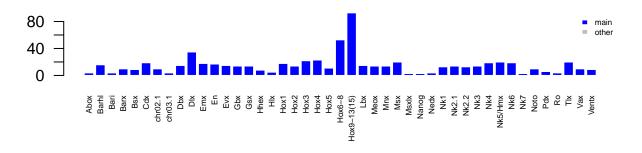


genes in best OG

n(best #=0.8 nlbs #5 (bef60)(c=0.999) n(best = all) = 44 (p=0.453 | r=1.000) n(best < all) = 1 (p=1.000 | r=0.957)

number of reference genes in main and other OGs



num OGs per refOG

