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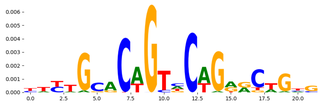
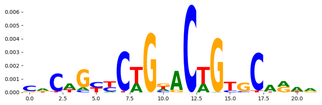
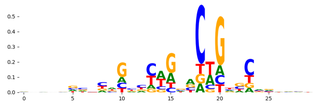
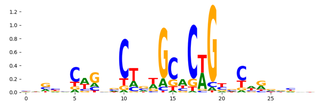
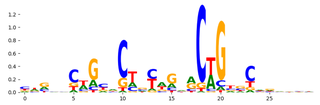
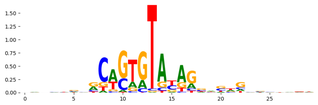
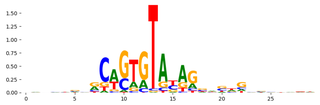
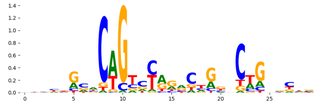
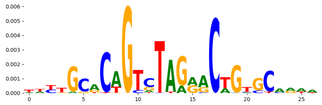
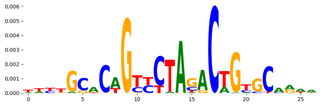
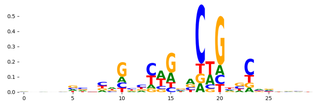
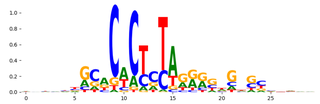
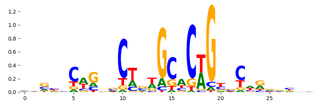
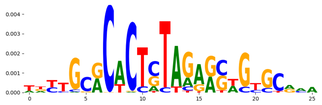
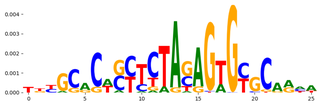
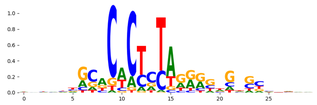
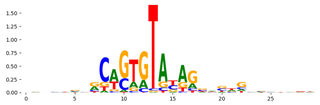
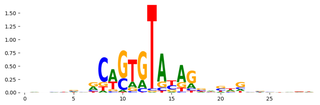
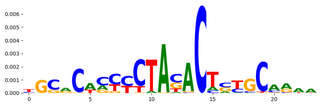
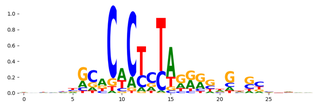
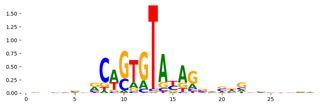
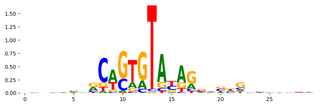
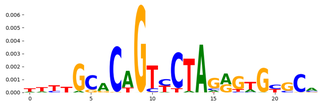
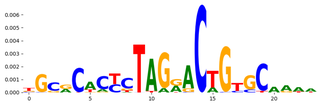
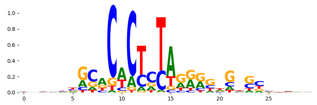
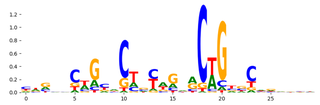
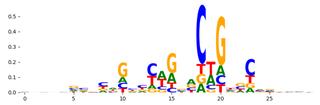
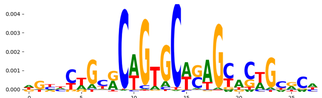
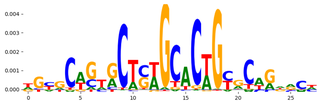
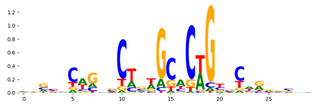
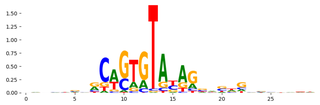
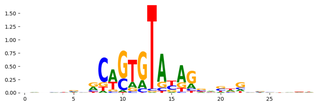
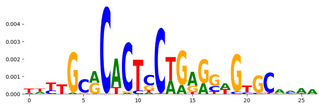
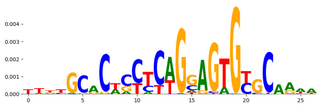
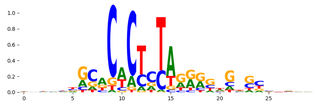
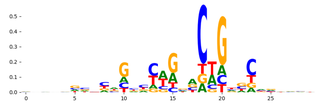
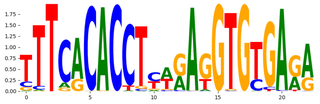
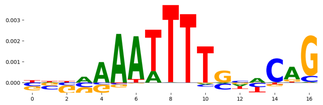
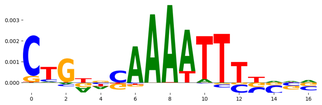
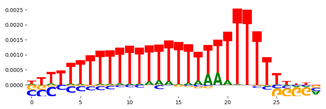
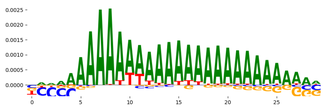
