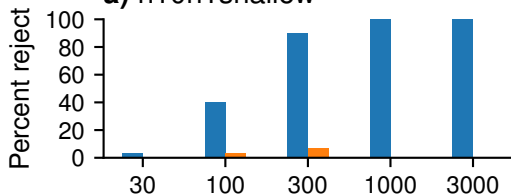
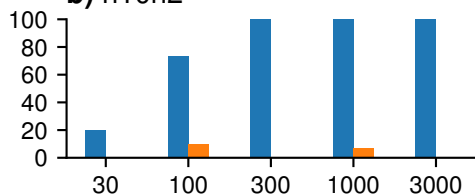


# Percent replicates TICR correctly rejects major tree

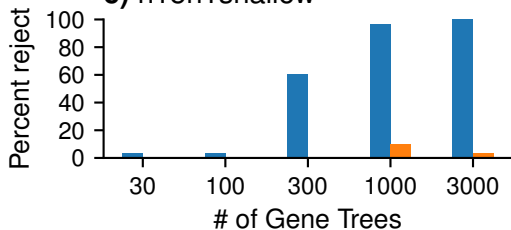
**a) n10h1shallow**



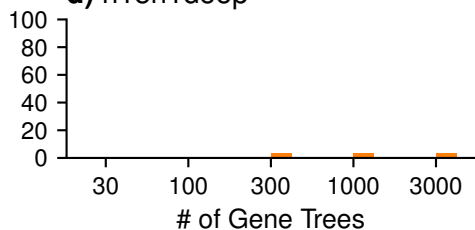
**b) n10h2**



**c) n15h1shallow**



**d) n15h1deep**



True gene trees      Estimated gene trees