
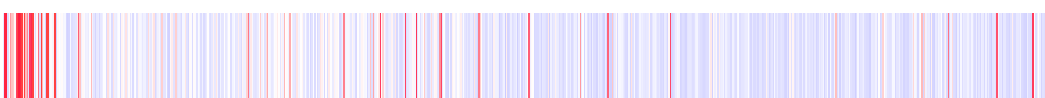
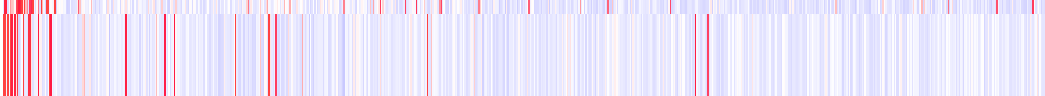
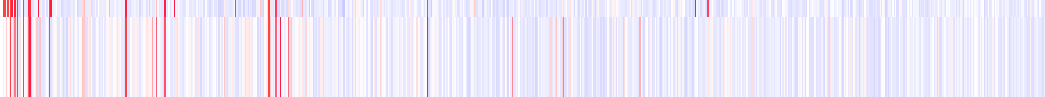
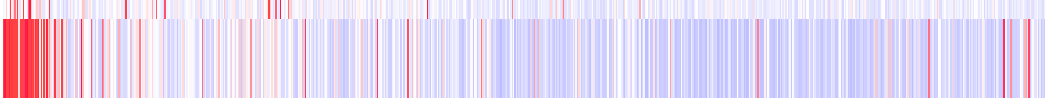
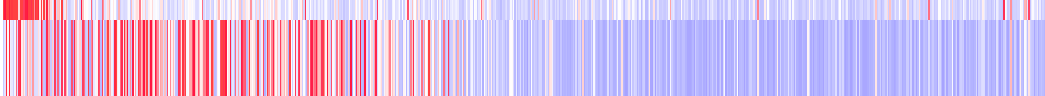
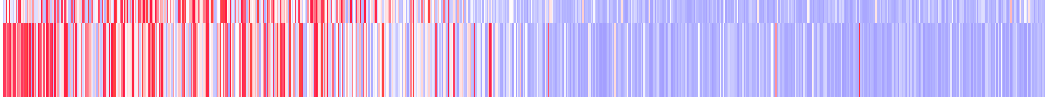
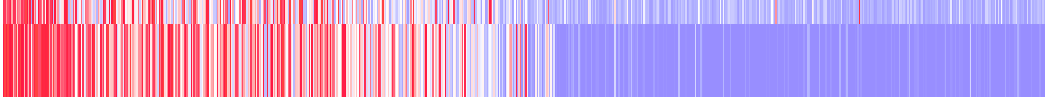
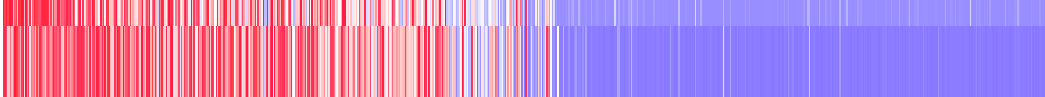
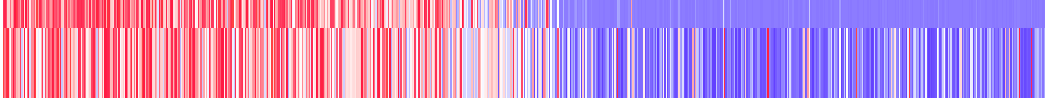
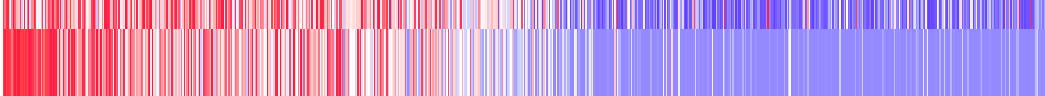
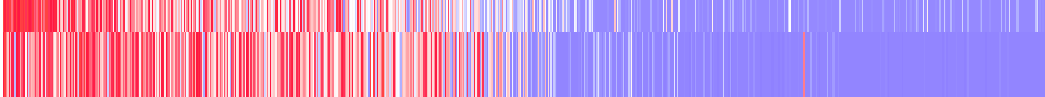
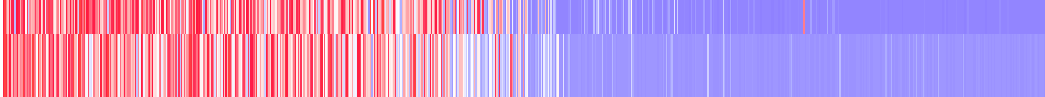
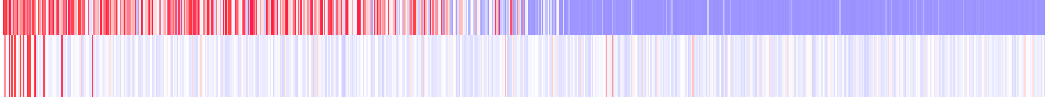
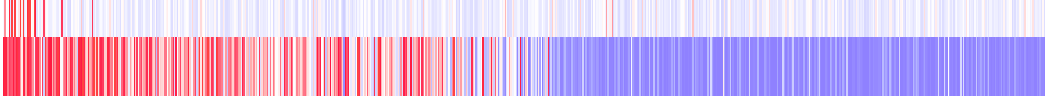
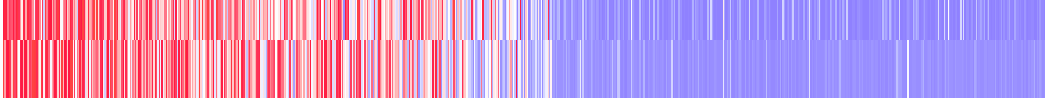
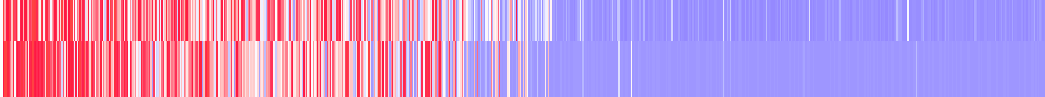
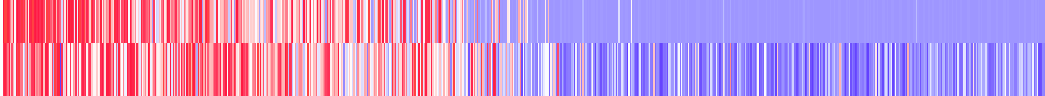
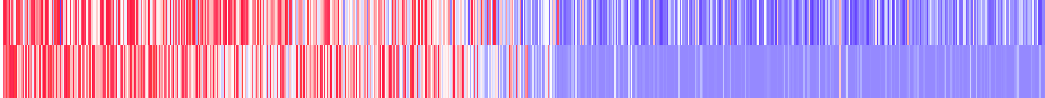
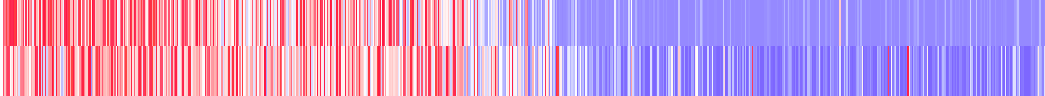
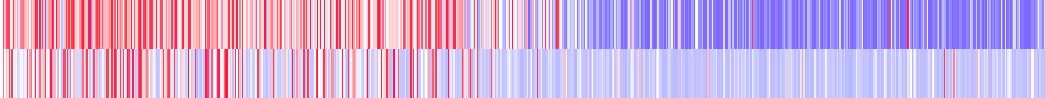
