
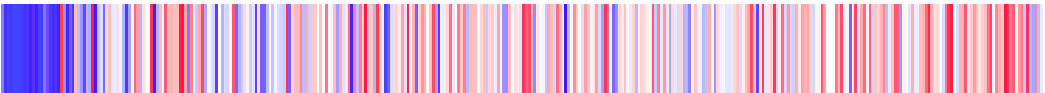
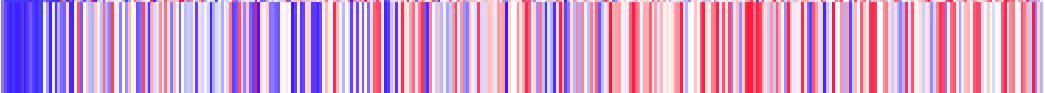
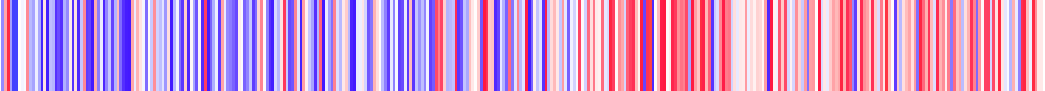
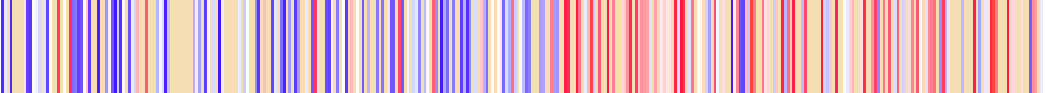
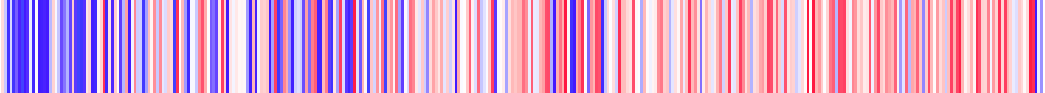
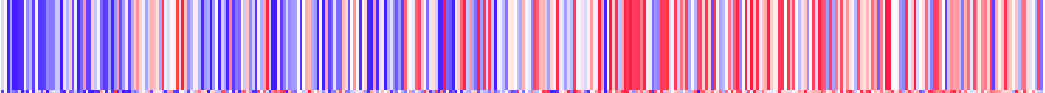

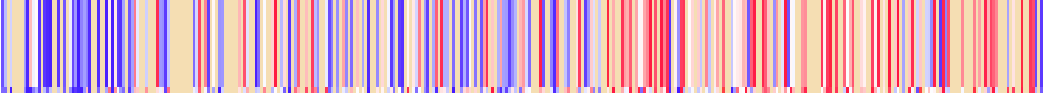
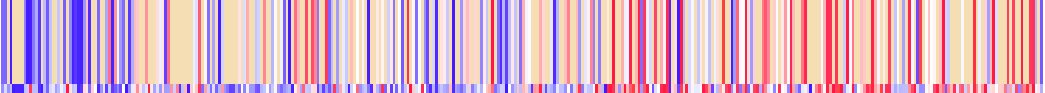
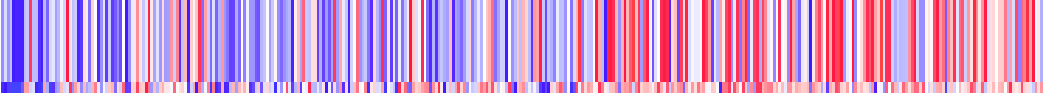
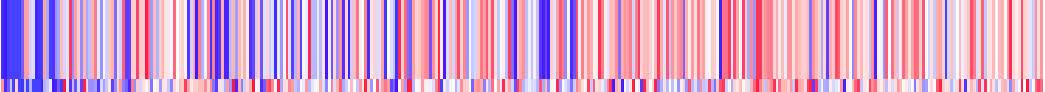
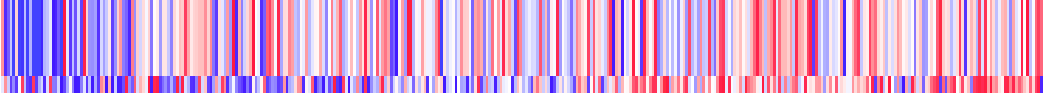
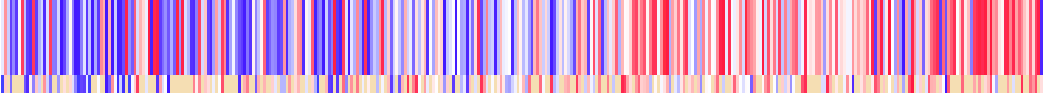
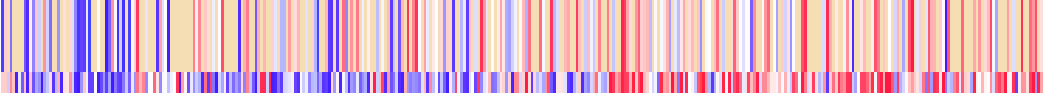
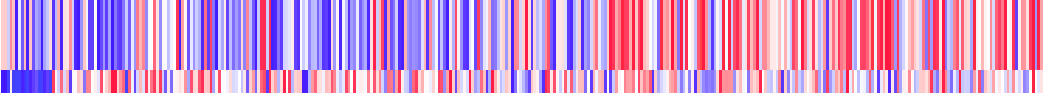
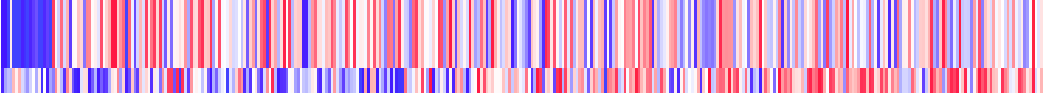
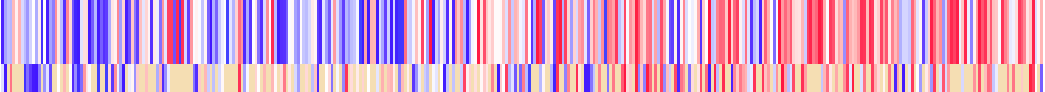
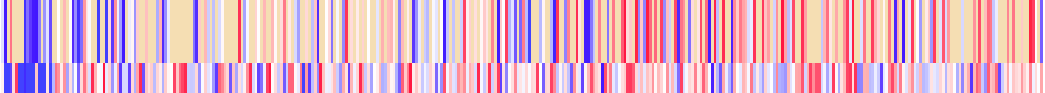
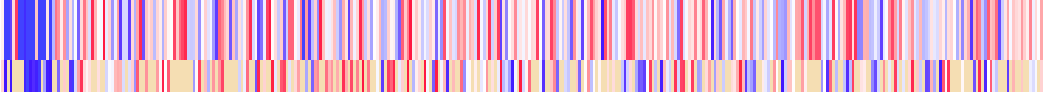
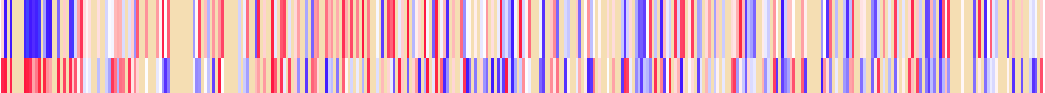
