
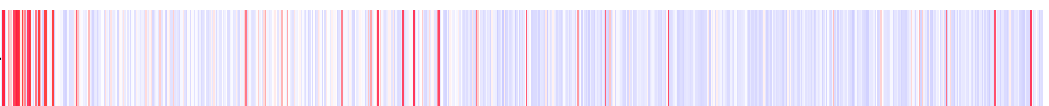
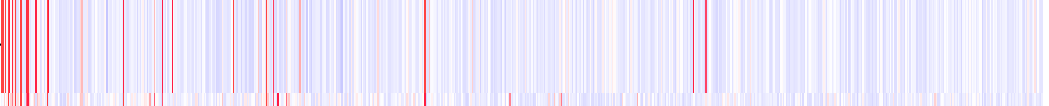
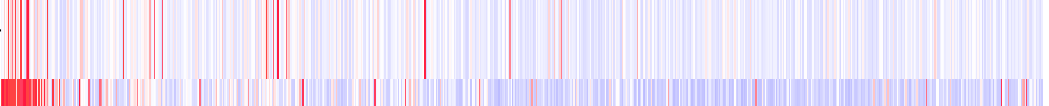
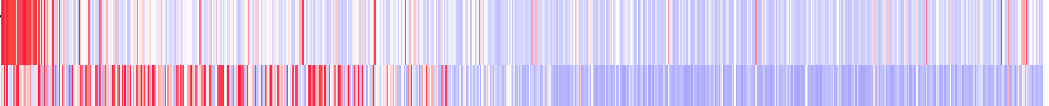
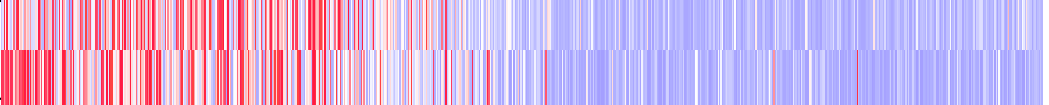
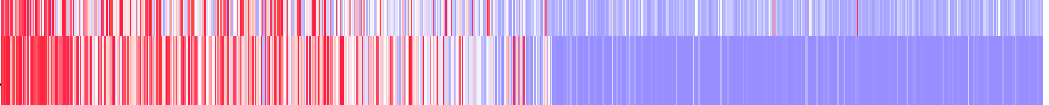
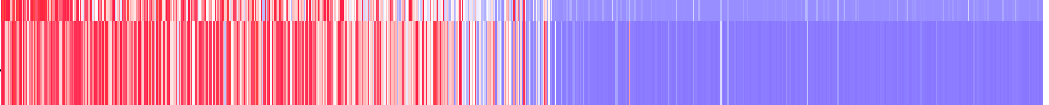
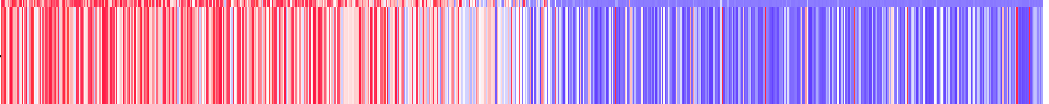
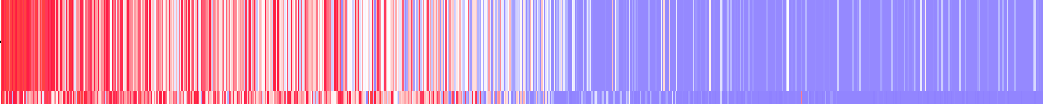
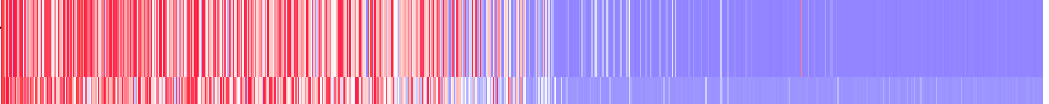
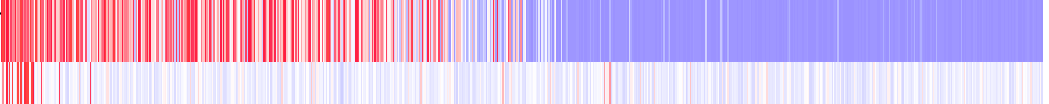
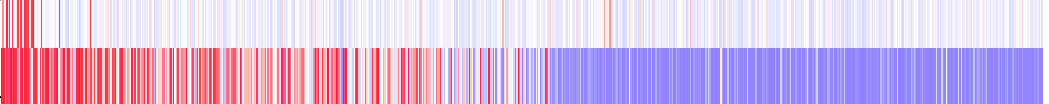
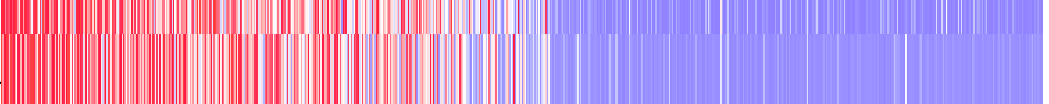
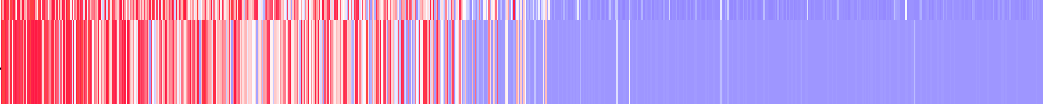
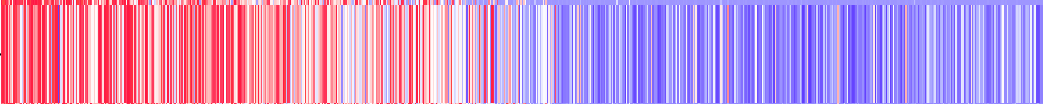
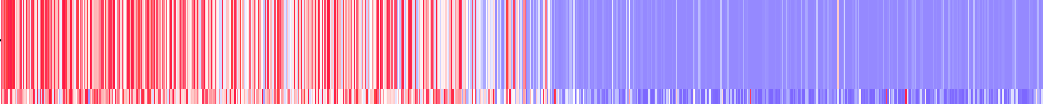
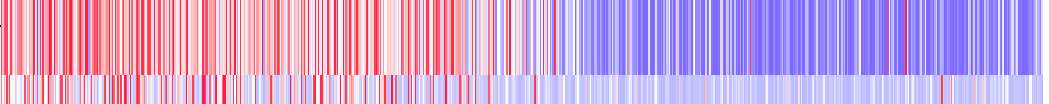
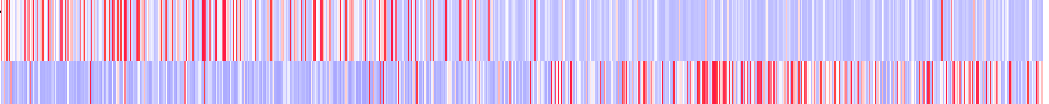
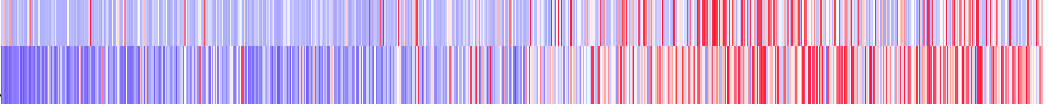
