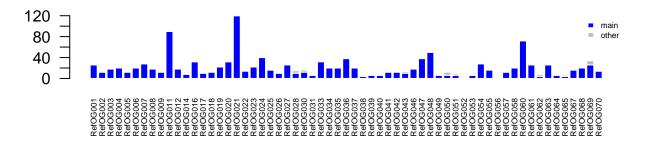


genes in best OG

n(best # 0.8 albs #7 (be115)(c=0.520) n(best = all) = 45 (p=0.911) r=0.513) n(best < all) = 16 (p=0.980 | r=0.479)

number of reference genes in main and other OGs



num OGs per refOG