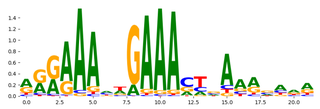
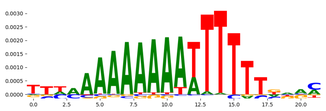
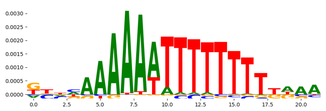
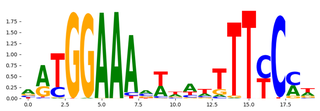
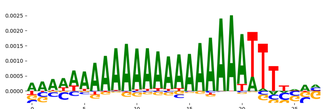
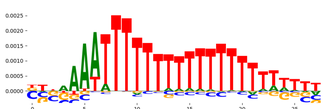
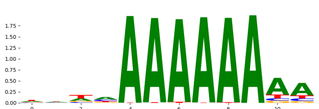
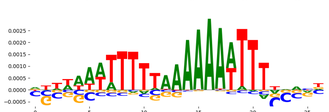
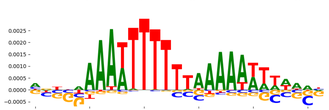
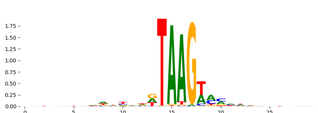
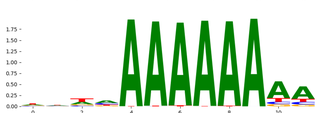
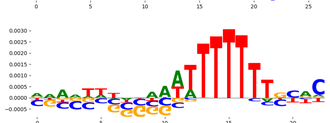
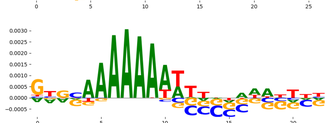
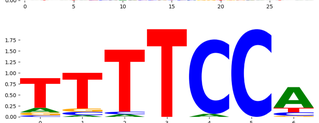
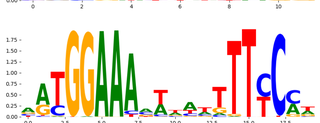
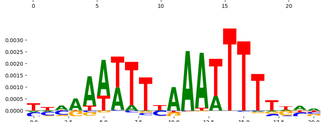
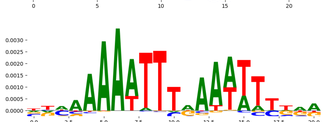
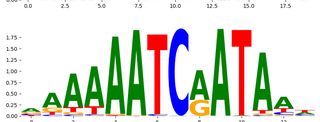
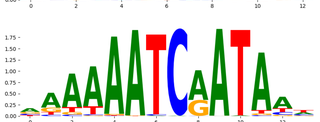
pos_patterns.pattern_0	8519			TN5_1	2.343550e-09		TN5_2	1.714920e-08		TN5_7	0.000100	
pos_patterns.pattern_1	6312			TN5_4	2.034240e-02		TN5_5	2.034240e-02		TN5_8	0.027068	
pos_patterns.pattern_2	4140			TN5_1	5.084640e-07		TN5_3	3.240360e-06		TN5_2	0.003034	
pos_patterns.pattern_3	3654			TN5_3	9.636770e-15		TN5_4	2.155130e-03		TN5_5	0.002155	
pos_patterns.pattern_4	3416			TN5_3	1.164160e-01		TN5_4	1.164160e-01		TN5_5	0.116436	
pos_patterns.pattern_5	2918			TN5_3	8.388180e-05		TN5_7	3.430450e-04		TN5_1	0.001952	
pos_patterns.pattern_6	2794			TN5_2	3.645640e-17		TN5_4	4.788040e-06		TN5_5	0.000005	
pos_patterns.pattern_7	796			TN5_3	4.789920e-07		TN5_1	9.746420e-04		TBX1_TBX_4	0.084708	
pos_patterns.pattern_8	223			POU3F4_POU_2	1.000000e-00		NFAC2_HUMAN.H11MO.0.B	1.000000e-00		NFAC2_MOUSE.H11MO.0.C	1.000000	
pos_patterns.pattern_9	146			ZNF384_MA1125.1	5.454130e-02		PRDM6_HUMAN.H11MO.0.C	6.609230e-02		STAT1_MOUSE.H11MO.0.A	0.114602	
