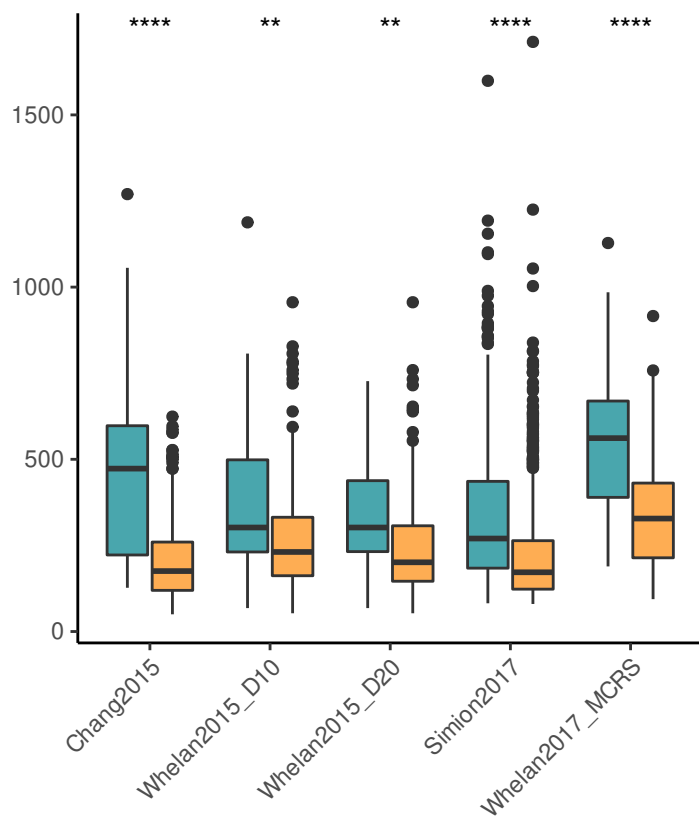
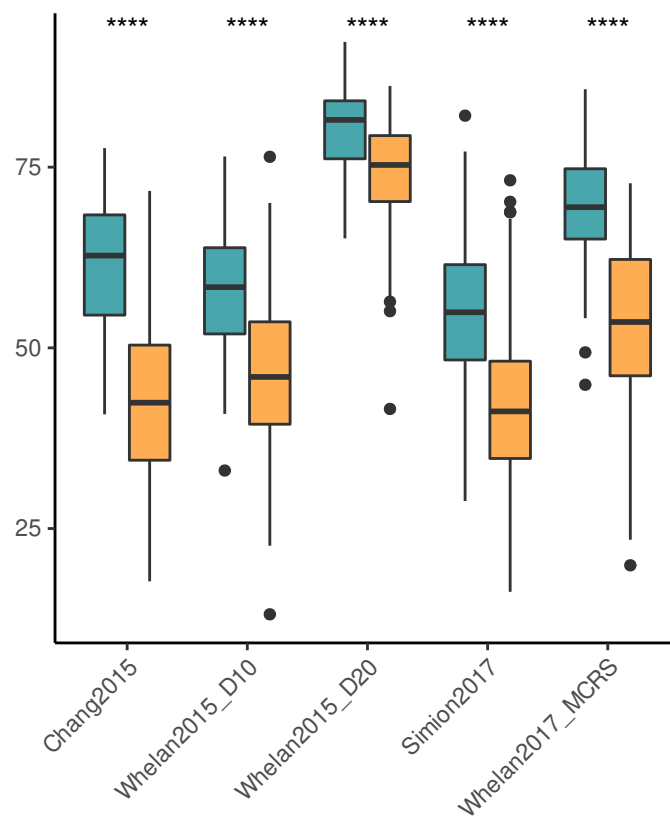


