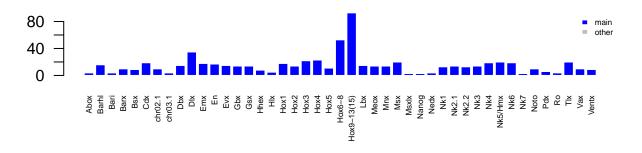


# genes in best OG

n(best #=0.8 albs #5 (be\$97)(5=0.999) n(best = all) = 44 (p=0.588 | r=1.000) n(best < all) = 1 (p=1.000 | r=0.957)

## number of reference genes in main and other OGs



## num OGs per refOG

