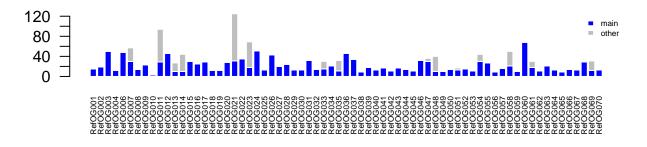


genes in best OG

n(best # 0.8 all) = \$1 (be\$100 (c=0.993) n(best = all) = 47 (p=1.000 | r=1.000) n(best < all) = 22 (p=0.994 | r=0.588)

number of reference genes in main and other OGs



num OGs per refOG

