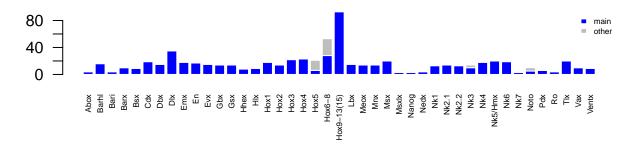


# genes in best OG

n(best #=0.8 albs 34 (be440)(c=0.998) n(best = all) = 37 (p=0.946 | r=1.000) n(best < all) = 6 (p=1.000 | r=0.655)

## number of reference genes in main and other OGs



## num OGs per refOG