pos_patterns.pattern_10	139			FOXD2_forkhead_1	1.000000e-00		NFATC1_NFAT_1	1.000000e-00		CPEB1_RRM_1	1.000000	
pos_patterns.pattern_11	125			ZNF384_MA1125.1	2.832720e-02		PRDM6_HUMAN.H11MO.0.C	2.832720e-02		FOXJ3_HUMAN.H11MO.0.A	0.165850	
pos_patterns.pattern_12	95			DNASE_2	2.608540e-01		ZNF384_MA1125.1	1.000000e-00		LHX3_HUMAN.H11MO.0.C	1.000000	
pos_patterns.pattern_13	85			NFATC2_MA0152.1	5.288450e-01		NFATC1_NFAT_1	5.288450e-01		PRDM6_HUMAN.H11MO.0.C	0.703003	
pos_patterns.pattern_14	74			LIN54_MA0610.1	1.000000e-00		ONECUT3_CUT_1	1.000000e-00		ONECUT3_MA0757.1	1.000000	

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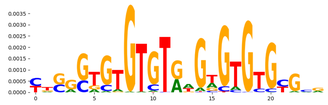
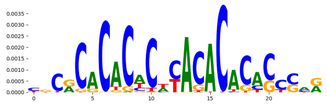
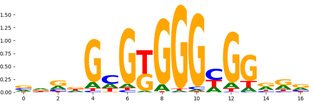
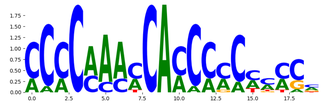
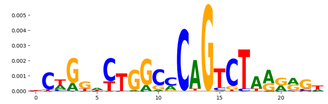
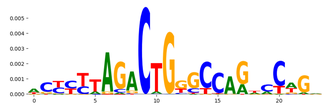
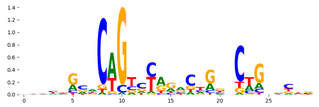
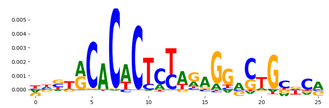
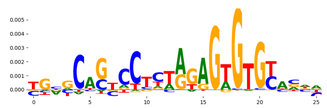
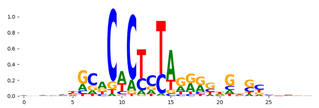
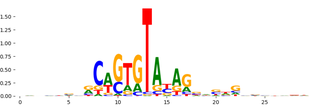
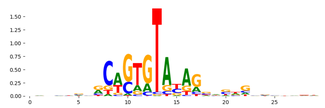
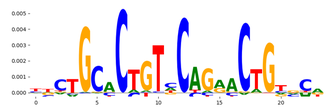
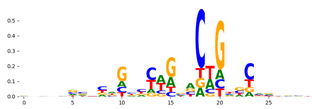
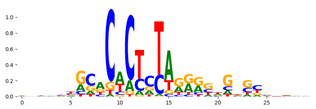
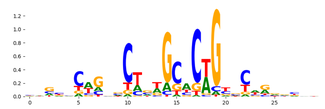
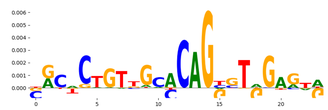
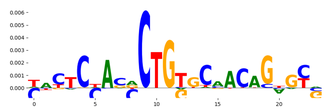
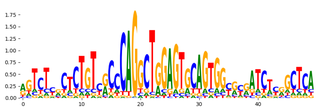
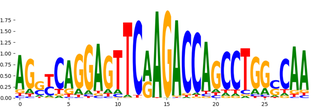
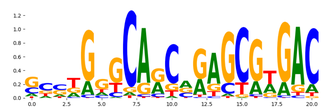
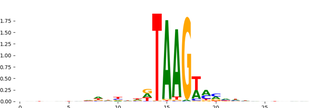
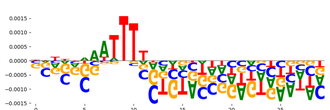
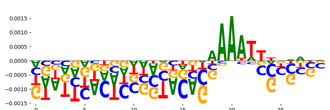
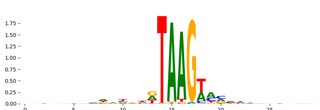
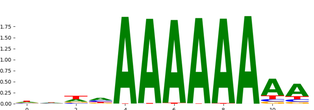
