

Supplemental Table 1

Genes that function in biofilm development				
Gene	Molecular function of gene product ^a	Mutant phenotype ^b	Filamentation	Reference
<i>ACE2</i>	Transcription factor	<i>ace2</i> ⁻ : severe biofilm defect; aberrant hyphae formation	Aberrant	118
<i>ADH1</i>	Alcohol dehydrogenase	<i>adh1</i> ⁻ : increased biofilm biomass	Normal	30
<i>ADH5</i>	Alcohol dehydrogenase	<i>ADH5</i> -oe: increased biofilm matrix	Normal	127
<i>ALS1</i>	Cell wall protein	<i>als1</i> - <i>als3</i> : severe biofilm defect. <i>ALS1</i> -oe: restores defect in <i>bcr1</i> -biofilm	Normal	22, 28
<i>ALS2</i>	Cell wall protein	<i>als2</i> ^{+/-} : defect in biofilm formation and hyphae formation. <i>ALS2</i> -oe: partially restores <i>als1</i> - <i>als3</i> - biofilm defect	Aberrant	22, 123
<i>ALS3</i>	Cell wall protein	<i>als3</i> ⁻ : reduced biofilm formation. <i>als1</i> - <i>als3</i> ⁻ : severe biofilm defect in vitro. <i>ALS3</i> -oe: restores biofilm in <i>bcr1</i> - and in <i>efg1</i> -	Normal	22, 28, 71
<i>ALS4</i>	Cell wall protein	<i>ALS4</i> -oe: partially restores defects in <i>als1</i> - <i>als3</i> -biofilm	Normal	22
<i>ALS5</i>	Cell wall protein	<i>ALS5</i> -oe: partially restores defects in <i>als1</i> - <i>als3</i> -biofilm	Normal	22
<i>ALS7</i>	Cell wall protein	<i>ALS7</i> -oe: partially restores defects in <i>als1</i> - <i>als3</i> -biofilm	Normal	22

<i>ALS9</i>	Cell wall protein	<i>ALS9</i> -oe: partially restores defects in <i>als1-</i> <i>als3</i> -biofilm	Normal	22
<i>BCR1</i>	Transcription factor	<i>bcr1-</i> : severe biofilm defect. <i>BCR1</i> -oe: partially restores biofilm in <i>tec1-</i>	Normal	27, 28
<i>CAT2</i>	Carnitine acetyltransferase	<i>cat2-</i> : reduced biofilm formation and biofilm thickness	Normal	131
<i>CBK1</i>	Protein Kinase	<i>cbk1-</i> : severe biofilm defect and defective hyphae formation	Defective	78
<i>CDR1</i>	Drug efflux pump	<i>cdr1-</i> : inhibited biofilm growth in the presence of azoles	Normal	89
<i>CDR2</i>	Drug efflux pump	<i>cdr2-</i> : inhibited biofilm growth in the presence of azoles	Normal	89
<i>CHK1</i>	Histidine kinase	<i>chk1-</i> : forms farnesol-resistant biofilm	Normal	129
<i>CPH1</i>	Transcription factor	<i>cph1-</i> <i>efg1-</i> : thin biofilm with reduce number of hyphae	Normal	119
<i>CSA1</i>	Cell wall protein	<i>csa1-</i> : defect in biofilm formation; additive with <i>pga10-</i> and <i>rbt5-</i>	Normal	74
<i>CSH1</i>	Aryl-alcohol dehydrogenase	<i>CSH1</i> -oe: decreased biofilm matrix in <i>zap1-</i>	Normal	29
<i>CZF1</i>	Transcription factor	<i>czf1-</i> : defect in biofilm formation. Aberrant hyphae	Aberrant	120
<i>EAP1</i>	Cell wall protein	<i>eap1-</i> : severe defect in biofilm formation	Normal	56

<i>ECE1</i>	Putative trans-membrane protein	<i>ECE1</i> -oe: restores defect in <i>bcr1</i> -biofilm	Normal	28
<i>EFG1</i>	Transcription factor	<i>efg1</i> -: severe biofilm defect and hyphae formation	Defective	14, 121
<i>FKS1</i>	Beta-1,3 glucan synthase	<i>fks1</i> / <i>FKS1</i> : reduced matrix production, reduce biofilm resistance to fluconazole	Normal	97
<i>FLO8</i>	Transcription factor	<i>czf1</i> -: defect in biofilm formation. Defective hyphae formation	Defective	122
<i>GCA1</i>	Glucosylase	<i>GCA1</i> -oe: increased biofilm matrix	Normal	29
<i>GCA2</i>	Glucosylase	<i>GCA2</i> -oe: increased biofilm matrix	Normal	29
<i>GCN4</i>	Transcription factor	<i>gcn4</i> -: decreased biofilm biomass	Normal	90
<i>GIN4</i>	Protein Kinase	<i>gin4</i> -: severe biofilm defect; defective hyphae	Defective	78
<i>HWP1</i>	Cell wall protein	<i>hwp1</i> -: defect in biofilm formation and hyphal formation. <i>HWP1</i> -oe: restores biofilm in <i>bcr1</i> -	Defective	23, 28
<i>HWP2</i>	Cell wall protein	<i>hwp2</i> -: decreased biomass	Defective	67
<i>IFD6</i>	Aryl-alcohol dehydrogenase	<i>IFD6</i> -oe: decreased biofilm matrix in <i>zap1</i> -	Normal	29
<i>IRE1</i>	Protein Kinase	<i>ire1</i> -: severe biofilm defect and hyphae formation	Defective	78
<i>KEM1</i>	Exo-RNase	<i>kem1</i> -: severe medium-dependent defect in biofilm formation	Defective	132

