

Search Details

Job Title/ Assay Name : glabrat(ex)	Amplicon Sequence: GCCTGTTTGAGCGTCATTTCTTCTCAAACACGTTGTGTTTGGTAGT GAGTGATACTCTCGTTTTTGAGTTAACTTGAAATTGTAGGCCATATC AGTATGTGGGACACGAGCGCAAGCTTCTCTATTAATCTGCTGCTCGT TTGCGCGAGCGGCGGGGGTTAATACTGTATTAGGTTTTACCAACTC GGTGTGATCTAGGGAGGGATAAGTGAGTGTTCTGTGCGTGCTGGG CAGACAGACGTCTTTAAGTTTGACCTCAAATCCGGTAGGGTTACCC GCTGAACTTAAGCATATCAATAAGCGGAGG
Effective Date : 2025-10-28	Forward: GCAGACAGACGTCTTTAAGTT
Database : core_nt	Reverse: CCTCCGCTTATTGATATGCTTA
Alignment algorithm : blastN	Probe: CCTCAAATCCGGTAGGGTTACCCGCTG
Exclusive Organism: Nakaseomyces glabratus (5478)	Inclusive Organism:
RID: G16K4FTV015	Total Result: 5000
App Version: 1.0	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
Candida albicans/5476 Analyzed Accessions: 12				
1	KT876205	Probe  	C10A	9.00    Detected
		Reverse   	C1-, C2-, +7C	
4	EF065107 EF065108 EF065104 EF065106	Probe  	C10A	9.00    Detected
		Reverse   	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-	
1	PP851215	Probe  	C10A	9.00    Detected
		Reverse   	C1-, C2-, T3-, C4-, A10T, +11A	
1	PP851213	Probe  	C10A	7.00  Not Detected
		Reverse 	T3C, G6C, A10T, G13-	
1	EF065103	Forward   	+251T	1.33    Not Detected
		Probe   	A5C, G12-, T13-, A14T	
		Reverse  	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-	
1	ON398749	Forward   	T12-, T20-, T21C	0.22    Not Detected
		Probe   	C10A, G11T, T18-, G27A	
		Reverse   	C1-, C2-, T3-, C4-, C5-, G6-, C19A	
1	JX912562	Forward   	G1-, G4A, A9G, C10A, G11A, T12C	1.67    Not Detected
		Probe  	C10A, G17A	
1	KJ706389	Forward   	G1-, G4A, A9G, C10G, G11A, T12C	5.67  Not Detected
		Probe  	C10A	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
		Reverse   	T3C	
1	KJ706866	Forward   	C2A, G4-, A9G, C10A, G11A, T12C	1.67    Not Detected
		Probe  	C10A, G17A	
1	PP851162	Probe  	A6C, C10A	4.67   limited coverage (excluded)
		Reverse 	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-, A14-	
1	PP851201	Probe   	C10A, C21-, C22-, C23-, G24-, C25-, T26-, G27-	0.34    limited coverage (excluded)
		Reverse   	Reverse Missing	
1	JN093144	Probe   	C9-, C10-, G11-, G12-, T13-, A14-, G15-, G16-, G17-, T18-, T19-, A20-, C21-, C22-, C23-, G24-, C25-, T26-, G27-	0.34    limited coverage (excluded)
		Reverse   	Reverse Missing	
1	OP042386	Forward   	T20-	0.22    limited coverage (excluded)
		Probe   	C10A, T13-, A14-, G15-, G16-, G17-, T18-, T19-, A20-, C21-, C22-, C23-, G24-, C25-, T26-, G27-	
		Reverse   	Reverse Missing	
1	PP851240	Forward   	T20-	0.22    limited coverage (excluded)
		Probe   	A7-, T8-, C9-, C10-, G11-, G12-, T13-, A14-, G15-, G16-, G17-, T18-, T19-, A20-, C21-, C22-, C23-, G24-, C25-, T26-, G27-	
		Reverse   	Reverse Missing	
1	PP851237	Forward   	T20-	0.22    limited coverage (excluded)
		Probe   	T3-, C4-, A5-, A6-, A7-, T8-, C9-, C10-, G11-, G12-, T13-, A14-, G15-, G16-, G17-, T18-, T19-, A20-, C21-, C22-, C23-, G24-, C25-, T26-, G27-	
		Reverse   	Reverse Missing	
1	MN419367	Forward   	A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-, A17-, A18-, G19-, T20-, T21-	0.22    limited coverage (excluded)
		Probe   	Probe Missing	
		Reverse   	Reverse Missing	
1	KY673190	Forward   	C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-, A17-, A18-, G19-, T20-, T21-	0.22    limited coverage (excluded)
		Probe   	Probe Missing	
		Reverse   	Reverse Missing	
8	KY673193 OP042382 KY673192 MN419375 LC639854 MN419379 MN419361 KY673191	Forward   	Forward Missing	0.22    limited coverage (excluded)
		Probe   	Probe Missing	
		Reverse   	Reverse Missing	
1	ON398728	Forward   	A9-, T12G	3.11   limited coverage (excluded)
		Probe   	C1-, C9G, C10G, T19A, G27-	
		Reverse   	Reverse Missing	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
<u>1</u>	MN419363	Forward XXX	C13T, +236A, +244TC, +251T	1.33 XXX limited coverage (excluded)
		Probe XXX	C10G, +4A, +9A, +14G, +20C	
		Reverse XXX	Reverse Missing	
Monosporozyma servazzii/27293 Analyzed Accessions: 22				
<u>10</u>	MW894863 OR964857 OR964858 OR964853 JQ808006 OR964859 JQ808010 OR964855 OR964854 OR964856	Forward XXX	A3G, G4A, A9T, C10G, G11T, +248T	0.34 XXX Not Detected
		Probe XXX	C10A, G17A	
<u>11</u>	MK044059 MK044060 MK044054 KX987246 MK044050 MK044044 MK044052 MK044053 MK044045 MK044065 MK044048	Forward XXX	A3G, G4A, A9T, C10G, G11T, +248T	0.34 XXX Not Detected
		Probe XXX	C10A, G17A, T18G	
<u>1</u>	KY103668	Forward XXX	A3G, G4A, A9T, C10G, G11T, +248T	3.22 XX Not Detected
		Probe XXX	C10A, G17A	
		Reverse ✓✓✓	C1-, C2-	
<u>2</u>	MH595423 MT645400	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Nakaseomyces nivariensis/418086 Analyzed Accessions: 19				
<u>15</u>	MW057254 OR690737 OR690735 NR_077073 MH545923 GU199442 MK394141 OK267572 GU199441 OR690738 MT731395 KP131740 PQ605688 KP131741 LR792666	Forward XXX	G1-, G4A, A9G, C10A, G11A, T12C	1.67 XXX Not Detected
		Probe XX	C10A, G17A	
<u>2</u>	FN298488 FM955316	Forward XXX	G1-, G4A, A9G, C10A, G11A, T12C	3.78 XX Not Detected
		Probe XX	C10A, G17A	
		Reverse ✓✓	A22T	
<u>1</u>	OK267677	Forward XXX	G1-, G4A, A9G, C10A, G11A, T12C	4.11 XX Not Detected
		Probe XX	C10A, G17A	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
		Reverse ✓✓✓	C1-, C2-, T3-	
1	PQ605689	Forward ✗✗✗	G1-, G4A, A9G, C10A, G11A, T12C	4.11 ✗✗ Not Detected
		Probe ✗✗	C10A, G17A	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-	
Lachancea thermotolerans/381046 Analyzed Accessions: 16				
14	MW384920 OQ137209 KX015903 KY207383 KF468194 KX015898 KY104005 PQ050730 OQ357445 KT253551 KY558362 KY207382 KY108259 KY108273	Forward ✗✗✗	G4A, A9T, C10A, G11T, T14A, T16A	1.67 ✗✗✗ Not Detected
		Probe ✗✗	C10A, G17A	
2	KY104004 KY104001	Forward ✗✗✗	G4A, A9T, C10A, G11T, T14A, T16A	4.11 ✗✗ Not Detected
		Probe ✗✗	C10A, G17A	
		Reverse ✓✓✓	C1-, C2-	
1	MN832581	Forward ✗✗✗	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	C10A, G17A	
1	GQ340447	Forward ✗✗✗	Forward Missing	0.22 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	Probe Missing	
		Reverse ✗✗✗	Reverse Missing	
Zygosaccharomyces rouxii/4956 Analyzed Accessions: 16				
1	AM943657	Forward ✗✗✗	G1C, C2T, G8A, C10A, G11T, T15C, T16A	0.67 ✗✗✗ Not Detected
		Probe ✗✗✗	C10A, G17A	
15	PV244003 MW979510 KM249341 ON710848 AM943656 PV244004 KY110271 MK044042 AM943655 OM905798 KY110285 KY106065 KY110254 KY110279 NR_165940	Forward ✗✗✗	G1-, C2-, A3-, G8A, C10A, G11T, T15C, T16A	0.67 ✗✗✗ Not Detected
		Probe ✗✗✗	C10A, G17A	
Maudiozyma exigua/34358 Analyzed Accessions: 15				
5	JQ808008 OP800236 MK394139	Forward ✗✗✗	G4A, A9T, C10A, G11C	0.34 ✗✗✗ Not Detected
		Probe ✗✗✗	C10A, G17A, T18A	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
	KY107912 KY107911			
1	KJ706449	Forward XXX	G4A, A9T, C10A, G11C	5.89 XX Not Detected
		Probe ✓✓✓	C10A, T18A	
		Reverse ✓✓✓	T3C, C7G, T8C	
1	KF728780	Forward XXX	G4A, A9T, C10A, G11C	5.89 XX Not Detected
		Probe ✓✓✓	C10A, G17A, T18A	
		Reverse ✓✓✓	+6G, T8C	
3	KY103635 KY103630 KY103634	Forward XXX	G4A, A9T, C10A, G11C, T20-	3.22 XX Not Detected
		Probe XXX	C10A, G17A, T18A	
		Reverse ✓✓✓	C1-, C2-	
1	D89894	Forward XXX	G4A, A9T, C10A, G11C	3.22 XX Not Detected
		Probe XXX	C10A, G17A, T18A	
		Reverse ✓✓✓	C1-, C2-, T3-	
1	FJ153216	Forward XXX	G4A, A9T, C10A, G11C	5.89 XX Not Detected
		Probe ✓✓✓	C10A, G17A, T18A	
		Reverse ✓✓✓	C7G, T8C	
2	KY103633 KY103632	Forward XXX	G4A, A9T, C10A, G11C	3.22 XX Not Detected
		Probe XXX	C10A, G17A, T18A	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-	
1	KJ808710	Forward XXX	G4A, A9T, C10A, G11C	5.89 XX Not Detected
		Probe ✓✓✓	C10A, G17A, T18A	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-	
30	30	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Kazachstania/71245 Analyzed Accessions: 14				
3	PQ314502 PQ309231 PQ309230	Forward X	G4A, A9T, C10A, G11C	3.33 XX Not Detected
		Probe XXX	C10A, G17A, T18A	
2	PQ309228 PQ309232	Forward X	A3G, G4A, G8A, A9T, C10A, G11C	3.33 XX Not Detected
		Probe XXX	C10A, G17A, T18A	
1	PP952090	Forward XXX	G4A, A9T, C10A, G11C	3.45 XX Not Detected
		Probe XXX	C10A, G17A, T18A	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-	
2	OM570227 MK322513	Forward XXX	A3G, G4A, A9T, C10G, G11T, +248T	0.34 XXX Not Detected
		Probe XXX	C10A, G17A, T18A	
3	JQ993382 PQ308368 OP740263	Forward XXX	A3G, G4A, A9T, C10G, G11T, +248T	0.34 XXX Not Detected
		Probe XXX	C10A, G17A	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	PP899636	Forward XXX	A3G, G4A, A9T, C10A, G11C	4.33
		Probe ✓✓✓	C10A, G17A, T18A	XX Not Detected
1	KX499461	Forward XXX	G4A, A9T, C10A, G11C	0.45
		Probe XXX	C10A, G11C, G17A, T18A	XXX
		Reverse XXX	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-	Not Detected
1	JQ993388	Forward XXX	A3G, G4A, A9T, C10G, G11T, +248T	3.22
		Probe XXX	C10A, G17A	XX
		Reverse ✓✓✓	T8C	Not Detected
18	18	Forward XXX	Forward Missing	0.22
		Probe XXX	Probe Missing	XXX
		Reverse XXX	Reverse Missing	limited coverage (excluded)
1	KX458459	Forward XXX	G4A, A9T, C10A, G11C	2.45
		Probe XXX	C10A, G17A, T18A	XXX
		Reverse X	Reverse Missing	limited coverage (excluded)
1	KT377034	Forward X	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-	3.33
		Probe XXX	C10A, G17A, T18A	XX limited coverage (excluded)
1	OR891688	Forward X	A3G, G4A, G8A, A9T, C10A, G11C	2.45
		Probe XXX	C10A, G17A, T18A	XXX
		Reverse XXX	Reverse Missing	limited coverage (excluded)
1	PQ531184	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T16A	2.00
		Probe XXX	C10A, G17A, T18A	XXX limited coverage (excluded)
Lachancea fermentati/4955 Analyzed Accessions: 13				
4	MK267548 MN907476 KP132361 KY103954	Forward XXX	A3G, G4A, A9T, C10A, G11T, T14A, T15C	1.67
		Probe XX	C10A, G17A	XXX Not Detected
4	KY103956 KY103955 KY103960 KY103957	Forward XXX	A3G, G4A, A9T, C10A, G11T, T14A, T15C	4.11
		Probe XX	C10A, G17A	XX
		Reverse ✓✓✓	C1-, C2-	Not Detected
1	GQ340438	Forward XXX	A3G, G4A, A9T, C10A, G11T, T14A, T15C	4.11
		Probe XX	C10A, G17A	XX
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-	Not Detected
3	OQ326712 OQ326710 OQ326709	Forward XXX	A3G, G4A, A9T, C10A, G11T, T14A, T15C	4.11
		Probe XX	C10A, G17A	XX
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-	Not Detected
1	OK050672	Forward XXX	A3G, G4A, A9T, C10A, G11T, T14A, T15C	4.11
		Probe XX	C10A, G17A, T26A	XX
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-	Not Detected

