortinarius citrinofulvescens/1032571 Analyzed Accessions: 0				
<u>3</u>	PQ652699 PV388800 PQ652962	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-	
Inocybe picrosma/1110399 Analyzed Accessions: 0				
<u>3</u>	OM439611 NR_121489 HQ201364	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-	
Panellus luxfilamentus/1524334 Analyzed Accessions: 0				
<u>3</u>	OM057666 OP985113 KJ206988	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22C, C26A	
Basidiomycota sp./1715256 Analyzed Accessions: 0				
<u>2</u>	PP840277 PP840293	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, C26G	
<u>1</u>	PP840267	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24G, T25A	
Agaricales sp./1756126 Analyzed Accessions: 0				
<u>1</u>	MH811171	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-	
<u>1</u>	MH811666	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, C26G	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	MH811061	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, C26T	
Pseudolaccaria pachyphylla/1766229 Analyzed Accessions: 0				
3	MK400231 PV388679 MK392074	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T19C, A22T, T23C, C26G	
Gymnopus foetidus/1928556 Analyzed Accessions: 0				
2	KY026731 KY026730	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, C26G	
1	PV683411	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T25G, C26A, A27C	
Calonarius mariekristinae/2558181 Analyzed Accessions: 0				
3	MK659910 MK659908 MK659909	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, C26T	
Hortiboletus engelii/2832327 Analyzed Accessions: 0				
3	PV030525 OR567836 PV094816	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
Inocybe somae/2874937 Analyzed Accessions: 0				
3	OK057157 OK057148 NR_176166	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, C26G	
Entoloma assiduum/2891379 Analyzed Accessions: 0				
3	OR755887 OL338089 OL338157	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, C26G	
Inocybe caesaraugustae/2892407 Analyzed Accessions: 0				
3	PX289273 OL352083 NR_176184	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, C26G	
Inocybe donata/3119109 Analyzed Accessions: 0				
3	PP274021 PP274019 NR_198728	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, T25G, C26A, A27C	
Ogataea pini/4923 Analyzed Accessions: 0				
2	HQ413282 HQ413279	Forward XXX	Forward Missing	0.34 XXX

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T19A, A22T, T24A	limited coverage (excluded)
Marasmius/34448 Analyzed Accessions: 0				
<u>2</u>	MZ027762 KP133212	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, C26G	
Armillaria gallica/47427 Analyzed Accessions: 0				
<u>2</u>	PV750360 MW418538	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T25G, C26A, A27C	
Gerronema/50968 Analyzed Accessions: 0				
<u>2</u>	MT537044 MT537041	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22G, C26G	
Chrysomphalina grossula/50976 Analyzed Accessions: 0				
<u>2</u>	PQ652544 MT636945	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, T23-, C26G	
Cyberlindnera amylophila/53489 Analyzed Accessions: 0				
<u>2</u>	KY107350 MK394132	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22C, T25-	
Cyberlindnera japonica/53492 Analyzed Accessions: 0				
<u>2</u>	KY107369 LC076106	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T20A, A22T, T25A	
Macrocybe titans/56473 Analyzed Accessions: 0				
<u>2</u>	MZ519068 LR992036	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, C26G	
Armillaria sinapina/64372 Analyzed Accessions: 0				
<u>2</u>	OR039429 MW871593	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T25G, C26A, A27C	
Amanita melleiceps/78362 Analyzed Accessions: 0				
<u>2</u>	ON131769 AB015688	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A22G, C26G	
Rhodotorula paludigena/86838 Analyzed Accessions: 0				
<u>1</u>	ON261353	Reverse XXX	Reverse Missing	0.00 XXX

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
				limited coverage (excluded)
<u>1</u>	OP077324	Probe XXX	Probe Missing	0.34 XXX
		Reverse XXX	Reverse Missing	limited coverage (excluded)
Naganishia liquefaciens/104408 Analyzed Accessions: 0				
<u>2</u>	MT303972 MT303956	Probe XXX	Probe Missing	0.34 XXX
		Reverse XXX	Reverse Missing	limited coverage (excluded)
Neohypochnicium geogenium/139119 Analyzed Accessions: 0				
<u>2</u>	DQ677509 PQ653225	Forward XXX	Forward Missing	0.34 XXX
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, C26G	limited coverage (excluded)
Ischnoderma benzoinum/139424 Analyzed Accessions: 0				
<u>2</u>	MG735352 MF782788	Forward XXX	Forward Missing	0.34 XXX
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, C26A	limited coverage (excluded)
Mycena pura/153505 Analyzed Accessions: 0				
<u>2</u>	MZ823603 MZ130507	Forward XXX	Forward Missing	0.34 XXX
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, C26G, +23A	limited coverage (excluded)
Panellus pusillus/160031 Analyzed Accessions: 0				
<u>1</u>	OQ680217	Forward XXX	Forward Missing	0.34 XXX
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22C, C26A	limited coverage (excluded)