		and hyphal formation		
<i>MDR1</i>	Drug efflux pump	<i>mdr1-</i> : inhibited biofilm growth in the presence of azoles	Normal	89
<i>MDS3</i>	Unknown	<i>mds3-</i> : severe medium-dependent defect in biofilm formation and hyphal formation	Defective	132
<i>MKC1</i>	MAP Kinase	<i>mkc1-</i> : defective biofilm, with reduced filamentation	Normal	64
<i>NDH51</i>	Subunit of nicotinamide adenine dinucleotide dehydrogenase complex I	<i>ndh51-</i> : decreased biofilm mass	Normal	133
<i>NRG1</i>	Transcription factor	<i>nrg1-</i> : decreased release of dispersal cells	Aberrant	99
<i>NUP85</i>	Nuclear pore protein	<i>nup85-</i> : severe medium-dependent defect in biofilm formation and hyphal formation	Defective	132
<i>OCH1</i>	Alpha-1,6-mannosyltransferase	<i>och1-</i> : defect in biofilm formation. Cellular aggregation, cell wall defects	Normal	38
<i>PBR1</i>	Unknown	<i>pbr1-</i> : forms a thin biofilm with reduced matrix production	Normal	68
<i>PES1</i>	Pescadillo homolog	<i>pes1-</i> : reduced cell dispersion <i>PES1-oe</i> : increased cell dispersion	Normal	98
<i>PDX1</i>	Pyruvate dehydrogenase	<i>pdx1-</i> : biofilm with reduced density	Normal	133

<i>PGA1</i>	Cell wall protein	<i>pga1-</i> : reduced metabolic activity	Normal	124
<i>PGA10</i>	Cell wall protein	<i>pga10-</i> : defect in biofilm formation; additive with <i>rbt5-</i> and <i>csa1-</i>	Aberrant	74
<i>PMT1</i>	Mannosyltransferase	<i>pmt1-</i> : biofilm with reduced biomass. Defective hyphae	Defective	125
<i>PMT2</i>	mannosyltransferase	<i>pmt2-/+</i> : biofilm with reduced biomass. Defective hyphae	Defective	125
<i>PMT4</i>	Mannosyltransferase	<i>pmt4-</i> : moderate defect in biofilm formation	Normal	125
<i>PMT6</i>	Mannosyltransferase	<i>pmt6-</i> : moderate defect in biofilm formation	Normal	125
<i>RBT1</i>	Cell wall protein	<i>rbt1-</i> : reduced biomass	Normal	67
<i>RBT5</i>	Cell wall protein	<i>rbt5-</i> : defect in biofilm formation; additive with <i>pga10-</i> and <i>csa1-</i> <i>RBT5-oe</i> : restores biofilm in <i>bcr1-</i>	Normal	28, 74
<i>RIX7</i>	AAA ATPase	<i>rix7+/-</i> : defect in biofilm formation	Normal	134
<i>SUN41</i>	Cell wall protein	<i>sun41-</i> : severe biofilm defect; aberrant hyphae	Aberrant	75
<i>SUV3</i>	Mito-chondrial RNA helicase	<i>suv3-</i> : severe medium-dependent defect in biofilm formation and hyphae formation	Defective	132
<i>TEC1</i>	Transcription factor	<i>tec1-</i> : severe biofilm defect and hyphae formation	Defective	27
<i>TOR1</i>	Phosphatidylinositol kinases	Rapamycin treatment inhibits <i>TOR1</i> resulting in decreased	Defective	130

		adhesion of cells and loss of filamentation		
<i>UME6</i>	Transcription factor	<i>ume6-</i> : reduced biomass biofilm and defective hyphae formation. Increased cell release from biofilm.	Defective	⁹⁸
<i>VAM3</i>	Vacuolar trafficking	<i>vam3-</i> : biofilm defect, fragile biofilm, reduced biomass and abnormal hyphae	Aberrant	¹³⁵
<i>VPS1</i>	Dynamain-family GTPase-related protein	<i>vps1-</i> : rudimentary biofilm composed primarily of yeast and pseudohyphal	Aberrant	¹³⁶
<i>YAK1</i>	Protein Kinase	<i>yak1-</i> : severe biofilm defect and hyphal formation	Defective	¹²⁸
<i>YWP1</i>	Cell wall protein	<i>ywp1-</i> : increased adhesion. Biofilm formation by only yeast-form cells	Normal	¹²⁶
<i>ZAP1</i>	Transcription factor	<i>zap1-</i> : increased production of biofilm matrix	Normal	²⁹

Footnotes:

a. Molecular functions have been inferred from protein sequence homology in most cases.

b. Mutant phenotype was observed with either homozygous loss-of-function alleles (e.g., *als1-* refers to an *als1/als1* homozygous mutant) or over-expression alleles (e.g., *ALS1-oe* refers to a strain in which one *ALS1* allele has been fused to a strong promoter). The strains which have *-/+* after the gene indicate a heterozygous genotype. The "*als2+/-*" is a heterozygous strain which has reduced but not abolished *ALS2* expression. Negative results (absence of a phenotype) are omitted.

c. The ability to filament was record as "Normal" for those mutants able to form hyphae, "Defective" for those unable to form hyphae, or "Aberrant" for those with an intermediate phenotype.