

Suppl. Table 3. Heterozygosity across chromosomes is not correlated with frequency of crossover events or nondisjunction (NDJ). Data are relative to *C. neoformans* H99 genome. Only one breakpoint for each gene conversion is considered.

Chr	Size	Num SNPs	SNP rate	Num. breaks	NDJ
1	2291499	72959	0.0318	1	1
2	1621675	39814	0.0246	1	29
3	1575141	50117	0.0318	5	6
4	1084805	30247	0.0279	2	0
5	1814975	54199	0.0299	6	0
6	1422463	45598	0.0321	3	0
7	1399503	42668	0.0305	7	0
8	1398693	42582	0.0304	1	0
9	1186808	37586	0.0317	1	12
10	1059964	29439	0.0278	5	1
11	1561994	47038	0.0301	4	0
12	774062	20441	0.0264	4	3
13	756744	20401	0.0270	3	1
14	926563	27207	0.0294	4	0