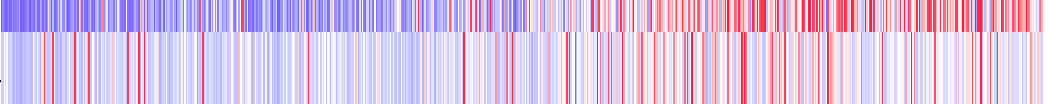
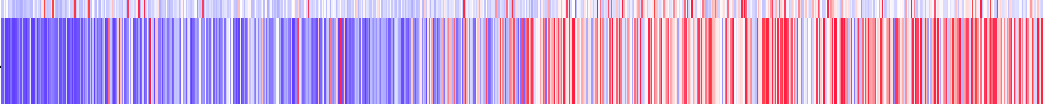
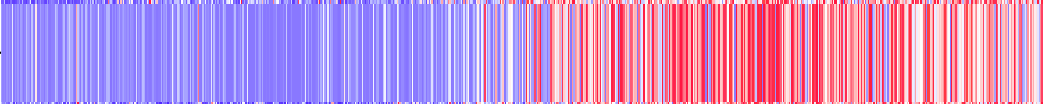
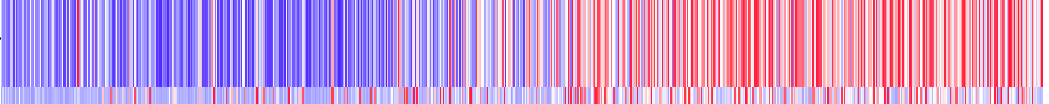
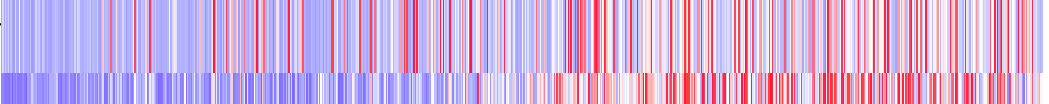
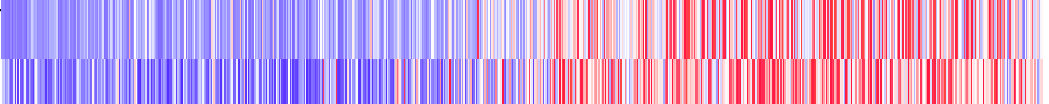
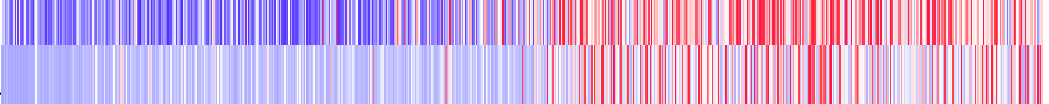
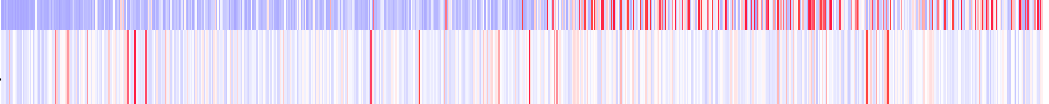
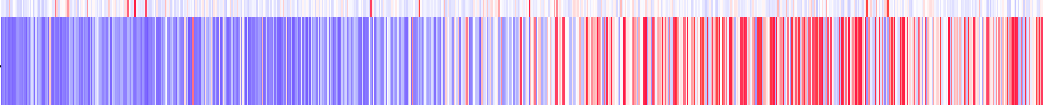
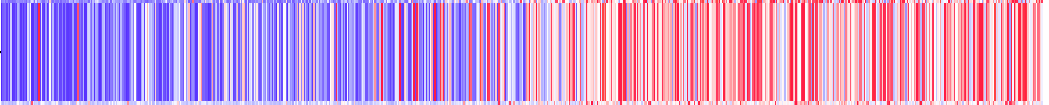
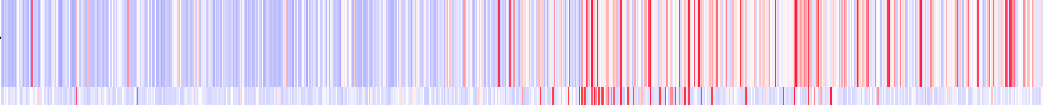
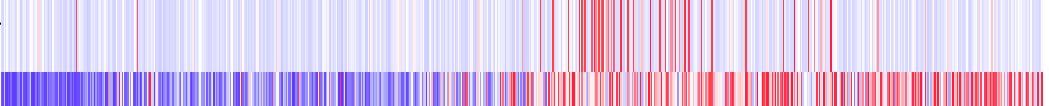
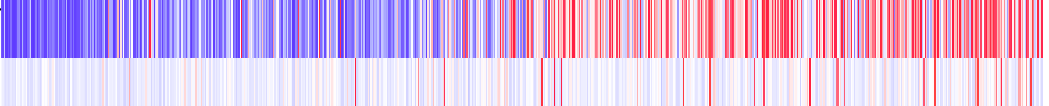
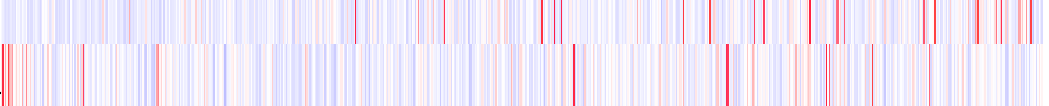
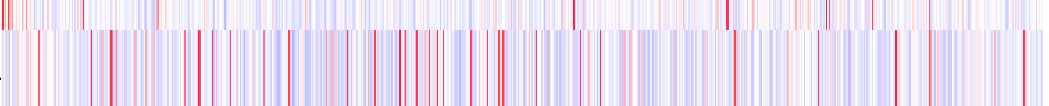
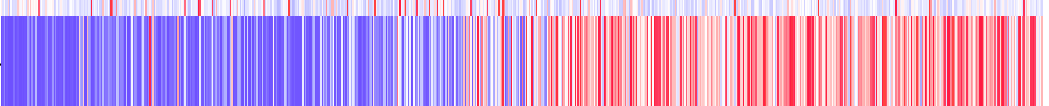



Gene Expression

	C1	IC(Δ)	p-value	FDR
				
RAB6C		0.940(0.045)	4.35e-07	1.00e-02
