

Fossil calibrations and comprehensive taxon sampling clarify the timescale of shorebird evolution

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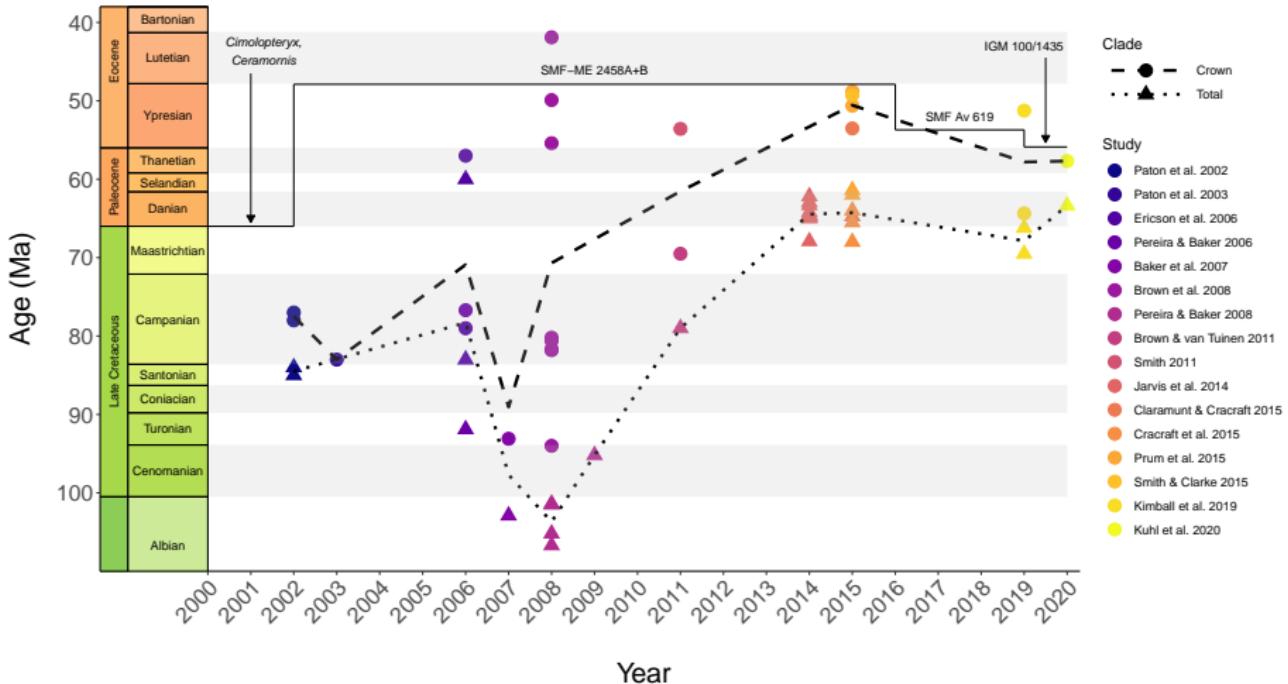
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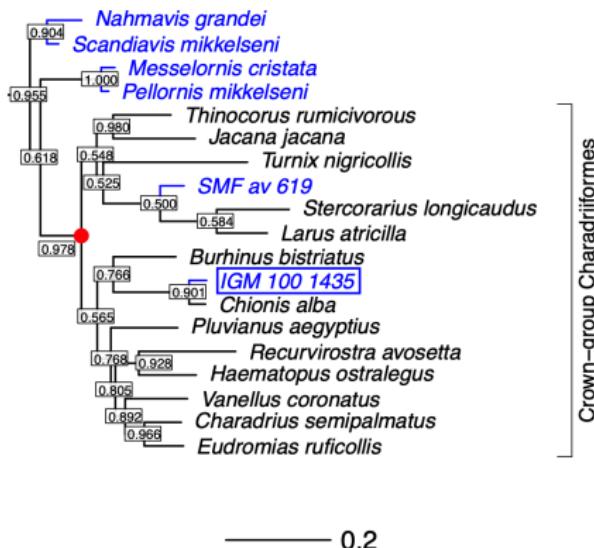
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Previous divergence time estimates

