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Figure 1. Stylized DateLife workflow. This shows the general workflows and analyses that can be performed with DateLife, via the R package or through the website. Details on the functions involved on each workflow are shown in **datelife**'s R package vignette.

Figure 2. Computation time of input processing and search across **datelife**s chronogram database.

Figure 3. Lineage through time (LTT) plots of source chronograms containing all or a subset of species from the bird family Fringillidae of true finches. Arrows indicate maximum age of each chronogram. Numbers reference to chronograms' original publications 1: Barker et al. (2012), 2: Barker et al. (2015), 3: Burns et al. (2014), 4: Claramunt and Cracraft (2015), 5: Gibb et al. (2015), 6: Hedges et al. (2015), 7: Hooper and Price (2017), 8: Jetz et al. (2012), 9: Price et al. (2014).

Figure 4. LTT plots of median and Supermatrix Distance Method (SDM) chronograms summarizing information from source chronograms found for the Fringillidae. Arrows indicate maximum age.

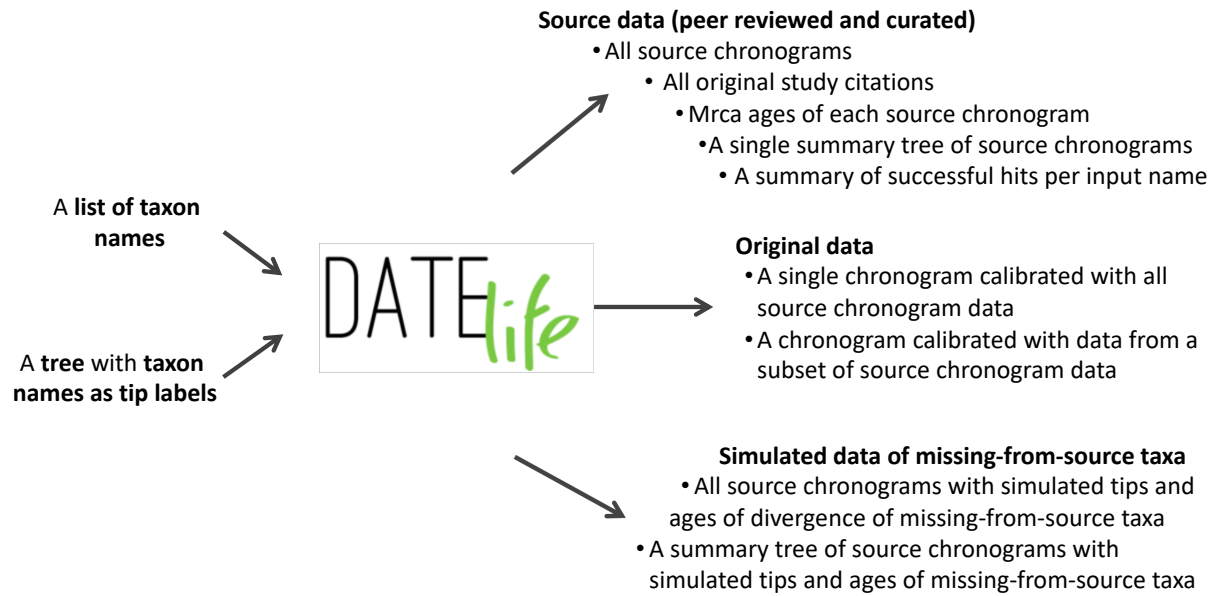


Figure 1

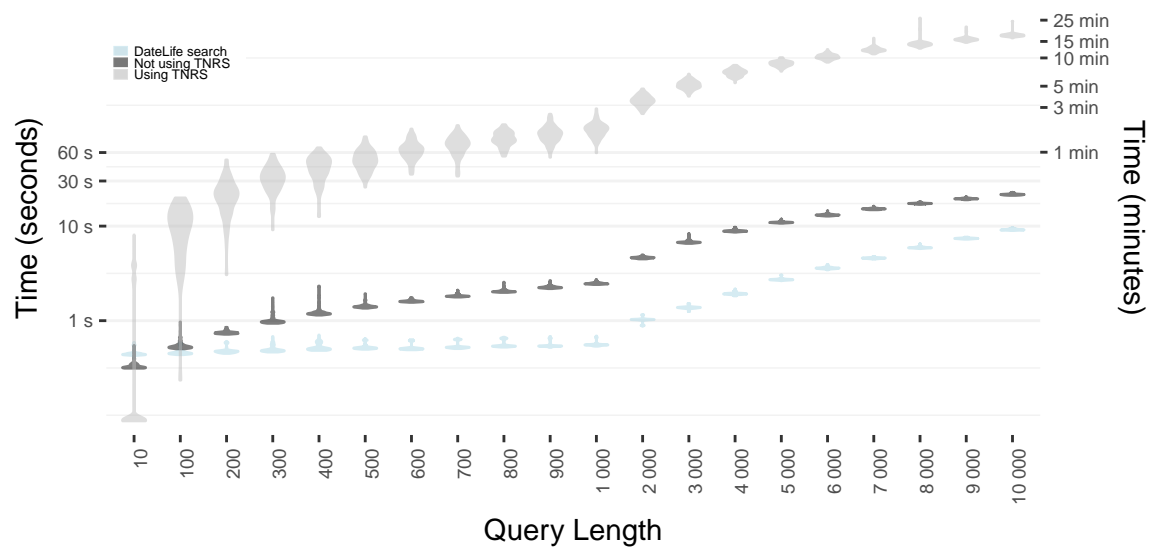


Figure 2

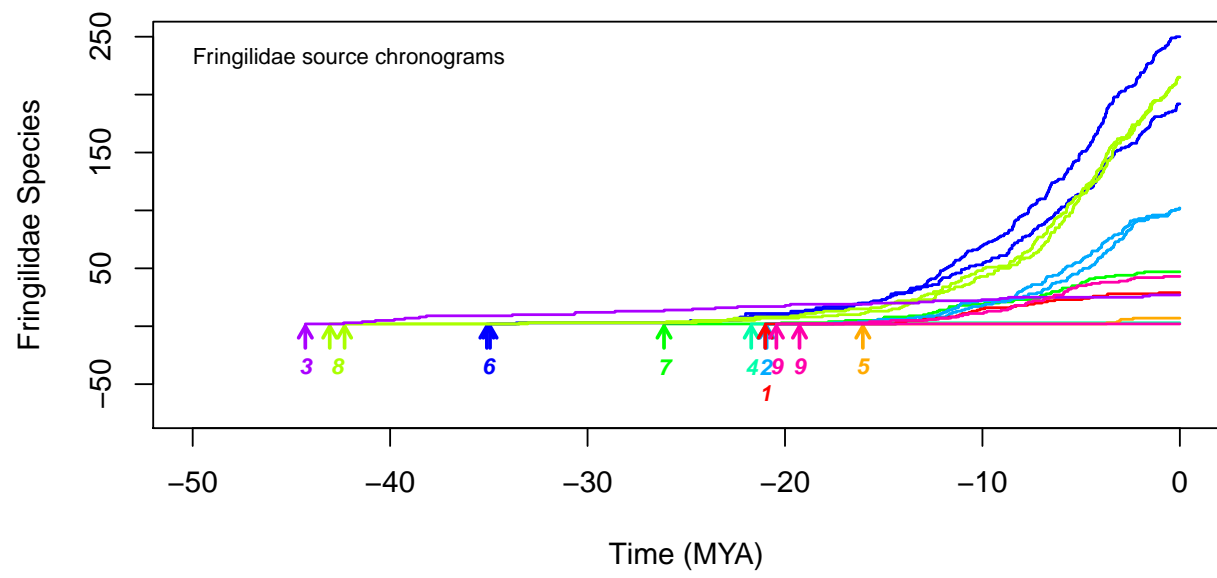


Figure 3

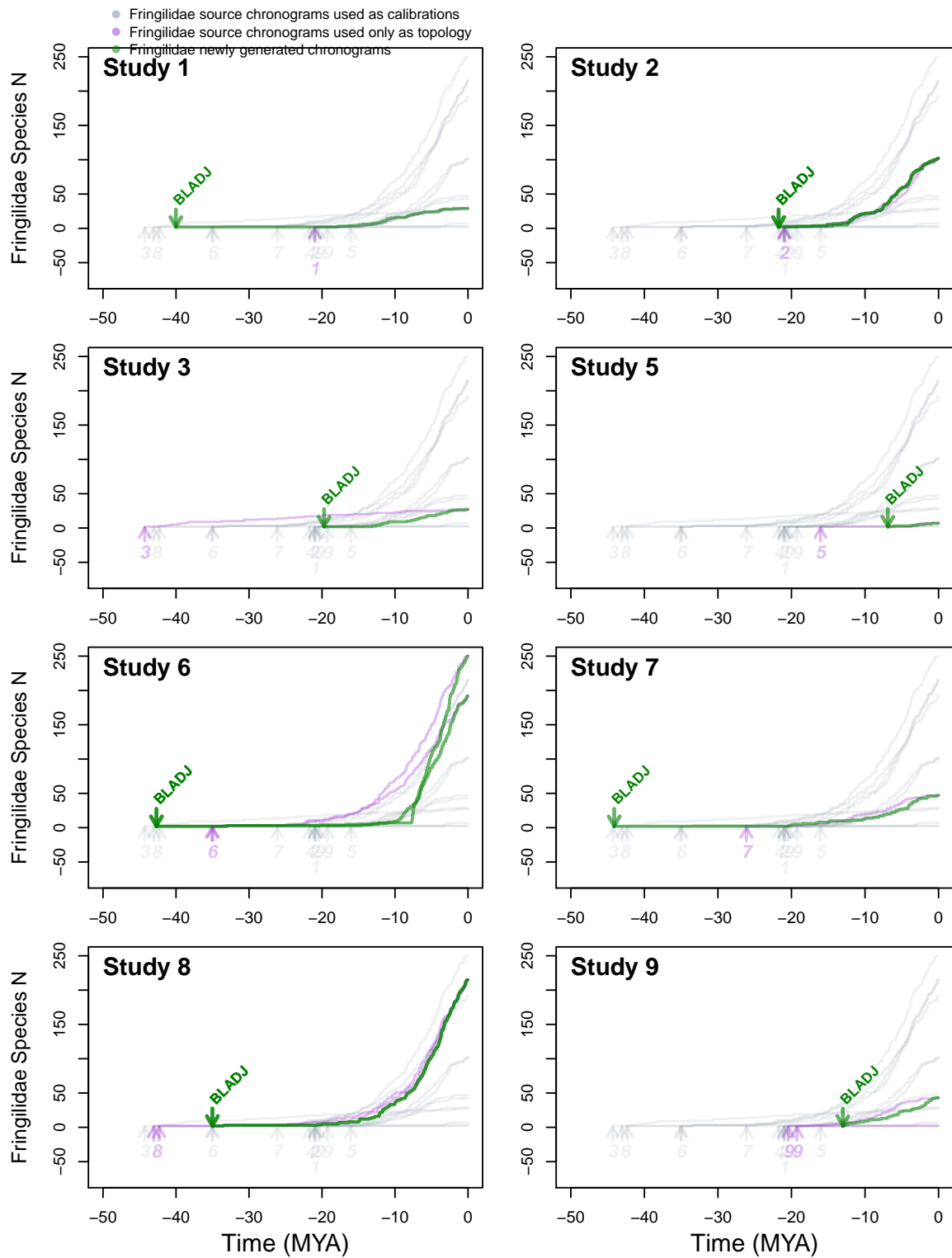


Figure 4

13 Barker F.K., Burns K.J., Klicka J., Lanyon S.M., Lovette I.J. 2012. Going to extremes: Contrasting rates of
14 diversification in a recent radiation of new world passerine birds. *Systematic biology*. 62:298–320.

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