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
64	64	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
1	KY108222	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-	2.00 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Nakaseomyces bracarensis/273131 Analyzed Accessions: 11				
3	GU199438 MF084287 GU199439	Forward XXX	G4A, C10A, G11A, T12C	4.11 XX Not Detected
		Probe XX	C10A, G17A	
		Reverse ✓✓✓	T8C	
1	KP131680	Forward XXX	G4A, C10A, G11A, T12C	0.34 XXX Not Detected
		Probe XXX	C9-, G11A, G17A	
1	KP674715	Forward XXX	G4A, C10A, G11A, T12C	4.11 XX Not Detected
		Probe XX	C10A, G17A	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, A10T	
2	MH545924 MK394142	Forward XXX	G4A, C10A, G11A, T12C, T20-	0.34 XXX Not Detected
		Probe XXX	A5-, C10A, G17A	
1	KP674833	Forward XXX	G4A, C10A, G11A, T12C	4.11 XX Not Detected
		Probe XX	C10A, G17A	
		Reverse ✓✓✓	+6G G, T12-	
1	KP674832	Forward XXX	G4A, C10A, G11A, T12C	4.11 XX Not Detected
		Probe XX	C10A, G17A	
		Reverse ✓✓✓	C1-, C2-, C7G, A10T, T11A	
1	MF187327	Forward XXX	G4A, C10A, G11A, T12C	4.11 XX Not Detected
		Probe XX	C10A, G17A	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, A10T	
1	OK267736	Forward XXX	G4A, C10A, G11A, T12C	4.11 XX Not Detected
		Probe XX	C10A, G17A, +23AG	
		Reverse ✓✓✓	C7G, +7G, A10T	
16	16	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
[Candida] sp. (uncertain placement)/1853550 Analyzed Accessions: 7				
4	OM802450 OM760527 OM802447 OM802403	Forward XXX	G4A, G8A, C10G, G11A, T12C	1.67 XXX Not Detected
		Probe XX	C10A, G17A	
1	OM480677	Forward XXX	G4A, G8A, C10G, G11A, T12C	4.11 XX Not Detected
		Probe XX	C10A, G17A	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
		Reverse   	C1-	
2	OM760525 OM570256	Forward 	G1-, G4A, G8A, C10G, G11A, T12C	3.33  
		Probe   	C10A, G17A	Not Detected
1	MH973205	Probe  	C10A	4.00  
		Reverse   	Reverse Missing	limited coverage (excluded)
1	ON907682	Forward   	T20-	0.22   
		Probe   	C10A, C25-, T26-, G27-	limited coverage (excluded)
		Reverse   	Reverse Missing	
2	ON907760 ON907759	Probe   	C10-, G11-, G12-, T13-, A14-, G15-, G16-, G17-, T18-, T19-, A20-, C21-, C22-, C23-, G24-, C25-, T26-, G27-	0.34   
		Reverse   	Reverse Missing	limited coverage (excluded)
6	MH973266 OM632680 MW895082 MW895096 MW895094 MW895081	Forward   	Forward Missing	0.22   
		Probe   	Probe Missing	limited coverage (excluded)
		Reverse   	Reverse Missing	
1	MW895095	Forward 	A3G, G4A, G8A, A9T, C10A, G11C	4.22  
		Probe   	C10A, G17A, T18A	limited coverage (excluded)
		Reverse 	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-	
1	LC272846	Forward   	G4A, A9T, C10A, G11A, T15C, T16A	2.22   
		Probe   	C10A, G17A, T18C	limited coverage (excluded)
		Reverse 	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-, A14-	
1	OM089792	Forward  	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-	2.00   
		Probe   	C10A, G17A	limited coverage (excluded)
1	PP962245	Forward 	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-	3.33  
		Probe   	C10A, G15A, G17A	limited coverage (excluded)
14	PQ032847 MG367321 OR582606 OM570241 MW764713 PP962240 LC815032 PP962241 MN299228 OM480687 OQ411075 OM570240 OP740281 LC815028	Forward   	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67   
		Probe   	C10A, G17A	limited coverage (excluded)
Maudiozyma barnettii/61262 Analyzed Accessions: 5				
1	FN394006	Forward 	G4A, A9T, C10A, G11C	4.33  

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
		Probe XXX	C10A, G17A, T18A	Not Detected
		Reverse ✓✓✓	C1-, C2-	
1	D89897	Forward XX	G4A, A9T, C10A, G11C	4.33
		Probe XXX	C10A, G17A, T18A	XX
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-	Not Detected
2	MW477701 FN667857	Forward X	G4A, A9T, C10A, G11C	3.33
		Probe XXX	C10A, G17A, T18A	XX Not Detected
1	MW477702	Forward XX	G4A, A9T, C10A, G11C, T16-	2.00
		Probe XXX	C10A, G17A, T18A	XXX Not Detected
5	KY103623 NR_155227 KY103624 AY046173 OQ275007	Forward XXX	Forward Missing	0.22
		Probe XXX	Probe Missing	XXX
		Reverse XXX	Reverse Missing	limited coverage (excluded)
Oxyporus populinus/139378 Analyzed Accessions: 5				
5	OL741476 PQ639264 PQ652838 OR475675 MF319116	Forward XXX	G1-, G11T, T20C, +239C, +250C	0.34
		Probe XXX	C10A, G17A, T18C	XXX Not Detected
Lachancea dasiensis/1072105 Analyzed Accessions: 5				
2	KY103942 FJ153187	Forward XXX	G4A, A9T, C10A, G11T, T14A, T15C, T16A	1.67
		Probe XX	C10A, G17A	XXX Not Detected
2	MK044068 MK044071	Forward XXX	A3G, G4A, A9T, C10A, G11T, T14A, T15C	1.67
		Probe XX	C10A, G17A	XXX Not Detected
1	MN648701	Forward XXX	A3G, G4A, A9T, C10A, G11T, T14A, T15C	4.11
		Probe XX	C10A, G17A	XX
		Reverse ✓✓✓	C1-, C2-, T3-	Not Detected
Lachancea/300275 Analyzed Accessions: 4				
2	ON864492 ON864497	Forward XXX	G4A, A9T, C10A, G11T, T14A, T16A	1.67
		Probe XX	C10A, G17A	XXX Not Detected
1	ON040667	Forward XXX	G4A, A9T, C10A, G11T, T14A, T15C, T16A	1.67
		Probe XX	C10A, G17A	XXX Not Detected
1	MZ701687	Forward XXX	G4A, A9T, C10A, G11T, T14A, T15C, T16A	4.11
		Probe XX	C10A, G17A	XX
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-	Not Detected
5	GQ340437 KJ451620 FJ666085 MZ701642 GQ340445	Forward XXX	Forward Missing	0.22
		Probe XXX	Probe Missing	XXX
		Reverse XXX	Reverse Missing	limited coverage (excluded)
Maudiozyma turicensis/432100 Analyzed Accessions: 4				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
2	MK044026 MK044040	Forward XXX	G4A, A9T, C10A, G11C	0.34 XXX Not Detected
		Probe XXX	C10A, G17A, T18A	
1	KY103679	Forward XXX	G4A, A9T, C10A, G11C	3.22 XX Not Detected
		Probe XXX	C10A, G17A, T18A	
		Reverse ✓✓✓	C1-, C2-	
1	MN648695	Forward XXX	G4A, A9T, C10A, G11C	3.22 XX Not Detected
		Probe XXX	C10A, G17A, T18A	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-	
3	NR_138171 AY046171 FJ178185	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
1	MN648699	Forward XXX	G4A, A9T, C10A, G11C	3.22 XX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18A	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-	
Monosporozyma aerobia/271218 Analyzed Accessions: 4				
1	OM523880	Forward XX	A3G, G4A, A9T, C10A, G11C	4.33 XX Not Detected
		Probe XXX	C10A, G17A, T18A	
		Reverse ✓✓✓	+7C	
2	NR_077087 MN328377	Forward XXX	A3G, G4A, A9T, C10G, G11T, +248T	0.34 XXX Not Detected
		Probe XXX	C10A, G17A	
1	MN328368	Forward XXX	A3G, G4A, A9T, C10G, G11T, +248T	5.89 X Not Detected
		Probe ✓✓✓	C10A, G17A	
		Reverse ✓✓✓	C1-, C2-, T3-	
1	KY103617	Forward XXX	A3G, G4A, A9T, C10G, G11T, +245T	4.89 XX limited coverage (excluded)
		Probe ✓✓✓	C10A, G17A	
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-	
1	PP589198	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Nakaseomyces/374468 Analyzed Accessions: 4				
2	MF975532 MK322511	Forward XXX	G4A, G8A, C10G, G11A, T12C	1.67 XXX Not Detected
		Probe XX	C10A, G17A	
1	OR052126	Forward XXX	G4A, G8A, C10G, G11A, T12C	4.11 XX Not Detected
		Probe XX	C10A, G17A	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-	
1	OM480721	Forward XXX	G4A, G8A, C10G, G11A, T12C	0.34 XXX Not Detected
		Probe XXX	C10A, G17A, T18C	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	KU218497	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Monosporozyma yasuniensis/1468137 Analyzed Accessions: 4				
3	HG934854 HG934853 NR_160315	Forward XXX	A3G, G4A, A9T, C10G, G11T, +245T	3.22 XX Not Detected
		Probe XXX	C10A, G17A, T18A	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-	
1	MW895952	Forward XXX	A3G, G4A, A9T, C10G, G11T, +245T	3.22 XX Not Detected
		Probe XXX	C10A, G17A, T18A	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-	
1	LN650650	Forward XXX	A3G, G4A, A9T, C10G, G11T, +248T	3.22 XX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18A	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-	
Nakaseomyces delphensis/51657 Analyzed Accessions: 4				
3	AY198400 NR_167682 MK394143	Forward XXX	C2A, T12A, +236GCCA, +240GAAT, +244A	4.33 XX Not Detected
		Probe ✓✓✓	C10A, G17A	
1	KP068958	Forward XXX	C2A, T12A, +236GCCA, +240GAAT, +244A	5.89 X Not Detected
		Probe ✓✓✓	C10A, G17A	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-	
Zygosaccharomyces siamensis/1074906 Analyzed Accessions: 4				
4	PV363964 PV363919 PV363929 LC547999	Forward XXX	G1C, C2T, A3T, C10A, G11T, T15C, T16A	1.67 XXX Not Detected
		Probe XX	C10A, G17A	
Saccharomyces cerevisiae/4932 Analyzed Accessions: 3				
3	OQ711757 LC576582 MT734684	Forward XXX	A3G, G4A, G8A, A9T, C10G, G11T, T16A	0.34 XXX Not Detected
		Probe XXX	C10A, G17A, T18G	
2070	2070	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
32	32	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T16A	2.00 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18G	
Maudiozyma humilis/51915 Analyzed Accessions: 3				
1	MH453885	Forward XXX	G4A, A9T, C10A, G11C	0.34 XXX Not Detected
		Probe XXX	C10A, G17A, T18A	
1	MH453888	Forward XXX	A3T, A9T, C10A, G11C	0.34 XXX Not Detected
		Probe XXX	C10A, G17A, T18A	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	KX951497	Forward XXX	A3T, G4A, A9T, C10A, G11C	0.34 XXX Not Detected
		Probe XXX	C10A, G17A, T18A	
10	MH453886 MH453887 PV826901 MN913450 MH453890 AY493349 MH453882 KY106506 KX951498 KX951499	Forward X	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-	3.33 XX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18A	
1	KX951500	Forward X	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-	5.56 X limited coverage (excluded)
		Probe XXX	C10A, G17A, T18A	
		Reverse ✓✓✓	C1-	
1	MT645398	Forward X	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-	5.56 X limited coverage (excluded)
		Probe XXX	C10A, G17A, T18A	
		Reverse ✓✓✓	C1-, C2-	
2	MT374224 MT374223	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Maudiozyma bulderi/112263 Analyzed Accessions: 3				
1	JQ808009	Forward XXX	G4A, A9T, C10A, G11C	0.34 XXX Not Detected
		Probe XXX	C10A, G17A, T18A	
2	MT645401 MT645402	Forward XXX	G4A, A9T, C10A, G11C	3.22 XX Not Detected
		Probe XXX	C10A, G17A, T18A	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-	
1	OM996016	Forward XXX	G4A, A9T, C10A, G11C	2.22 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18A	
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-, A14-, T15-, A16-	
4	MW624470 KY103628 NR_138198 PP466950	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Jamesozyma lodderae/1089531 Analyzed Accessions: 3				
1	KY103654	Forward XXX	A3G, G4A, G8A, A9T, C10A, G11C	3.45 XX Not Detected
		Probe XXX	C10A, G17A, T18A	
		Reverse ✓✓✓	C1-, C2-	
1	KY103650	Forward XXX	A3G, G4A, G8A, A9T, C10A, G11C	3.45 XX Not Detected
		Probe XXX	C10A, G17A, T18A	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	KY103652	Forward XXX	A3G, G4A, G8A, A9T, C10A, G11C, T20-	3.45 XX Not Detected
		Probe XXX	C10A, G17A, T18A	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-	
4	KY103651 MH595324 NR_138196 MH595321	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Yueomyces sinensis/58632 Analyzed Accessions: 3				
2	NR_159012 KP866235	Forward XXX	A3G, G4A, G8A, A9T, C10A, G11T, T15C, T16A	1.67 XXX Not Detected
		Probe XX	C10A, T18C	
1	KY105247	Forward XXX	A3G, G4A, G8A, A9T, C10A, G11T, T15C, T16A	4.11 XX Not Detected
		Probe XX	C10A, T18C	
		Reverse ✓✓✓	C1-, C2-	
1	AB056132	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Monosporozyma unispora/27294 Analyzed Accessions: 3				
3	EU794732 FN394008 MK268124	Forward XXX	A3G, G4A, A9T, C10G, G11T, +248T	0.34 XXX Not Detected
		Probe XXX	C10A, G17A, T18G	
Hanseniaspora osmophila/56408 Analyzed Accessions: 3				
3	MK044002 FR774545 KJ880970	Forward XX	A3C, G4A, A9T, C10A, G11T, T14A, T16A	2.00 XXX Not Detected
		Probe XXX	C10A, G17A, T18G	
Zygosaccharomyces pseudorouxii (nom. inval.)/227369 Analyzed Accessions: 3				
1	AM279696	Forward XXX	G1C, C2T, A3T, C10A, G11T, T15C, T16A	1.67 XXX Not Detected
		Probe XX	C10A, G17A	
1	AM279464	Forward XXX	G1C, C2T, G8A, C10A, G11T, T15C, T16A	0.67 XXX Not Detected
		Probe XXX	C10A, G17A	
1	AM279465	Forward XXX	G1-, C2-, A3-, G8A, C10A, G11T, T15C, T16A	0.67 XXX Not Detected
		Probe XXX	C10A, G17A	
Maudiozyma pseudohumilis/640839 Analyzed Accessions: 3				
3	NR_137720 KY102343 FJ888526	Forward XXX	A3T, G4A, A9T, C10A, G11C	0.34 XXX Not Detected
		Probe XXX	C10A, G17A, T18A	
fungal sp./1709941 Analyzed Accessions: 2				
1	OL998449	Forward XX	A3G, G4A, A9T, C10A, G11C	4.33 XX Not Detected
		Probe XXX	C10A, G17A, T18A	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	MH812033	Forward XXX	G1-, C2-, A3-, G8A, C10A, G11T, T15C, T16A	0.67
		Probe XXX	C10A, G17A	XXX Not Detected
1	MK040496	Probe ✓✓	C10A	4.00
		Reverse XXX	Reverse Missing	XX limited coverage (excluded)
1	MT912769	Forward ✓✓✓	T12G	3.11
		Probe XXX	C10A, A14T	XX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
3	MT596299 OP537717 MK040529	Forward XXX	Forward Missing	0.22
		Probe XXX	Probe Missing	XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
3	MH811239 MH811904 MH812499	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	2.00
		Probe XX	A5G, C10A	XXX limited coverage (excluded)
Zygosaccharomyces bisporus/4957 Analyzed Accessions: 2				
1	DQ167569	Forward XXX	G4A, G8A, A9T, C10A, G11C, T15C, T20C	0.34
		Probe XXX	C10A, G17A	XXX Not Detected
1	KY106028	Forward XXX	G4A, G8A, A9T, C10A, G11C, T15C, T20-, T21C	3.22
		Probe XXX	C10A, G17A	XX Not Detected
		Reverse ✓✓✓	C1-, C2-	
13	PX058846 MK044037 NR_138188 KY106029 MK592835 PX058848 AY046192 PX058847 ON908978 PQ038114 MK044023 KY106030 KC881074	Forward XXX	Forward Missing	0.22
		Probe XXX	Probe Missing	XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
Zygosaccharomyces/4953 Analyzed Accessions: 2				
1	MN268785	Forward XXX	G1C, C2T, A3T, C10A, G11T, T15C, T16A	0.34
		Probe XXX	C10A, G17A, A20-	XXX Not Detected
1	HE984156	Forward XXX	G1C, C2T, G8A, C10A, G11T, T15C, T16A	0.67
		Probe XXX	C10A, G17A	XXX Not Detected
4	OR481161 KU350363 KU350462 OR481213	Forward XXX	Forward Missing	0.22
		Probe XXX	Probe Missing	XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
2	PV244009 PV244008	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67
				XXX