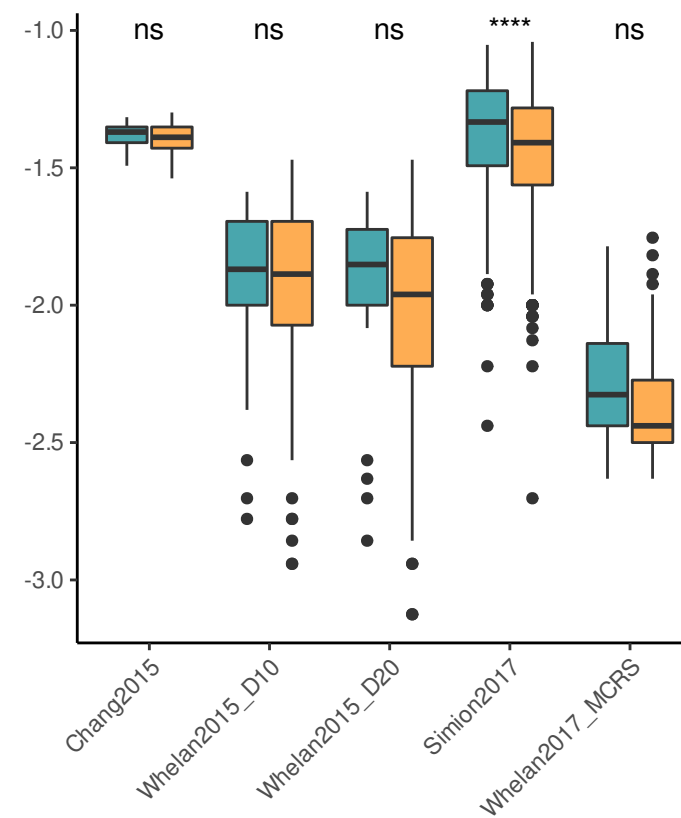
Alignment length



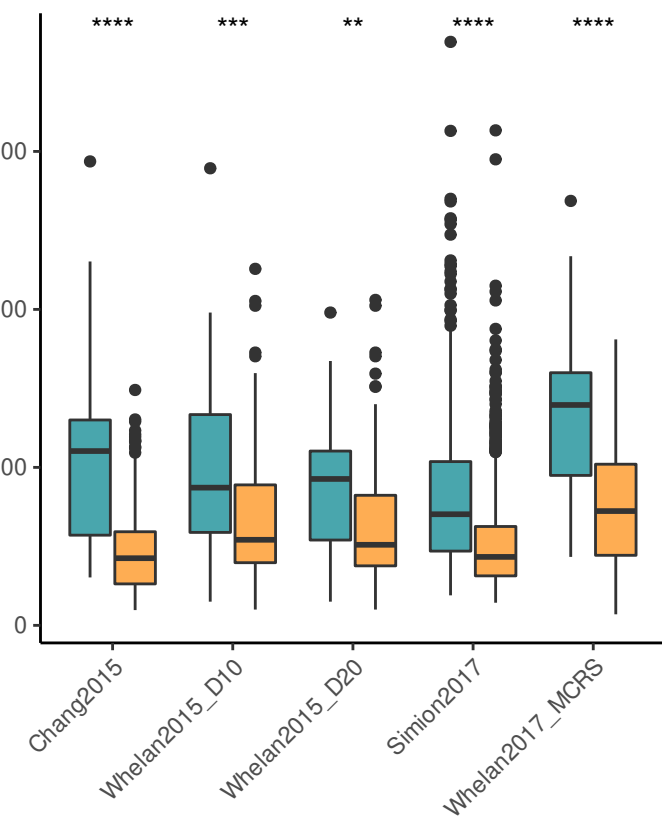
Mean bipartition support



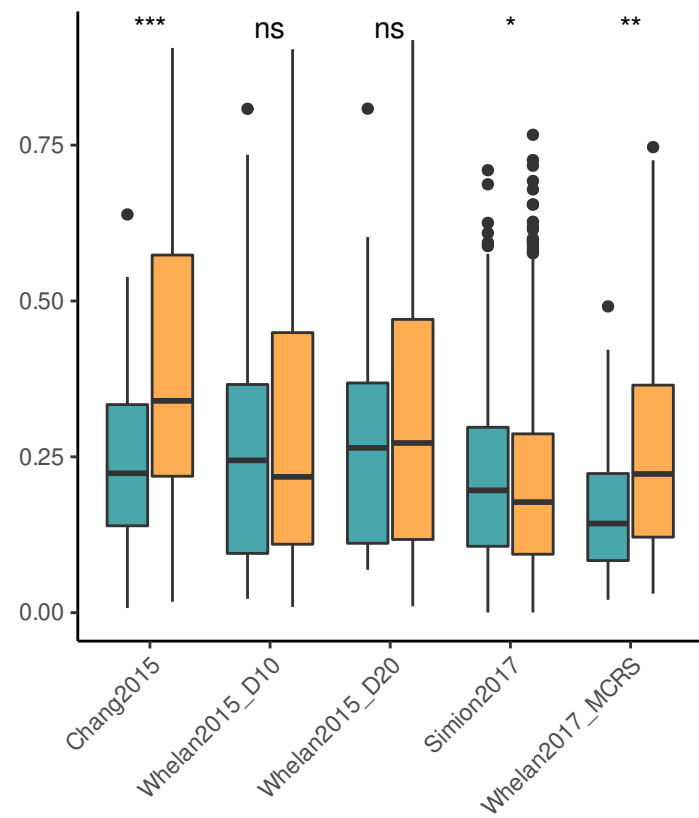
Mean long branch score



