

**Table 4.** Percentages of loci in different retention classes between *S. cerevisiae* and the real *K. polysporus* genome, and in 100 pseudo-genomes where the tracking of *K. polysporus* single-copy genes was randomized.

Locus class ( <i>K. pol.</i> : <i>S. cer.</i> )	Real genome		Pseudo-genomes			
	Number of loci	Percent	Number of loci		Percent	
			Mean	S.D.	Mean	S.D.
2:2	212	6.52%	209.17	1.81	7.60%	0.87%
2:1	238	7.32%	234.90	1.65	8.53%	0.70%
1:2	221	6.80%	183.27	5.00	6.66%	2.73%
1:1 orthologs	1455	44.74%	1195.72	26.65	43.42%	2.23%
1:1 paralogs	1126	34.62%	930.61	24.61	33.80%	2.64%
Total	3252		2753.67			
Proportion of paralogs among 1:1 loci		44%			44%	1%