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
		Probe XXX	C10A, G17A	limited coverage (excluded)
Kazachstania kunashirensis/488016 Analyzed Accessions: 2				
1	NR_138158	Forward XXX	G4A, A9T, C10G, G11T, +235TGGCG	3.22 XX Not Detected
		Probe XXX	C10A, G17A, T18A	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-	
1	KY103649	Forward XXX	A5G, A7G, G8A, +235T, +242A, +244GTTC	1.33 XXX Not Detected
		Probe XXX	C10A, G17A, T18A	
		Reverse XX	C1-, C2-	
6	AY046163 OK052128 KM384184 OK051061 KM384185 OK051060	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Sungouiozyma intestinalis/935788 Analyzed Accessions: 2				
2	NR_155231 KY103647	Forward XXX	A3G, G4A, G8A, A9T, C10A, G11C	3.22 XX Not Detected
		Probe XXX	C10A, G15A, G17A, T18A	
		Reverse ✓✓✓	C1-, C2-	
4	MN194592 HQ263350 HQ260334 HQ999969	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Naumovozyma dairenensis/27289 Analyzed Accessions: 2				
1	D89893	Forward XXX	A3G, G4A, A9T, C10A, G11C	3.45 XX Not Detected
		Probe XXX	C10A, G17A, T18A	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-	
1	NR_165980	Forward XXX	A3G, G4A, A9T, C10A, G11C, T20-	3.45 XX Not Detected
		Probe XXX	C10A, G17A, T18A	
		Reverse ✓✓✓	C1-, C2-	
1	KY108664	Forward X	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-	3.33 XX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18A	
2	AJ229072 AY046181	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Jamesozyma spencerorum/1089533 Analyzed Accessions: 2				
1	D89898	Forward XX	A3G, G4C, G8A, A9T, C10A, G11C, +237A	4.33 XX Not Detected
		Probe XXX	C10A, G17A, T18A	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-	
1	KY103674	Forward XXX	A3G, G4A, G8T, T20-, +238A, +245A	3.22 XX Not Detected
		Probe XXX	C10A, G17A, T18A	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-	
<u>2</u>	NR_138197 KY103673	Forward ✗✗✗	Forward Missing	0.22 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	Probe Missing	
		Reverse ✗✗✗	Reverse Missing	
<u>1</u>	MH595421	Forward ✗	A3G, G4A, G8A, A9T, C10A, G11C	2.22 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	C10A, G17A, T18A	
		Reverse ✗✗✗	Reverse Missing	
Maudiozyma wufongensis/561021 Analyzed Accessions: 2				
<u>1</u>	KY103689	Forward ✗✗✗	G4A, G11T, +240T	4.33 ✗✗ Not Detected
		Probe ✓✓✓	C10A, G17A, T18A	
<u>1</u>	FJ153181	Forward ✗✗✗	G4A, A9T, C10A, G11C	4.33 ✗✗ Not Detected
		Probe ✓✓✓	C10A, G17A, T18A	
<u>1</u>	OW987344	Forward ✗✗✗	G4A, A9T, C10A, G11C	5.11 ✗ limited coverage (excluded)
		Probe ✓✓✓	C10A, G17A, T18A	
		Reverse ✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-	
<u>1</u>	HQ999975	Forward ✗✗✗	Forward Missing	0.22 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	Probe Missing	
		Reverse ✗✗✗	Reverse Missing	
Naumovozyma baii/1209964 Analyzed Accessions: 2				
<u>2</u>	NR_155462 JX087438	Forward ✗✗✗	A3G, G4A, G8A, A9T, C10A, G11C	0.67 ✗✗✗ Not Detected
		Probe ✗✗✗	C10A, G17A, T18A	
Lachancea mirantina/1230905 Analyzed Accessions: 2				
<u>2</u>	NR_155330 KY103990	Forward ✗✗✗	G4A, A9T, C10A, G11A, T16A	3.22 ✗✗ Not Detected
		Probe ✗✗✗	C10A, G17A, T18C	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-	
Candida parapsilosis/5480 Analyzed Accessions: 1				
<u>1</u>	OM523864	Forward ✗✗✗	G8A, C10T, G11A, T16C, G19A, T20A, +246C	1.33 ✗✗✗ Not Detected
		Probe ✗✗✗	T3A, C9A, C10G, T13A, A14G, +20C	
		Reverse ✗✗	C1-, C2-, A10C, T12C, G13T, G18T, C19T, A22G	
<u>1</u>	KX268323	Forward ✗✗✗	T20-	0.22 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	C10-, G11-, G12-, T13-, A14-, G15-, G16-, G17-, T18-, T19-, A20-, C21-, C22-, C23-, G24-, C25-, T26-, G27-	
		Reverse ✗✗✗	Reverse Missing	
<u>1</u>	MF767671	Forward ✗✗✗	Forward Missing	0.22 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	Probe Missing	
		Reverse ✗✗✗	Reverse Missing	
<u>1</u>	MF767639	Forward ✗✗✗	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-, G19A	4.00 ✗✗

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
		Probe  	C10A	limited coverage (excluded)
Saccharomyces paradoxus/27291 Analyzed Accessions: 1				
1	KP960515	Forward   	G4A, A9T, C10A, G11T, T14A, T16A	1.67
		Probe  	C10A, G17A	   Not Detected
281	281	Forward   	Forward Missing	0.22
		Probe   	Probe Missing	   limited coverage (excluded)
		Reverse   	Reverse Missing	
5	KY109448 KY109455 KY109443 FN667999 FN868260	Forward  	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T16A	2.00
		Probe   	C10A, G17A, T18G	   limited coverage (excluded)
Zygosaccharomyces mellis/42258 Analyzed Accessions: 1				
1	LT963551	Forward   	G1C, C2T, A3T, C10A, G11T, T15C, T16A	1.67
		Probe  	C10A, G17A	   Not Detected
9	PV244013 PV243998 PV243999 PV244002 PV244001 PV244015 PV244000 PV244012 PV244014	Forward   	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67
		Probe   	C10A, G17A	   limited coverage (excluded)
1	NR_197461	Forward   	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	3.45
		Probe   	C10A, G17A	  limited coverage (excluded)
		Reverse   	C1-, C2-	
Yueomyces/1723649 Analyzed Accessions: 1				
1	OM845233	Forward   	G4A, G8A, A9T, C10A, G11C, T15C, T16A	1.67
		Probe  	C10A, G17A	   Not Detected
10	PP589140 OP886715 OM095318 PP589139 PP589145 OP886717 OP886713 OP886716 MT009486 OP886714	Forward   	Forward Missing	0.22
		Probe   	Probe Missing	   limited coverage (excluded)
		Reverse   	Reverse Missing	
Wickerhamomyces/599737 Analyzed Accessions: 1				
1	PQ309225	Forward   	A3C, G4T, C6A, A7C, G8A, G11T, T14A, T16A, +242CTA	0.34
		Probe   	C10A, G17A, T18C	   Not Detected

