
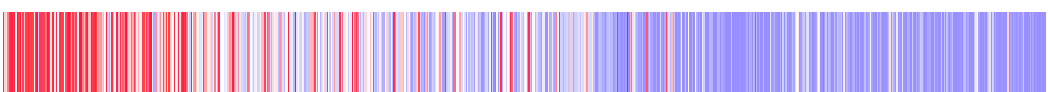
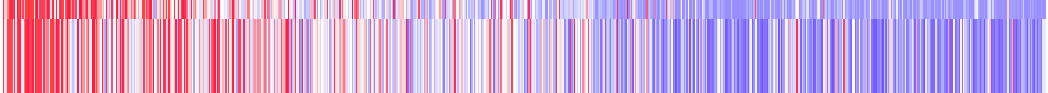
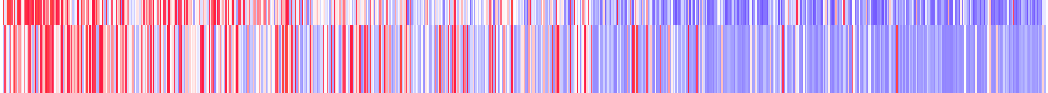
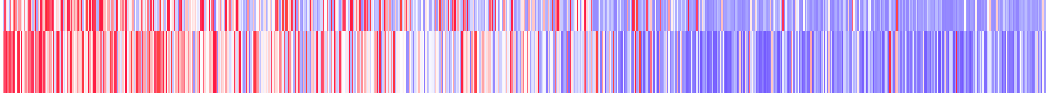
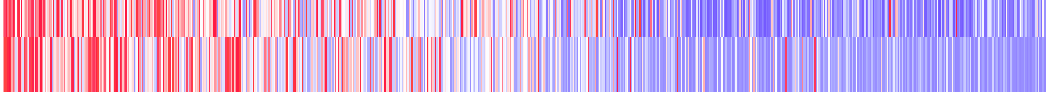
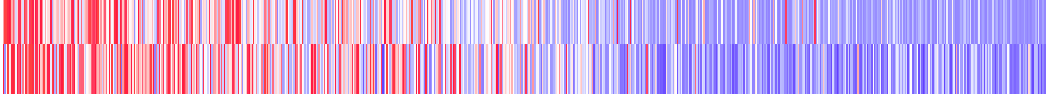
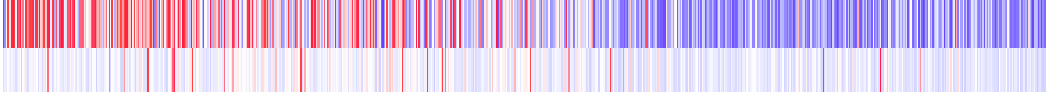
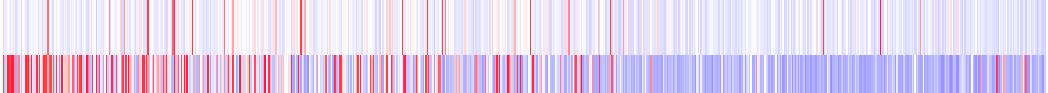
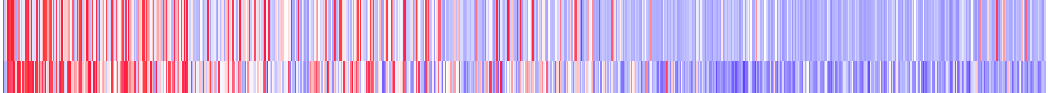
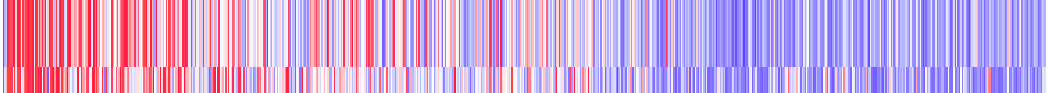
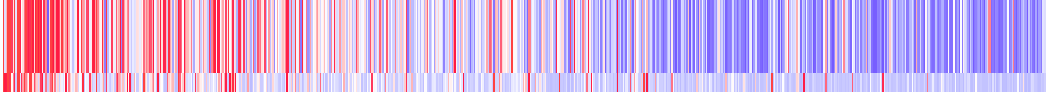
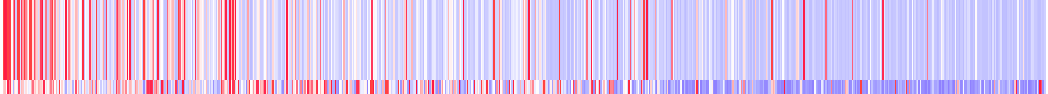
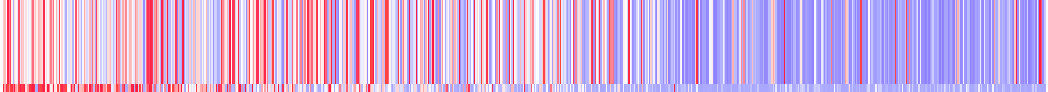
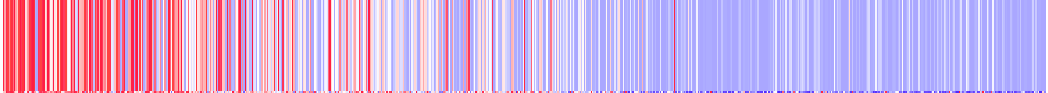
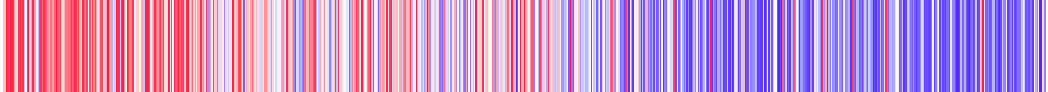
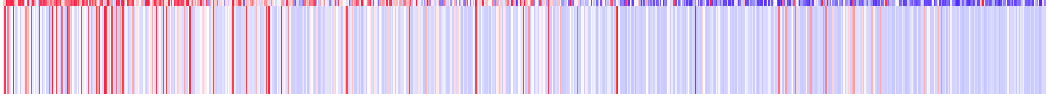
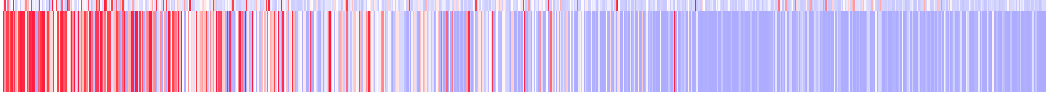
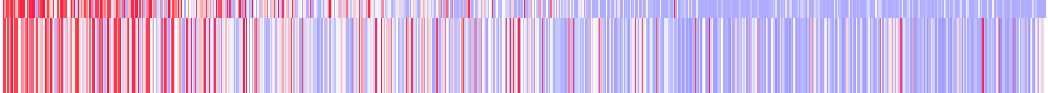
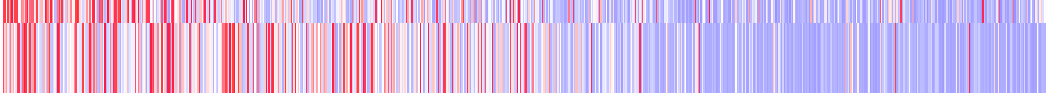
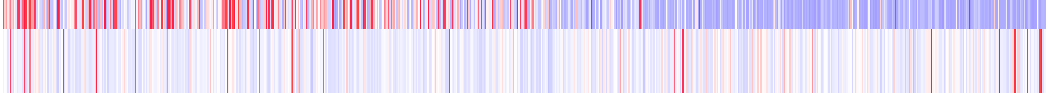