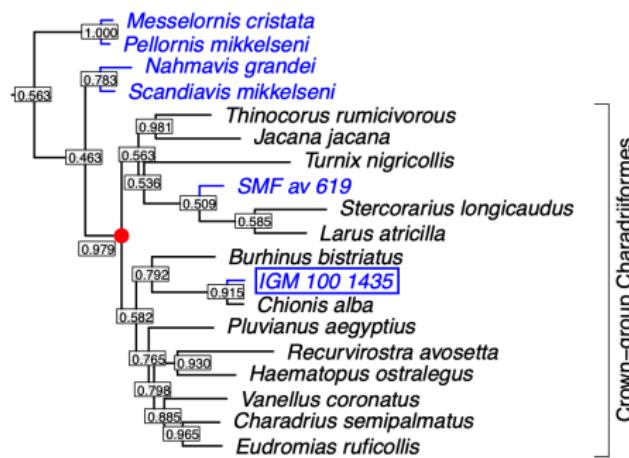




Kimball et al. (2019) backbone



Reddy et al. (2017) backbone

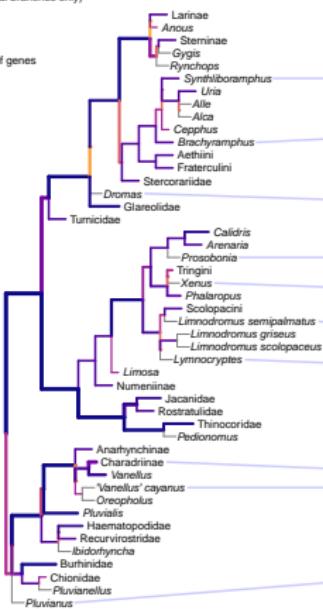


a) ASTRAL

Coalescent units
(non-terminal branches only)

Effective # of genes
— 2
— 4
— 6
— 8

Local PP
0.75
0.50
0.25

**b) RAxML**

Expected substitutions
per site

Locus coverage

Bootstrap

Alle

Uria

Synthliboramphus

Cephus

Brachyramphus

Aethini

Fraternalini

Stercorariidae

Glareolidae

Dromas

Turonicidae

Calidris

Arenaria

Prosobonia

Tringini

Xenus

Phalaropus

Scolopacini

Limnodromus semipalmatus

Limnodromus griseus

Limnodromus scolopaceus

Limnodromus semipalmatus

Lymnocryptes

Numeninae

Jacanidae

Rostratulidae

Threskiornithidae

Pedionomus

Anarhynchinae

Charadrinae

Vanellus

'Vanellus' cayanus

Oreopholus

Haematopodidae

Recurvirostridae

Ibdorhyncha

Burhinidae

Chionidae

Pluvianellus

c) ExaBayes

Expected substitutions
per site

Locus coverage

Posterior probability

Alle

Uria

Synthliboramphus

Cephus

Brachyramphus

Aethini

Fraternalini

Stercorariidae

Glareolidae

Dromas

Turonicidae

Calidris

Arenaria

Prosobonia

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Charadrinae

Vanellus

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Oreopholus

Haematopodidae

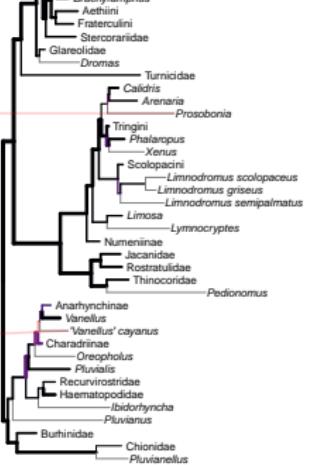
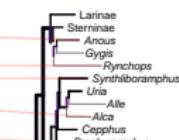
Recurvirostridae

