














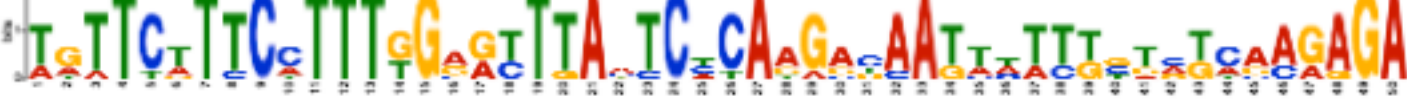










Conserved sequences - COLEC/10

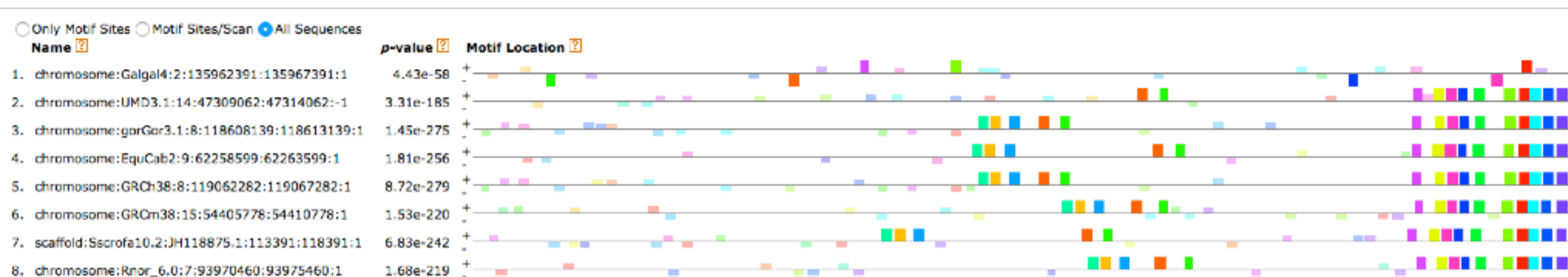
		Logo	E-value <a href="#">?</a>	Sites <a href="#">?</a>	Width <a href="#">?</a>	More <a href="#">?</a>	Submit/Download <a href="#">?</a>
1.			4.4e-083	8	50	<a href="#">I</a>	<a href="#">...</a>
2.			2.6e-065	7	50	<a href="#">I</a>	<a href="#">...</a>
3.			2.1e-048	8	50	<a href="#">I</a>	<a href="#">...</a>
4.			1.6e-045	7	50	<a href="#">I</a>	<a href="#">...</a>
5.			3.1e-043	6	50	<a href="#">I</a>	<a href="#">...</a>
6.			1.1e-037	7	50	<a href="#">I</a>	<a href="#">...</a>
7.			7.1e-038	7	48	<a href="#">I</a>	<a href="#">...</a>
8.			6.3e-031	8	50	<a href="#">I</a>	<a href="#">...</a>
9.			1.1e-030	8	50	<a href="#">I</a>	<a href="#">...</a>
10.			4.6e-030	7	49	<a href="#">I</a>	<a href="#">...</a>

Sites ?

TTGAGATAAG	TTTCAATAAACTGTTGTC	ACTTGAAAATAAACAGTCTGGACAATGTATGG	TTCATTTGGA
CAGAGATAAT	TTTCAATAAACTGTTGTC	ACTTGAAAATAAACAGTCTGGACAATGTATGG	TCCATTTGGA
CAGAGATAAT	TTTCAATAAACTGTTGTC	ACTTGAAAATAAACAGTCTGGACAATGTATGG	TCCATTTGGA
CAGAGATAAT	TTTCAATAAACTGTTGTC	ACTTGAAAATAAACAGTCTGGACAATGTATGG	TCCATTTGGA
TAGAGATAAT	TTTCAATAAACTGTTGTC	ATTGGAAAAGAAACAGTCTGGACAATGTATGG	TCCATTTGGG
TAGAGATAAT	TTTCAATAAACTGTTGTC	ATTTGAAAAGAAACGGTCTGGACAATGTATGG	TCCATTTGGA
CAGAGATAAT	TTTCAATAAACTCTTGTC	ACTTGAAAGTAAACAGTCTGGAAAATGTACGG	CTCATTTGGA
AAGAGATAAA	TTGGAATAAACAGCTGTC	AGTGTAATAATAATATCCTGGATAATGTGCAG	GACAGAAATA

# Conserved sequences align across species

## *COLEC10*





# Conserved sequences - COLEC10



## Sites

TTGAGATAAG TTTCAATAAACTGTTGTCACTTGAATAAAACAGTCTGGACAATGTATGG TTCATTTGGA  
CAGAGATAAT TTTCAATAAACTGTTGTCACTTGAATAAAACAGTCTGGACAATGTATGG TCCATTTGGA  
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