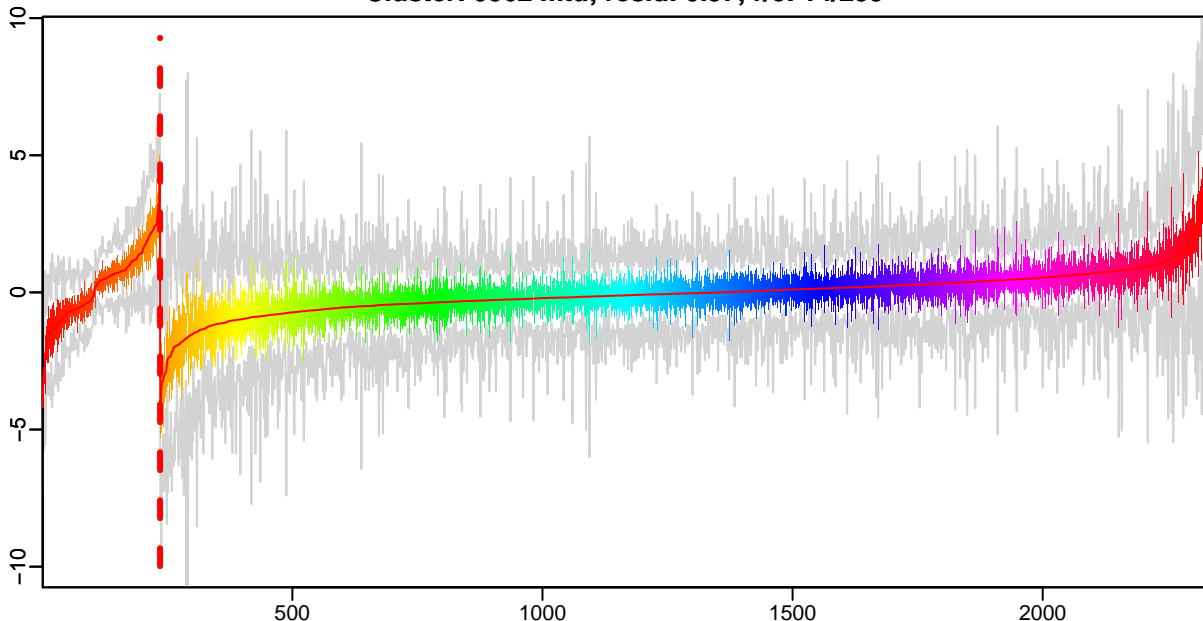
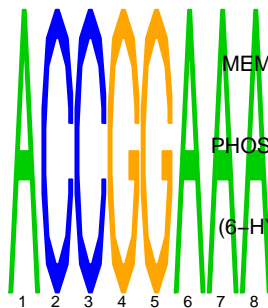
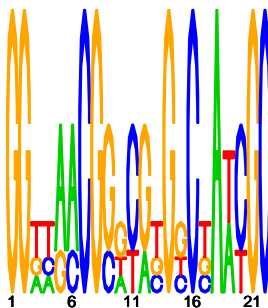


Cluster: 0562 mtu; resid: 0.57; r/c: 14/235

13 Oct 25 14:25:18 iter=2000
cMonkey Version 4.9.8 mtu



upstream meme PSSM #1; e=1.5e+00 upstream meme PSSM #2; e=6.4e+03



log10(P) upstream meme

log10(P.clust)=-6.46; 14 seqs; 4 uniq

MEMBRANE-BOUND PROTEASE FTSH (CELL DIVISION PROTEIN): RV3610C

-7.10 GTP cyclohydrolase I: RV3609C

PHOSPHORYLASE 1 (DIHYDROPTEROATE DIPHOSPHORYLASE 1): RV3608C

PROBABLE DIHYDRONEOPTERIN ALDOLASE FOLB (DHNA): RV3607C

(6-HYDROXYMETHYL-7,8-DIHYDROPTERIN DIPHOSPHOKINASE): RV3606C

-7.10 PROBABLE CONSERVED SECRETED PROTEIN: RV3605C

-7.10 hypothetical protein: RV2097C

-7.10 hypothetical protein: RV2096C

-6.50 hypothetical protein: RV3627C

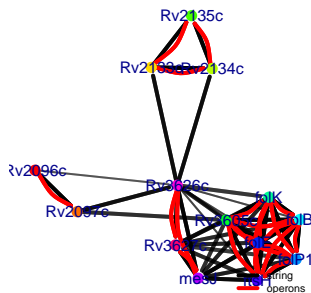
-6.50 hypothetical protein: RV3626C

-6.50 POSSIBLE CELL CYCLE PROTEIN MESJ: RV3625C

-4.70 hypothetical protein: RV2135C

-4.70 hypothetical protein: RV2134C

-4.70 hypothetical protein: RV2133C



-200 -100 -1