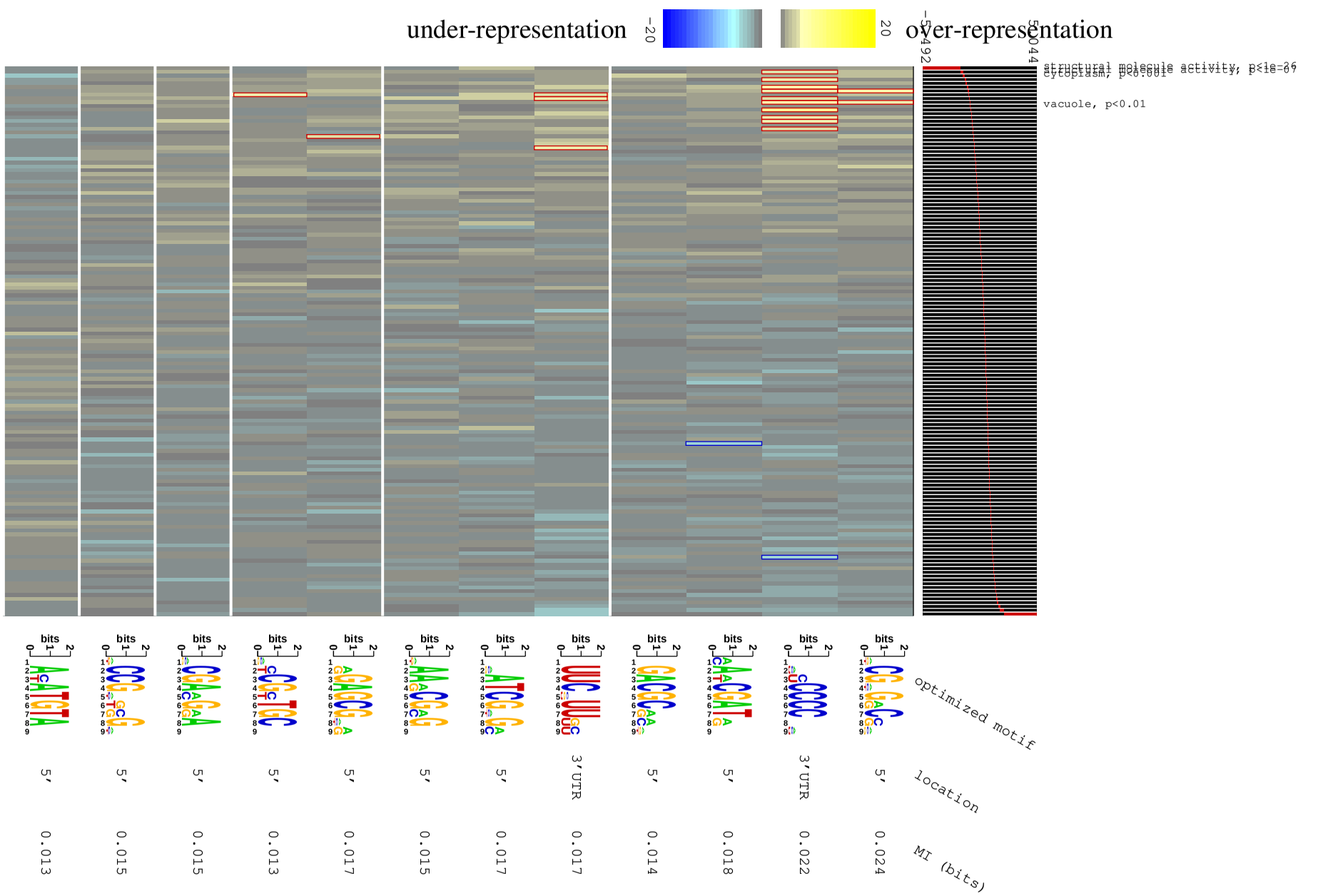
under-representation
-20
20
over-representation
-5.492
5.044
structural molecule activity, p<1e-06
cytoplasm, p<0.001
vacuole, p<0.01
optimized motif
location
MI (bits)
z-score
robustness
position bias
orientation bias
conservation index
seed
motif name
0.024 19.8 10/10 - - 1.00 CGAGACC -
0.022 17.4 10/10 Y → 1.00 UUCCCCC -
0.018 13.1 10/10 - ← 0.99 ATCGATA -
0.014 8.2 8/10 - ← 0.96 GACGCGC -
0.017 12.0 10/10 - → 0.99 UUCGUUU -
0.017 11.1 10/10 Y - 0.71 AATCGCG -
0.015 9.7 10/10 - ← 0.92 AAACGAG -
0.017 10.8 10/10 - ← 1.00 AGAGCGC -
0.013 7.2 6/10 - ← 0.87 TCGTTGC -
0.015 9.1 10/10 - ← 0.93 CGAAGAA -
0.015 8.7 9/10 - ← 0.95 CCGTTCG -
0.013 6.7 7/10 - ← 0.79 ATATGTA -