**S6 Table:** Summary of parameters used for the coalescent simulations.

Species pair	Shape	Scale	μ (average) <sup>a</sup>	N <sub>a</sub>	t-split
					(generations)
Human-chimp	16.82	1.7X10 <sup>-10</sup>	2.9X10 <sup>-9</sup>	70000	240000
GERP 10					
Human-chimp	15.68	1.8X10 <sup>-10</sup>	2.8X10 <sup>-9</sup>	71000	280000
GERP 25					
Human-mouse	$4.0X10^3$	5.0X10 <sup>-12</sup>	2.0X10 <sup>-8</sup>	940000	N/A
GERP 10					
Human-mouse	$1.6X10^3$	1.7X10 <sup>-11</sup>	2.7X10 <sup>-8</sup>	375000	N/A
GERP 25					

Note that shape and scale refer to the shape and scale parameters used for the gamma distribution of mutation rates.

<sup>a</sup>We filtered all AT→GC changes between the human and chimp sequences as they could be affected by biased gene conversion. Thus, the mutation rates used here are substantially lower than commonly used values.