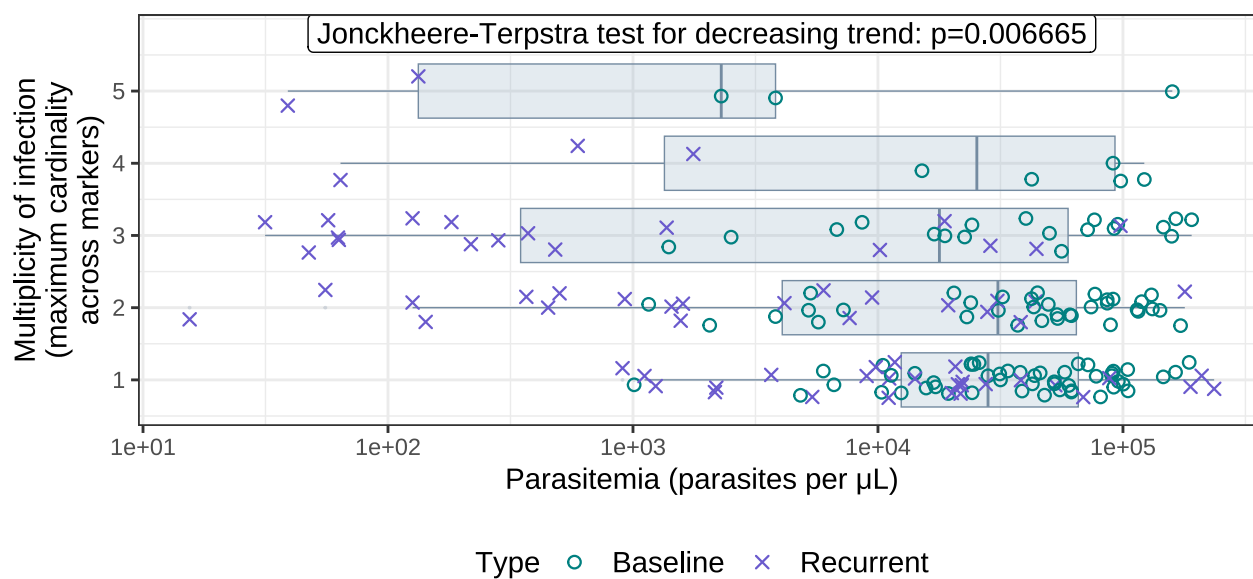
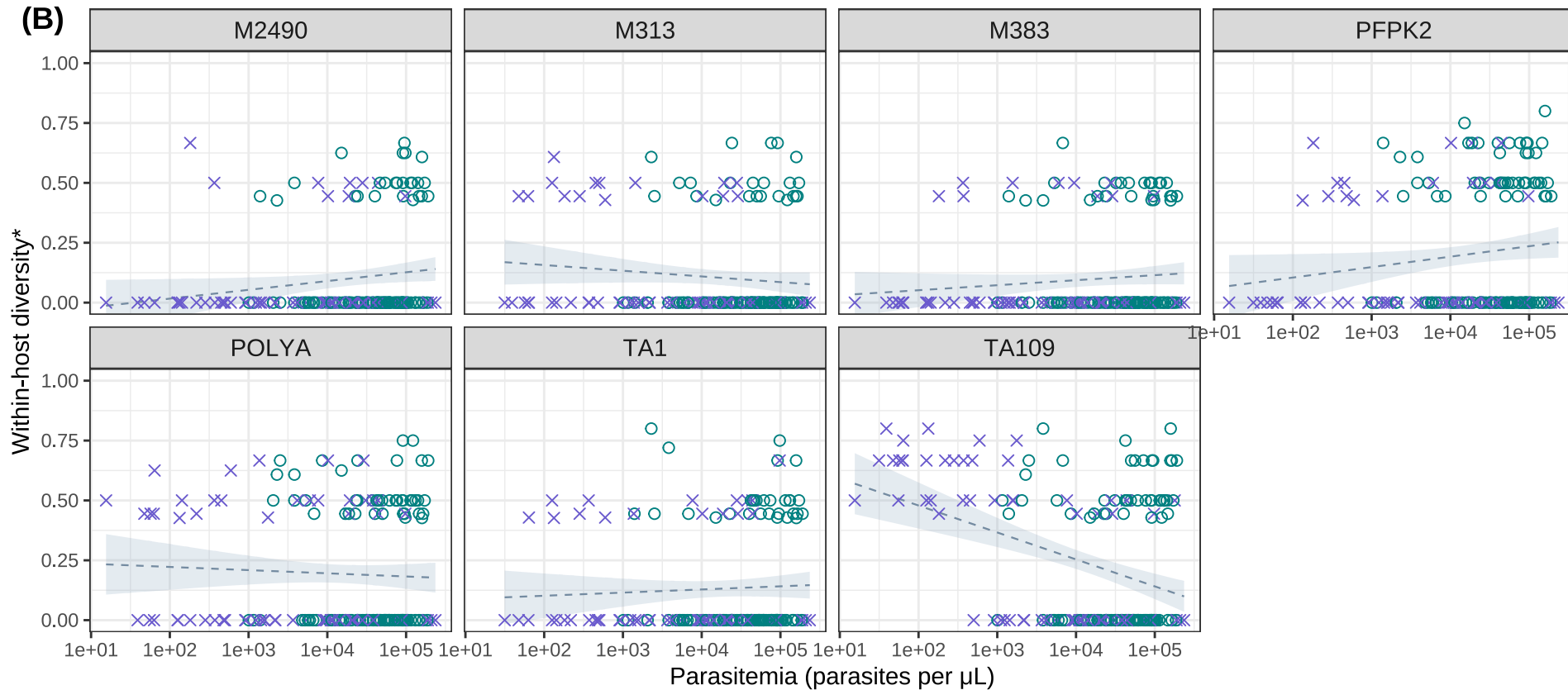


**(A)****(B)**

\*Expected value of 1-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)