

Table S13. Summary statistics for the log of polymorphism/divergence ratio in human-mouse conserved sequences (HMCS) versus unfiltered data.

Data	Min	1 st Quartile	Median	Mean	3 rd Quartile	Max.
HMCS	-3.1350	-1.5040	-0.9808	-1.0220	-0.5108	2.4850
Unfiltered	-3.4340	-1.6090	-1.0990	-1.1040	-0.5878	2.8900