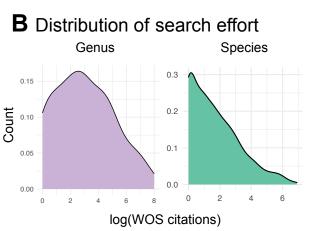
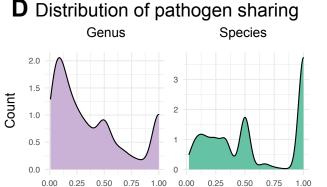
**A** Data collection process (workflow) **B** Data summary (counts)



C Distribution of phylogenetic distance



**E** Distribution of PSR

**F** Host search effort vs. parasite species richness

% Shared Pathogens