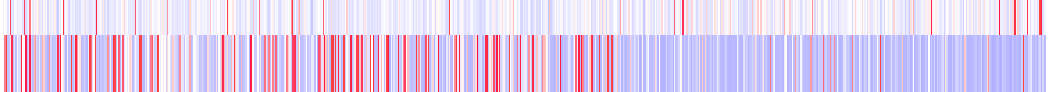
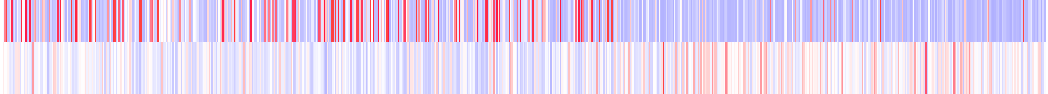
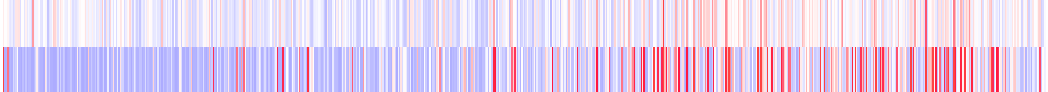
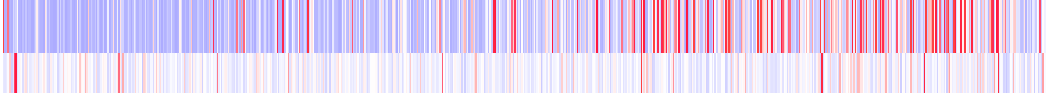
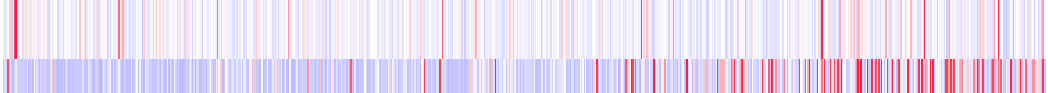
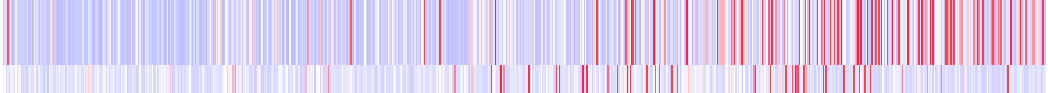
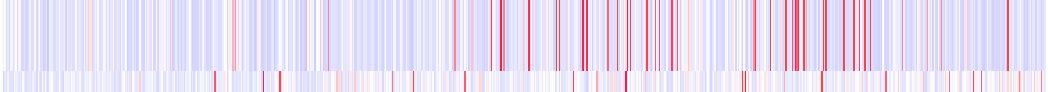
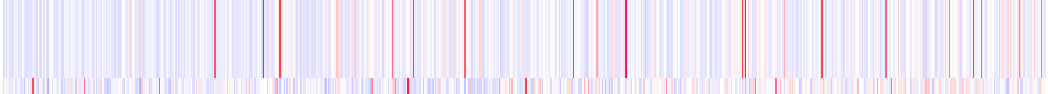
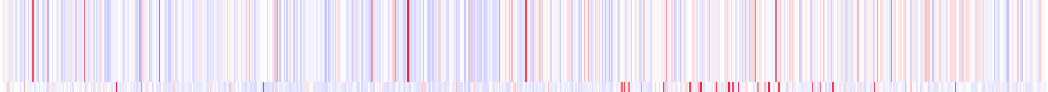
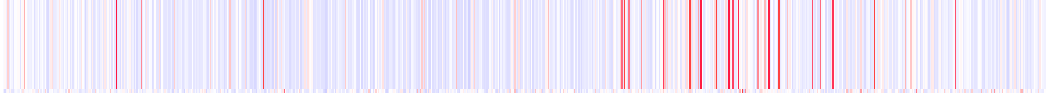
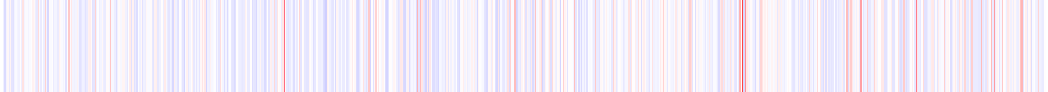
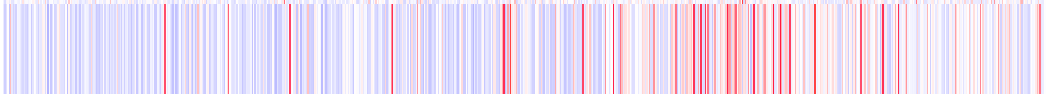
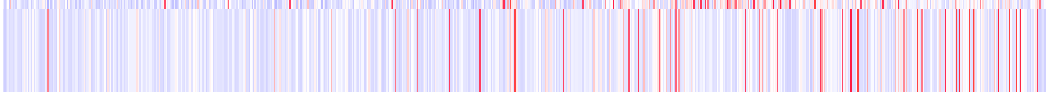
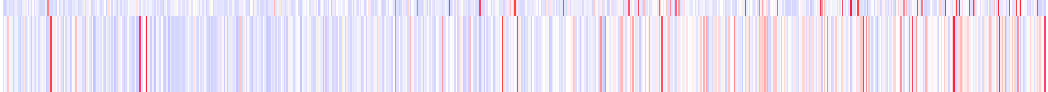
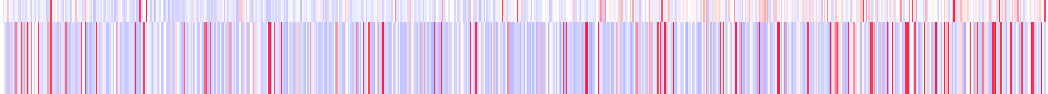
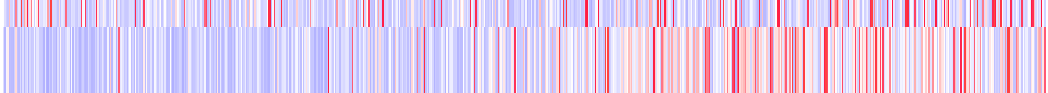
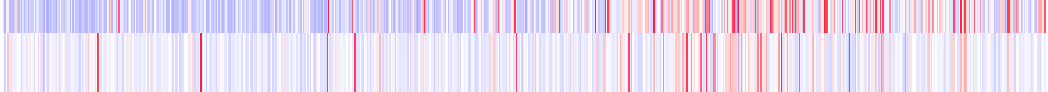
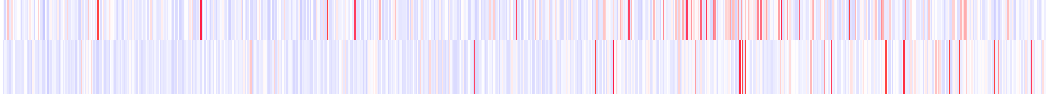
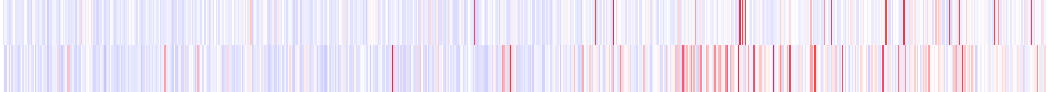
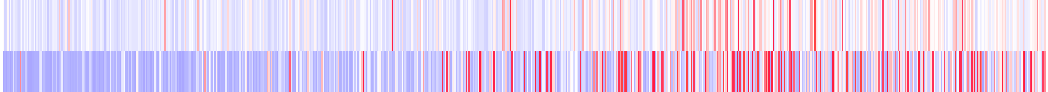