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
2	OP292297 JF781478	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
4	MF737105 MF737104 MF737103 MF737102	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, T18C	
1	PQ309223	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T15A	2.00 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18C	
1	PQ533516	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Maudiozyma serrabonitensis/1392788 Analyzed Accessions: 1				
1	MW895939	Forward X	C2T, G4A, A9T, C10A, G11C	5.22 X Not Detected
		Probe XXX	C10A, G17A, T18A	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-	
1	NR_164524	Forward X	G4A, A9T, C10A, G11C	4.22 XX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18A	
		Reverse X	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-	
6	LT158247 LT158249 OK051505 OK051519 OK051517 OK051514	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Jamesozyma piceae/56483 Analyzed Accessions: 1				
1	FR716598	Forward XX	G4A, G8A, A9T, C10A, G11C	2.00 XXX Not Detected
		Probe XXX	C10A, G17A, T18A	
2	KY107939 KY107938	Forward X	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-	3.33 XX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18A	
4	HF558661 NR_155232 KY103660 AY046159	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Lachancea waltii/4914 Analyzed Accessions: 1				
1	OQ326708	Forward XXX	G4A, A9T, C10A, G11T, T14A, T15C, T16A	4.11 XX Not Detected
		Probe XX	C10A, G17A	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-	
1	PP589011	Forward XXX	G4A, A9T, C10A, G11T, T14A, T15C, T16A	4.11 XX limited coverage (excluded)
		Probe XX	C10A, G17A	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-, G13-	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
4	NR_138205 KY104011 KF057702 OK051390	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Lachancea cidri/29831 Analyzed Accessions: 1				
1	NR_165978	Forward XXX	A3G, G4A, A9T, C10A, G11T, T14A, T15C	4.11 XX Not Detected
		Probe XX	C10A, G17A	
		Reverse ✓✓✓	C1-, C2-	
5	KY103940 AY046205 PP906256 KY103939 KY103938	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Tetrapisipora/113604 Analyzed Accessions: 1				
1	MW895328	Forward XXX	A3G, G4A, A9T, C10G, G11T, +248T	0.34 XXX Not Detected
		Probe XXX	C10A, G17A, T18A	
5	MW895948 MW895023 MW895950 MW895947 MW895247	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Vanderwaltozyma verrucispora/907289 Analyzed Accessions: 1				
1	KY105796	Forward XXX	A3G, G4A, G8T, G11T, T15C, T16A, +238A	1.67 XXX Not Detected
		Probe XX	C10A, G17A	
5	OQ255923 NR_137559 HM461673 HM461674 EU515872	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Naumovozyma/278028 Analyzed Accessions: 1				
1	PQ309241	Forward XXX	A3G, G4A, A9T, C10A, G11C	3.45 XX Not Detected
		Probe XXX	C10A, G17A, T18A	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-	
3	PQ309239 PQ309240 MK713457	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Lachancea kluyveri/4934 Analyzed Accessions: 1				
1	OQ357476	Forward XXX	A3G, G4A, A9T, C10A, G11T, T14A, T16A	1.67 XXX Not Detected
		Probe XX	C10A, G17A	
2	KY108233 KY108232	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Oligophagozymba castelii/51914 Analyzed Accessions: 1				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	NR_154958	Forward XXX	G4A, G8A, T15C, T16A, T20-	3.22 XX Not Detected
		Probe XXX	C10A, G17A, T18C	
		Reverse ✓✓✓	C1-, C2-	
2	AY046196 OW983542	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Lachancea nothofagi/679534 Analyzed Accessions: 1				
1	GQ411403	Forward XXX	G4A, A9T, C10A, G11T, T14A, T15C, T16A	1.67 XXX Not Detected
		Probe XX	C10A, G17A	
1	KF826533	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Maudiozyma saulgeensis/1789683 Analyzed Accessions: 1				
1	NR_164574	Forward XX	G4A, A9T, C10A, G11C	2.00 XXX Not Detected
		Probe XXX	C10A, G17A, T18A	
1	LT158244	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Pinus wallichiana/3341 Analyzed Accessions: 1				
1	AY430074	Forward XXX	G1C, C2T, G8A, C10A, G11T, T15C, T16A	0.67 XXX Not Detected
		Probe XXX	C10A, G17A	
Tortispora caseinolytica/51930 Analyzed Accessions: 1				
1	KY109824	Forward X	G1-, G4A, A9G, C10A, G11A, T12C	3.33 XX Not Detected
		Probe XXX	C10A, G17A	
Hanseniaspora vineae/56409 Analyzed Accessions: 1				
1	MN371905	Forward XX	A3C, G4A, A9T, C10A, G11T, T14A, T16A	2.00 XXX Not Detected
		Probe XXX	C10A, G17A, T18G	
Pinus albicaulis/71622 Analyzed Accessions: 1				
1	AY430073	Forward XXX	G1C, C2T, G8A, C10A, G11T, T15C, T16A	0.67 XXX Not Detected
		Probe XXX	C10A, G17A	
Ramaria rubrievanescens/113070 Analyzed Accessions: 1				
1	JX310408	Forward XXX	G1-, C2-, A3-, G8A, C10A, C13G, T16G, A18C	0.34 XXX Not Detected
		Probe XXX	C10A, T18C	
Fomitiporia mediterranea/208960 Analyzed Accessions: 1				
1	OQ979759	Forward XXX	G8A, A9C, C10A, G11T, A17G, A18C	0.34 XXX Not Detected
		Probe XXX	C10A, G17A	
Kluyveromyces hubeiensis/237925 Analyzed Accessions: 1				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	MW709994	Forward XX	A3G, G4A, G8A, A9T, C10A, G11C	4.33 XX Not Detected
		Probe XXX	C10A, G17A, T18A	
		Reverse ✓✓✓	+7C, A10T	
Lachancea meyersii/300276 Analyzed Accessions: 1				
1	NR_077162	Forward XXX	G4A, C6T, A9T, C10A, G11T, T14A, T16A	1.67 XXX Not Detected
		Probe XX	C10A, G17A	
Trigonopsis californica/378053 Analyzed Accessions: 1				
1	NR_154533	Forward X	G1-, C2-, A3-, G4-, A5-, C10A, G11C, C13T	5.56 X Not Detected
		Probe XXX	C10A, T18A	
		Reverse ✓✓✓	C1-, C2-	
Monosporozyma solicola/1089532 Analyzed Accessions: 1				
1	AY582128	Forward XXX	A3G, G4A, A9T, C10G, G11T, +248T	4.33 XX Not Detected
		Probe ✓✓✓	C10A, G17A	
Maudiozyma rupicola/1198442 Analyzed Accessions: 1				
1	JQ861272	Forward X	G4A, A9T, C10A, G11C	3.33 XX Not Detected
		Probe XXX	C10A, G17A, T18A	
[Candida] sp. VVT-2012/1214789 Analyzed Accessions: 1				
1	JX406304	Forward XXX	C2-, A3-, G4-, A5-, G8C, G19-, T20-	0.22 XXX Not Detected
		Probe XXX	A5-, C10-, G11-, G12-, T18-, A20C	
		Reverse XXX	C1-, C2-, T8-, T9-, A16-, T21A	
Rigidoporus piceicola/1765029 Analyzed Accessions: 1				
1	PQ653016	Forward XXX	G1-, G11T, T20C, +239C, +250C	0.34 XXX Not Detected
		Probe XXX	C10A, G17A, T18C	
Fomitopsis flavimontis/3151108 Analyzed Accessions: 1				
1	PV749276	Forward XXX	G1-, C2-, A3-, G8-, A17G, G19C, +251C	0.34 XXX Not Detected
		Probe XXX	C10A, G17A	
[Candida] glabrosa/45554 Analyzed Accessions: 0				
1	PP544253	Forward XXX	T20-	0.22 XXX limited coverage (excluded)
		Probe XXX	C10A, G15C, G17-, T18-, C25-, T26-, G27-	
		Reverse XXX	Reverse Missing	
5	MK394113 ON261242 KY102112 MW959737 FM178351	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Candida/5475 Analyzed Accessions: 0				
1	LT799386	Probe XXX	C10A, T18-, T19-, A20-, C21-, C22-, C23-, G24-, C25-, T26-, G27-	0.34 XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
Candida africana/241526 Analyzed Accessions: 0				
1	MF769545	Forward XXX	T20-	0.22 XXX limited coverage (excluded)
		Probe XXX	C10A, G17-, T18-, T19-, A20-, C21-, C22-, C23-, G24-, C25-, T26-, G27-	
		Reverse XXX	Reverse Missing	
Debaryomyces hansenii/4959 Analyzed Accessions: 0				
129	129	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Saccharomyces cf. cerevisiae/paradoxus/1911677 Analyzed Accessions: 0				
95	95	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
8	KY109324 KY109344 KY109359 KY109341 KY109332 KY109329 KY109343 KY109346	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T16A	2.00 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18G	
Ogataea uvarum/2126558 Analyzed Accessions: 0				
98	98	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Cryptococcus neoformans/5207 Analyzed Accessions: 0				
91	91	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, A17T, A18T	0.34 XXX limited coverage (excluded)
		Probe XXX	C10A, T18C	
Saccharomyces/4930 Analyzed Accessions: 0				
1	HQ623547	Forward XXX	A3G, G4A, G8A, A9T, C10A, G11C, T15C, T16C	4.11 XX limited coverage (excluded)
		Probe XX	C10A, G17A	
		Reverse ✓✓✓	C1-, C2-, T3-, C4-, C5-, G6-, C7-, T8-, T9-, A10-, T11-, T12-	
69	69	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Saccharomyces bayanus/4931 Analyzed Accessions: 0				
52	52	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
5	KY109213 KY109195 KY109202	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T16A	2.00 XXX
		Probe XXX	C10A, G17A, T18G	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
	KY109198 KY109212			limited coverage (excluded)
Zygosaccharomyces bailii/4954 Analyzed Accessions: 0				
54	54	Forward XXX	Forward Missing	0.22
		Probe XXX	Probe Missing	XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
Debaryomyces/4958 Analyzed Accessions: 0				
49	49	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67
		Probe XXX	C10A, G17A	XXX limited coverage (excluded)
Hanseniaspora uvarum/29833 Analyzed Accessions: 0				
48	48	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67
		Probe XXX	C10A, G17A	XXX limited coverage (excluded)
Saccharomyces uvarum/230603 Analyzed Accessions: 0				
41	41	Forward XXX	Forward Missing	0.22
		Probe XXX	Probe Missing	XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
1	KY109468	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T16A	2.00
		Probe XXX	C10A, G17A, T18G	XXX limited coverage (excluded)
Saccharomyces kudriavzevii/114524 Analyzed Accessions: 0				
39	39	Forward XXX	Forward Missing	0.22
		Probe XXX	Probe Missing	XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
Cyberlindnera/604195 Analyzed Accessions: 0				
2	PQ309206 PQ309205	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, +247A	2.00
		Probe XXX	C10A, G17A	XXX limited coverage (excluded)
19	19	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-	2.00
		Probe XXX	C10A, G17A	XXX limited coverage (excluded)
16	16	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67
		Probe XXX	C10A, G17A	XXX limited coverage (excluded)
Saccharomyces pastorianus/27292 Analyzed Accessions: 0				
29	29	Forward XXX	Forward Missing	0.22
		Probe XXX	Probe Missing	XXX limited coverage (excluded)
		Reverse XXX	Reverse Missing	
Zygorulaspora florentina/48255 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
29	29	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Debaryomyces fabryi/58627 Analyzed Accessions: 0				
29	29	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Saccharomyces eubayanus/1080349 Analyzed Accessions: 0				
28	28	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
1	KF826534	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T16A	2.00 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18G	
[Candida] subhashii/561895 Analyzed Accessions: 0				
28	28	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Yamadazyma/766765 Analyzed Accessions: 0				
11	MK130967 LC076105 OP941484 OP941480 PQ309190 PP849692 PP849693 MK130964 MF631784 PV866810 OR064519	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-	2.00 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
17	17	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Hanseniaspora opuntiae/211096 Analyzed Accessions: 0				
26	26	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Scheffersomyces/766733 Analyzed Accessions: 0				
25	25	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] boidinii/5477 Analyzed Accessions: 0				
23	23	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, T16A	2.00 XXX

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
		Probe XXX	C10A, G17A	limited coverage (excluded)
[Candida] palmiophila/45574 Analyzed Accessions: 0				
23	23	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX
		Probe XXX	C10A, G17A	limited coverage (excluded)
Saccharomyces cf. bayanus/pastorianus/1911676 Analyzed Accessions: 0				
17	17	Forward XXX	Forward Missing	0.22 XXX
		Probe XXX	Probe Missing	limited coverage (excluded)
		Reverse XXX	Reverse Missing	
3	KY109222 KY109218 KY109224	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T16A	2.00 XXX
		Probe XXX	C10A, G17A, T18G	limited coverage (excluded)
Doratomyces/186351 Analyzed Accessions: 0				
19	19	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T15C	2.00 XXX
		Probe XXX	A5G, A6G, C10A	limited coverage (excluded)
Jamesozyma gamospora/308752 Analyzed Accessions: 0				
3	KY107922 KY107919 KY107920	Forward X	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-	3.33 XX
		Probe XXX	C10A, G17A	limited coverage (excluded)
1	KY103642	Forward X	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-	5.56 X
		Probe XXX	C10A, G17A	limited coverage (excluded)
		Reverse ✓✓✓	C1-, C2-	
14	ON306659 KY103643 KY363436 MH595441 KY103638 KY363445 PQ868069 PQ868067 NR_137688 PQ868068 KY103639 KY363444 KY103641 KY103640	Forward XXX	Forward Missing	0.22 XXX
		Probe XXX	Probe Missing	limited coverage (excluded)
		Reverse XXX	Reverse Missing	
Cyberlindnera fabianii/36022 Analyzed Accessions: 0				
17	17	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-	2.00 XXX
		Probe XXX	C10A, G17A	limited coverage (excluded)
Naumovozyma castellii/27288 Analyzed Accessions: 0				
1	KY108663	Forward X	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-	3.33 XX
		Probe XXX	C10A, G17A, T18A	limited coverage (excluded)