<u>1</u>	PP525847	Forward XXX	Forward Missing	0.34 XXX
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T19C, A22C, C26A	limited coverage (excluded)
Favolaschia calocera/181918 Analyzed Accessions: 0				
<u>2</u>	OP985987 KX447791	Forward XXX	Forward Missing	0.34 XXX
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, C26G	limited coverage (excluded)
Mycetinis scorodonius/182058 Analyzed Accessions: 0				
<u>2</u>	PV388814 PV248389	Forward XXX	Forward Missing	0.34 XXX
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T25G, C26A, A27C	limited coverage (excluded)
Cortinarius bolaris/183875 Analyzed Accessions: 0				
<u>2</u>	AY669596 KY964797	Forward XXX	Forward Missing	0.34 XXX
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22T	limited coverage (excluded)
Cortinarius decipiens/190316 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
2	PX139223 PX220874	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24-, T25-	
Inocybe nitidiuscula/217426 Analyzed Accessions: 0				
1	HQ604090	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, C26G	
1	HQ604214	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T19A, T21G, T24G, +26G	
Laetisaria/228945 Analyzed Accessions: 0				
2	OK053440 OK053453	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T25A, A27T	
Peniophorella/391237 Analyzed Accessions: 0				
1	MW507051	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-	
1	MW507055	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, T23-	
Candida frijolesensis/434044 Analyzed Accessions: 0				
1	EF658667	Probe XXX	G2C, C3G, G6-, T7A, G9A, C16T, T19-, T20-, A22C, T25G	0.34 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
1	NR_136989	Probe XXX	G2C, C3G, G6-, T7A, G9A, C16T, T19-, T20-, A22C, T25C	0.34 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
Ogataea/461281 Analyzed Accessions: 0				
2	MW764715 OP602957	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T19A, A22T, T24A	
Calonarius elotoides/477510 Analyzed Accessions: 0				
2	NR_137120 EU056948	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, C26T	
Candidozyma auris/498019 Analyzed Accessions: 0				
1	OR791935	Reverse XXX	Reverse Missing	0.00 XXX limited coverage (excluded)

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	OR791934	Probe XXX	Probe Missing	0.34 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
Inocybe aff. antillana BZ4124/512473 Analyzed Accessions: 0				
2	EU555441 PQ844550	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, C26G	
Tricholoma leucophyllum/519974 Analyzed Accessions: 0				
2	PP375511 PP375510	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24A, C26G	
Cortinarius huronensis/545998 Analyzed Accessions: 0				
1	EU821663	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-	
1	OQ549975	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22-, T23-	
Cortinarius colus/560664 Analyzed Accessions: 0				
2	HM240522 FJ157129	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24C, +24A	
Inocybe moelleri/674014 Analyzed Accessions: 0				
2	OR419907 FN550913	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, A22G, T23-, T24-, T25-, C26-	
Calonarius lavandulochlorus/983295 Analyzed Accessions: 0				
2	ON322986 HQ843177	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, C26T	
Rhizoctonia/1322061 Analyzed Accessions: 0				
2	KC176298 KM488558	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T25A, A27T	
Auricularia nigricans/1331584 Analyzed Accessions: 0				
2	JX065172 KM396803	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T19C, C26T	
Cercopomyces crocodilinus/1334636 Analyzed Accessions: 0				
2	JX409899 NR_175602	Forward XXX	Forward Missing	0.34 XXX

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C16A, T19A, C26G	limited coverage (excluded)
Osmoporus protractus/1349983 Analyzed Accessions: 0				
<u>2</u>	OP980890 JQ518280	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-	
Psathyrella gordonii/1365038 Analyzed Accessions: 0				
<u>2</u>	PQ849528 KC992924	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, C26G	
Psathyrella sulcatotuberculosa/1506512 Analyzed Accessions: 0				
<u>2</u>	OL518973 MW375696	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, C26G	
Butyriboletus/1524449 Analyzed Accessions: 0				
<u>2</u>	KM388727 KM388725	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, A27G	
Cortinarius angustisporus/1573779 Analyzed Accessions: 0				
<u>2</u>	KP165561 NR_131884	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, T24A	
Pseudomicrostroma glucosiphilum/1684307 Analyzed Accessions: 0				
<u>2</u>	MZ098659 OM337566	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-	
Fonsecazyma/1909748 Analyzed Accessions: 0				
<u>2</u>	PQ803299 PQ803290	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, T23C	
Entolomataceae sp./1935319 Analyzed Accessions: 0				
<u>1</u>	MH812941	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, C26G	
<u>1</u>	MH810651	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
Collybiopsis obscuroides/2007092 Analyzed Accessions: 0				