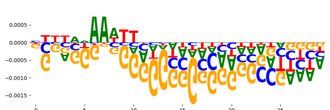
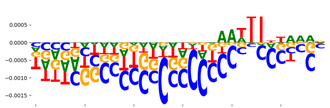
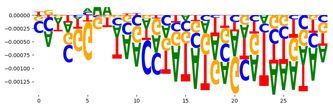
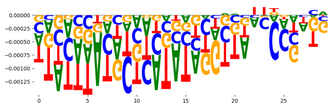
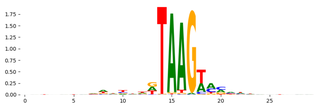
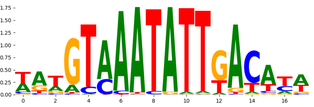
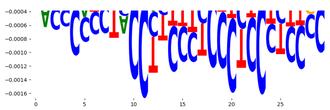
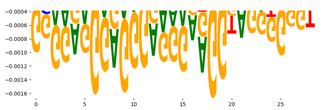
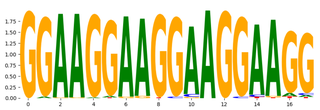
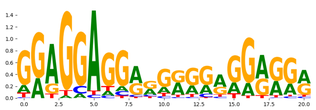
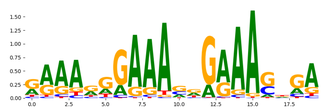
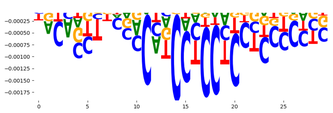
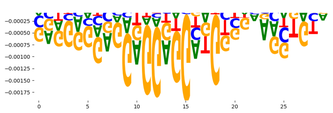
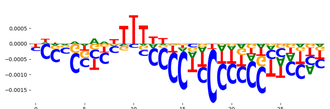
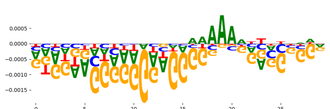
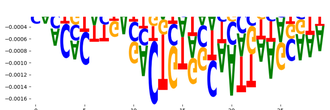
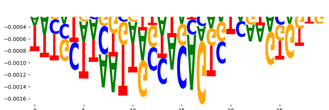
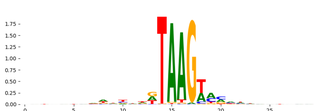
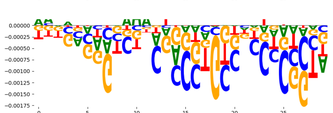
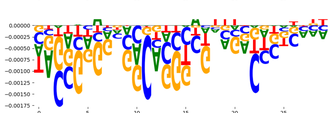
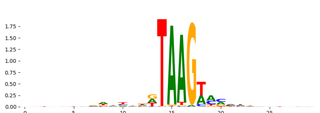
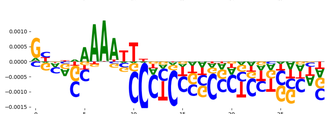
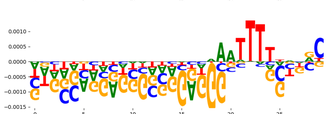
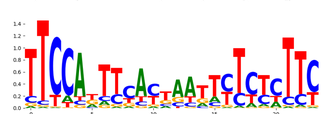
pos_patterns.pattern_15	72			EGR2_HUMAN.H11MO.0.A	1.000000e-00		EGR1_HUMAN.H11MO.0.A	1.000000e-00		RREB1_MA0673.1	1.000000	
pos_patterns.pattern_16	64			TN3_8	9.999990e-01		RORA_HUMAN.H11MO.0.C	9.999990e-01		RORA_MOUSE.H11MO.0.C	0.999999	
pos_patterns.pattern_17	42			TN5_3	4.137570e-04		TN5_4	2.010910e-01		TN5_5	0.201091	
pos_patterns.pattern_18	33			TN5_1	1.252380e-01		TN5_3	1.252380e-01		TN5_2	0.125238	
pos_patterns.pattern_19	20			TN3_6	4.466370e-08		DNASE_5	3.970190e-01		PAX5_HUMAN.H11MO.0.A	0.419224	
neg_patterns.pattern_0	98			ZNF384_MA1125.1	1.310690e-02		DNASE_2	2.011360e-02		ONECUT1_CUT_1	0.030979	
neg_patterns.pattern_1	85			DNASE_2	8.722660e-04		ZNF384_MA1125.1	1.020160e-01		FOXJ3_HUMAN.H11MO.0.A	0.569880	
