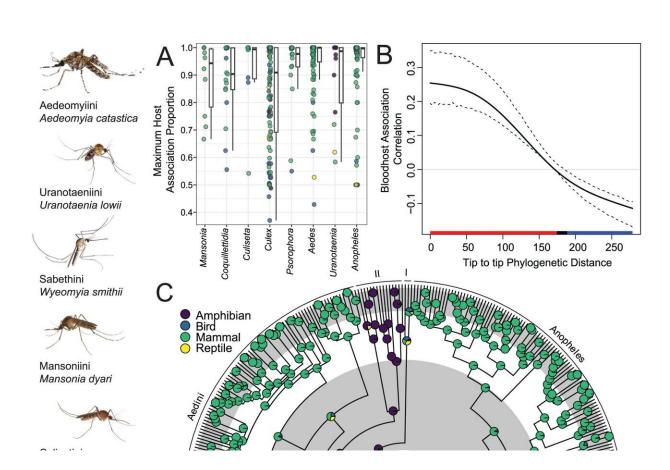


Study elucidates evolution of mosquitoes and their hosts



October 18 2023, by Mick Kulikowski

Macroevolutionary analyses of host associations for blood-feeding in mosquitoes based on 293,308 bloodmeal records from 422 different mosquito species. A The maximum host association score among several genera indicating that many genera have a high degree of class-level host association, although there are exceptions. Each point is a species in a given genus. Boxplots for each genus show median max host association values as black bars, along with interquartile ranges as boxes. Whiskers are drawn to +/-1.5 times the interquartile range. **B** Vertebrate host association contains significant phylogenetic signal as estimated



by a phylogenetic Mantel test (shown) and multivariate Blomberg's K = 0.06, P

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