<u>2</u>	KX958398 KX958399	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, A22T, C26G	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
Inocybe ionochlora/2033263 Analyzed Accessions: 0				
<u>2</u>	OP164054 OP164061	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, C26G	
Cortinarius quercoconicus/2044529 Analyzed Accessions: 0				
<u>2</u>	PQ652876 MT935365	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, T24A	
Inocybe derbschii/2291540 Analyzed Accessions: 0				
<u>2</u>	MW845868 PV487616	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, C26G	
Candolleomyces aberdarensis/2316362 Analyzed Accessions: 0				
<u>2</u>	MH880928 MK421517	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, C26G	
Pseudolaccaria fellea/2585278 Analyzed Accessions: 0				
<u>2</u>	PV440447 PV750474	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T19C, A22T, T23C, C26G	
Marasmius tricystidiatus/2733409 Analyzed Accessions: 0				
<u>2</u>	MT441866 NR_182345	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22-, C26G	
Candolleomyces rogueianus/2799848 Analyzed Accessions: 0				
<u>2</u>	MW412407 NR_173216	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, C26G	
Peniophorella aspersa/2805651 Analyzed Accessions: 0				
<u>2</u>	NR_172775 MN062097	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, T23-	
Inocybe xantholeuca/2822544 Analyzed Accessions: 0				
<u>2</u>	MW845940 NR_173446	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, T21C, C26G	
Candolleomyces eurysporus/2828524 Analyzed Accessions: 0				
<u>2</u>	NR_172427 MT651560	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, C26G	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
Entoloma leptopus/2891367 Analyzed Accessions: 0				
<u>2</u>	OL338108 OL338043	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
Entoloma ventricosum/2891560 Analyzed Accessions: 0				
<u>2</u>	PQ653170 PX273841	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
Dendrosporomyces prolifer/2923081 Analyzed Accessions: 0				
<u>2</u>	OM906797 OM906796	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C	
Gymnopus strigosipes/2925462 Analyzed Accessions: 0				
<u>2</u>	NR_182585 OM970867	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T25G, C26A, A27C	
Tricholoma cf. viridilutescens/2949213 Analyzed Accessions: 0				
<u>2</u>	OQ832780 OQ832779	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, C26T, +24A	
Inocybe soliana/2949346 Analyzed Accessions: 0				
<u>2</u>	ON003425 NR_184496	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, C14A, T20A, C26G, +22A	
Inocybe sepiana/2970770 Analyzed Accessions: 0				
<u>2</u>	OP164056 OP164100	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, +26TG	
Inocybe telluris/3023422 Analyzed Accessions: 0				
<u>2</u>	OQ324783 NR_189410	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, T20A, C26G	
Meyerozyma guilliermondii/4929 Analyzed Accessions: 0				
<u>1</u>	OK090876	Probe XXX	Probe Missing	0.34 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
Histoplasma capsulatum/5037 Analyzed Accessions: 0				
<u>1</u>	OR447425	Probe XXX	G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, T23-, T24-, T25-, C26-, A27-	0.34 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
Penicillium/5073 Analyzed Accessions: 0				
1	AM176725	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T19C, A22C, T25C, A27-	0.34 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
Penicillium chrysogenum/5076 Analyzed Accessions: 0				
1	FJ345355	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1C, C3G, G6A, T7C, G9T, A17G, A18-, T20A, +23C	
Ganoderma/5314 Analyzed Accessions: 0				
1	KR155124	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, T24G, T25A	
Coprinus/5345 Analyzed Accessions: 0				
1	KR155053	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, C26G	
Boletus/5369 Analyzed Accessions: 0				
1	KY548850	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-	
Colletotrichum acutatum/27357 Analyzed Accessions: 0				
1	MG561705	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, C26G	
Ogataea henricii/36024 Analyzed Accessions: 0				
1	DQ414540	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T19A, A22T, T24A	
Omphalina/39675 Analyzed Accessions: 0				
1	PX273843	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, C26-, A27-	
Gloeophyllum sepiarium/40444 Analyzed Accessions: 0				
1	AB733139	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-	
Mycena/41247 Analyzed Accessions: 0				
1	PV750438	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, C26G	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
Amanita muscaria/41956 Analyzed Accessions: 0				
1	EU071889	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22G, C26G, +23C	
Cortinarius miniatopus/47326 Analyzed Accessions: 0				
1	MG597353	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24C, +24A	
Thaxterogaster vibratilis/47332 Analyzed Accessions: 0				
1	OK239906	Forward XXX	Forward Missing	3.22 XX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, C26T	
		Reverse ✓✓✓	+7C	
