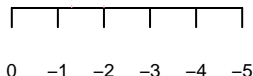


## PBStop domain

CaMKII_AD	p = 0.00281 n = 1
DUF2358	p = 0.00281 n = 1
DUF4414	p = 0.00281 n = 1
IFT43	p = 0.00281 n = 1
IMS_HHH	p = 0.00281 n = 1
MCRS_N	p = 0.00281 n = 1
RBM39linker	p = 0.00281 n = 1
REV1_C	p = 0.00281 n = 1
SIP1	p = 0.00281 n = 1
IMS	p = 0.00841 n = 1
IMS_C	p = 0.00841 n = 1
LysM	p = 0.00841 n = 1
Myosin_TH1	p = 0.00841 n = 1
S-methyl_trans	p = 0.00841 n = 1
Tim44	p = 0.00841 n = 1
Bioplerin_H	p = 0.0112 n = 1
CIDE-N	p = 0.0112 n = 1
Methyltransf_16	p = 0.0112 n = 1
zf-Tim10_DDP	p = 0.0112 n = 1
BRCT_2	p = 0.014 n = 1
DFRP_C	p = 0.014 n = 1
IRS	p = 0.014 n = 1
adh_short	p = 0.0148 n = 2
Hairy_orange	p = 0.0167 n = 1
IBR	p = 0.0195 n = 1
RWD	p = 0.0195 n = 1
Ribosomal_L7Ae	p = 0.0223 n = 1
Amidase_2	p = 0.025 n = 1
Cyclin_C	p = 0.025 n = 1
CD36	p = 0.0359 n = 1



log(p)

n = 39 genes | p adjust BH FALSE