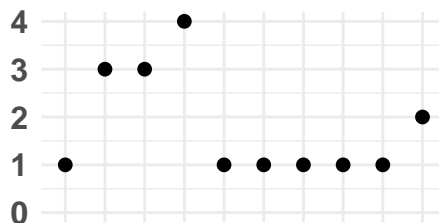
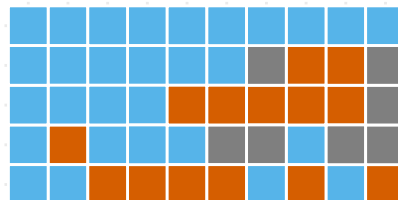


HD AOO GWAS hits



genetic_evidence
cross_repeat_expansion
hd_exome_support
hpsc_msn_evidence
omim_phenotypes



Favorable
Unfavorable
Unfavorable.25

knockdown_beneficial
druggable_any
pli_tolerance
rna_tissue_specificity
interacting_partners



Favorable
Unfavorable
Unfavorable.25

PMS1
LIG1
MSH3
FAN1
PMS2
RRM2B
TCERG1
MLH1
CCDC82
HTT