

	Centromere linkage				Overall linkage (control)							
	Significant		Not significant		All chromosomes		All chrs., excl. 4 smallest		All chromosomes		All chrs., excl. 4 smallest	
	<i>R</i>	<i>p-val</i>	<i>R</i>	<i>p-val</i>	<i>R</i>	<i>p-val</i>	<i>R</i>	<i>p-val</i>	<i>R</i>	<i>p-val</i>		
This study	0.77	0.01	0.66	0.03	0.74	0.001	0.44	0.15				
SGD	0.70	0.002	0.42	0.17	0.64	0.008	0.05	0.87				
Gerton <i>et al.</i> , 2000	0.78	0.0003	0.74	0.006	0.46	0.08	-0.07	0.82				
Borde <i>et al.</i> , 2004	0.79	0.001	0.92	0.0002	0.25	0.34	0.04	0.89				
Buhler <i>et al.</i> , 2007	0.63	0.04	0.72	0.03	0.33	0.22	0.15	0.64				
Blitzblau <i>et al.</i> , 2007	0.59	0.02	0.67	0.02	0.62	0.01	0.11	0.74				

Figure S5 Pearson correlation coefficients and the corresponding significance p-values were computed between chromosome size and centromere-proximal recombination reported by several published studies, as described in Methods.