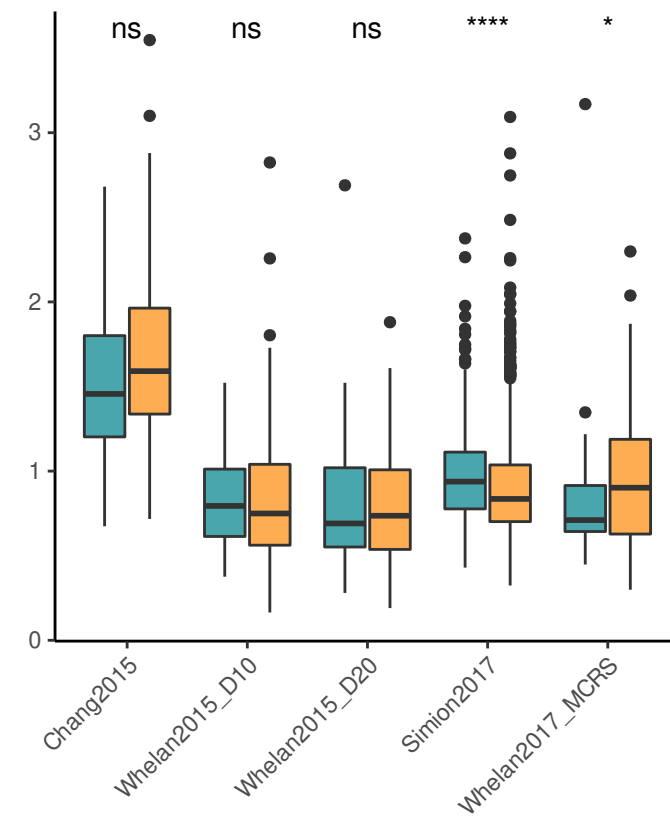
Parsimony informative sites



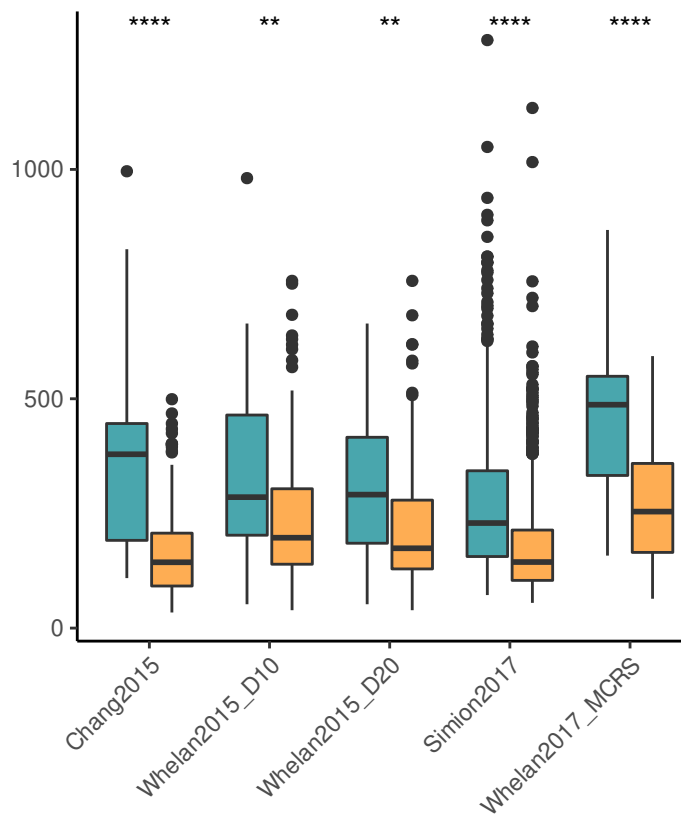
Saturation



Treeness by RCV



Variable sites



Passed clan_check filter
Failed clan_check filter