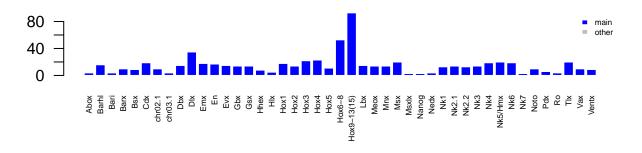


genes in best OG

n(best # 6.8 Albs 15 (bef20)(5-1.000) n(best = all) = 44 (p=0.428 | r=1.000) n(best < all) = 1 (p=0.084 | r=1.000)

number of reference genes in main and other OGs



num OGs per refOG

