

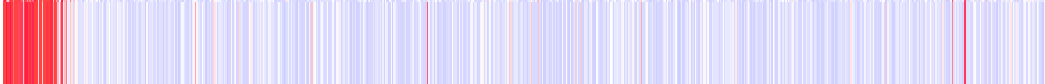
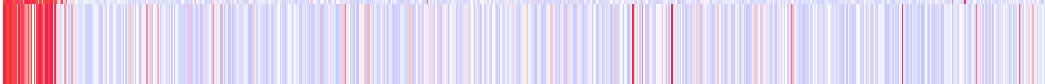
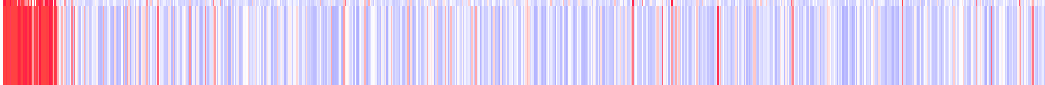
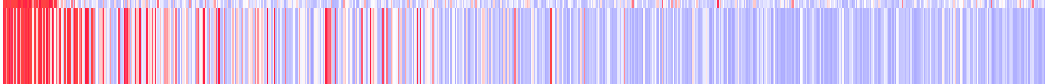
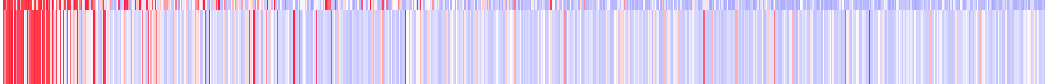
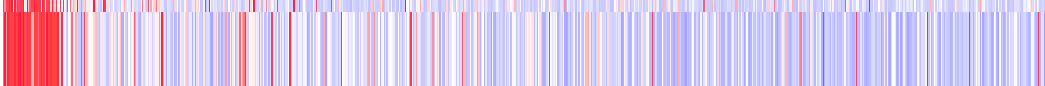
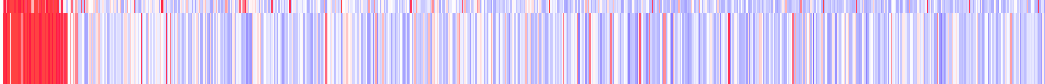
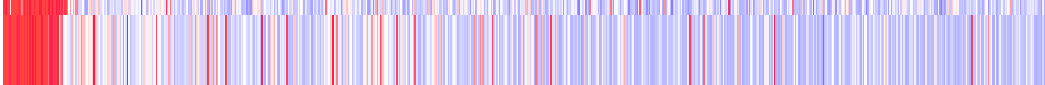
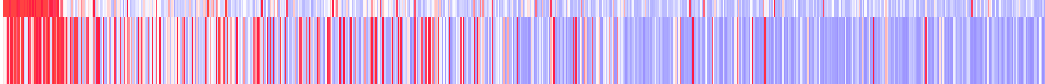
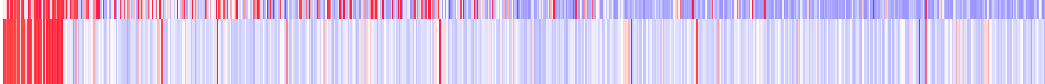
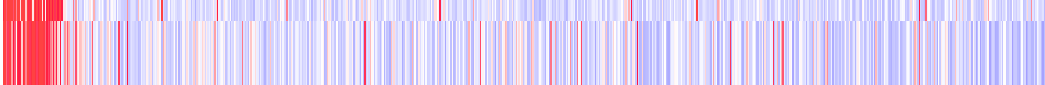
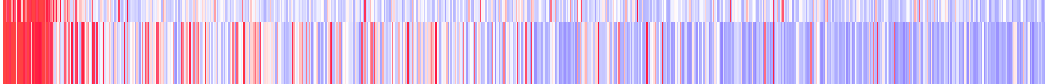
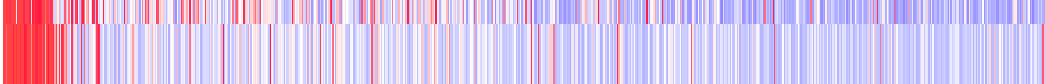
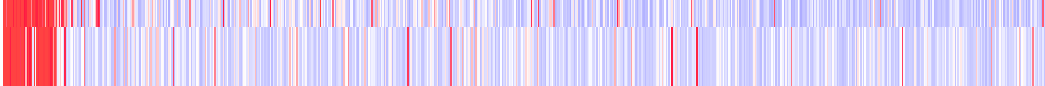
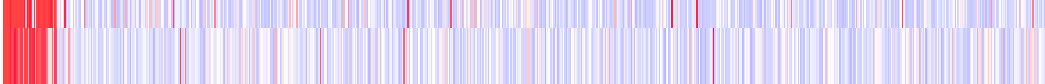
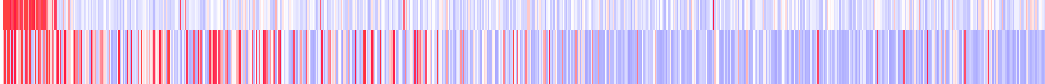
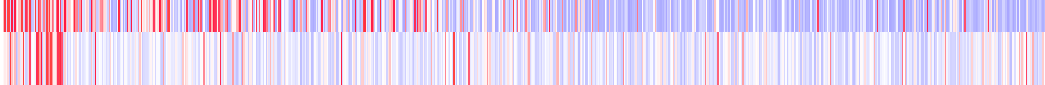
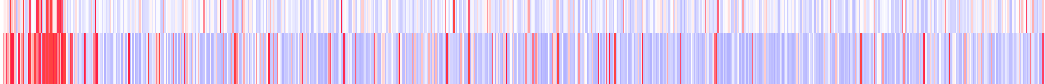
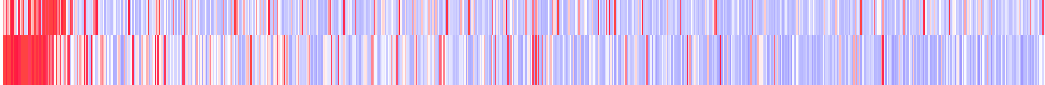
