

Error and Error Mitigation in Low-Coverage Genomes

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Table S3: Estimates of d_N/d_S for chr22 genes and four primates

branch	full data	high-quality only	SEM
tarsier	0.179	0.164	0.165
mouse lemur	0.173	0.166	0.162
bushbaby	0.189	0.179	0.171
tree shrew	0.135	0.123	0.124
<i>internal branch</i>	0.161	0.142	0.156