Aureonarius limonius/47422 Analyzed Accessions: 0				
1	FJ039667	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-	
Armillaria cepistipes/47426 Analyzed Accessions: 0				
1	MW418553	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T25G, C26A, A27C	
Tricholoma intermedium/54539 Analyzed Accessions: 0				
1	OQ987878	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24A, C26G	
Microsporium canis/63405 Analyzed Accessions: 0				
1	PV706791	Reverse XXX	Reverse Missing	0.00 XXX limited coverage (excluded)
Stropharia/68745 Analyzed Accessions: 0				
1	OL678095	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24G, T25A	
Clavulina/68756 Analyzed Accessions: 0				
1	OM337549	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, T23C	
Conocybe/71669 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	PQ531150	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, T23-, T24-, T25-, C26-	
Coprinosia atramentaria/71694 Analyzed Accessions: 0				
1	MT451919	Forward XXX	Forward Missing	3.22 XX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C13A, C16A, T19A, C26G	
		Reverse ✓✓✓	C1-, C2-	
Tricholoma sejunctum/76327 Analyzed Accessions: 0				
1	HM240544	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24G, C26G, +18C	
Chamonixia/80632 Analyzed Accessions: 0				
1	MK307840	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C26A	
Stropharia coronilla/83508 Analyzed Accessions: 0				
1	MT908250	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24G, T25A	
Moesziomyces aphidis/84754 Analyzed Accessions: 0				
1	PQ677007	Forward XXX	G1-, C2-, C3-, T4-, G5-, T6-, T7-, T14G, C15A, A16C	0.22 XXX limited coverage (excluded)
		Probe XXX	A5G, G6C, T7A, C26G	
		Reverse XXX	Reverse Missing	
Clavaria/104242 Analyzed Accessions: 0				
1	PQ579499	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
Crepidotus/104351 Analyzed Accessions: 0				
1	PV682888	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, C26G	
Thanatephorus cucumeris/107832 Analyzed Accessions: 0				
1	KJ489354	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, C14A, C26-	
Tricholoma ustale/113601 Analyzed Accessions: 0				
1	MT622855	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24G, T25A	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
Favolaschia/139071 Analyzed Accessions: 0				
1	OP985989	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, C26G	
Pseudotomentella nigra/148297 Analyzed Accessions: 0				
1	MK290718	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, T23-, C26G	
Cortinarius meleagris/160473 Analyzed Accessions: 0				
1	OR434403	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22-, T23-	
Auricularia delicata/160860 Analyzed Accessions: 0				
1	JX065159	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-	
Thaxterogaster/165407 Analyzed Accessions: 0				
1	OR035547	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22-	
Inocybe lanuginosa/165588 Analyzed Accessions: 0				
1	HQ604525	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, C26T, A27G	
Inocybe sindonia/166043 Analyzed Accessions: 0				
1	OK057131	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, T23-, C26G	
Calonarius corrosus/171475 Analyzed Accessions: 0				
1	EU056947	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, C26T	
Cortinarius corrugatus/171476 Analyzed Accessions: 0				
1	EU056949	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, C26T	
Calonarius flavaurora/171478 Analyzed Accessions: 0				
1	EU056946	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, C26T	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
Cortinarius laniger/171481 Analyzed Accessions: 0				
1	MT935187	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24-, T25-	
Callistosporium xanthophyllum/171504 Analyzed Accessions: 0				
1	JF907781	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T20A, A22T, T24C, C26G	
Collybia odora/181985 Analyzed Accessions: 0				
1	OL455051	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24G, T25A	
Cortinarius alnetorum/190313 Analyzed Accessions: 0				
1	FJ039534	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, T24A	
Micropsalliota/201197 Analyzed Accessions: 0				
1	PQ456422	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, C26G	
Rhodocollybia butyracea/206335 Analyzed Accessions: 0				
1	PV259563	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T25G, C26A, A27C	
Thaxterogaster myxoclaricolor/213137 Analyzed Accessions: 0				
1	AF539733	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22-	
Pseudosperma rimosum/217425 Analyzed Accessions: 0				
1	HQ604207	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, C26G	
Inocybe teraturgus/226591 Analyzed Accessions: 0				
1	MH310765	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, C26G	
Laetisaria arvalis/228946 Analyzed Accessions: 0				
1	ON965011	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T25A, A27T	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
Cortinarius croceus/239117 Analyzed Accessions: 0				
1	KY287714	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22-, T23-	