STARD3		0.928(0.061)	4.35e-07	1.00e-02
ORMDL3		0.912(0.057)	2.17e-06	3.00e-02
CREB3L4		0.909(0.017)	2.61e-06	3.00e-02
MKRN2-AS1		0.888(0.023)	9.57e-06	3.11e-02
RP11-517C16.2		0.884(0.022)	1.09e-05	3.11e-02
RP11-532F12.5		0.883(0.005)	1.09e-05	3.11e-02
C1orf172		0.883(0.005)	1.13e-05	3.11e-02
INADL		0.878(0.015)	1.22e-05	3.11e-02
ESRP2		0.874(0.005)	1.39e-05	3.11e-02
ESRP1		0.873(0.006)	1.44e-05	3.11e-02
MARVELD3		0.870(0.005)	1.48e-05	3.11e-02
SERHL2		0.869(0.043)	1.48e-05	3.11e-02
CDC42BPG		0.868(0.006)	1.52e-05	3.11e-02
PRRG2		0.866(0.006)	1.57e-05	3.11e-02
GRHL2		0.865(0.006)	1.57e-05	3.11e-02
HOOK2		0.865(0.010)	1.57e-05	3.11e-02
SPINT1		0.864(0.006)	1.65e-05	3.11e-02
F11R		0.864(0.011)	1.65e-05	3.11e-02
C9orf169		0.864(0.040)	1.65e-05	3.11e-02
NEK10		-0.761(0.026)	7.74e-05	9.19e-02
