

Table 1: The overlap between the hotspots we identified in *M. m. castaneus* and the locations of DSB hotspots in wild-derived strains obtained by Smagulova et al. (2016). The corrected overlap is the number of overlapping hotspots, above the null expectation, over the total.

Strain ID	Sub-species	# DSB Hotspots	# Overlaps	% Overlap Uncorrected	Null Expectation	% Overlap Corrected
13R	<i>domesticus</i>	14744	1202	8.2	1169	0.2
B6	<i>domesticus</i>	19455	1533	7.9	1505	0.1
C3H	<i>domesticus</i>	14635	1399	9.6	1308	0.6
CAST	<i>castaneus</i>	15061	1831	12.2	1221	4.1
MOL	<i>molossinus</i>	15718	1559	9.9	1351	1.3
PWD	<i>musculus</i>	14483	1569	10.8	1205	2.5