

Motif	Enrichment score
vfl_SANGER_5_FBgn0259789	12.759135723067
CAGGTAG	11.1380836470664
PF0076.1-CAGGTA	11.0717022210421
vfl_SOLEXA_5_FBgn0259789	10.2502212575165
CASGTAR	9.74366411671557
TACCTGC	8.85525369244997
MA0340.1-MOT3	4.24295891331343
GAGAGAG	3.46645717797162
CTCTCTC	3.46645717797162
M00043-I-DL_01	3.35453118268841
dl_FlyReg_FBgn0000462	3.24723752294856
dl	3.24723752294856
RAAMGGRTTA-Kruppel	3.24664910672761
M00774-V-NFKB_Q6_01	3.19416468733339
Gsc Cell_FBgn0010323	3.17917161127211
bcd_FlyReg_FBgn0000166	3.13986886898448
bcd	3.13986886898448
Trl_FlyReg_FBgn0013263	3.09551843852693
Trl	3.09551843852693
MA0205.1-Trl	3.09551843852693
M00723-I-GAGAFACITOR_Q6	3.04762020438423
Oe SOLEXA_FBgn0004102	3.00892895346348
Ptx1 SOLEXA_FBgn0020912	2.98415778431874
TIFDMEM0000009	2.93688834790237
CGTGNGAA	2.93685373518348
Bcd SOLEXA_FBgn0000166	2.93221563085364
MA0022.1-dl_1	2.92012425439175
M01111-V-RBPJK_Q4	2.9091693288664
M00412-V-AREB6_01	2.86589189269411
Bcd Cell_FBgn0000166	2.85202949878289
AGAGAGCG	2.83936124367301
Kr FlyReg_FBgn0001325	2.79895089438123
Kr	2.79895089438123
PF0086.1-GGGNNTTTC	2.79666645493519
MA0306.1-GIS1	2.7539658974109
MA0372.1-RPH1	2.73781906405363
AGAGAGAG	2.73031964162975
M01112-V-RBPJK_01	2.65071615699347
Kr NAR_FBgn0001325	2.60858670933069
M00234-I-SUH_01	2.59301098583493
MA0023.1-dl_2	2.58319251124613
Gsc SOLEXA_FBgn0010323	2.5705069497768
M00935-V-NFAT_Q4_01	2.56416705343538
PH0035.1-Gsc	2.53477508632023

Logos

Recovery curves