Inocybe fuscodisca/240309 Analyzed Accessions: 0				
1	PX307095	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, C26G	
Inocybe pusio/240321 Analyzed Accessions: 0				
1	HQ604521	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, T20A, A22C, C26G	
Onnia orientalis/271455 Analyzed Accessions: 0				
1	AY558606	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, T20C, C26T	
[Candida] sp. 36M/273132 Analyzed Accessions: 0				
1	AY589575	Probe XXX	C3G, G6-, T7C, G8A, G9A, C10A, A12T, C16A, T21-, A22-, T23C, A27-	0.34 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
Ogataea falcaomoraisii/281045 Analyzed Accessions: 0				
1	KY108672	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T19A, A22T, T24A	
Mycetinis copelandii/297704 Analyzed Accessions: 0				
1	MK234198	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T25G, C26A, A27C	
Rhodocollybia asema/298344 Analyzed Accessions: 0				
1	PQ653023	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T25G, C26A, A27C	
saccharomycete sp. jbra529/301054 Analyzed Accessions: 0				
1	AY796118	Probe XXX	G2T, C3A, T19C, A22-, C26T	0.34 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
Cortinarius aprinus/304809 Analyzed Accessions: 0				
1	AJ889942	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24A	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
Cortinarius psammocephalus/304815 Analyzed Accessions: 0				
1	AY669672	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, T24A	
Cortinarius cf. saniosus HL04-506O/342227 Analyzed Accessions: 0				
1	DQ102682	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, T24A	
[Candida] sp. BCMU BX02/413944 Analyzed Accessions: 0				
1	AB285025	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22C, T25-	
Candolleomyces leucotephrus/415551 Analyzed Accessions: 0				
1	KC992885	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, C26G	
Nolanea strictior/445260 Analyzed Accessions: 0				
1	PV248527	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
Fomitiporia langloisii/447965 Analyzed Accessions: 0				
1	EF437947	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, C26G	
Rhizoctonia solani/456999 Analyzed Accessions: 0				
1	KJ170365	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, C14A, C26-	
Inocybe griseoscabrosa/467846 Analyzed Accessions: 0				
1	FN550928	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, C26G	
Inocybe fuscidula/467874 Analyzed Accessions: 0				
1	HQ604301	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, C26G	
Inocybe phaeodisca/467877 Analyzed Accessions: 0				
1	MW845950	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, A18T, A22-, T23-, C26G	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
Inocybe hirtella/467885 Analyzed Accessions: 0				
1	AM882934	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, C26G	
Inocybe cf. flocculosa EL10605/467933 Analyzed Accessions: 0				
1	AM882992	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, C26G	
Calonarius elotus/477511 Analyzed Accessions: 0				
1	EU056951	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, C26T	
Cortinarius brunneifolius/498358 Analyzed Accessions: 0				
1	OL602057	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24-, T25-	
Cortinarius leiocastaneus/498359 Analyzed Accessions: 0				
1	PX316310	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24-, T25-	
Inocybe luteifolia/502799 Analyzed Accessions: 0				
1	OP879234	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, A22C, C26G	
Inocybe whitei/507466 Analyzed Accessions: 0				
1	FN550915	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C14A, C16A, T19A, T21C, C26G	
Daedaleopsis rubescens/528340 Analyzed Accessions: 0				
1	EU661889	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, C26G	
Cortinarius illibatus/545999 Analyzed Accessions: 0				
1	EU821685	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22T	
Cortinarius cf. azureus PK3036/560639 Analyzed Accessions: 0				
1	FJ039650	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22T, C26T	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
Cortinarius cf. anomalus F16129/561903 Analyzed Accessions: 0				
1	FJ157010	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22T, C26T	
Cercospora sophorae/578111 Analyzed Accessions: 0				
1	AB470876	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, C26G	
Cyberlindnera/604195 Analyzed Accessions: 0				
1	KY010353	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, C26T	
Cortinarius fasciatus/653840 Analyzed Accessions: 0				
1	GQ159913	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, T24A	
Inocybe argenteolutea/673979 Analyzed Accessions: 0				
1	FN550889	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, C26G	
Inocybe cf. griseolilacina EL12006/673990 Analyzed Accessions: 0				
1	FN550929	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A18T, A22T, C26G, +22CT	
Inocybe cf. hirtelloides EL9506/673993 Analyzed Accessions: 0				
1	FN550903	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, C26G	
Inocybe squarrosa/674023 Analyzed Accessions: 0				