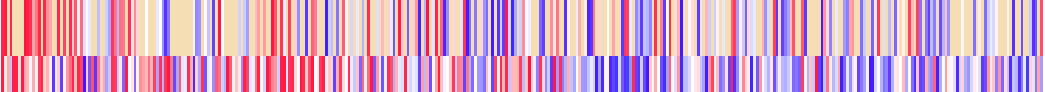
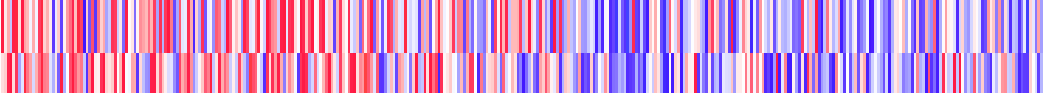
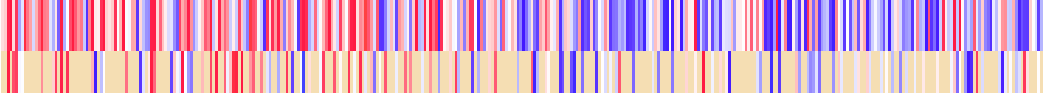
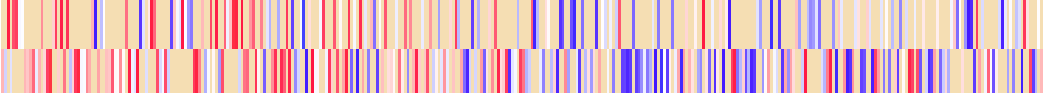
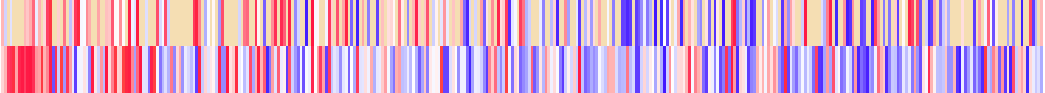
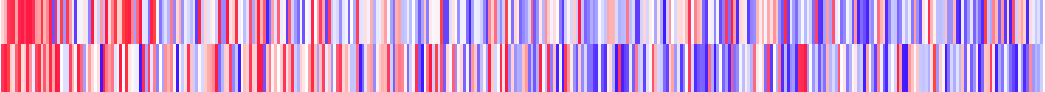
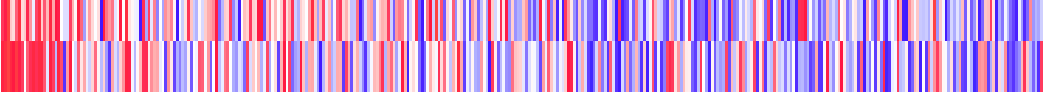
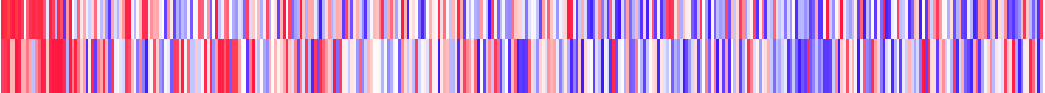
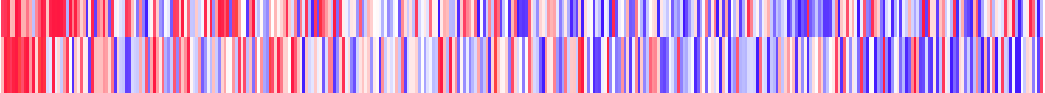
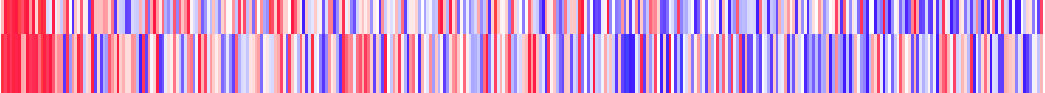
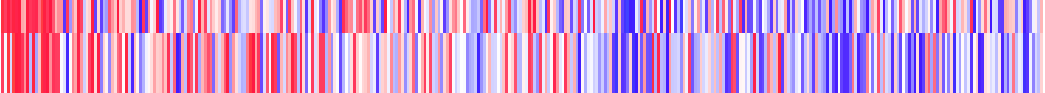
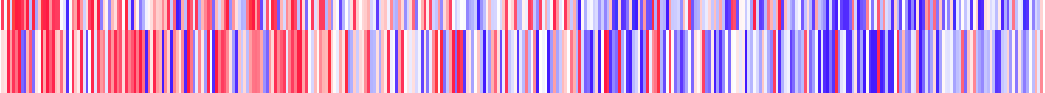
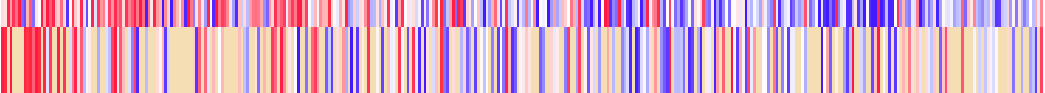
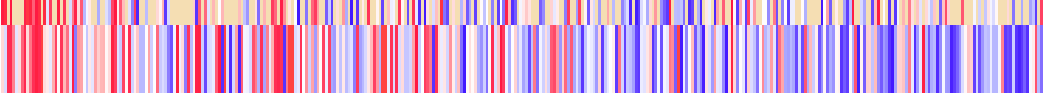
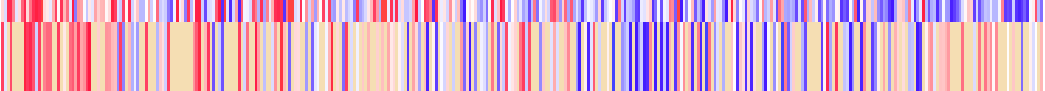
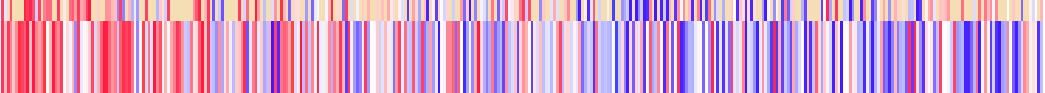
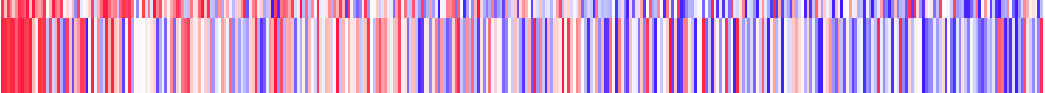
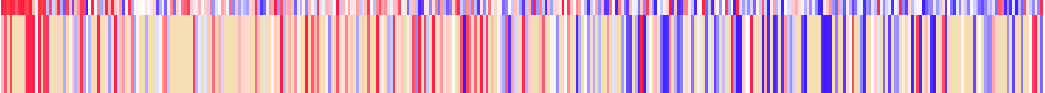
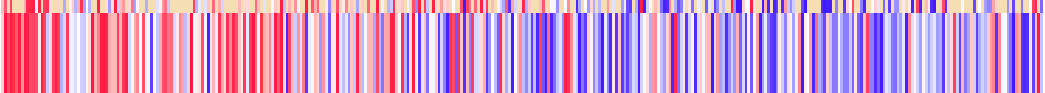



C1 vs Gene Dependency

