

6. Maximum likelihood bootstrap values for individual partitions within linkage groups

Grouping	Mitochondrial DNA			18s+28s		H3 (codon positions)		EF1- α (codon positions)	
	12s	16s	COII	18s	28s	1 st + 2 nd	3 rd	1 st + 2 nd	3 rd
Palaeoptera	0.24	0.03	0.53	0.32	0.07	0.17	0.21	0.25	0.49
Chiaatomyaria	0.46	0.78	0.32	0.68	0.27	0.48	0.74	0.11	0.14
Metapterygota	0.30	0.19	0.15	0.00	0.66	0.35	0.05	0.64	0.37

Note. All results are shown for our 38-species dataset 2, with corrections for heterogeneous base content.