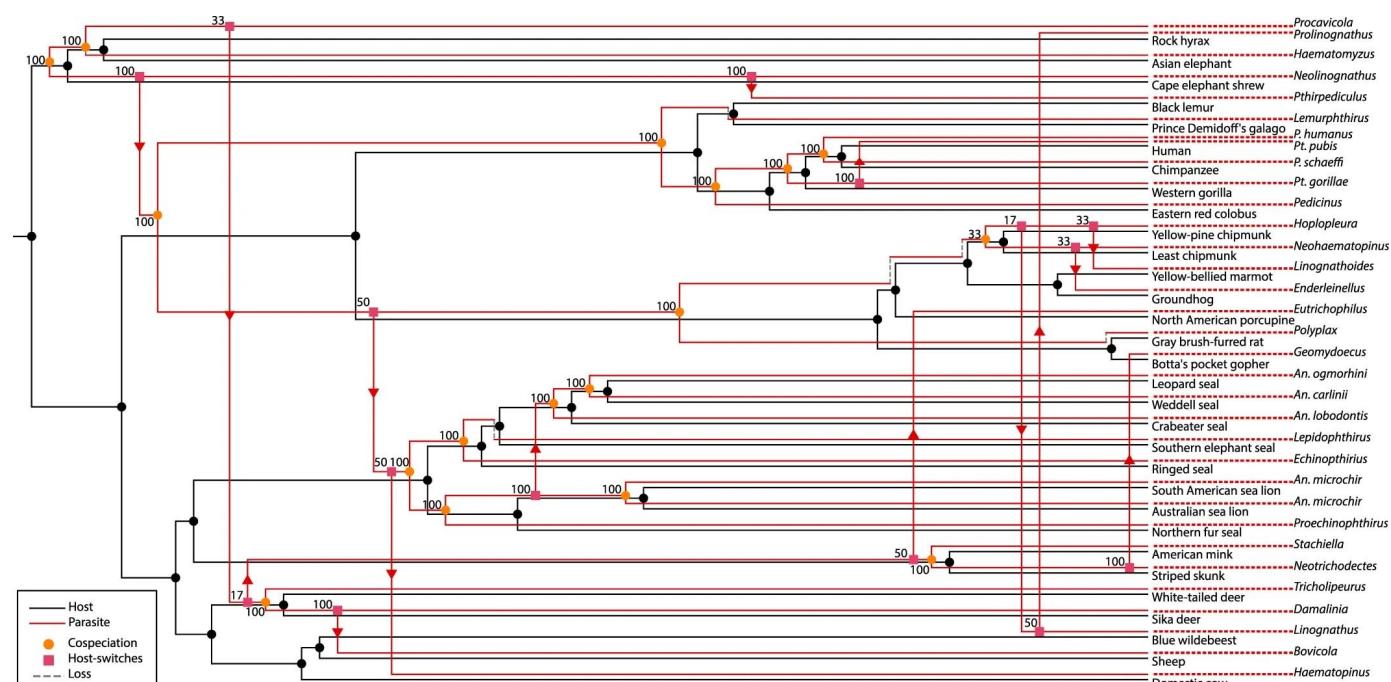




# Extended Data Fig. 1: Summary of cophylogenetic reconstruction of optimal MPRs from eMPReSS comparison (cost scheme duplication: 1, sorting: 1, and host-switching: 2) of the louse (concatenated) tree with the mammal host tree.

From: [Phylogenomics reveals the origin of mammal lice out of Afrotheria](#)



Arrows indicate direction of host-switches. Numbers associated with events are the percentage of MPRs with that event.

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