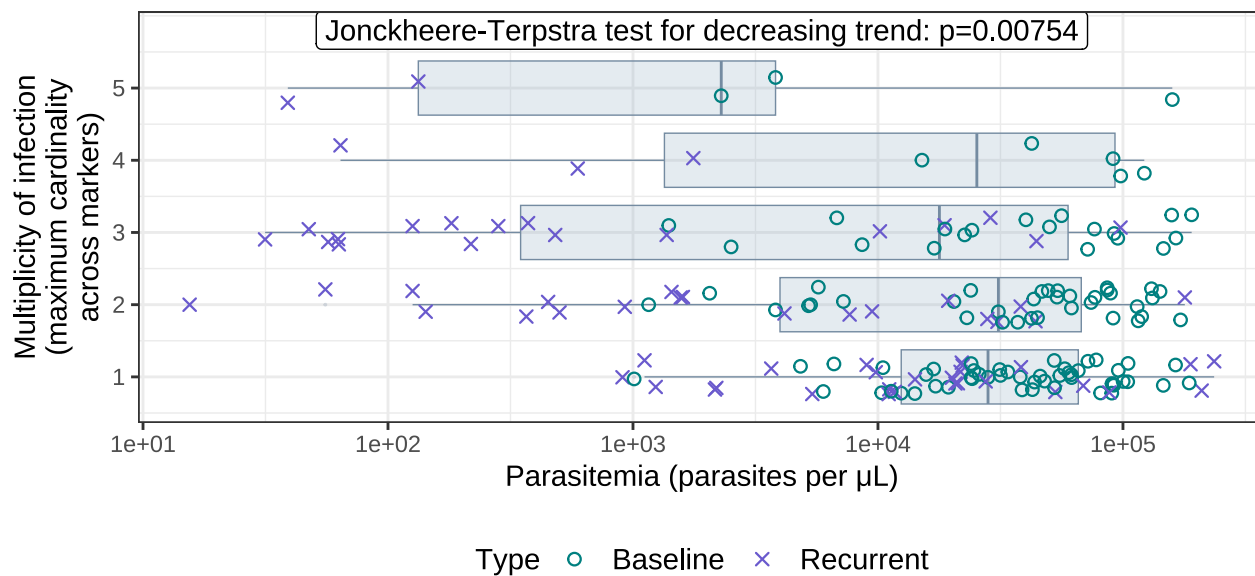
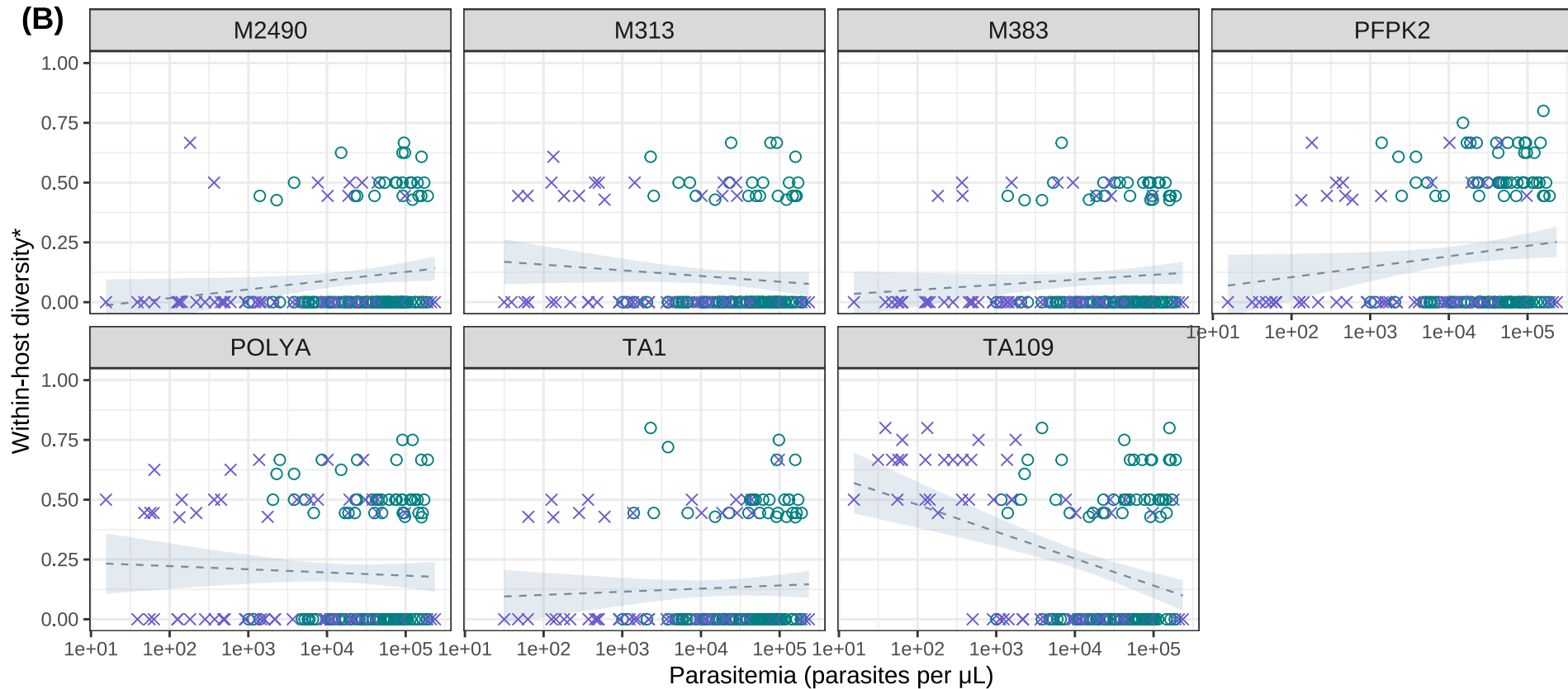


(A)**(B)**

*Expected value of $1 - \text{Nei's gene identity metric}$, assuming M equiprequent clones (where M is the maximum cardinality across all loci) exhibit A distinct alleles (where A is the locus cardinality)