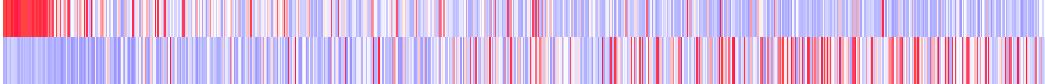
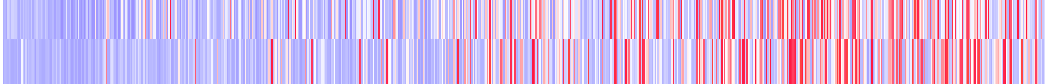
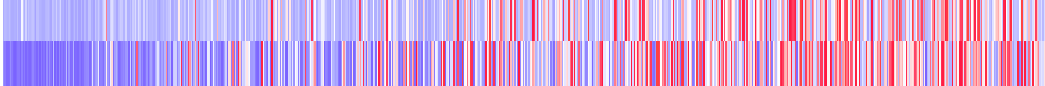
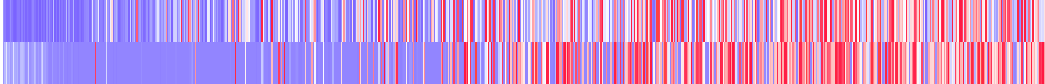
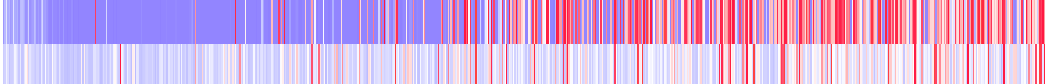
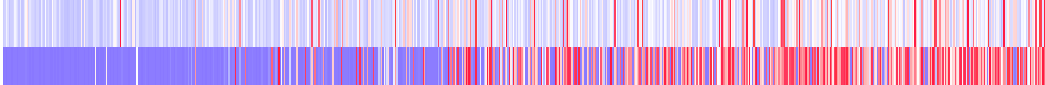
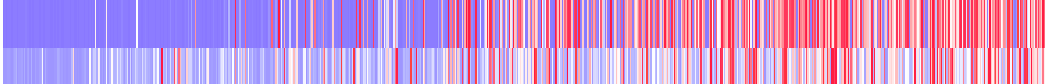
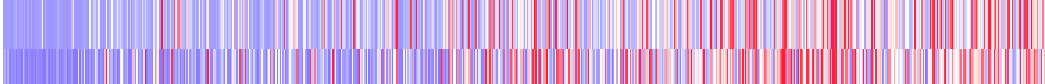

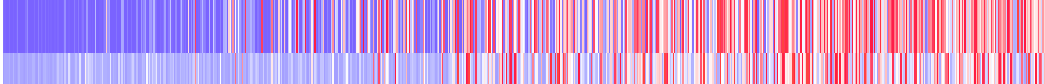
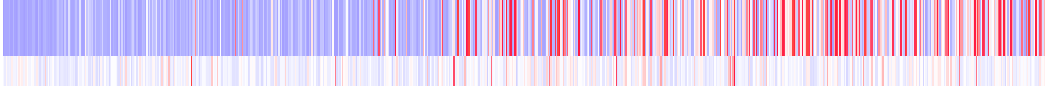
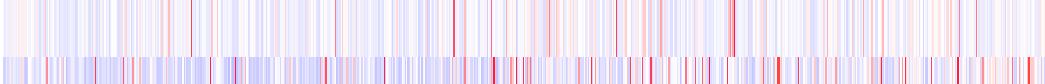
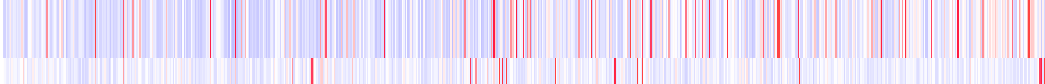
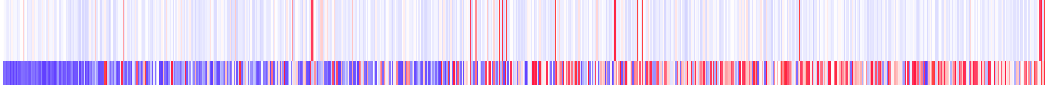
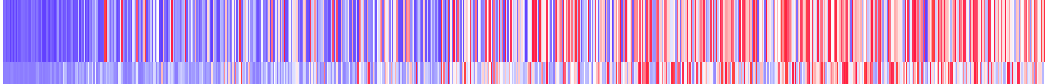
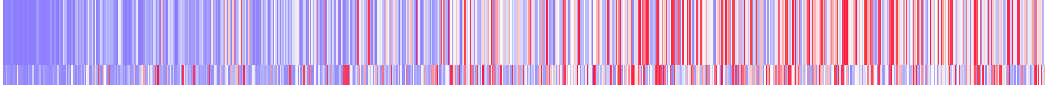
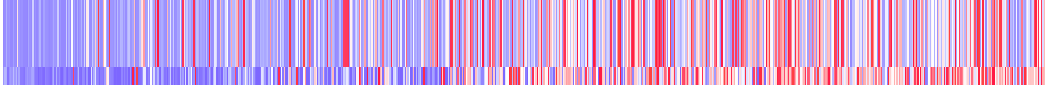
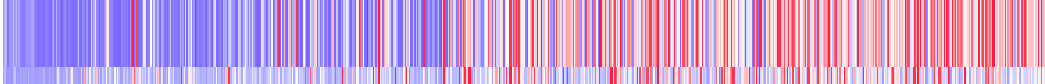
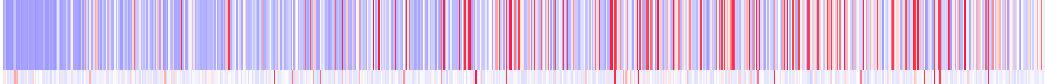
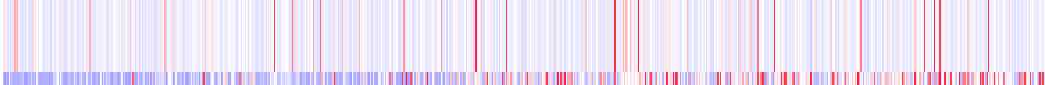


C6 vs Gene Expression

C6		IC(Δ)	P-val	FDR
