

positive regulation of synapse assembly

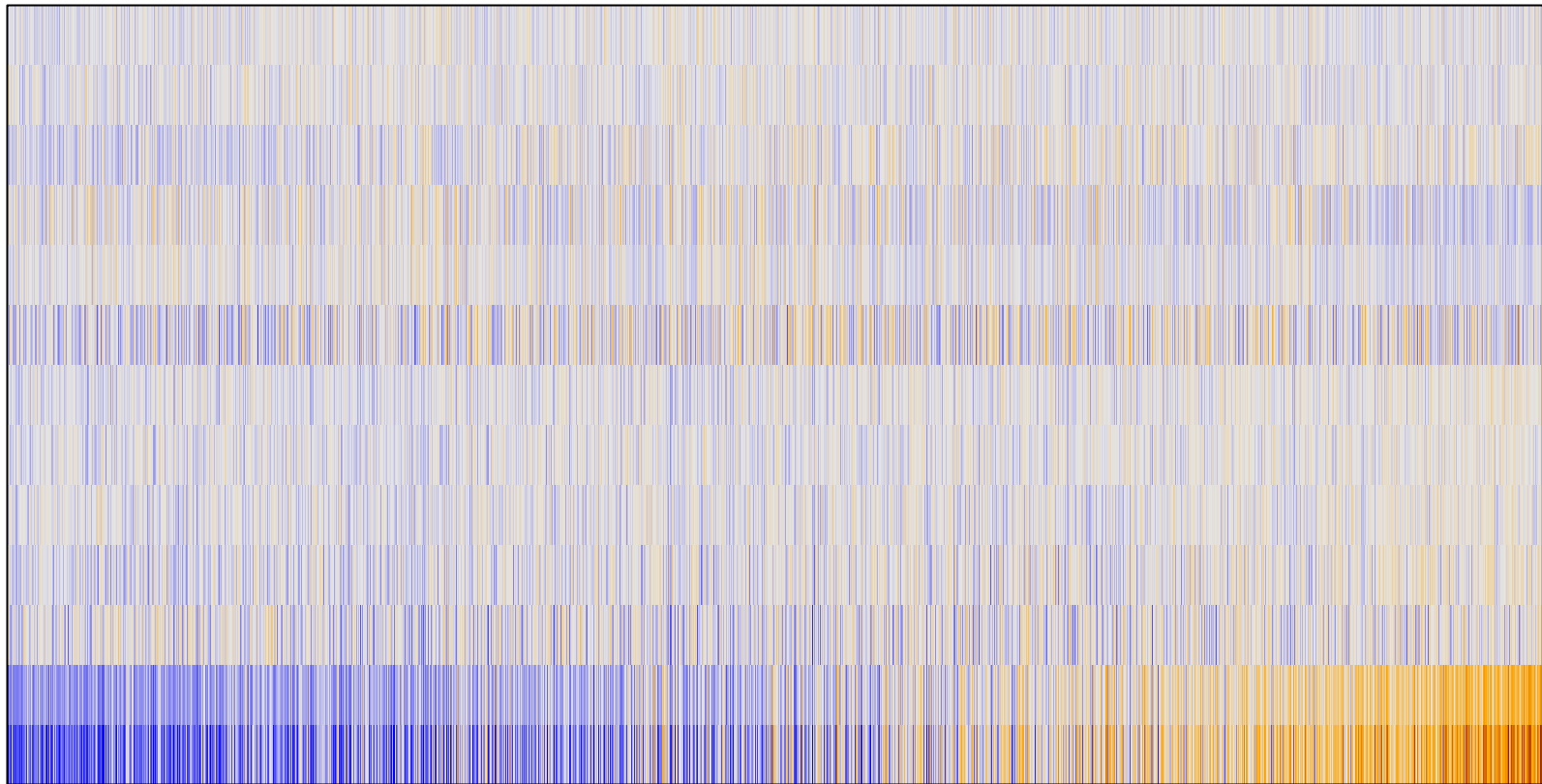
GSZ

2.91
1.18
-0.55
-2.27

- A*
- AC*
- ACF*
- AF*
- CF*
- F*
- FJ*
- J*
- N*

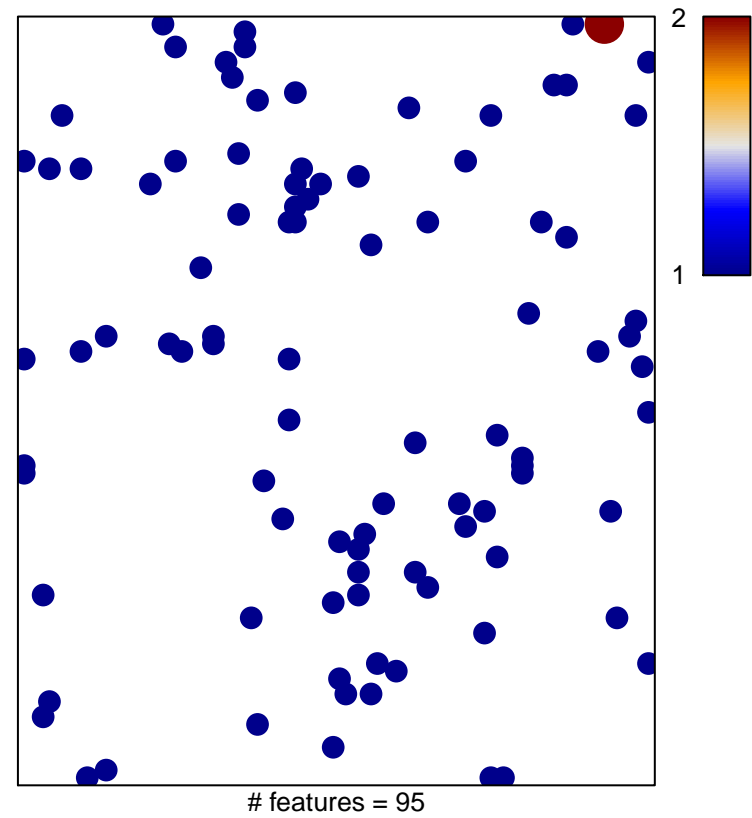