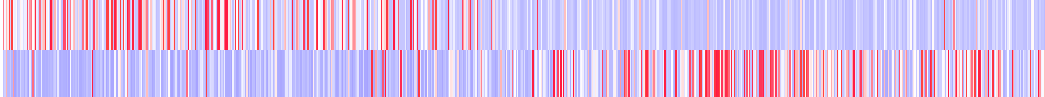
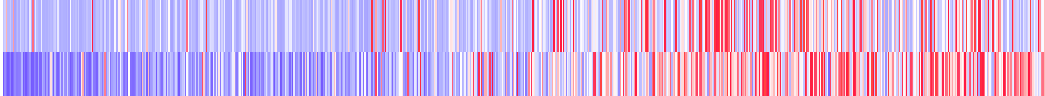
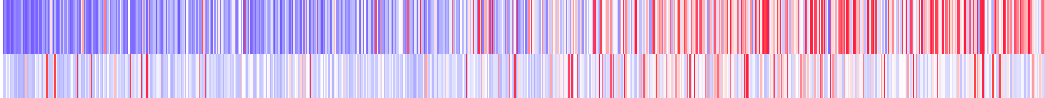
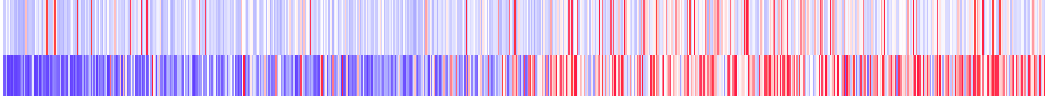
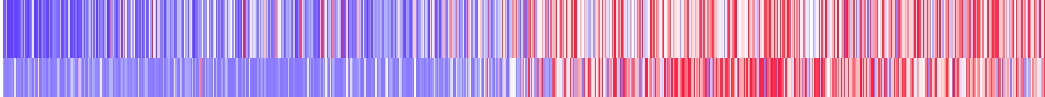
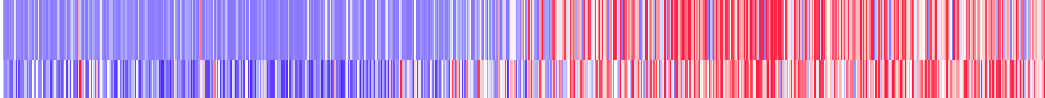
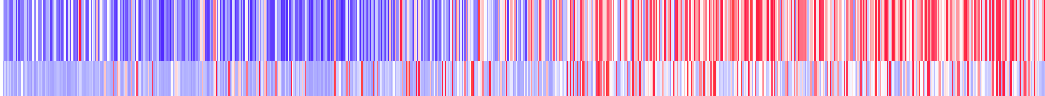
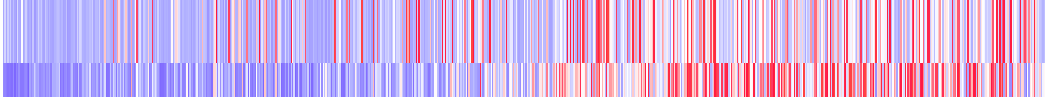
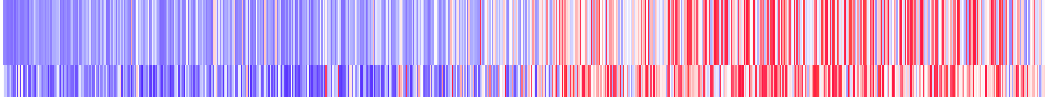
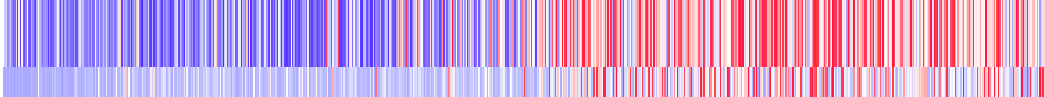
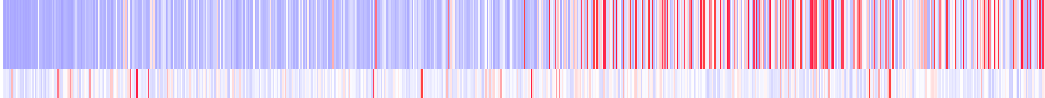
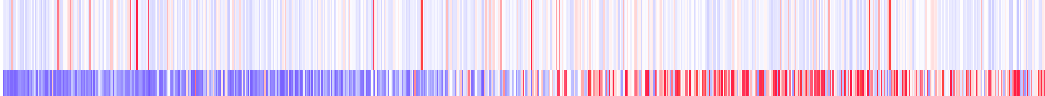
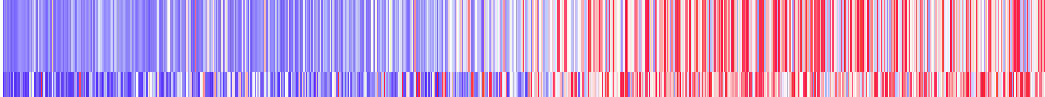
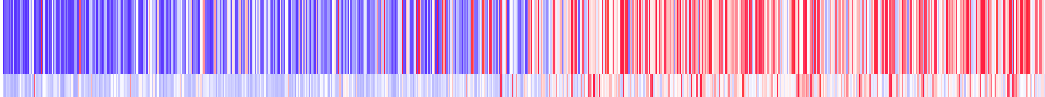
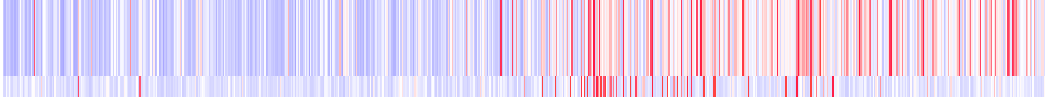
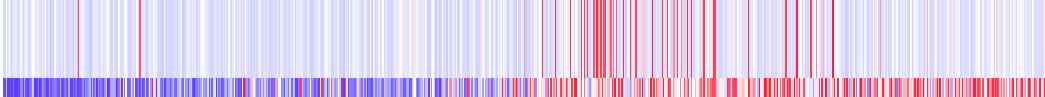
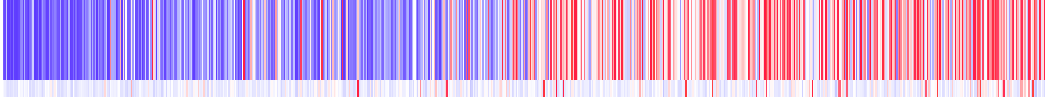
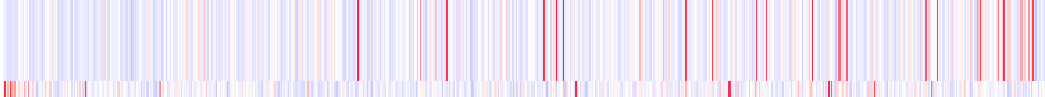
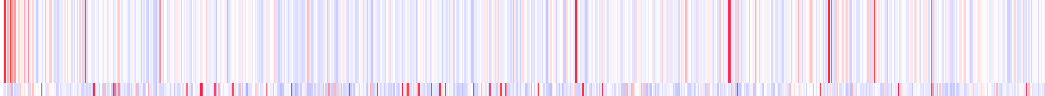
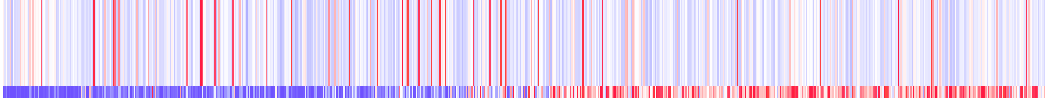