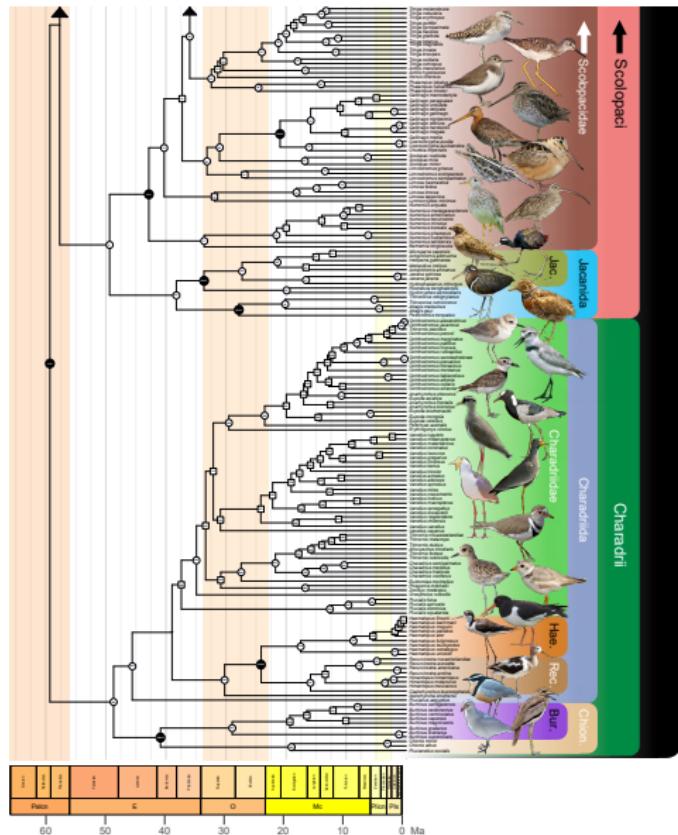
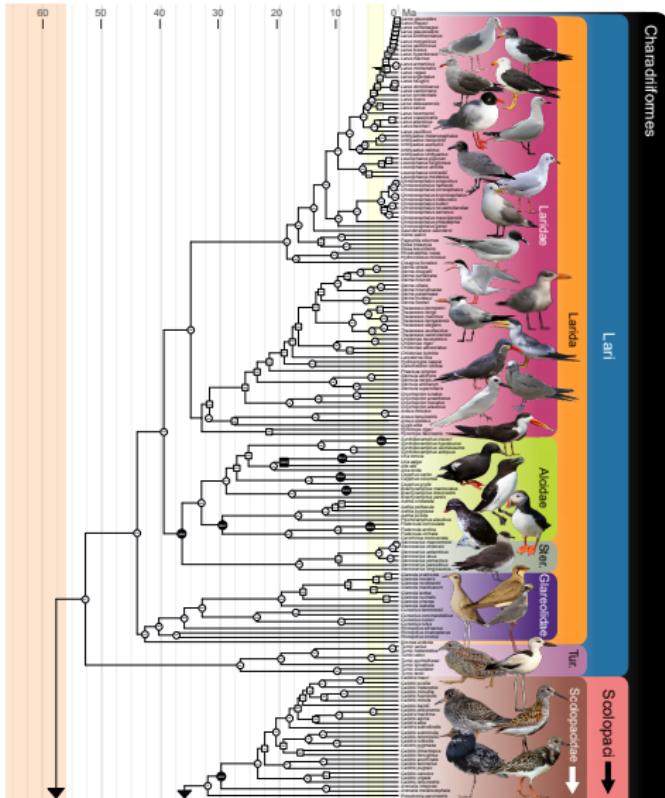
Ibdorhyncha