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
15	ON935715 KY104379 KY104380 KY104382 D89604 D89892 HQ026730 MW710017 KY104383 NR_144905 MH595422 D89602 AJ229062 MH595212 OP018314	Forward XXX Probe XXX Reverse XXX	Forward Missing Probe Missing Reverse Missing	0.22 XXX limited coverage (excluded)
Scheffersomyces shehatae/45589 Analyzed Accessions: 0				
16	16	Forward XXX Probe XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16- C10A, G17A	0.67 XXX limited coverage (excluded)
Zygosaccharomyces parabailii/1365886 Analyzed Accessions: 0				
16	16	Forward XXX Probe XXX Reverse XXX	Forward Missing Probe Missing Reverse Missing	0.22 XXX limited coverage (excluded)
Kurtzmaniella quercitrusa/2763128 Analyzed Accessions: 0				
16	16	Forward XXX Probe XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16- C10A, G17A	0.67 XXX limited coverage (excluded)
Saccharomycodes ludwigii/36035 Analyzed Accessions: 0				
14	MT937310 XR_006829194 ON763770 XR_006829210 KY109478 EU556322 XR_006829191 MT937312 PQ677536 XR_006829197 XR_006829200 FM199957 KY109476 MT937305	Forward XX Probe XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T15A C10A, G17A, T18A	2.00 XXX limited coverage (excluded)
Nakazawaea/460521 Analyzed Accessions: 0				
14	PQ411191 PP809586 PQ568186 PQ309155 MH844420 PQ568166 PQ568167 ON306382 PQ309154 PQ568201 PQ411190 OM480689	Forward XXX Probe XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16- C10A, G17A	0.67 XXX limited coverage (excluded)

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
	PQ568193 PQ568169			
Priceomyces carsonii/28549 Analyzed Accessions: 0				
12	PQ677527 PQ677524 PQ677526 PQ677528 PQ677525 PQ677533 PQ677535 PQ677530 PQ677534 PQ677531 PQ677529 PQ677532	Forward XXX Probe XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16- C10A, T18C	0.67 XXX limited coverage (excluded)
Amanita/41955 Analyzed Accessions: 0				
10	KY435392 OR296876 KY774702 OL414952 KU248100 KP662539 PV070461 OL555792 KX061528 OR296885	Forward XXX Probe XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, G19A C10A, G17A, T18C	0.34 XXX limited coverage (excluded)
2	ON817223 ON817224	Forward XXX Probe ✓✓	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-, G19C C10A	4.00 XX limited coverage (excluded)
Saccharomyces mikatae/114525 Analyzed Accessions: 0				
12	KY105198 NR 111354 KY105199 AJ229064 KY105200 AY046149 PP551941 KY105202 MH595374 KY105201 KY105203 PP551945	Forward XXX Probe XXX Reverse XXX	Forward Missing Probe Missing Reverse Missing	0.22 XXX limited coverage (excluded)
Chroogomphus rutilus/85976 Analyzed Accessions: 0				
11	PQ653344 ON692958 PQ653096 PQ641299 ON869350 PQ641304 PQ653039 MG457862 DQ367894 HM049562 HM049561	Forward XXX Probe XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, A18T C10A, G17A, T18C	0.34 XXX limited coverage (excluded)
Vanderwaltozyma/374469 Analyzed Accessions: 0				
10	LC191730 LC191744	Forward XXX	Forward Missing	0.22 XXX

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
	LC191731 FJ873456 OM095313 OM095311 OP099834 OM095315 OM095316 FJ873444	Probe XXX Reverse XXX	Probe Missing Reverse Missing	limited coverage (excluded)
Nakazawaea molendinolei/1538181 Analyzed Accessions: 0				
3	KJ706898 KJ706421 KJ706262	Forward XXX Probe ✓✓	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16- C10A	4.33 XX limited coverage (excluded)
7	PV092539 JN688668 OK051845 HE799678 JN688669 JN688670 NR_172220	Forward XXX Probe XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16- C10A, G17A	0.67 XXX limited coverage (excluded)
Saccharomyces cariocanus/114526 Analyzed Accessions: 0				
9	MH595281 MH595331 NR_144772 MH595418 KY104990 AJ229061 KY104988 KY104991 KY104989	Forward XXX Probe XXX Reverse XXX	Forward Missing Probe Missing Reverse Missing	0.22 XXX limited coverage (excluded)
Debaryomyces subglobosus/522693 Analyzed Accessions: 0				
8	MZ227346 JN851033 KP132014 FN675240 MK156303 KP132015 JN850992 KJ706916	Forward XXX Probe XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16- C10A, G17A	0.67 XXX limited coverage (excluded)
1	KJ706439	Forward XXX Probe ✓✓ Reverse ✓✓✓	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16- C10A T3C	5.89 X limited coverage (excluded)
Chroogomphus mediterraneus/1811971 Analyzed Accessions: 0				
9	PQ764697 MG457868 MG457867 PQ653340 PQ653257 PQ653457 MG457857 PQ653461 MG457839	Forward XXX Probe XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, A18T C10A, G17A, T18C	0.34 XXX limited coverage (excluded)
Grigorovia yakushimaensis/432099 Analyzed Accessions: 0				
8	NR_155237 KY103692	Forward XXX	Forward Missing	0.22 XXX

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
	KY103693 AB467304 KY103691 KY103690 AB097400 AB467303	Probe XXX Reverse XXX	Probe Missing Reverse Missing	limited coverage (excluded)
Zygorulasporea/1196389 Analyzed Accessions: 0				
8	MN721359 OR554094 OR554070 PQ314506 OR481445 PQ314510 MN721358 PQ314503	Forward XXX Probe XXX Reverse XXX	Forward Missing Probe Missing Reverse Missing	0.22 XXX limited coverage (excluded)
Kluyveromyces marxianus/4911 Analyzed Accessions: 0				
1	MK027128	Forward XXX Probe XXX Reverse XXX	Forward Missing Probe Missing Reverse Missing	0.22 XXX limited coverage (excluded)
6	KM921926 KM921929 KM921928 KM921927 KM921933 KM921930	Forward XXX Probe XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16- C10A, T18G	0.67 XXX limited coverage (excluded)
Hanseniaspora/29832 Analyzed Accessions: 0				
1	KU350325	Forward XXX Probe XXX Reverse XXX	Forward Missing Probe Missing Reverse Missing	0.22 XXX limited coverage (excluded)
6	MW057255 OP044349 KR185333 OP043928 MT988220 OP043932	Forward XXX Probe XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16- C10A, G17A	0.67 XXX limited coverage (excluded)
Vanderwaltozyma polyspora/36033 Analyzed Accessions: 0				
7	HM461668 AJ229076 HM461667 HM461672 HM461669 NR_111124 KY105794	Forward XXX Probe XXX Reverse XXX	Forward Missing Probe Missing Reverse Missing	0.22 XXX limited coverage (excluded)
Cephalotrichum stemonitis/186354 Analyzed Accessions: 0				
7	LC177641 MH854674 MF782693 FJ914696 FJ914658 FR799473 PV491314	Forward XX Probe XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T15C A5G, A6G, C10A	2.00 XXX limited coverage (excluded)
Torulaspora maleeae/200334 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
7	AB304160 KY105663 MT131815 NR_136949 AB304159 MT138578 AB304156	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Ogataea/461281 Analyzed Accessions: 0				
7	PQ149074 PQ032851 PQ149073 PQ380279 HE799687 PQ309161 ON303737	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Cephalotrichum asperulum/1658437 Analyzed Accessions: 0				
7	OR514928 PQ527657 PP947297 PQ109189 PP947299 PP947298 PP851738	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T15C	2.00 XXX limited coverage (excluded)
		Probe XXX	A5G, A6G, C10A	
Cora hawksworthiana/1924445 Analyzed Accessions: 0				
7	OR543616 OR543590 OR543545 OR543549 OR543546 OR543547 OR543592	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Nakazawaea ambrosiae/2518133 Analyzed Accessions: 0				
7	MK562509 MK508963 MK562507 MK562510 NR_165549 MK562506 MK562508	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Pichia/4919 Analyzed Accessions: 0				
1	MF662379	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
5	AB513333 AB513335 AB513336 AB513332 FJ153190	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Grigorovia transvaalensis/36916 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
6	OW983946 KY103678 AB011808 AY046156 Z80093 KX018479	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Cryptococcus deneoformans/40410 Analyzed Accessions: 0				
6	KT067745 AF219003 AF218975 AF444324 PV828365 PP994669	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, A17T, A18T	0.34 XXX limited coverage (excluded)
		Probe XXX	C10A, T18C	
Zygosaccharomyces lentus/85125 Analyzed Accessions: 0				
6	KY106033 KY106036 MH252540 KY106035 AY046194 NR_156001	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Cyberlindnera mycetangii/130814 Analyzed Accessions: 0				
1	KY106591	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-	2.00 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
5	NR_152475 OK052441 KY102221 OK052205 OK052442	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Cephalotrichum microsporum/186352 Analyzed Accessions: 0				
6	FJ914695 GU566278 MT447483 LC177639 OR514918 HQ607849	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T15C	2.00 XXX limited coverage (excluded)
		Probe XXX	A5G, A6G, C10A	
Xylaria enterogena/477508 Analyzed Accessions: 0				
6	PP662562 GU324736 KP133357 KP133368 KP133364 KP133365	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T16A, T20G	1.67 XXX limited coverage (excluded)
		Probe XX	A5G, C10A	
Cephalotrichum verrucisporum/644588 Analyzed Accessions: 0				
6	MF448345 MF002121 MF002122 FJ914678 LC177638 MF448346	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T15C	2.00 XXX limited coverage (excluded)
		Probe XXX	A5G, A6G, C10A	
Cryptococcus gattii/37769 Analyzed Accessions: 0				
5	EU402437 JN939461 KY102787	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, A17T, A18T	0.34 XXX