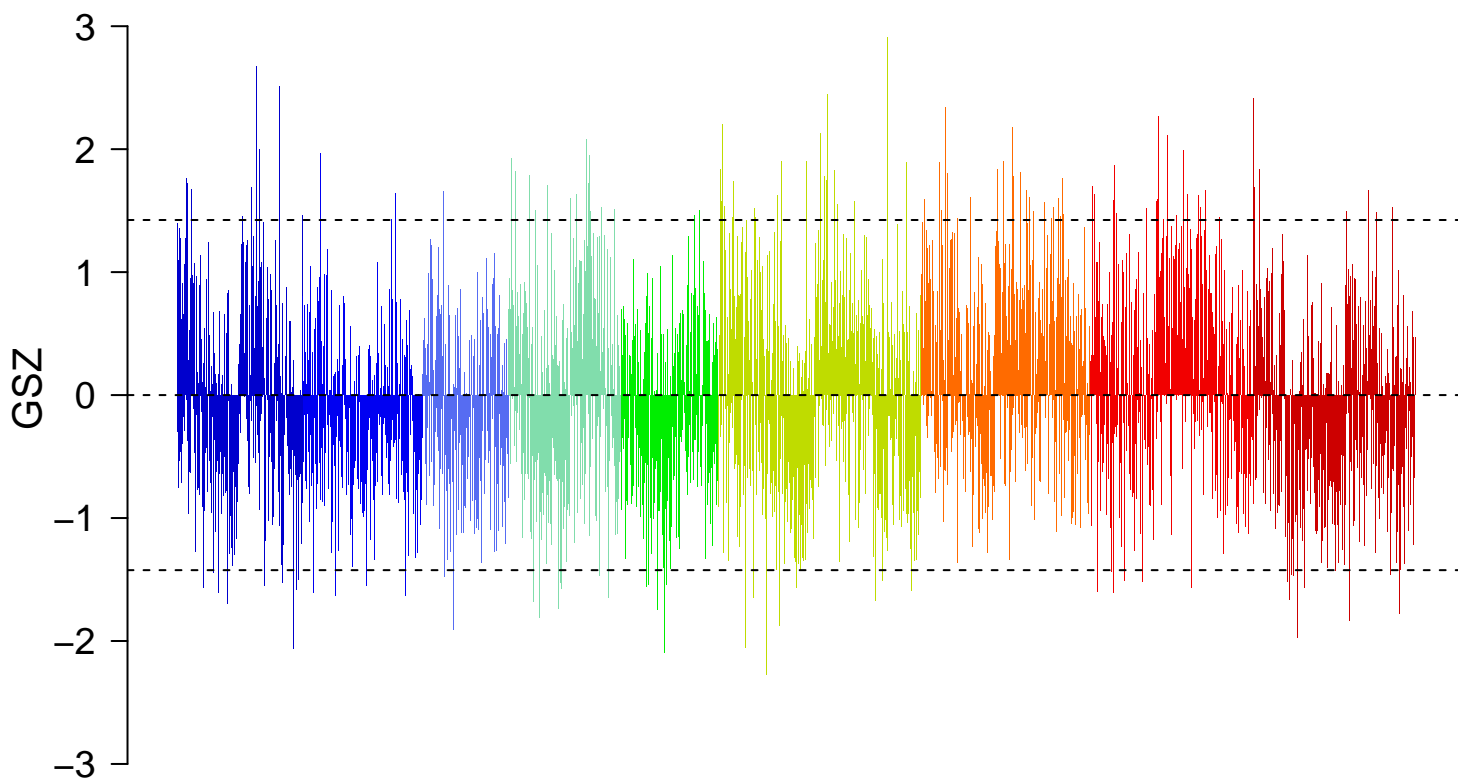


OXTR
MECP2
EPHB1
SLITRK1
BDNF
IL1RAP
ADGRL1
LOC101930123
UBE2V2
ADNP
CLSTN1
LRRN3
LRRN3



-1.2 Δe 1.8

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