C1		IC(Δ)	P-val	FDR
FOXA1		-0.604(0.023)	1.17e-06	2.22e-03
KDM1A		-0.531(0.023)	1.17e-06	2.22e-03
MYBL2		-0.523(0.023)	1.17e-06	2.22e-03
PRSS3P2		-0.516(0.019)	1.17e-06	2.22e-03
ADSL		-0.507(0.022)	1.17e-06	2.22e-03
SRRM1		-0.501(0.021)	1.17e-06	2.22e-03
COPS8		-0.501(0.024)	1.17e-06	2.22e-03
SPC25		-0.498(0.023)	1.17e-06	2.22e-03
GRPEL1		-0.493(0.015)	1.17e-06	2.22e-03
NUP93		-0.483(0.022)	2.34e-06	2.86e-03
PIK3CA		-0.483(0.033)	2.34e-06	2.86e-03
TFAP2C		-0.482(0.024)	2.34e-06	2.86e-03
CLTC		-0.476(0.022)	2.34e-06	2.86e-03
NUP43		-0.474(0.021)	2.34e-06	2.86e-03
HNRNPM		-0.473(0.021)	3.51e-06	3.20e-03
SPDEF		-0.472(0.028)	3.51e-06	3.20e-03
NSMCE1		-0.472(0.023)	3.51e-06	3.20e-03
NARS2		-0.472(0.017)	3.51e-06	3.20e-03
ESR1		-0.469(0.023)	4.69e-06	3.20e-03
