

Table S3. Summary of genomic variations in the clinical, environmental and fluconazole-resistant strains

Genome Comparisons	Deleted/sequence diverged genes ^a	Amplified genes ^b
	WM276 tiling array	
WM276gfp2 vs WM276	24	0
R794 vs WM276	59	2
KB3864 vs WM276	58	7
	R265 tiling array	
R272 vs R265	18	0
RB28 vs R265	16	3
	R265 tiling array	
R-1402F vs R-1402	0	276^c
R-1401F vs R-1401	5	340^d
R-1346F vs R-1346	0	0
R-1347F vs R-1347	65^e	2
	WM276 tiling array	
R-1413F vs R-1413	1	75^f
R-1412F vs R-1412	231^g	12

^adeletion/sequence divergence designated at \log_2 ratio \leq 1.0

^bamplification designated at \log_2 ratio \geq 1.0

^c involves amplification of chromosomes 1 and 10

^dinvolves amplification of a major portion of chromosome 1

^einvolves deletion/sequence divergence of chromosomes 6, 11, 12

^finvolves amplification of chromosome 10

^ginvolves amplification of chromosome 14