C7 vs Gene Expression

C7		IC( $\Delta$ )	P-val	FDR
FOSL1		0.770(0.008)	1.35e-05	6.06e-02
ELK3		0.744(0.009)	2.13e-05	6.06e-02
DCBLD2		0.738(0.015)	2.26e-05	6.06e-02
BCAR3		0.738(0.012)	2.35e-05	6.06e-02
PLAUR		0.726(0.014)	2.87e-05	6.06e-02
DRAP1		0.718(0.008)	3.17e-05	6.06e-02
SEC61G		0.704(0.047)	3.78e-05	6.06e-02
SMURF2		0.700(0.010)	4.09e-05	6.06e-02
C11orf68		0.695(0.008)	4.31e-05	6.06e-02
RP11-394J1.2		0.695(0.009)	4.31e-05	6.06e-02
AC137932.6		0.687(0.036)	4.91e-05	6.06e-02
ELL2		0.686(0.019)	4.96e-05	6.06e-02
RAC2		0.684(0.008)	5.18e-05	6.06e-02
CAPN2		0.683(0.009)	5.18e-05	6.06e-02
MOK		0.681(0.050)	5.35e-05	6.06e-02
Z82188.1		0.681(0.008)	5.35e-05	6.06e-02
ERRFI1		0.679(0.015)	5.39e-05	6.06e-02
SLC22A4		0.677(0.020)	5.65e-05	6.06e-02
GPAM		0.676(0.150)	5.65e-05	6.06e-02
SELPLG		0.675(0.045)	5.83e-05	6.06e-02
