Percent replicates TICR correctly rejects major tree a) n5h2 **b)** n10h2 c) n10h1shallow 100 · 100 -Percent reject True gene trees Estimated gene trees 300 1000 3000 300 1000 3000 300 1000 3000 d) n15h1shallow e) n15h1intermediate f) n15h1deep **g)** n15h3 100 -100 -Percent reject 40 -1000 3000 300 1000 3000 300 1000 3000 300 1000 3000 # of Gene Trees # of Gene Trees # of Gene Trees # of Gene Trees