neg_patterns.pattern_2	83			SPIB_HUMAN.H11MO.0.A	4.822550e-02		SPI1_HUMAN.H11MO.0.A	4.822550e-02		BC11A_HUMAN.H11MO.0.A	0.219973	
neg_patterns.pattern_3	80			DNASE_2	2.098000e-01		FOXB1_forkhead_2	2.098000e-01		FOXB1_MA0845.1	0.226153	
neg_patterns.pattern_4	76			EWSR1-FLI1_MA0149.1	1.836460e-02		ZNF263_MA0528.1	6.070330e-02		IRF3_HUMAN.H11MO.0.B	0.065934	
neg_patterns.pattern_5	76			ELF5_MOUSE.H11MO.0.A	4.641330e-02		SPI1_HUMAN.H11MO.0.A	5.171870e-02		SPIB_HUMAN.H11MO.0.A	0.051719	
neg_patterns.pattern_6	68			PRDM6_HUMAN.H11MO.0.C	2.897410e-02		ETS2_HUMAN.H11MO.0.B	4.946930e-02		ETS2_MOUSE.H11MO.0.A	0.049469	
neg_patterns.pattern_7	55			DNASE_2	3.608670e-01		POU3F3_POU_3	3.608670e-01		POU6F2_POU_2	0.360867	
neg_patterns.pattern_8	53			DNASE_2	8.982450e-01		ZNF85_HUMAN.H11MO.0.C	8.982450e-01		ONECUT1_CUT_1	0.898245	
neg_patterns.pattern_9	44			SPI1_HUMAN.H11MO.0.A	1.280920e-01		SPIB_HUMAN.H11MO.0.A	1.280920e-01		ZFP82_HUMAN.H11MO.0.C	0.195097	

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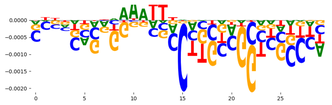
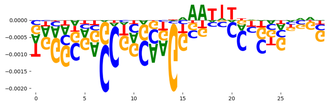
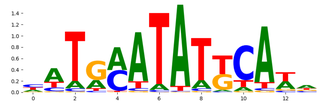
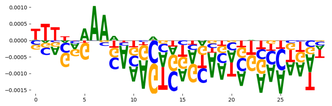
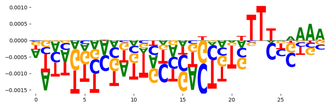
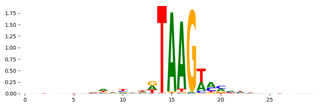
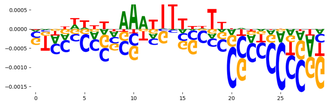
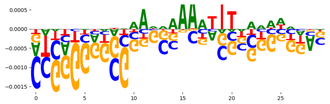
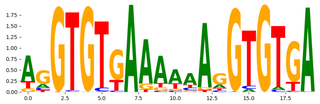
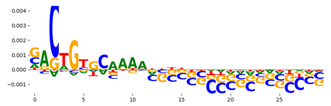
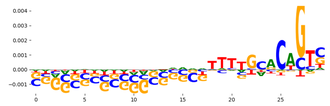
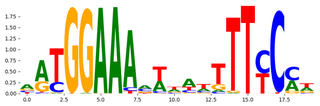
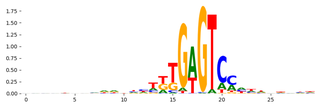
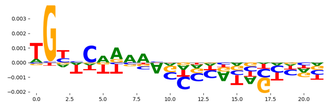
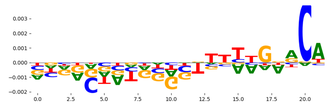
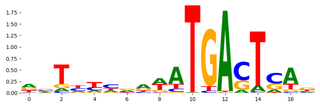
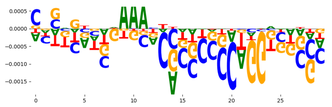
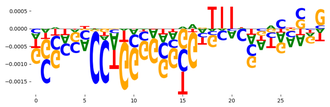