DDX51		-0.633(0.037)	5.57e-05	4.93e-02
AC011747.7		-0.642(0.015)	5.00e-05	4.93e-02
KIAA0196-AS1		-0.651(0.099)	4.48e-05	4.93e-02
RP4-659J6.2		-0.652(0.018)	4.44e-05	4.93e-02
C19orf71		-0.654(0.039)	4.22e-05	4.93e-02
ORMDL3		-0.659(0.053)	3.96e-05	4.93e-02
AC019201.1		-0.659(0.042)	3.96e-05	4.93e-02
EYS		-0.664(0.050)	3.52e-05	4.93e-02
RECQL5		-0.668(0.041)	3.48e-05	4.93e-02
PI4KA		-0.672(0.038)	3.39e-05	4.93e-02
RAB6C		-0.672(0.054)	3.35e-05	4.93e-02
LRP6		-0.675(0.032)	3.26e-05	4.93e-02
AB019441.29		-0.686(0.067)	2.96e-05	4.93e-02
PLA2G6		-0.694(0.026)	2.74e-05	4.93e-02
FRS3		-0.697(0.077)	2.57e-05	4.93e-02
SERHL2		-0.702(0.062)	2.35e-05	4.93e-02
ZNF74		-0.719(0.049)	1.65e-05	4.93e-02
BTG2		-0.722(0.021)	1.48e-05	4.93e-02
STARD3		-0.746(0.082)	9.57e-06	4.93e-02
HSD17B1		-0.794(0.051)	2.17e-06	4.93e-02