

	HAR1	HAR2	HAR3	HAR4	HAR5
<i>4d sites</i>					
All	$3.3e^{-14}$	$4.7e^{-11}$	$8.3e^{-5}$	$1.2e^{-3}$	$5.5e^{-4}$
Last Band	$2.4e^{-12}$	$1.8e^{-9}$	$4.7e^{-4}$	$4.9e^{-3}$	$4.8e^{-3}$

Table S6. Significance of human acceleration. Wright-Fisher independent sites model p -values for human substitutions compared to the genome-wide human-chimp background rate based on ENCODE region³⁹ 4d sites and the human background rate in 4d sites of ENCODE regions in final chromosome bands. Unadjusted p -values are reported.