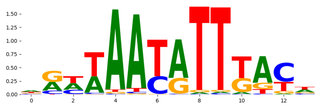
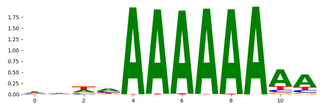
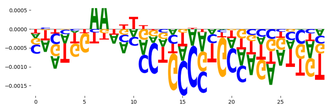
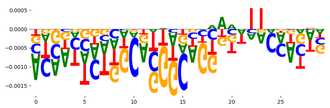
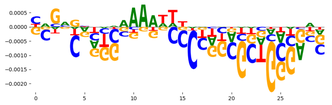
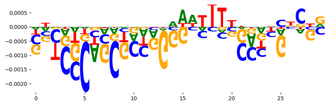
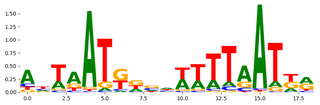
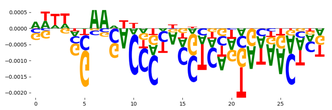
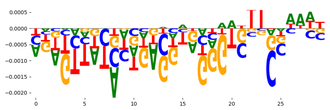
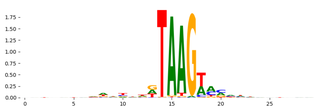
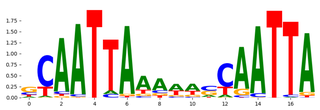
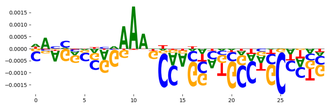
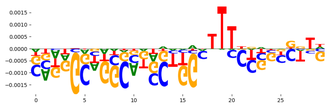
neg_patterns.pattern_10	44			PBX2_HUMAN.H11MO.0.C	2.971310e-01		PBX2_MOUSE.H11MO.0.C	2.971310e-01		POU2F1_POU_2	0.420332	
neg_patterns.pattern_11	39			DNASE_2	5.112270e-02		ZNF384_MA1125.1	3.196580e-01		PRD86_HUMAN.H11MO.0.C	0.989106	
neg_patterns.pattern_12	36			ZNF384_MA1125.1	8.221930e-01		TBX1_TBX_1	8.221930e-01		LHX3_HUMAN.H11MO.0.C	0.822193	
neg_patterns.pattern_13	35			NFATC1_NFAT_1	2.918110e-01		DNASE_4	4.694330e-01		FOXA2_HUMAN.H11MO.0.A	1.000000	
neg_patterns.pattern_14	31			AridB_MA0601.1	1.000000e-00		BATF_MOUSE.H11MO.0.A	1.000000e-00		ZNF435_C2H2_1	1.000000	
neg_patterns.pattern_15	29			FOXD2_forkhead_1	7.085750e-01		FOXD3_forkhead_1	7.085750e-01		ZNF384_MA1125.1	0.708575	
neg_patterns.pattern_16	27			ZNF384_MA1125.1	1.310330e-01		HOXC10_homeodomain_2	1.310330e-01		Hoxc10.mouse_homeodomain_2	0.131033	
neg_patterns.pattern_17	25			NKX61_MOUSE.H11MO.0.A	5.028620e-01		ONECUT3_CUT_1	5.028620e-01		ONECUT3_MA0757.1	0.502862	
neg_patterns.pattern_18	24			DNASE_2	5.609400e-01		MSX2_homeodomain_1	9.847780e-01		LHX3_HUMAN.H11MO.0.C	0.984778	
neg_patterns.pattern_19	23			NR2E3_HUMAN.H11MO.0.C	1.000000e-00		NR2E3_MOUSE.H11MO.0.C	1.000000e-00		ZN317_HUMAN.H11MO.0.C	1.000000	