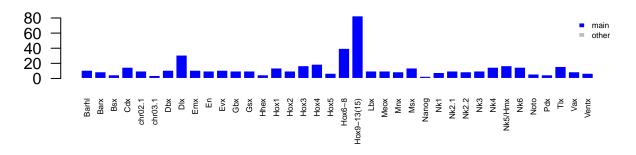


genes in best OG

n(best # 6.8 Albs 3A (best 30)(c=1.000) n(best = all) = 38 (p=0.630 | r=1.000) n(best < all) = 0 (p=NaN | r=NaN)

number of reference genes in main and other OGs



num OGs per refOG

