

Scale
chr1:
--->

0 bases | hg38_2

G

W

variants derived from NyuWa project

NyuWa variants

blocks of chr22

Blocks_chr22

Recombination map of 200 individuals

Recombination Map

NCBI RefSeq genes, curated subset (NM_*, NR_*, NP_* or YP_*)

MTOR