ZNF652		-0.469(0.071)	4.69e-06	3.20e-03
PABPC1L2A		0.433(0.021)	7.73e-05	6.48e-02
CERKL		0.434(0.026)	7.61e-05	6.48e-02
IRF5		0.435(0.026)	7.38e-05	6.48e-02
MANBA		0.435(0.025)	7.38e-05	6.48e-02
MREG		0.436(0.022)	7.15e-05	6.48e-02
OR5T3		0.441(0.027)	5.51e-05	6.27e-02
PPP3R2		0.443(0.027)	4.57e-05	5.57e-02
RCN2		0.446(0.027)	3.98e-05	5.23e-02
GBGT1		0.447(0.031)	3.75e-05	5.23e-02
NTRK3		0.453(0.024)	2.93e-05	4.55e-02
ELAVL2		0.458(0.024)	1.29e-05	2.20e-02
AOC1		0.459(0.032)	1.17e-05	2.20e-02
TPH2		0.462(0.024)	1.05e-05	2.20e-02
PABPC1L2B		0.465(0.024)	9.37e-06	2.20e-02
ZIM3		0.480(0.024)	1.17e-06	3.33e-03
MSTO1		0.486(0.017)	1.17e-06	3.33e-03
ZSCAN32		0.491(0.017)	1.17e-06	3.33e-03
KPNA1		0.497(0.026)	1.17e-06	3.33e-03
APOF		0.506(0.018)	1.17e-06	3.33e-03
CXCL13		0.516(0.024)	1.17e-06	3.33e-03