1	FN550924	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, C26G	
Entoloma hebes/684104 Analyzed Accessions: 0				
1	OL338327	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
Aphanobasidium pseudotsugae/693291 Analyzed Accessions: 0				
1	KP814259	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T19G, T25-	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
Marasmius fusicystidiosus/707263 Analyzed Accessions: 0				
1	FJ917624	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, C26G	
Psathyrella maculata/752688 Analyzed Accessions: 0				
1	KC992935	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22-, C26G	
Inocybe abjecta/935512 Analyzed Accessions: 0				
1	HQ604302	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, C26G	
Inocybe cf. hirtella UBC F19257/935521 Analyzed Accessions: 0				
1	HQ604179	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, C26G	
Inocybe cf. sambucina UBC F19091/935537 Analyzed Accessions: 0				
1	HQ604287	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, C26G	
Inocybe cf. sambucina UBC F19157/935538 Analyzed Accessions: 0				
1	HQ604288	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, C26G	
fungal sp. BMP3018/986089 Analyzed Accessions: 0				
1	HQ833000	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, C26G, +22C	
Inocybe semifulva/993620 Analyzed Accessions: 0				
1	PV487638	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, T23-, C26G	
Inocybe grammopodia/1032817 Analyzed Accessions: 0				
1	OK057203	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, C26G	
Pluteus galeroideis/1033748 Analyzed Accessions: 0				
1	MH010877	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T21C, C26G	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
Phellodon ellisianus/1086382 Analyzed Accessions: 0				
1	JN135204	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, A22C, C26G	
Steccherinum aridum/1130391 Analyzed Accessions: 0				
1	JN710583	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24A, A27T	
Pleurotus giganteus/1156456 Analyzed Accessions: 0				
1	OQ916940	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-	
Inocybe polytrichi-norvegici/1169576 Analyzed Accessions: 0				
1	PV074833	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, C26G	
Agaricaceae sp. 347 A/1198530 Analyzed Accessions: 0				
1	JQ312188	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, C26G	
Agaricaceae sp. 377/1198531 Analyzed Accessions: 0				
1	JQ312189	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, C26G	
Agaricaceae sp. 325/1198532 Analyzed Accessions: 0				
1	JQ312190	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, C26G	
Phlegmacium vernicifer/1199181 Analyzed Accessions: 0				
1	MG019352	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, T19A, A22T	
Entoloma kristiansenii/1217149 Analyzed Accessions: 0				
1	PQ652815	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
Entoloma minutum/1217151 Analyzed Accessions: 0				
1	OL338210	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, C26G	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
Entoloma nitens/1217152 Analyzed Accessions: 0				
1	OL338321	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
Entoloma olivaceohebes/1217153 Analyzed Accessions: 0				
1	OL338158	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
Entoloma fuscohebes/1218271 Analyzed Accessions: 0				
1	OL338339	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, C26G	
Microstromatales sp. CHAM-8/1221400 Analyzed Accessions: 0				
1	JX290153	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-	
[Candida] sp. TH1/1221642 Analyzed Accessions: 0				
1	JX129175	Probe XXX	T4-, G9C, A18G, T19C, T21-, A22-, A27-, +7CG	0.34 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
[Candida] sp. TH4/1221643 Analyzed Accessions: 0				
1	JX129176	Probe XXX	T1G, G6T, T7G, A12C, C16-, A18T	0.34 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
fungal sp. MS578/1264013 Analyzed Accessions: 0				
1	JQ919949	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, C26G	
Basidiomycota sp. 28g/1264978 Analyzed Accessions: 0				
1	JX243956	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, T23C	
Candolleomyces badhyzensis/1365029 Analyzed Accessions: 0				
1	KC992883	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, C26G	
Candolleomyces trinitatensis/1365065 Analyzed Accessions: 0				
1	KC992882	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, C26G	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
Entoloma brunneosericeum/1365595 Analyzed Accessions: 0				
1	ON041428	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, C26G	
[Candida] sp. D141_4/1412056 Analyzed Accessions: 0				
1	KC349927	Probe XXX	Probe Missing	0.34 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
fungal sp. XXR-C/1429447 Analyzed Accessions: 0				
1	KF572128	Forward XXX	+8C, +11G, +14T, +17C	0.22 XXX limited coverage (excluded)
		Probe XXX	G6A, G8A, C10A, C11G, T23G, +5T, +13ACTACA, +22AT	
		Reverse XXX	Reverse Missing	
