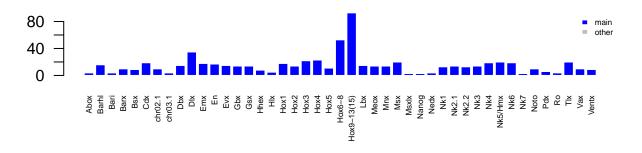


# genes in best OG

n(best # 6.8 Albs 15 (best 50)(5-1.000) n(best = all) = 44 (p=0.656 | r=1.000) n(best < all) = 1 (p=0.523 | r=1.000)

## number of reference genes in main and other OGs



## num OGs per refOG

