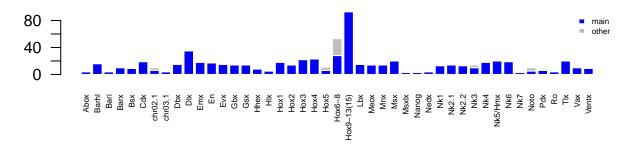


genes in best OG

n(best # 0.8 nlbs 14 (best 20)(5-1.000) n(best = all) = 38 (p=0.921 | r=1.000) n(best < all) = 7 (p=1.000 | r=1.000)

number of reference genes in main and other OGs



num OGs per refOG