Phlegmacium mediterraneense/1440159 Analyzed Accessions: 0				
1	PQ587305	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22-, T23-	
Entoloma graphitipes/1502248 Analyzed Accessions: 0				
1	PQ527616	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, C26G	
Cortinarius coracis/1526703 Analyzed Accessions: 0				
1	MW862310	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24-, T25-	
Cortinarius anfractoides/1538812 Analyzed Accessions: 0				
1	NR_132060	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T19A, A22T	
Cortinarius persoonianus/1538907 Analyzed Accessions: 0				
1	KJ421118	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22T	
Auricularia villosula/1579976 Analyzed Accessions: 0				
1	OP345202	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-	
Tricholomataceae sp. 8PBL11/1585262 Analyzed Accessions: 0				
1	LC014889	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, C26G	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
Volvanarius chlorophanus/1586121 Analyzed Accessions: 0				
1	KJ421096	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22T, +23A	
Phlegmacium xantho-ochraceum/1586131 Analyzed Accessions: 0				
1	MK965073	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22-, T23-	
Cortinarius uliginobtusus/1619579 Analyzed Accessions: 0				
1	NR_175037	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22-, T23-	
Cortinarius tillamookensis/1619605 Analyzed Accessions: 0				
1	PQ653144	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22-, T23-	
Calonarius anetholens/1658746 Analyzed Accessions: 0				
1	NR_153048	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, C26T	
Amanita subparvipantherina/1660236 Analyzed Accessions: 0				
1	MG030644	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22G, T24C, C26G	
Cyanoboletus/1708618 Analyzed Accessions: 0				
1	KC152076	Forward XXX	Forward Missing	3.22 XX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, A27G	
		Reverse ✓✓✓	C1-, C2-	
Entoloma exile/1710133 Analyzed Accessions: 0				
1	PV388690	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-	
Tricholoma viridilutescens/1779462 Analyzed Accessions: 0				
1	OM417577	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, C26T, +24A	
Inocybe tiliae/1881966 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	OP164070	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, +26TG	
Vishniacozyma/1891946 Analyzed Accessions: 0				
1	PQ536259	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-	
Cortinarius birenensis/1930068 Analyzed Accessions: 0				
1	NR_153081	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24-, T25-	
Inocybe fuscithurnata/1978405 Analyzed Accessions: 0				
1	PP346048	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, C26G	
Pseudomicrostoma/1981958 Analyzed Accessions: 0				
1	OK053499	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, T23-	
Inocybaceae sp./2011741 Analyzed Accessions: 0				
1	MN096203	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, C26G	
Cortinarius subionophyllus/2020524 Analyzed Accessions: 0				
1	MT935497	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24-, T25-	
Cortinarius subminiatus/2044531 Analyzed Accessions: 0				
1	MT935502	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24C, +24A	
Inocybe muricellatoides/2053744 Analyzed Accessions: 0				
1	KY402220	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, C26G	
Cyberlindnera tropicalis/2057510 Analyzed Accessions: 0				
1	KT377032	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, C26T	
Cortinarius mycenarum/2136233 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	NR_156300	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22-, T23-	
Fulvoderma australe/2282104 Analyzed Accessions: 0				
1	MW742567	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, A18C, T25-, C26-	
Thaxterogaster emollitus/2493877 Analyzed Accessions: 0				
1	PP695561	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, C26T	
Lentinus brumalis/2498619 Analyzed Accessions: 0				
1	JX009175	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, T23-	
Cortinarius friesianus/2681340 Analyzed Accessions: 0				
1	PX220850	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24-, T25-	
Cortinarius pseudotorvus/2691840 Analyzed Accessions: 0				
1	NR_173297	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24-, T25-	
Inocybe neorufula/2712091 Analyzed Accessions: 0				
1	MT101876	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, C26G	
Tricholoma cf. equestre/2724809 Analyzed Accessions: 0				
1	MT303144	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24A, C26G	
Zhuliangomyces/2738246 Analyzed Accessions: 0				
1	MT424868	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T25G, C26-	
Coprinellus aff. xanthothrix/2740687 Analyzed Accessions: 0				
1	MT537064	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, C26G	
Fomitiporia aff. australiensis/2740694 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	MT537079	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, T21C, C26G	
Cortinarius distortus/2764271 Analyzed Accessions: 0				