C19orf71		0.944(0.033)	4.35e-07	1.00e-02
MFSD12		0.918(0.032)	4.35e-07	1.00e-02
LYST-IT1		0.830(0.022)	5.22e-06	3.68e-02
LYST		0.822(0.016)	6.52e-06	3.68e-02
SH2B3		0.813(0.011)	8.26e-06	3.68e-02
PFKFB4		0.805(0.032)	8.70e-06	3.68e-02
KAT2B		0.792(0.015)	1.13e-05	3.68e-02
BIN3		0.788(0.016)	1.13e-05	3.68e-02
AP1S2		0.785(0.016)	1.17e-05	3.68e-02
ZCCHC24		0.776(0.022)	1.39e-05	3.68e-02
INPP5F		0.773(0.018)	1.44e-05	3.68e-02
ATOX1		0.770(0.017)	1.48e-05	3.68e-02
OSTM1		0.764(0.016)	1.65e-05	3.68e-02
SDC3		0.762(0.017)	1.70e-05	3.68e-02
LDHAP2		0.759(0.019)	1.83e-05	3.68e-02
TBC1D7		0.755(0.018)	1.87e-05	3.68e-02
FRMD4A		0.754(0.026)	1.91e-05	3.68e-02
PRR4		0.754(0.045)	1.91e-05	3.68e-02
RCAN1		0.752(0.024)	2.00e-05	3.68e-02
ZNF697		0.750(0.018)	2.09e-05	3.68e-02
RAB11FIP1		-0.620(0.009)	7.74e-05	7.27e-02