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
	EU402436 JN939462	Probe XXX	C10A, T18C	limited coverage (excluded)
Vanderwaltozyma yarrowii/51620 Analyzed Accessions: 0				
4	KY105798 NR_155996 KY105799 AY046183	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
1	KY110004	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Tetrapispora arboricola/113605 Analyzed Accessions: 0				
5	NR_155927 KY105600 AY046178 KY105601 KY105599	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Debaryomyces prosopidis/148098 Analyzed Accessions: 0				
5	PQ530893 NR_077067 PQ044656 MT187616 MZ227312	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Cephalotrichum nanum/644570 Analyzed Accessions: 0				
5	LC177640 MN535086 OP961970 FJ914706 FJ914668	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T15C	2.00 XXX limited coverage (excluded)
		Probe XXX	A5G, A6G, C10A	
Cephalotrichum dendrocephalum/644594 Analyzed Accessions: 0				
5	FJ914659 MG886313 FJ914660 FJ914691 KY379554	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T15C	2.00 XXX limited coverage (excluded)
		Probe XXX	A5G, A6G, C10A	
Saccharomyces arboricola/706196 Analyzed Accessions: 0				
5	KY104944 EF580917 MZ068118 NR_153296 KY104943	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Torulaspora indica/1056991 Analyzed Accessions: 0				
5	MW895345 HE575668 HE575669 NR_155891 HE574639	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] adriatica/1171601 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
5	HE799686 JN637464 HE799682 NR_152453 HE574653	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T16A	2.00 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Cora/1399800 Analyzed Accessions: 0				
5	KJ780628 KX772631 KJ780630 MK010863 KX772535	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Scheffersomyces coipomensis/1788519 Analyzed Accessions: 0				
5	LC120359 NR_111424 MF417396 KY109557 MT277139	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Saccharomyces eubayanus x Saccharomyces uvarum/1911678 Analyzed Accessions: 0				
4	KY105191 KY105193 KY105192 KY105190	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
1	KY109433	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T16A	2.00 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18G	
Yamadazyma tenuis/2315449 Analyzed Accessions: 0				
5	KY106801 HE612104 KY106799 HE612107 MH844426	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Vanderwaltozyma molinica/3393988 Analyzed Accessions: 0				
5	HM461683 HM461684 FJ873443 HM461680 HM461681	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Kluyveromyces/4910 Analyzed Accessions: 0				
4	LC229722 LC191810 LC155264 MN244369	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Torulaspora/4948 Analyzed Accessions: 0				
4	OM480730 MZ254721 LC486365 FJ873436	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Eremothecium coryli/33173 Analyzed Accessions: 0				
4	KY107678 LC663179	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
	LC663180 KY107673	Probe XXX	C10A, G17A	limited coverage (excluded)
[Candida] saitoana/45582 Analyzed Accessions: 0				
4	JX188109 JX188110 FN667843 OQ238757	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX
		Probe XXX	C10A, G17A	limited coverage (excluded)
Kurtzmaniella/549703 Analyzed Accessions: 0				
4	OM760529 MK130974 OM760530 PP936020	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX
		Probe XXX	C10A, G17A	limited coverage (excluded)
Trichurus terrophilus/644595 Analyzed Accessions: 0				
4	OM372923 FJ914684 OM372942 ON127887	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T15C	2.00 XXX
		Probe XXX	A5G, A6G, C10A	limited coverage (excluded)
Nakaseomyces uthaithaninus/1040648 Analyzed Accessions: 0				
4	AB610144 NR_154963 AB610143 AB610145	Forward XXX	Forward Missing	0.22 XXX
		Probe XXX	Probe Missing	limited coverage (excluded)
		Reverse XXX	Reverse Missing	
Scheffersomyces insectosa/1111082 Analyzed Accessions: 0				
4	PP808761 PP410361 LC120362 KY109564	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX
		Probe XXX	C10A, G17A	limited coverage (excluded)
Ogataea histrianica/1171602 Analyzed Accessions: 0				
4	HE799681 NR_155556 HE799680 HF559223	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX
		Probe XXX	C10A, G17A	limited coverage (excluded)
Zygosaccharomyces pseudobailii/1365887 Analyzed Accessions: 0				
4	OW847531 JX458105 JX458108 JX458107	Forward XXX	Forward Missing	0.22 XXX
		Probe XXX	Probe Missing	limited coverage (excluded)
		Reverse XXX	Reverse Missing	
Nakazawaea ishiwadae/1538179 Analyzed Accessions: 0				
4	KX078419 KX078416 KX078417 KX078418	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX
		Probe XXX	C10A, G17A	limited coverage (excluded)
Grigorovia jiainica (nom. inval.)/1734212 Analyzed Accessions: 0				
4	KY103648 EF612201 FJ196780 AB467302	Forward XXX	Forward Missing	0.22 XXX
		Probe XXX	Probe Missing	limited coverage (excluded)
		Reverse XXX	Reverse Missing	
Cryptococcus deuterogettii/1859096 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
4	PV938973 PV828368 KY102648 MN660280	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, A17T, A18T	0.34 XXX limited coverage (excluded)
		Probe XXX	C10A, T18C	
Eukaryota sp./1928008 Analyzed Accessions: 0				
4	MK208486 MK208485 MK208484 MK208495	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Zygosaccharomyces favi/2024901 Analyzed Accessions: 0				
4	KJ782415 KJ782412 KJ782413 KJ782416	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Rhodocollybia maculata/2074862 Analyzed Accessions: 0				
4	KU314929 OR822111 KU314930 MW862377	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, A18T	0.34 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18C	
Yamadazyma cocois/2681892 Analyzed Accessions: 0				
4	NR_173296 MN764370 MN764371 MN764369	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Vanderwaltozyma meishanica/3393987 Analyzed Accessions: 0				
4	HM461679 HM461678 HQ623545 NR_199198	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Eremothecium koelreuteriae/3454402 Analyzed Accessions: 0				
4	LC835708 LC835690 LC835692 LC835691	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Scheffersomyces stipitis/4924 Analyzed Accessions: 0				
3	JQ026363 DQ409167 MW165504	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Eremothecium gossypii/33169 Analyzed Accessions: 0				
3	LC835689 AF113137 LC835688	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Arxiozyma telluris/36907 Analyzed Accessions: 0				
3	OP642538 AJ871396 KY107954	Forward X	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-	3.33 XX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18G	
Scheffersomyces segobiensis/45512 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
3	LC120358 HG737349 NR_111217	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Kurtzmaniella natalensis/45571 Analyzed Accessions: 0				
3	MF927606 MF927607 ON797475	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Wickerhamia fluorescens/45790 Analyzed Accessions: 0				
3	KY110053 KY110054 FM178333	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Sarcosoma globosum/47207 Analyzed Accessions: 0				
3	OP023316 PP756121 FJ499393	Forward X	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-	3.33 XX limited coverage (excluded)
		Probe XXX	A5G, A6G, C10A, T18A	
Ogataea naganishii/50265 Analyzed Accessions: 0				
3	KY104411 MZ098645 KY108685	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] vartiovaarae/51918 Analyzed Accessions: 0				
3	FR774538 HF934034 KY106865	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] sonorensis/51926 Analyzed Accessions: 0				
3	KY106767 KY106770 KY106769	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Ogataea nonfermentans/53937 Analyzed Accessions: 0				
3	NR_155607 KY108687 MF574035	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Saccharomyces bayanus x Saccharomyces cerevisiae/89981 Analyzed Accessions: 0				
3	LT577618 EU719073 OP168927	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Tetrapisipora iriomotensis/113606 Analyzed Accessions: 0				
3	KY105605 NR_155929 AY046179	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Cephalotrichum purpureofuscum/186353 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
<u>3</u>	FJ914690 LN851031 FJ914688	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T15C	2.00 XXX limited coverage (excluded)
		Probe XXX	A5G, A6G, C10A	
Gomphidius smithii/186657 Analyzed Accessions: 0				
<u>3</u>	PV750447 MT703790 EF530941	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, A18T	0.34 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18C	
Gomphidius oregonensis/283674 Analyzed Accessions: 0				
<u>3</u>	KC581306 DQ533976 PP799164	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, A18T	0.34 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18C	
Blastobotrys serpentis/456365 Analyzed Accessions: 0				
<u>3</u>	KM056340 KM056339 AM410670	Forward X	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-	3.33 XX limited coverage (excluded)
		Probe XXX	C10A, G15A, G17A	
Jamesozyma hellenica/473794 Analyzed Accessions: 0				
<u>3</u>	NR_111415 EF620862 KY103644	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Ogataea kolombanensis/1171605 Analyzed Accessions: 0				
<u>3</u>	HF913624 NR_155557 HF559222	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Kazachstania molopis/1286812 Analyzed Accessions: 0				
<u>3</u>	NR_165506 HM627145 OR346357	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Saccharomyces cerevisiae x Saccharomyces eubayanus x Saccharomyces uvarum/1470621 Analyzed Accessions: 0				
<u>3</u>	MG641177 KF530462 MK049898	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Nakazawaea pomicola/1538183 Analyzed Accessions: 0				
<u>1</u>	KJ706397	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	5.89 X limited coverage (excluded)
		Probe ✓✓	C10A	
		Reverse ✓✓✓	T3C	
<u>2</u>	KJ706874 NR_138216	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Nakazawaea wickerhamii/1538186 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
<u>3</u>	HE799683 NR_155483 KY104377	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Saccharomycetales sp./1891166 Analyzed Accessions: 0				
<u>3</u>	MH811278 MW299120 MH811196	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Eremothecium leptocoridis/2841763 Analyzed Accessions: 0				
<u>3</u>	LC835753 LC835716 LC835713	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Pichia membranifaciens/4926 Analyzed Accessions: 0				
<u>2</u>	OR186608 OR186609	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Gomphidius glutinosus/5393 Analyzed Accessions: 0				
<u>2</u>	MT644863 LN714548	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, A18T	0.34 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18C	
Alloascoidea africana/27309 Analyzed Accessions: 0				
<u>2</u>	MH859161 NR_159607	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Saccharomycodes/36034 Analyzed Accessions: 0				
<u>2</u>	KP866237 KP866236	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T15A	2.00 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18A	
Ambrosiozyma/36917 Analyzed Accessions: 0				
<u>2</u>	PQ568195 PQ568179	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Eremothecium sinecaudum/45286 Analyzed Accessions: 0				
<u>2</u>	NR_155081 FJ422506	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Priceomyces medius/45504 Analyzed Accessions: 0				
<u>2</u>	KY108942 DQ409170	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] silvicultrix/51917 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
<u>2</u>	NR_165969 MK394131	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Cyberlindnera amylophila/53489 Analyzed Accessions: 0				
<u>2</u>	KY107350 MK394132	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-	2.00 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Cyberlindnera veronae/53496 Analyzed Accessions: 0				
<u>2</u>	DQ414543 AF335966	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Ambrosiozyma angophorae/53648 Analyzed Accessions: 0				
<u>2</u>	KF826536 KY106094	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Ogataea methylovora/53649 Analyzed Accessions: 0				
<u>2</u>	KY108683 KY108682	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Hanseniaspora guilliermondii/56406 Analyzed Accessions: 0				
<u>2</u>	MG367319 KY107803	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Ogataea ramenticola/104402 Analyzed Accessions: 0				
<u>2</u>	KY108713 KY108714	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Moniliophthora/221102 Analyzed Accessions: 0				
<u>2</u>	AY916754 MT162719	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-, A17-, A18-	4.00 XX limited coverage (excluded)
		Probe ✓✓	C10A	
Arxiozyma pintolopesii/278025 Analyzed Accessions: 0				
<u>2</u>	KP132307 KP132309	Forward X	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-	3.33 XX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18G	
Arxiozyma slooffiae/278027 Analyzed Accessions: 0				
<u>2</u>	OP642535 MT457819	Forward X	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-	3.33 XX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18G	
Jamesozyma zonata/308750 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
<u>2</u>	KY107970 KY107972	Forward ✖	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-	3.33 ✖✖ limited coverage (excluded)
		Probe ✖✖✖	C10A, G17A	
Kuraishia/317045 Analyzed Accessions: 0				
<u>2</u>	OP856837 PQ380277	Forward ✖✖✖	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 ✖✖✖ limited coverage (excluded)
		Probe ✖✖✖	C10A, G17A	
Kuraishia capsulata/317047 Analyzed Accessions: 0				
<u>1</u>	MH844414	Forward ✖✖	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, T16A	2.00 ✖✖✖ limited coverage (excluded)
		Probe ✖✖✖	C10A, G17A	
<u>1</u>	HQ413277	Forward ✖✖✖	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 ✖✖✖ limited coverage (excluded)
		Probe ✖✖✖	C10A, G17A	
Hanseniaspora pseudoguilliermondii/332229 Analyzed Accessions: 0				
<u>2</u>	OR295855 OQ357464	Forward ✖✖✖	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 ✖✖✖ limited coverage (excluded)
		Probe ✖✖✖	C10A, G17A	
Monosporozyma siamensis/382647 Analyzed Accessions: 0				
<u>2</u>	KY103669 NR_111356	Forward ✖✖✖	Forward Missing	0.22 ✖✖✖ limited coverage (excluded)
		Probe ✖✖✖	Probe Missing	
		Reverse ✖✖✖	Reverse Missing	
Ogataea polymorpha/460523 Analyzed Accessions: 0				
<u>2</u>	MK038753 KY102443	Forward ✖✖✖	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 ✖✖✖ limited coverage (excluded)
		Probe ✖✖✖	C10A, G17A	
Kuraishia ogatae/537810 Analyzed Accessions: 0				
<u>2</u>	EU722412 EU722411	Forward ✖✖✖	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 ✖✖✖ limited coverage (excluded)
		Probe ✖✖✖	C10A, G17A	
[Candida] vrieseae/622359 Analyzed Accessions: 0				
<u>2</u>	NR_155991 KY102517	Forward ✖✖	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-	2.00 ✖✖✖ limited coverage (excluded)
		Probe ✖✖✖	C10A, G17A	
Kazachstania taianensis/655095 Analyzed Accessions: 0				
<u>2</u>	GQ181169 KY103675	Forward ✖✖✖	Forward Missing	0.22 ✖✖✖ limited coverage (excluded)
		Probe ✖✖✖	Probe Missing	
		Reverse ✖✖✖	Reverse Missing	
Hannaella/663591 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
<u>1</u>	KX067813	Forward ✗	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-	3.33 ✗✗ limited coverage (excluded)
		Probe ✗✗✗	C10A, T18C	
<u>1</u>	OP754975	Forward ✗✗	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, T16A	2.00 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	C10A, G17A	
Hanseniaspora thailandica (nom. inval.)/689838 Analyzed Accessions: 0				
<u>2</u>	MG367285 MG251411	Forward ✗✗✗	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	C10A, G17A	
[Candida] sp. LCF-39/907295 Analyzed Accessions: 0				
<u>2</u>	HM461638 HM461639	Forward ✗✗✗	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	C10A, G17A	
Debaryomyces vindobonensis/1156046 Analyzed Accessions: 0				
<u>2</u>	MZ227344 HE681106	Forward ✗✗✗	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	C10A, G17A	
Maudiozyma bozae/1286813 Analyzed Accessions: 0				
<u>2</u>	KC118125 KX369579	Forward ✗✗✗	Forward Missing	0.22 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	Probe Missing	
		Reverse ✗✗✗	Reverse Missing	
Nakazawaea populi/1538184 Analyzed Accessions: 0				
<u>2</u>	NR_138241 KY104375	Forward ✗✗✗	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	C10A, G17A	
Xylaria aff. enterogena RV-2015/1636134 Analyzed Accessions: 0				
<u>2</u>	KP133314 KP133315	Forward ✗✗✗	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T16A, T20G	1.67 ✗✗✗ limited coverage (excluded)
		Probe ✗✗	A5G, C10A	
Pezizomycetes sp./1711683 Analyzed Accessions: 0				
<u>1</u>	KP991842	Forward ✗	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-	3.33 ✗✗ limited coverage (excluded)
		Probe ✗✗✗	A5G, C10A, T18A	
<u>1</u>	KP991657	Forward ✗	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-	3.33 ✗✗ limited coverage (excluded)
		Probe ✗✗✗	A5G, C10A, T18A	
Saccharomycetaceae sp./1769366 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
<u>1</u>	OQ708375	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
<u>1</u>	MH812850	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Spathaspora hagerdaliae/1774706 Analyzed Accessions: 0				
<u>2</u>	NR_155800 KU556168	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Spathaspora gorwiae/1852163 Analyzed Accessions: 0				
<u>2</u>	KC959474 NR_155784	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Kernia columnaris/2020669 Analyzed Accessions: 0				
<u>2</u>	FJ914707 FJ914671	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T15C	2.00 XXX limited coverage (excluded)
		Probe XXX	A5G, A6G, C10A	
Cephalotrichum gorgonifer/2041049 Analyzed Accessions: 0				
<u>2</u>	OW988170 NR_190248	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T15C	2.00 XXX limited coverage (excluded)
		Probe XXX	A5G, A6G, C10A	
Saccharomyces cf. cerevisiae/2069377 Analyzed Accessions: 0				
<u>2</u>	MK044014 MG757427	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Zygorulasporea chibaensis/2116648 Analyzed Accessions: 0				
<u>2</u>	NR_160344 MH055359	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Zygorulasporea dagestanica/2775096 Analyzed Accessions: 0				
<u>2</u>	NR_182852 MW036827	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Hanseniaspora terricola/3031818 Analyzed Accessions: 0				
<u>2</u>	MN706596 MN706599	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Klebsiella pneumoniae/573 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	CP096810	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T16A	2.00 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18G	
Cyberlindnera jadinii/4903 Analyzed Accessions: 0				
1	MG550146	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-	2.00 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Pachysolen tannophilus/4918 Analyzed Accessions: 0				
1	KY108725	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Dioszegia crocea/4974 Analyzed Accessions: 0				
1	MN065520	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Cryptococcus/5206 Analyzed Accessions: 0				
1	PV828378	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, A17T, G19T	0.34 XXX limited coverage (excluded)
		Probe XXX	C10A, T18C	
Cryptococcus depauperatus/5208 Analyzed Accessions: 0				
1	PV828385	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, A17T, A18T	0.34 XXX limited coverage (excluded)
		Probe XXX	C10A, T18C	
Suillus/5379 Analyzed Accessions: 0				
1	MK509801	Forward X	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-	7.00 ✓ limited coverage (excluded)
		Probe ✓✓	C10A	
Chroogomphus/5390 Analyzed Accessions: 0				
1	MG597339	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, A18T	0.34 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18C	
Gomphidius/5392 Analyzed Accessions: 0				
1	KU314926	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, A18T	0.34 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18C	
Candida tropicalis/5482 Analyzed Accessions: 0				
1	KY673188	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Collybia/5624 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	HM639274	Forward ✗	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-	3.33 ✗✗ limited coverage (excluded)
		Probe ✗✗✗	C10A, G17A, T18C	
Schwanniomyces etchellsii/28550 Analyzed Accessions: 0				
1	JQ726605	Forward ✗✗✗	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	C10A, G17A	
Debaryomyces udonii/28551 Analyzed Accessions: 0				
1	NR_077068	Forward ✗✗✗	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	C10A, G17A	
Eremothecium/33170 Analyzed Accessions: 0				
1	LC769456	Forward ✗✗✗	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	C10A, G17A	
Eremothecium ashbyi/33171 Analyzed Accessions: 0				
1	MH857759	Forward ✗✗✗	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	C10A, G17A	
Nakazawaea holstii/36025 Analyzed Accessions: 0				
1	AB449811	Forward ✗✗✗	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	C10A, G17A	
Ogataea minuta/36026 Analyzed Accessions: 0				
1	LC580394	Forward ✗✗✗	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	C10A, G17A	
Ambrosiozyma platypodis/36918 Analyzed Accessions: 0				
1	NR_153573	Forward ✗✗✗	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	C10A, G17A	
Mycena/41247 Analyzed Accessions: 0				
1	LC327056	Forward ✗✗✗	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-, G19T	4.00 ✗✗ limited coverage (excluded)
		Probe ✓✓	C10A	
[Candida] fluviatilis/45551 Analyzed Accessions: 0				
1	MT502785	Forward ✗✗✗	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	C10A, G17A	
[Candida] multigemmis/45569 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	NR_111434	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Wickerhamia/45789 Analyzed Accessions: 0				
1	PP692555	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Torulaspora globosa/48254 Analyzed Accessions: 0				
1	MG817611	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Ogataea kodamae/50263 Analyzed Accessions: 0				
1	KY108676	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
basidiomycete from Bashania fargesii/51124 Analyzed Accessions: 0				
1	U65615	Forward X	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-	3.33 XX limited coverage (excluded)
		Probe XXX	C10A, T18C	
[Candida] succiphila/51927 Analyzed Accessions: 0				
1	KY102419	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] pini/51934 Analyzed Accessions: 0				
1	KY106694	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Kurtzmaniella fragi/52254 Analyzed Accessions: 0				
1	FM178329	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Cyberlindnera mississippiensis/53607 Analyzed Accessions: 0				
1	KY103067	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-	2.00 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Hanseniaspora valbyensis/53685 Analyzed Accessions: 0				
1	MK044019	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Vibrio anguillarum/55601 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	CP011464	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T16A	2.00 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18G	
Hanseniaspora lindneri/56411 Analyzed Accessions: 0				
1	KY107973	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Jamesozyma rosinii/56413 Analyzed Accessions: 0				
1	KY103665	Forward X	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-	5.56 X limited coverage (excluded)
		Probe XXX	C10A, G17A, T18A	
		Reverse ✓✓✓	C1-, C2-	
[Candida] karawaiewii/58622 Analyzed Accessions: 0				
1	FR774536	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Saccharomyces chevalieri/62270 Analyzed Accessions: 0				
1	AF005714	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Saccharomyces ellipsoideus/62271 Analyzed Accessions: 0				
1	AF005716	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Pichia sporocuriosa/87101 Analyzed Accessions: 0				
1	OR346353	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Nowakowskiella/109880 Analyzed Accessions: 0				
1	PV921549	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Grigorovia humatica/121233 Analyzed Accessions: 0				
1	AB097397	Forward XXX	G4A, A9T, C10A, G11C	2.22 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18A	
		Reverse X	Reverse Missing	
Kurtzmaniella cleridarum/121466 Analyzed Accessions: 0				
1	KY108193	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
[Candida] tammaniensis/130808 Analyzed Accessions: 0				
1	KY106792	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-	2.00 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] caryicola/147333 Analyzed Accessions: 0				
1	MN861537	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Kuraishia hungarica/157291 Analyzed Accessions: 0				
1	EU722410	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Melanophyllum haematospermum/181865 Analyzed Accessions: 0				
1	PQ653280	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Trichurus/186363 Analyzed Accessions: 0				
1	MK442919	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T15C	2.00 XXX limited coverage (excluded)
		Probe XXX	A5G, A6G, C10A	
Trichurus spiralis/186364 Analyzed Accessions: 0				
1	MK463989	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T15C	2.00 XXX limited coverage (excluded)
		Probe XXX	A5G, A6G, C10A	
Gomphidius subroseus/186658 Analyzed Accessions: 0				
1	DQ384576	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, A18C	0.34 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18C	
Hanseniaspora meyeri/211097 Analyzed Accessions: 0				
1	KY107811	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] heliconiae/221911 Analyzed Accessions: 0				
1	KY106498	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] temnochilae/234530 Analyzed Accessions: 0				
1	FJ153211	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
Yamadazyma paraphyllophila/267649 Analyzed Accessions: 0				
1	NR_111164	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] michaelii/271314 Analyzed Accessions: 0				
1	KY102212	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-	2.00 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] cerambycidarum/271315 Analyzed Accessions: 0				
1	NR_111392	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-	2.00 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] endomychidarum/271317 Analyzed Accessions: 0				
1	NR_111393	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Arxiozyma bovina/278024 Analyzed Accessions: 0				
1	KP132306	Forward X	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-	3.33 XX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18G	
Teunomyces lycoperdinae/297653 Analyzed Accessions: 0				
1	NR_111400	Forward X	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-	3.33 XX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18C	
[Candida] lessepsii/297795 Analyzed Accessions: 0				
1	NR_111395	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-	2.00 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] pattaniensis/298207 Analyzed Accessions: 0				
1	NR_144776	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
saccharomycete sp. jbra611/301062 Analyzed Accessions: 0				
1	AY796126	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
saccharomycete sp. MVCH-J15/301080 Analyzed Accessions: 0				
1	AY796191	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
saccharomycete sp. SCH-3/301082 Analyzed Accessions: 0				
1	AY796193	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
saccharomycete sp. MVCH-J18/301083 Analyzed Accessions: 0				
1	AY796194	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
saccharomycete sp. jbra2913/306905 Analyzed Accessions: 0				
1	AY842480	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Naganishia randhawae/312146 Analyzed Accessions: 0				
1	MF662353	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
saccharomycete sp. NZSBMV10-B6/317868 Analyzed Accessions: 0				
1	AY942697	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Saccharomyces cerevisiae x Saccharomyces kudriavzevii/332112 Analyzed Accessions: 0				
1	PQ898003	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Teunomyces tritomae/348312 Analyzed Accessions: 0				
1	NR 111399	Forward X	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-	3.33 XX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18C	
[Candida] tibetensis/356546 Analyzed Accessions: 0				
1	MN861536	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Saccharomycetaceae sp. BC41/364392 Analyzed Accessions: 0				
1	DQ317356	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Hyphoderma heterocystidia/394044 Analyzed Accessions: 0				
1	DQ677495	Forward X	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-	3.33 XX

