

	<b>HAR1</b>	<b>HAR2</b>	<b>HAR3</b>	<b>HAR4</b>	<b>HAR5</b>	<b>Total</b>
<i>Substitution Rates</i>						
<b>Human:Chimp</b>	3.56	12.25	12.93	17.33	10.21	7.24
<b>Human:(Chimp-Mouse)</b>	76.45	50.32	46.08	20.17	9.27	25.87

**Table S5. Comparison of estimated substitution rates in HAR1-HAR5.** Substitution rates are posterior expected value of the number of substitutions using the method described in Siepel *et al.*<sup>38</sup> with the CONS model as prior. The human:chimp ratio is the estimated number of substitutions per site in human compared to chimp. The human:(chimp-mouse) ratio is the estimated number of substitutions per site per million years in human compared to the chimp-mouse phylogeny. Note that because rodent generation times are much shorter than human, we expect the rates in years to be *smaller* in human than in the chimp-mouse if the regions are evolving at the same rate per generation. Branch lengths for scaling rates in millions of years: human=5my, chimp-mouse=150my. Totals computed by concatenating the five elements.