

Table S1 : Fossil calibrations

Dataset	Node	Maximum age (My)	Minimum age (My)	References
Birds	Paleognathae / Neognathae	86.5	66	Benton et al. (2009)
	<i>Acanthisitta</i> / other Passeriformes	65	–	Benton et al. (2009)
	Sphenisciformes / Procellariiformes	–	62	Slack et al. 2006
Mammals	<i>Pan</i> / <i>Homo</i>	10	5.7	Benton et al. (2009)
	<i>Pan</i> / <i>Hylobates</i>	33.7	11.2	Benton et al. (2009)
	Cercopithecoidea / Hominoidea	34	23.5	Benton et al. (2009)
	lemuriforms / other Primates	65.8	55.6	Benton et al. (2009)
	Feliformia / Caniformia	65.8	39.7	Benton et al. (2009)
	Suidae / Cetacea	65.8	52.4	Benton et al. (2009)
	Eulipotyphlans / other Laurasiatheria	131.5	62.5	Benton et al. (2009)
	Ochotoma / other Lagomorpha	65.8	48.6	Benton et al. (2009)
	sciromorphs / Caviomorpha +	65.8	55.6	Benton et al. (2009)
	Myomorpha			
	Caviomorpha /	58.9	52.5	Benton et al. (2009)
	Myomorpha			
	<i>Rattus</i> / <i>Mus</i>	14	10.4	Benton et al. (2009)
	Macroscelidea / Proboscidea	131.5	48.4	Benton et al. (2009)
	Afrotheria / other placentalia	131.5	62.5	Benton et al. (2009)

Table S2 : Covariance between Kr/Kc and LHTs using the alternative birds topology of Hackett et al. 2008. The correlations without control (–) or controlling for base composition, . Change in polarity and volume are considered as radical substitutions.

Control	Body mass	Longevity	Sexual maturity
–	0.39 (0.99)**	0.32 (0.91)	0.11 (0.62)
π_C and π_T	0.45 (>0.99)**	0.35 (0.92)	0.23 (0.80)
GC bias, AT/GC skew	0.43 (>0.99)**	0.33 (0.91)	0.23 (0.80)

* indicates pp > 0.95 or < 0.05

** indicates pp < 0.025 or > 0.975

Table S3 : Covariance between Kr/Kc and LHTs excluding ND6. The correlations without control (–) or controlling for base composition. Change in polarity and volume are considered as radical substitutions.

Datasets	Control	Body mass	Longevity	Sexual maturity
–	–	0.35 (>0.99)**	0.42 (>0.99)**	0.63 (>0.99)**
Placentals (All)				
pc and p _T	–	0.30 (>0.99)**	0.44 (>0.99)**	0.51 (0.99)**
GC bias, AT/GC skew	–	0.32 (>0.99)**	0.34 (0.98)**	0.530 (>0.99)**
–	–	0.36 (0.98)**	0.29 (0.89)	0.08 (0.60)
Birds				
pc and p _T	–	0.40 (0.99)**	0.32 (0.90)	0.20 (0.77)
GC bias, AT/GC skew	–	0.37 (0.98)**	0.29 (0.88)	0.19 (0.77)

* indicates pp > 0.95 or < 0.05

** indicates pp < 0.025 or > 0.975

References

- Benton MJ, Donoghue PCJ, Asher RJ. 2009. Calibrating and constraining molecular clocks. *Timetree Life*:35–86.
- Slack KE, Jones CM, Ando T, Harrison GLA, Fordyce RE, Arnason U, Penny D. 2006. Early penguin fossils, plus mitochondrial genomes, calibrate avian evolution. *Mol. Biol. Evol.* 23:1144–1155.