N20R1 propQuartets = 0.5 propQuartets = 0.7 propQuartets = 0.9 propQuartets = 1 6 probQR = Average Hardwired Cluster Distance probQR **SNaQ Version** v1.0 = 0.5 v1.1 probQR 0 -3001000 3001000 3000 3001000 3000 3001000 3000 3000 Number of gene trees