Table S2. The proportions of enhancers that underwent selection based on mouse-human alignments (neutral reference: intron sites)

	Total	Under selection ^a	Positively selected ^b	Under selection ^a /Total	Positively selected ^b /Under selection ^a	Positively selected ^b /Total
HT	2273	946	415	41.61%	43.86%	18.25%
FB	1183	724	79	61.20%	10.91%	6.67%
MB	1186	656	142	55.31%	21.64%	11.97%
LB	2368	1257	274	53.08%	21.79%	11.57%

FB: forebrain; MB: midbrain; LB: limb; HT: heart.

^a A significantly unequal proportion of substituted sites were observed between enhancers and the intron sites of the neighboring gene by Fisher's exact test.

^b Enhancers that underwent selection and had a higher substitution rate than the intron sites of the neighboring gene.