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
		Probe XXX	C10A, G17A, T18C	limited coverage (excluded)
Saccharomyces cerevisiae/Candida shehatae fusant/410624 Analyzed Accessions: 0				
1	EF042285	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal endophyte/410823 Analyzed Accessions: 0				
1	PQ418041	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Spathaspora/412764 Analyzed Accessions: 0				
1	LC762486	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] sp. BCMU BX01/413943 Analyzed Accessions: 0				
1	AB285024	Forward X	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T14A	3.33 XX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] sp. BCMU BX02/413944 Analyzed Accessions: 0				
1	AB285025	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-	2.00 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] sp. BCMU BX03/413945 Analyzed Accessions: 0				
1	AB285026	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-	2.00 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] sp. BCMU BX04/427486 Analyzed Accessions: 0				
1	AB294738	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Saccharomycetaceae sp. KS2-2007/447692 Analyzed Accessions: 0				
1	EF620029	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
[Candida] sp. EJ2M02/452172 Analyzed Accessions: 0				
1	FJ873562	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] nonsorbophila/456248 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	KY102232	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Pichia rarassimilans/477177 Analyzed Accessions: 0				
1	KY104644	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Xylaria telfairii/477509 Analyzed Accessions: 0				
1	OR088130	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T16A, T20G	1.67 XXX limited coverage (excluded)
		Probe XX	A5G, C10A	
Saccharomyces cerevisiae x Saccharomyces uvarum/489137 Analyzed Accessions: 0				
1	KY105189	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Candidozyma auris/498019 Analyzed Accessions: 0				
1	KY657043	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
[Candida] sp. SG6L02/511688 Analyzed Accessions: 0				
1	FJ153162	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
fungal endophyte sp. R78_2/548639 Analyzed Accessions: 0				
1	EU561626	Forward X	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-	3.33 XX limited coverage (excluded)
		Probe XXX	C10A, G15A, G17A	
[Candida] sp. SM1S01/561015 Analyzed Accessions: 0				
1	FJ153160	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Saccharomyces cerevisiae x Saccharomyces paradoxus/595493 Analyzed Accessions: 0				
1	OP243461	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
[Candida] sp. GJ15M15/639351 Analyzed Accessions: 0				
1	FJ873586	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] sp. GY7M03/639365 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	FJ873600	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Cephalotrichum castaneum/644568 Analyzed Accessions: 0				
1	FJ914681	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T15C	2.00 XXX limited coverage (excluded)
		Probe XXX	A5G, A6G, C10A	
Grigorovia aff. yakushimaensis IMUFRJ 51944/655874 Analyzed Accessions: 0				
1	FN424113	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Saccharomycetes sp. HZ140/700463 Analyzed Accessions: 0				
1	GU213441	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Saccharomycetes sp. HZ178/700465 Analyzed Accessions: 0				
1	GU213443	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Saccharomycetes sp. OHZ67/747488 Analyzed Accessions: 0				
1	GU931321	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. ARIZ AZ0283/757750 Analyzed Accessions: 0				
1	HM123030	Forward X	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-	3.33 XX limited coverage (excluded)
		Probe XXX	A5G, C10A, T18A	
fungal sp. ARIZ AZ0659/758095 Analyzed Accessions: 0				
1	HM123375	Forward X	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-	3.33 XX limited coverage (excluded)
		Probe XXX	A5G, C10A, T18A	
[Candida] sp. TCJ249/765288 Analyzed Accessions: 0				
1	HM044870	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Priceomyces/766759 Analyzed Accessions: 0				
1	LC815030	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
[Candida] sphagnicola/766783 Analyzed Accessions: 0				
1	FN868154	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
fungal endophyte sp. g19/859732 Analyzed Accessions: 0				
1	HM537029	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Scheffersomyces queiroziae/880181 Analyzed Accessions: 0				
1	HM566445	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] sp. ZIM 2279/884009 Analyzed Accessions: 0				
1	HE574656	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] sp. AZ004/933230 Analyzed Accessions: 0				
1	HQ413283	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] sp. 1 TMS-2011/942759 Analyzed Accessions: 0				
1	HQ631020	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Ogataea parapolymorpha/1005962 Analyzed Accessions: 0				
1	XR_001314120	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Kazachstania bromeliacearum/1006466 Analyzed Accessions: 0				
1	KY103627	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
[Candida] sp. LCF-105/1041462 Analyzed Accessions: 0				
1	HQ623566	Forward X	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-	3.33 XX limited coverage (excluded)
		Probe XXX	C10A, G17A, T18C	
Wickerhamomyces ciferrii/1041607 Analyzed Accessions: 0				
1	FJ153191	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
Cladosporium halotolerans/1052096 Analyzed Accessions: 0				
1	MH171918	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T20G	4.00 XX limited coverage (excluded)
		Probe ✓✓	C10A	
[Candida] sp. ZIM 2329/1056998 Analyzed Accessions: 0				
1	HE574658	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] sp. K-708/1095004 Analyzed Accessions: 0				
1	HE602532	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] sp. KBP Y-3971/1095006 Analyzed Accessions: 0				
1	HE602534	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] sp. LCF963/1095152 Analyzed Accessions: 0				
1	JN637463	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T16A	2.00 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Scheffersomyces lignosus/1111081 Analyzed Accessions: 0				
1	LC120360	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Saccharomycetales sp. WB/1117288 Analyzed Accessions: 0				
1	JN084124	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
[Candida] sp. KP-2012/1130972 Analyzed Accessions: 0				
1	HQ916868	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] sp. DMKU-RK512/1177565 Analyzed Accessions: 0				
1	AB714251	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] sp. K29/1177566 Analyzed Accessions: 0				
1	AB714255	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
[Candida] sp. ST-236/1177569 Analyzed Accessions: 0				
1	AB714252	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] sp. LN06/1177570 Analyzed Accessions: 0				
1	AB714253	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Petrozyma/1196393 Analyzed Accessions: 0				
1	KX766419	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Starmerella bacillaris/1247836 Analyzed Accessions: 0				
1	MH595413	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Ogataea wickerhamii/1249076 Analyzed Accessions: 0				
1	KY108721	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
fungal sp. F271/1312607 Analyzed Accessions: 0				
1	JQ747698	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Priceomyces vitoshaensis/1331796 Analyzed Accessions: 0				
1	PP481707	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] aff. saitoana VKPM Y3988/1343201 Analyzed Accessions: 0				
1	HG007840	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Ambrosiozyma kamigamensis/1382202 Analyzed Accessions: 0				
1	KY101638	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Ambrosiozyma neoplatypodis/1382203 Analyzed Accessions: 0				
1	KY101645	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
Psathyrella ornatispora/1395785 Analyzed Accessions: 0				
1	KC992968	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Ogataea mangiferae/1428343 Analyzed Accessions: 0				
1	KF585022	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
fungal sp. AT1350/1437062 Analyzed Accessions: 0				
1	KF725624	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Nakazawaea siamensis/1449382 Analyzed Accessions: 0				
1	MG367290	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Kuraishia piskuri/1538145 Analyzed Accessions: 0				
1	NR_155218	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Kuraishia cidri/1538149 Analyzed Accessions: 0				
1	KY108183	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Nakazawaea anatomiae/1538176 Analyzed Accessions: 0				
1	KM065947	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Nakazawaea ernobii/1538178 Analyzed Accessions: 0				
1	NR_165955	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Nakazawaea laoshanensis/1538180 Analyzed Accessions: 0				
1	FJ613523	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Nakazawaea wyomingensis/1538187 Analyzed Accessions: 0				
1	KY108657	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
fungal sp. 169 MCS-2014/1543510 Analyzed Accessions: 0				
1	KM374215	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 17 MCS-2014/1543511 Analyzed Accessions: 0				
1	KM374167	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 173 MCS-2014/1543515 Analyzed Accessions: 0				
1	KM374217	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 174 MCS-2014/1543516 Analyzed Accessions: 0				
1	KM374118	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 176 MCS-2014/1543518 Analyzed Accessions: 0				
1	KM374219	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 177 MCS-2014/1543519 Analyzed Accessions: 0				
1	KM374119	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 178 MCS-2014/1543520 Analyzed Accessions: 0				
1	KM374220	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 179 MCS-2014/1543521 Analyzed Accessions: 0				
1	KM374221	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 18 MCS-2014/1543522 Analyzed Accessions: 0				
1	KM374168	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 182 MCS-2014/1543525 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	KM374224	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 185 MCS-2014/1543528 Analyzed Accessions: 0				
1	KM374120	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 186 MCS-2014/1543529 Analyzed Accessions: 0				
1	KM374227	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 188 MCS-2014/1543530 Analyzed Accessions: 0				
1	KM374228	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 190 MCS-2014/1543533 Analyzed Accessions: 0				
1	KM374122	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 192 MCS-2014/1543534 Analyzed Accessions: 0				
1	KM374124	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 193 MCS-2014/1543535 Analyzed Accessions: 0				
1	KM374125	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 198 MCS-2014/1543537 Analyzed Accessions: 0				
1	KM374231	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 509 MCS-2014/1543581 Analyzed Accessions: 0				
1	KM374082	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 510 MCS-2014/1543582 Analyzed Accessions: 0				
1	KM374250	Forward XXX	Forward Missing	0.22 XXX
		Probe XXX	Probe Missing	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
		Reverse XXX	Reverse Missing	limited coverage (excluded)
fungal sp. 511 MCS-2014/1543583 Analyzed Accessions: 0				
1	KM374251	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 514 MCS-2014/1543586 Analyzed Accessions: 0				
1	KM374253	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 516 MCS-2014/1543588 Analyzed Accessions: 0				
1	KM374255	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 517 MCS-2014/1543589 Analyzed Accessions: 0				
1	KM374256	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 520 MCS-2014/1543592 Analyzed Accessions: 0				
1	KM374259	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 521 MCS-2014/1543593 Analyzed Accessions: 0				
1	KM374260	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 522 MCS-2014/1543594 Analyzed Accessions: 0				
1	KM374261	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 523 MCS-2014/1543595 Analyzed Accessions: 0				
1	KM374262	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 539 MCS-2014/1543596 Analyzed Accessions: 0				
1	KM374263	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
fungal sp. 540 MCS-2014/1543597 Analyzed Accessions: 0				
1	KM374264	Forward XXX	Forward Missing	0.22
		Probe XXX	Probe Missing	XXX
		Reverse XXX	Reverse Missing	limited coverage (excluded)
fungal sp. 541 MCS-2014/1543598 Analyzed Accessions: 0				
1	KM374265	Forward XXX	Forward Missing	0.22
		Probe XXX	Probe Missing	XXX
		Reverse XXX	Reverse Missing	limited coverage (excluded)
fungal sp. 547 MCS-2014/1543604 Analyzed Accessions: 0				
1	KM374271	Forward XXX	Forward Missing	0.22
		Probe XXX	Probe Missing	XXX
		Reverse XXX	Reverse Missing	limited coverage (excluded)
fungal sp. 551 MCS-2014/1543608 Analyzed Accessions: 0				
1	KM374275	Forward XXX	Forward Missing	0.22
		Probe XXX	Probe Missing	XXX
		Reverse XXX	Reverse Missing	limited coverage (excluded)
fungal sp. 554 MCS-2014/1543609 Analyzed Accessions: 0				
1	KM374276	Forward XXX	Forward Missing	0.22
		Probe XXX	Probe Missing	XXX
		Reverse XXX	Reverse Missing	limited coverage (excluded)
fungal sp. 555 MCS-2014/1543610 Analyzed Accessions: 0				
1	KM374277	Forward XXX	Forward Missing	0.22
		Probe XXX	Probe Missing	XXX
		Reverse XXX	Reverse Missing	limited coverage (excluded)
fungal sp. 558 MCS-2014/1543612 Analyzed Accessions: 0				
1	KM374279	Forward XXX	Forward Missing	0.22
		Probe XXX	Probe Missing	XXX
		Reverse XXX	Reverse Missing	limited coverage (excluded)
fungal sp. 560 MCS-2014/1543613 Analyzed Accessions: 0				
1	KM374280	Forward XXX	Forward Missing	0.22
		Probe XXX	Probe Missing	XXX
		Reverse XXX	Reverse Missing	limited coverage (excluded)
fungal sp. 561 MCS-2014/1543614 Analyzed Accessions: 0				
1	KM374281	Forward XXX	Forward Missing	0.22
		Probe XXX	Probe Missing	XXX
		Reverse XXX	Reverse Missing	limited coverage (excluded)
fungal sp. 562 MCS-2014/1543615 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	KM374282	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 563 MCS-2014/1543616 Analyzed Accessions: 0				
1	KM374283	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 565 MCS-2014/1543618 Analyzed Accessions: 0				
1	KM374284	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 567 MCS-2014/1543620 Analyzed Accessions: 0				
1	KM374286	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 568 MCS-2014/1543621 Analyzed Accessions: 0				
1	KM374287	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 605 MCS-2014/1543642 Analyzed Accessions: 0				
1	KM374292	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 613 MCS-2014/1543648 Analyzed Accessions: 0				
1	KM374084	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 614 MCS-2014/1543649 Analyzed Accessions: 0				
1	KM374297	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 619 MCS-2014/1543654 Analyzed Accessions: 0				
1	KM374300	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 622 MCS-2014/1543656 Analyzed Accessions: 0				
1	KM374085	Forward XXX	Forward Missing	0.22 XXX
		Probe XXX	Probe Missing	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
		Reverse XXX	Reverse Missing	limited coverage (excluded)
fungal sp. 625 MCS-2014/1543659 Analyzed Accessions: 0				
1	KM374157	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 627 MCS-2014/1543661 Analyzed Accessions: 0				
1	KM374301	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 628 MCS-2014/1543662 Analyzed Accessions: 0				
1	KM374302	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 629 MCS-2014/1543663 Analyzed Accessions: 0				
1	KM374159	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 15h/1629110 Analyzed Accessions: 0				
1	KJ956018	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
fungal sp. 23h/1629113 Analyzed Accessions: 0				
1	KJ956021	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, T16A	2.00 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Cephalotrichum/1658452 Analyzed Accessions: 0				
1	PQ422811	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T15C	2.00 XXX limited coverage (excluded)
		Probe XXX	A5G, A6G, C10A	
Amanita kalamundae/1705360 Analyzed Accessions: 0				
1	KP898378	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-, G19T	4.00 XX limited coverage (excluded)
		Probe ✓✓	C10A	
Sordariomycetes sp./1709940 Analyzed Accessions: 0				
1	KY965434	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T16G	3.33 XX limited coverage (excluded)
		Probe XX	A5G, C10A	
Yamadazyma endophytica/1714400 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	KT307981	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] sp. 1 AK-2015/1739950 Analyzed Accessions: 0				
1	LN890936	Forward X	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-	3.33 XX limited coverage (excluded)
		Probe XXX	C10A, G15A, G17A	
[Candida] sp. DspF14/1745351 Analyzed Accessions: 0				
1	KT958535	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-	2.00 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
[Candida] sp. DspF15/1745352 Analyzed Accessions: 0				
1	KT958536	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-	2.00 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Helotiales sp./1755440 Analyzed Accessions: 0				
1	MF331014	Forward XX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-	2.00 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Ascomycota sp. L210/1796131 Analyzed Accessions: 0				
1	KU535740	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	2.00 XXX limited coverage (excluded)
		Probe XX	A5G, C10A	
Saccharomyces uvarum x Saccharomyces eubayanus/1862628 Analyzed Accessions: 0				
1	LT594194	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Clavariaceae sp./1935309 Analyzed Accessions: 0				
1	MH811373	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T20C	0.34 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Yamadazyma laniorum/1944661 Analyzed Accessions: 0				
1	KY588136	Forward XXX	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 XXX limited coverage (excluded)
		Probe XXX	C10A, G17A	
Saccharomyces jurei/1987369 Analyzed Accessions: 0				
1	NR_197987	Forward XXX	Forward Missing	0.22 XXX limited coverage (excluded)
		Probe XXX	Probe Missing	
		Reverse XXX	Reverse Missing	
Dictyosporiaceae sp./2012064 Analyzed Accessions: 0				

