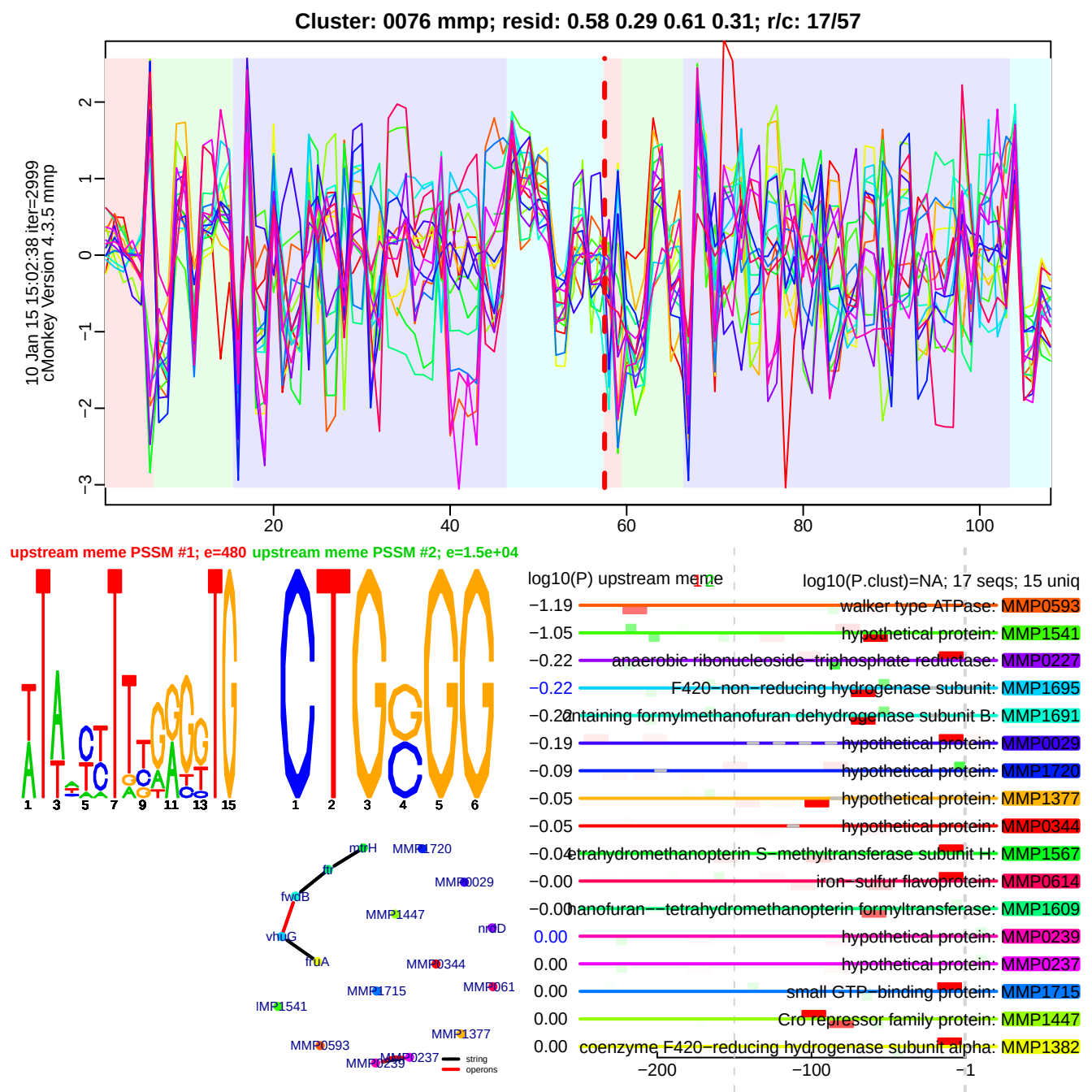
Cluster: 0076 mmp; resid: 0.58 0.29 0.61 0.31; r/c: 17/57
10 Jan 15 15:02:38 iter=2999
cMonkey Version 4.3.5 mmp
upstream meme PSSM #1; e=480
upstream meme PSSM #2; e=1.5e+04
log10(P) upstream meme
log10(P.clust)=NA; 17 seqs; 15 uniq
−1.19 walker type ATPase: MMP0593
−1.05 hypothetical protein: MMP1541
−0.22 anaerobic ribonucleoside−triphosphate reductase: MMP0227
−0.22 F420−non−reducing hydrogenase subunit: MMP1695
−0.20 ntaining formylmethanofuran dehydrogenase subunit B: MMP1691
−0.19 hypothetical protein: MMP0029
−0.09 hypothetical protein: MMP1720
−0.05 hypothetical protein: MMP1377
−0.05 hypothetical protein: MMP0344
−0.04 etrahydromethanopterin S−methyltransferase subunit H: MMP1567
−0.00 iron−sulfur flavoprotein: MMP0614
−0.00 hanofuran−−tetrahydromethanopterin formyltransferase: MMP1609
0.00 hypothetical protein: MMP0239
0.00 hypothetical protein: MMP0237
0.00 small GTP−binding protein: MMP1715
0.00 Cro repressor family protein: MMP1447
0.00 coenzyme F420−reducing hydrogenase subunit alpha: MMP1382
−200
−100
−1
mdH
MMP1720
MMP0029
fwdB
MMP1447
vhuG
fruA
MMP0344
MMP1715
MMP0061
MMP1541
MMP1377
MMP0593
MMP0237
MMP0239
string
operons