SACS		-0.763(0.010)	7.61e-05	9.19e-02
PPAP2A		-0.767(0.037)	7.00e-05	9.19e-02
RP11-529K1.4		-0.768(0.011)	6.83e-05	9.19e-02
SYDE1		-0.773(0.009)	6.35e-05	9.19e-02
DENND5A		-0.774(0.011)	6.13e-05	9.19e-02
TUBB2A		-0.774(0.026)	6.13e-05	9.19e-02
LEPRE1		-0.778(0.011)	5.91e-05	9.19e-02
AP1M1		-0.780(0.011)	5.52e-05	9.19e-02
SH2B3		-0.784(0.019)	5.05e-05	9.19e-02
PABPC4		-0.788(0.070)	4.61e-05	9.19e-02
IKBIP		-0.788(0.011)	4.57e-05	9.19e-02
STX2		-0.788(0.012)	4.48e-05	9.19e-02
RAB8B		-0.789(0.019)	4.39e-05	9.19e-02
C19orf71		-0.791(0.050)	4.26e-05	9.19e-02
ST3GAL2		-0.793(0.010)	4.04e-05	9.19e-02
GPAM		-0.799(0.051)	3.61e-05	9.19e-02
KIAA0196		-0.816(0.192)	2.09e-05	9.19e-02
MDM1		-0.821(0.182)	1.91e-05	9.19e-02
LIX1L		-0.835(0.007)	1.61e-05	9.19e-02