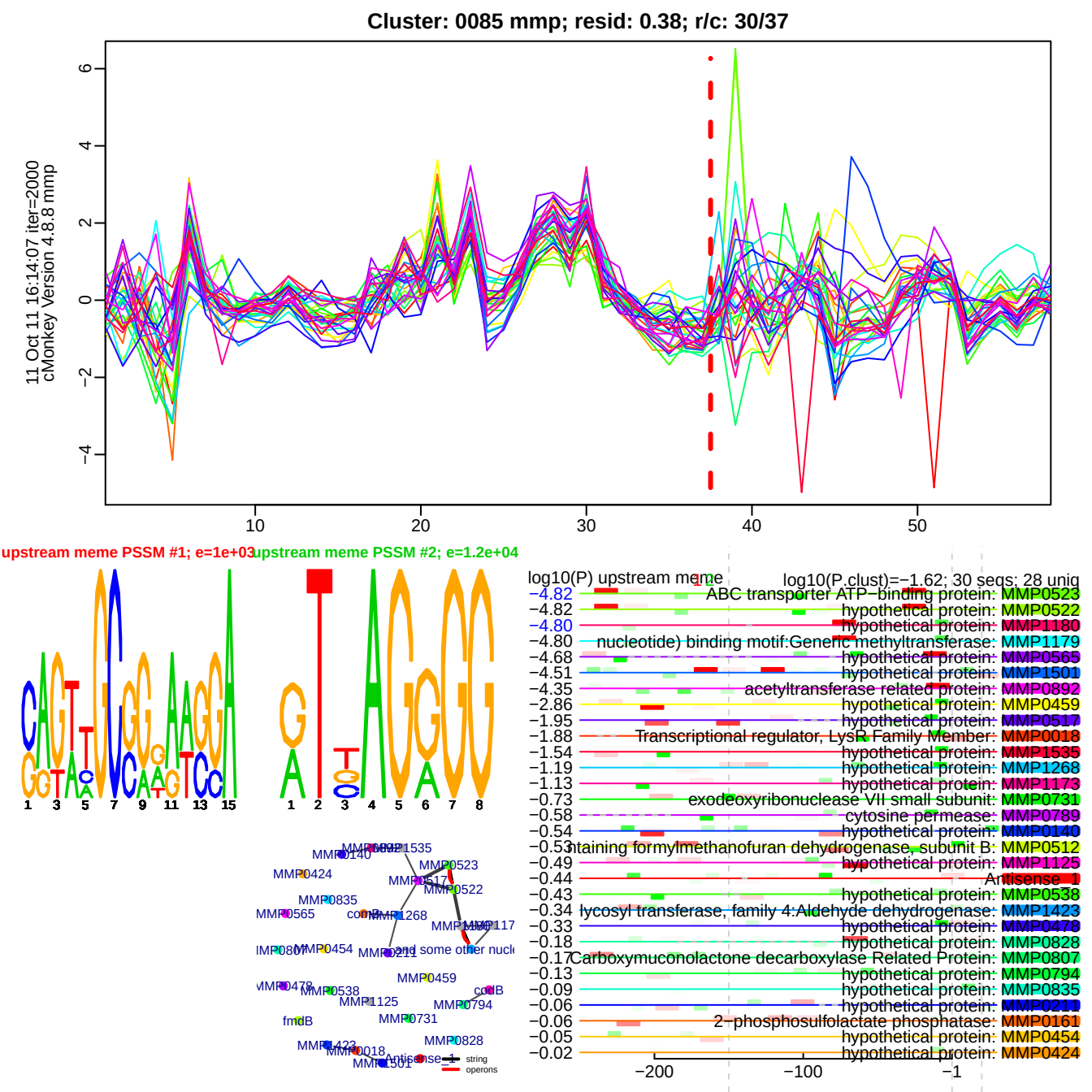
Cluster: 0085 mmp; resid: 0.38; r/c: 30/37
11 Oct 11 16:14:07 iter=2000
cMonkey Version 4.8.8 mmp
upstream meme PSSM #1; e=1e+03
upstream meme PSSM #2; e=1.2e+04
log10(P) upstream meme
log10(P.clust)=−1.62; 30 seqs; 28 uniq
−4.82 ABC transporter ATP−binding protein: MMP0523
−4.82 hypothetical protein: MMP0522
−4.80 hypothetical protein: MMP1180
−4.80 nucleotide) binding motif:Generic methyltransferase: MMP1179
−4.68 hypothetical protein: MMP0565
−4.51 hypothetical protein: MMP1501
−4.35 acetyltransferase related protein: MMP0892
−2.86 hypothetical protein: MMP0459
−1.95 hypothetical protein: MMP0517
−1.88 Transcriptional regulator, LysR Family Member: MMP0018
−1.54 hypothetical protein: MMP1535
−1.19 hypothetical protein: MMP1268
−1.13 hypothetical protein: MMP1173
−0.73 exodeoxyribonuclease VII small subunit: MMP0731
−0.58 cytosine permease: MMP0789
−0.54 hypothetical protein: MMP0140
−0.53 ntaining formylmethanofuran dehydrogenase, subunit B: MMP0512
−0.49 hypothetical protein: MMP1125
−0.44 Antisense_1
−0.43 hypothetical protein: MMP0538
−0.34 lycosyl transferase, family 4:Aldehyde dehydrogenase: MMP1423
−0.33 hypothetical protein: MMP0478
−0.18 hypothetical protein: MMP0828
−0.17 Carboxymuconolactone decarboxylase Related Protein: MMP0807
−0.13 hypothetical protein: MMP0794
−0.09 hypothetical protein: MMP0835
−0.06 hypothetical protein: MMP0211
−0.06 2−phosphosulfolactate phosphatase: MMP0161
−0.05 hypothetical protein: MMP0454
−0.02 hypothetical protein: MMP0424
−200
−100
−1
MMP0140
MMP0523
MMP0424
MMP0517
MMP0522
MMP0835
MMP0565
MMP1268
MMP0454
MMP0211
MMP0459
MMP0538
MMP1125
MMP0794
fmdB
MMP0731
MMP0828
MMP1423
MMP0018
MMP1501
Antisense_1
string
operons