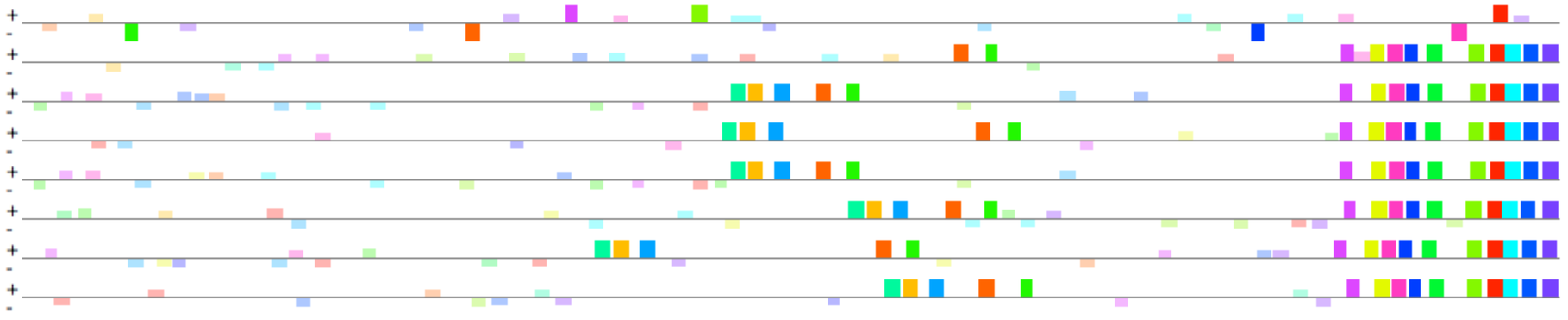
Conserved sequences align across species

Only Motif Sites | Motif Sites/Scan | All Sequences

	Name	*p*-value	Motif Location
1.	chromosome:Galgal4:2:135962391:135967391:1	4.43e-58	
2.	chromosome:UMD3.1:14:47309062:47314062:-1	3.31e-185	
3.	chromosome:gorGor3.1:8:118608139:118613139:1	1.45e-275	
4.	chromosome:EquCab2:9:62258599:62263599:1	1.81e-256	
5.	chromosome:GRCh38:8:119062282:119067282:1	8.72e-279	
6.	chromosome:GRCm38:15:54405778:54410778:1	1.53e-220	
7.	scaffold:Sscrofa10.2:JH118875.1:113391:118391:1	6.83e-242	
8.	chromosome:Rnor_6.0:7:93970460:93975460:1	1.68e-219	



Upstream analysis

Gene X
5kb upstream

CIS-BP v.1.02
PWM- LogOdds > 8
Bos taurus

"All" potential TFBS

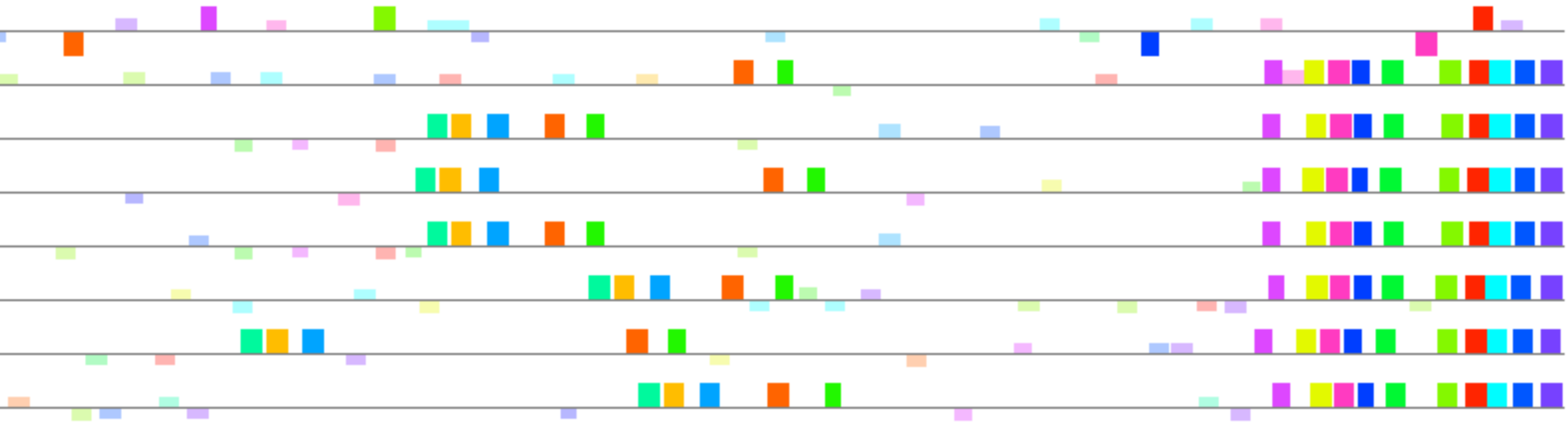
Gene X
5kb upstream

Horse | Cow | Mouse | Rat
Human | Gorilla | Pig | Chicken

MEME v.4.10.1

Conserved sequences

"Conserved" potential TFBS

TFBS with SNVs

Conserved sequences align across species