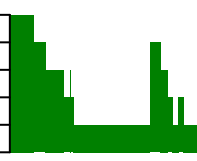


5
4
3
2
1
0



0 50 150 250



27%



TCGA.A2.A4S0.01A(26



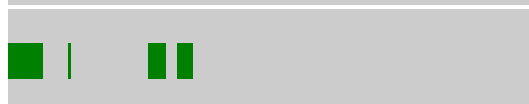
17%



TCGA.PE.A5DE.01A(7



14%



TCGA.A1.A0SK.01A(8



13%



TCGA.A1.A0SI.01A(93



10%



TCGA.D8.A27M.01A(9



ER

positive
negative
indeterminate