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
1	MH812580	Forward ✗	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-	3.33 ✗✗ limited coverage (excluded)
		Probe ✗✗✗	A5G, C10A, T18A	
Chroogomphus britannicus/2052982 Analyzed Accessions: 0				
1	MG457841	Forward ✗✗✗	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, A18T	0.34 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	C10A, G17A, T18C	
Veluticeps africana/2075487 Analyzed Accessions: 0				
1	MH873336	Forward ✗✗✗	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T16G, A18C, G19A	4.00 ✗✗ limited coverage (excluded)
		Probe ✓✓	C10A	
Saccharomycotina sp./2100775 Analyzed Accessions: 0				
1	MH001970	Forward ✗✗✗	Forward Missing	0.22 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	Probe Missing	
		Reverse ✗✗✗	Reverse Missing	
Cora boleslia/2770919 Analyzed Accessions: 0				
1	KF443230	Forward ✗✗✗	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-, T16-	0.67 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	C10A, G17A	
Jamesozyma jinghongensis/2862417 Analyzed Accessions: 0				
1	NR_173271	Forward ✗	A3G, G4A, G8A, A9T, C10A, G11C	2.45 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	C10A, G17A, T18A	
		Reverse ✗✗✗	Reverse Missing	
Cyberlindnera maritima/2878918 Analyzed Accessions: 0				
1	KY106562	Forward ✗✗	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-, C13-, T14-, T15-	2.00 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	C10A, G17A	
Gomphidius cf. glutinosus/2912858 Analyzed Accessions: 0				
1	OM401944	Forward ✗✗✗	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, A18T	0.34 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	C10A, G17A, T18C	
Vanderwaltozyma huisunica/3393986 Analyzed Accessions: 0				
1	NR_199199	Forward ✗✗✗	Forward Missing	0.22 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	Probe Missing	
		Reverse ✗✗✗	Reverse Missing	
Torulaspora jiuxiensis/3398717 Analyzed Accessions: 0				
1	MK682817	Forward ✗✗✗	Forward Missing	0.22 ✗✗✗ limited coverage (excluded)
		Probe ✗✗✗	Probe Missing	
		Reverse ✗✗✗	Reverse Missing	

#Accessions	#Accession Details	Oligo	Mutation	BDS Analysis Performance Prediction
Hannaella pteridophytorum/3421687 Analyzed Accessions: 0				
1	OP470253	Forward ✗	G1-, C2-, A3-, G4-, A5-, C6-, A7-, G8-, A9-, C10-, G11-, T12-	3.33 ✗✗ limited coverage (excluded)
		Probe ✗✗✗	C10A, T18C	