C1 vs Gene Expression

C1		IC(Δ)	P-val	FDR
RAB6C		0.940(0.061)	4.35e-07	1.00e-02
STARD3		0.928(0.064)	4.35e-07	1.00e-02
ORMDL3		0.912(0.058)	8.70e-07	1.33e-02
CREB3L4		0.909(0.017)	1.30e-06	1.50e-02
MKRN2-AS1		0.888(0.027)	7.83e-06	3.03e-02
RP11-517C16.2		0.884(0.022)	7.83e-06	3.03e-02
RP11-532F12.5		0.883(0.004)	8.26e-06	3.03e-02
C1orf172		0.883(0.005)	8.26e-06	3.03e-02
INADL		0.878(0.012)	9.57e-06	3.03e-02
ESRP2		0.874(0.005)	1.00e-05	3.03e-02
ESRP1		0.873(0.006)	1.04e-05	3.03e-02
MARVELD3		0.870(0.006)	1.39e-05	3.03e-02
SERHL2		0.869(0.049)	1.39e-05	3.03e-02
CDC42BPG		0.868(0.006)	1.44e-05	3.03e-02
PRRG2		0.866(0.005)	1.57e-05	3.03e-02
GRHL2		0.865(0.007)	1.61e-05	3.03e-02
HOOK2		0.865(0.010)	1.61e-05	3.03e-02
SPINT1		0.864(0.006)	1.65e-05	3.03e-02
F11R		0.864(0.011)	1.65e-05	3.03e-02
