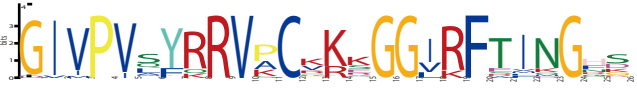
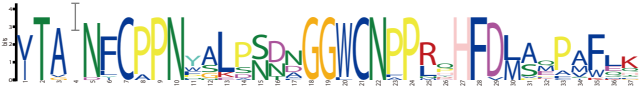
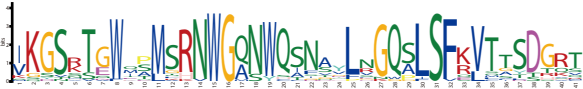
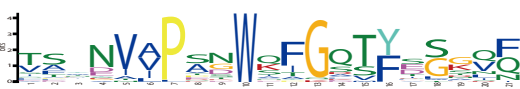


motif	width	sites	Logo	consensus sequence	e-value
1	29	30		QGYGTNTAALSTALFNNGLSCGACYZIRC	1.5e-408
2	26	20		GIVPVSYRRVPCKKKGGIRFTINGHS	4.1e-305
3	37	18		VTATNFCPPNYALPSBNGWCNPPRQHFDLAQPAFLK	2.0e-429
4	41	22		IKGSRTGWMPMsrnwgqNwqSNaylNgqSLsFKVtTSDGRT	2.1e-471
5	29	24		GWQSAHATFYGGDASGTMGACGYGNLY	1.5e-329
6	21	30		TSYNVAPSNWQFGQTYESGQF	1.4e-193
7	15	23		YFNLVLITNVGGAGD	1.1e-152
8	32	9		FMALARPGMAPQLMNLGTVDIEYKRIPCEYPG	1.7e-076
9	49	7		NLTVKVEESSQKPSNLAILILYQGQTDILAVDVAQVGSSDWKYMTRSY	1.9e-096
10	11	18		DPRWCLEGSIV	6.7e-054