## N10R1 propQuartets = 0.5 propQuartets = 0.7 propQuartets = 0.9 propQuartets = 1 6probQR = Average Hardwired Cluster Distance probQR = 0.5**SNaQ Version** v1.0 v1.1 6 probQR = 0 -3001000 3000 3001000 3000 3001000 3000 3001000 3000 Number of gene trees