1	PQ639037	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24-, T25-	
Candolleomyces/2791032 Analyzed Accessions: 0				
1	PV751186	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, C26G	
Cortinarius rufulus/2795720 Analyzed Accessions: 0				
1	MT271791	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22-, T23-	
Tricholoma atrofibrillosum/2807643 Analyzed Accessions: 0				
1	MW597242	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24G, C26G, +18C	
Cortinarius modestus/2812699 Analyzed Accessions: 0				
1	MZ580446	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22-, T23-	
Inocybe patibilis/2822546 Analyzed Accessions: 0				
1	MW845951	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, C26G	
Inocybe tigrinella/2822547 Analyzed Accessions: 0				
1	MW845953	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, C26G	
Psathyrella subcacao/2843179 Analyzed Accessions: 0				
1	PQ527795	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, C26G	
Entoloma belouvense/2861840 Analyzed Accessions: 0				
1	MZ611628	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
Inocybe ovilla/2874935 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	OK057191	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, T23-, C26G	
Entoloma cuneatum/2891365 Analyzed Accessions: 0				
1	OL338199	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
Entoloma transitionisporum/2891377 Analyzed Accessions: 0				
1	NR_177610	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
Amanitaceae sp. OMS-016/2905774 Analyzed Accessions: 0				
1	LC667046	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
Inocybe cf. pusio/2919401 Analyzed Accessions: 0				
1	OL352057	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, C26G	
Inocybe aff. lacera/2928573 Analyzed Accessions: 0				
1	PQ658724	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, C26G	
Gymnopus longisterigmaticus/2935534 Analyzed Accessions: 0				
1	OM030282	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T25G, C26A, A27C	
Inocybe tenuiorparietalis/2952547 Analyzed Accessions: 0				
1	ON754369	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, +26TG	
Calonarius pulcher/2955213 Analyzed Accessions: 0				
1	PX094299	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, T21C, C26T	
Tricholoma yatesii/2994527 Analyzed Accessions: 0				
1	PV650144	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24A, C26G	
Infundibulicybe bispora/3003539 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	OP983927	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T25G, C26A, A27C	
Inocybe amygdaliolens/3025631 Analyzed Accessions: 0				
1	PQ099327	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, C26G	
Hymenochaetales sp. IV-2023a/3043945 Analyzed Accessions: 0				
1	QQ913425	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T23C, A27T	
Infundibulicybe cf. gibba/3064479 Analyzed Accessions: 0				
1	OR419909	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T25G, C26A, A27C	
Inocybe maremmiana/3088919 Analyzed Accessions: 0				
1	OR765951	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, C26G	
Candolleomyces singeri/3095788 Analyzed Accessions: 0				
1	PQ613312	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T23C, C26G	
Paragymnopus perforans/3103146 Analyzed Accessions: 0				
1	PQ653032	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T25G, C26A, A27C	
Inocybe naekkii/3119111 Analyzed Accessions: 0				
1	PP274025	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, A17-, A18-, T19-, T20-, T21-, A22-, C26G	
Cortinarius urbis-veteris/3122135 Analyzed Accessions: 0				
1	PP357035	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24A	
Gymnopus brunneostipitatus/3138791 Analyzed Accessions: 0				
1	PP657586	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T25G, C26A, A27C	
Gymnopus biyangensis/3383057 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	PQ651936	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T25G, C26A, A27C	
Omphalina cf. rivulicola/3383740 Analyzed Accessions: 0				
1	PQ675388	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, C26-, A27-	
Cortinarius infractitenebricus/3384554 Analyzed Accessions: 0				
1	OR873435	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A22-	
Calonarius magnificus/3384560 Analyzed Accessions: 0				
1	HQ997909	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, A15-, C16-, C26T	
Hortiboletus hershenzoniae/3396226 Analyzed Accessions: 0				
1	PV030532	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
Inocybe favoris/3415765 Analyzed Accessions: 0				
1	PV487617	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T20G, A22G, C26G	
Inocybe schweyckertii/3415766 Analyzed Accessions: 0				
1	PV487636	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T19C, C26G, +21C	
Cortinarius subdeciapiens/3457080 Analyzed Accessions: 0				
1	PX139152	Forward XXX	Forward Missing	0.34 XXX limited coverage (excluded)
		Probe XXX	T1-, G2-, C3-, T4-, A5-, G6-, T7-, G8-, G9-, C10-, C11-, A12-, C13-, C14-, T24-, T25-	