C9orf169		0.864(0.033)	1.65e-05	3.03e-02
NEK10		-0.761(0.034)	7.92e-05	9.16e-02
SACS		-0.763(0.012)	7.65e-05	9.16e-02
PPAP2A		-0.767(0.027)	7.13e-05	9.16e-02
RP11-529K1.4		-0.768(0.013)	6.96e-05	9.16e-02
SYDE1		-0.773(0.008)	6.31e-05	9.16e-02
DENND5A		-0.774(0.010)	6.22e-05	9.16e-02
TUBB2A		-0.774(0.025)	6.22e-05	9.16e-02
LEPRE1		-0.778(0.013)	6.00e-05	9.16e-02
AP1M1		-0.780(0.010)	5.65e-05	9.16e-02
SH2B3		-0.784(0.014)	5.22e-05	9.16e-02
PABPC4		-0.788(0.128)	4.83e-05	9.16e-02
IKBIP		-0.788(0.013)	4.78e-05	9.16e-02
STX2		-0.788(0.014)	4.65e-05	9.16e-02
RAB8B		-0.789(0.020)	4.52e-05	9.16e-02
C19orf71		-0.791(0.042)	4.44e-05	9.16e-02
ST3GAL2		-0.793(0.013)	4.22e-05	9.16e-02
GPAM		-0.799(0.065)	3.74e-05	9.16e-02
KIAA0196		-0.816(0.179)	2.35e-05	9.16e-02
MDM1		-0.821(0.171)	2.22e-05	9.16e-02
LIX1L		-0.835(0.007)	1.74e-05	9.16e-02