MKRN2-AS1		-0.622(0.029)	7.48e-05	7.27e-02
RP11-195E2.1		-0.623(0.015)	7.44e-05	7.27e-02
ESRP1		-0.625(0.006)	7.26e-05	7.27e-02
TOB1		-0.626(0.022)	7.26e-05	7.27e-02
C1orf172		-0.626(0.006)	7.22e-05	7.27e-02
CASZ1		-0.628(0.015)	6.96e-05	7.27e-02
RP4-537K17.2		-0.628(0.019)	6.96e-05	7.27e-02
MARVELD2		-0.628(0.006)	6.92e-05	7.27e-02
RP11-517C16.2		-0.632(0.020)	6.52e-05	7.27e-02
SERPINE3		-0.634(0.080)	6.44e-05	7.27e-02
TRAF3IP3		-0.635(0.073)	6.35e-05	7.27e-02
SERHL2		-0.639(0.046)	5.96e-05	7.27e-02
INADL		-0.647(0.015)	5.39e-05	7.27e-02
EPB41L4B		-0.647(0.012)	5.39e-05	7.27e-02
PITPNM3		-0.653(0.022)	4.83e-05	7.27e-02
F11R		-0.654(0.011)	4.83e-05	7.27e-02
KIAA1161		-0.656(0.014)	4.70e-05	7.27e-02
HSD17B1		-0.665(0.048)	4.22e-05	7.27e-02
REM2		-0.679(0.020)	3.70e-05	7.27e-02