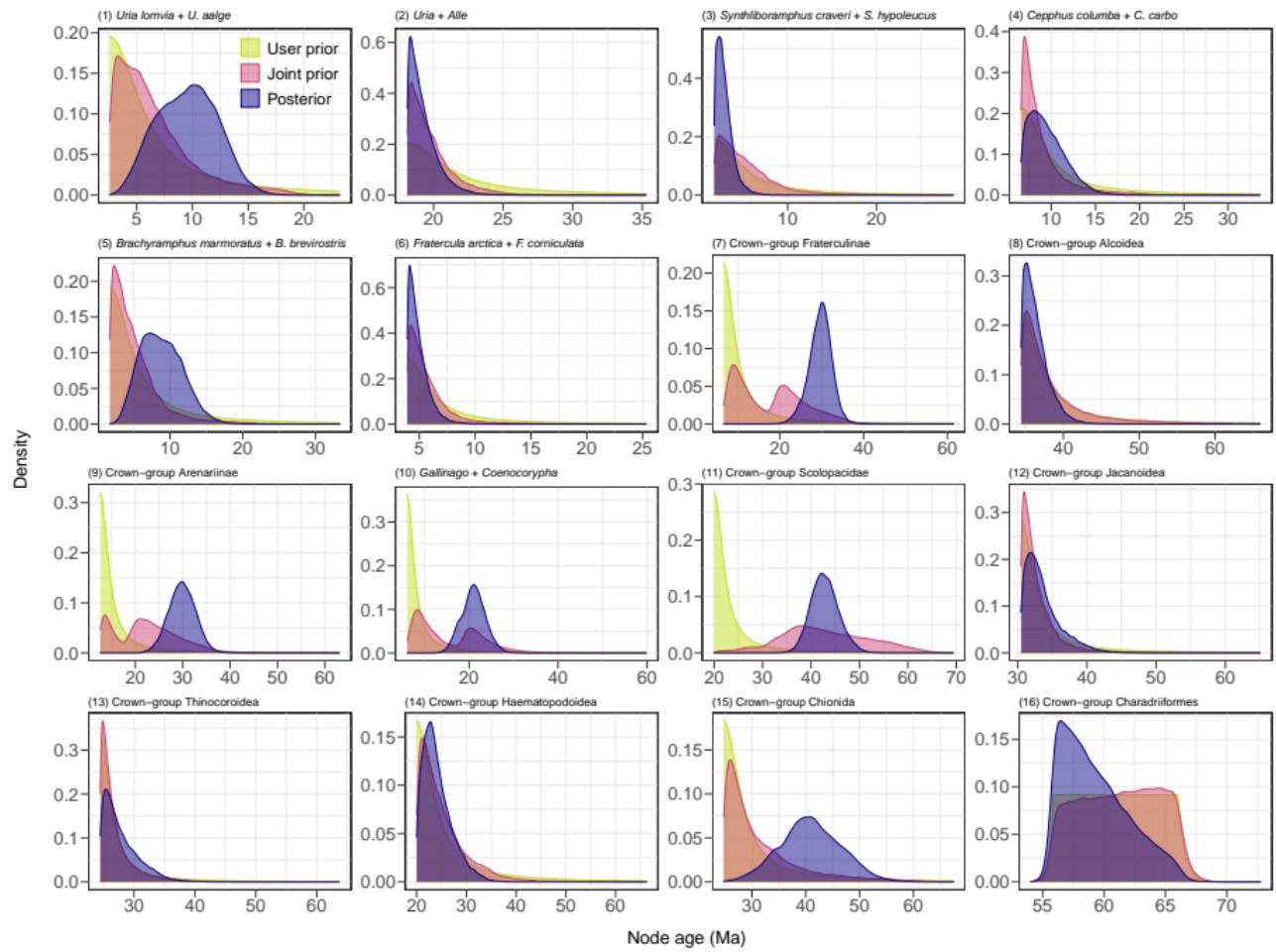
Burhinidae

Chionidae

Pluvianellus







Relaxed-clock model comparisons

Dataset	Model	Bayes factor	Posterior probability
19 species	Autocorrelated rates	72.64	0.986
	Uncorrelated rates	1	0.014
40 species	Autocorrelated rates	1	0.130
	Uncorrelated rates	6.67	0.870

