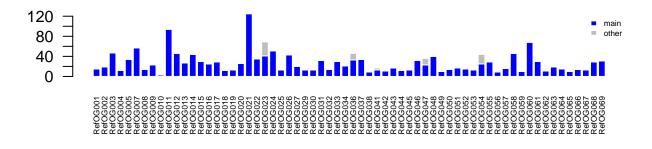


genes in best OG

n(best # 6.8 all) = 56 (p=0.857 | r=0.996) n(best = all) = 56 (p=0.857 | r=0.996) n(best < all) = 8 (p=0.818 | r=1.000)

number of reference genes in main and other OGs



num OGs per refOG

