

cluster	SDs	motif finders	sequence logo
Wasserman genes	(not better than randoms)	AA, MD, ME	A A A A A A C
	2.3	AA, BP	G G G G G G G G
	0.54	AA	G G G A G G A G G A
	0.34	ME	G C G G C C C
	1.3	ME	G G G C G G G
skeletal muscle	4.8	AA	T C A C G A A A A
	2.1	AA	A A A A A A A A A A A A
	1.3	AA, BP	A A G C A G A
	0.93	ME	G G G G G G G G G G G G
	2.3	AA	I A A G A A A A A
	0.63	BP	I G G G G G A G G G

Additional Figure 2

Examples of motifs found by MultiFinder that failed the block filtering criteria.