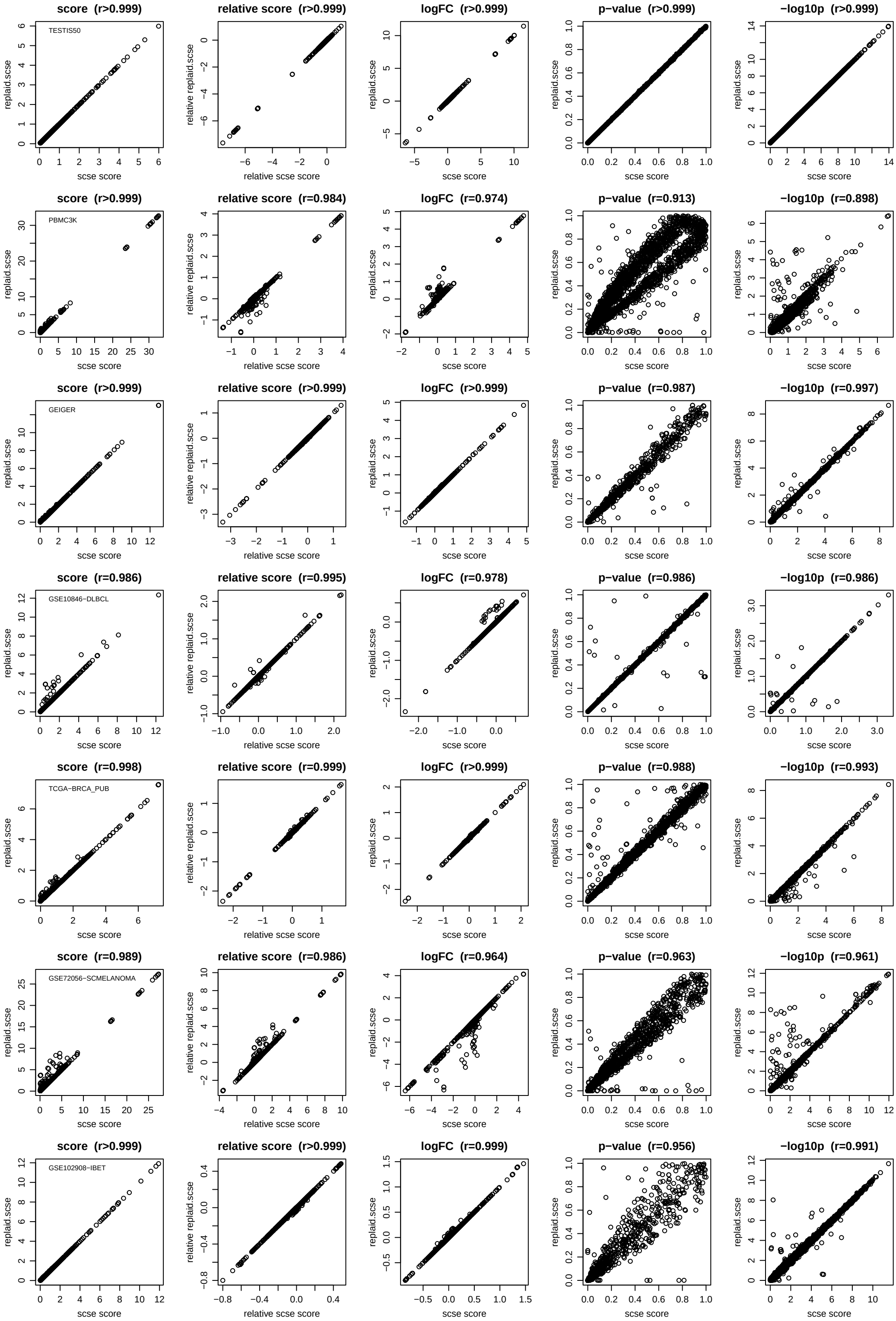
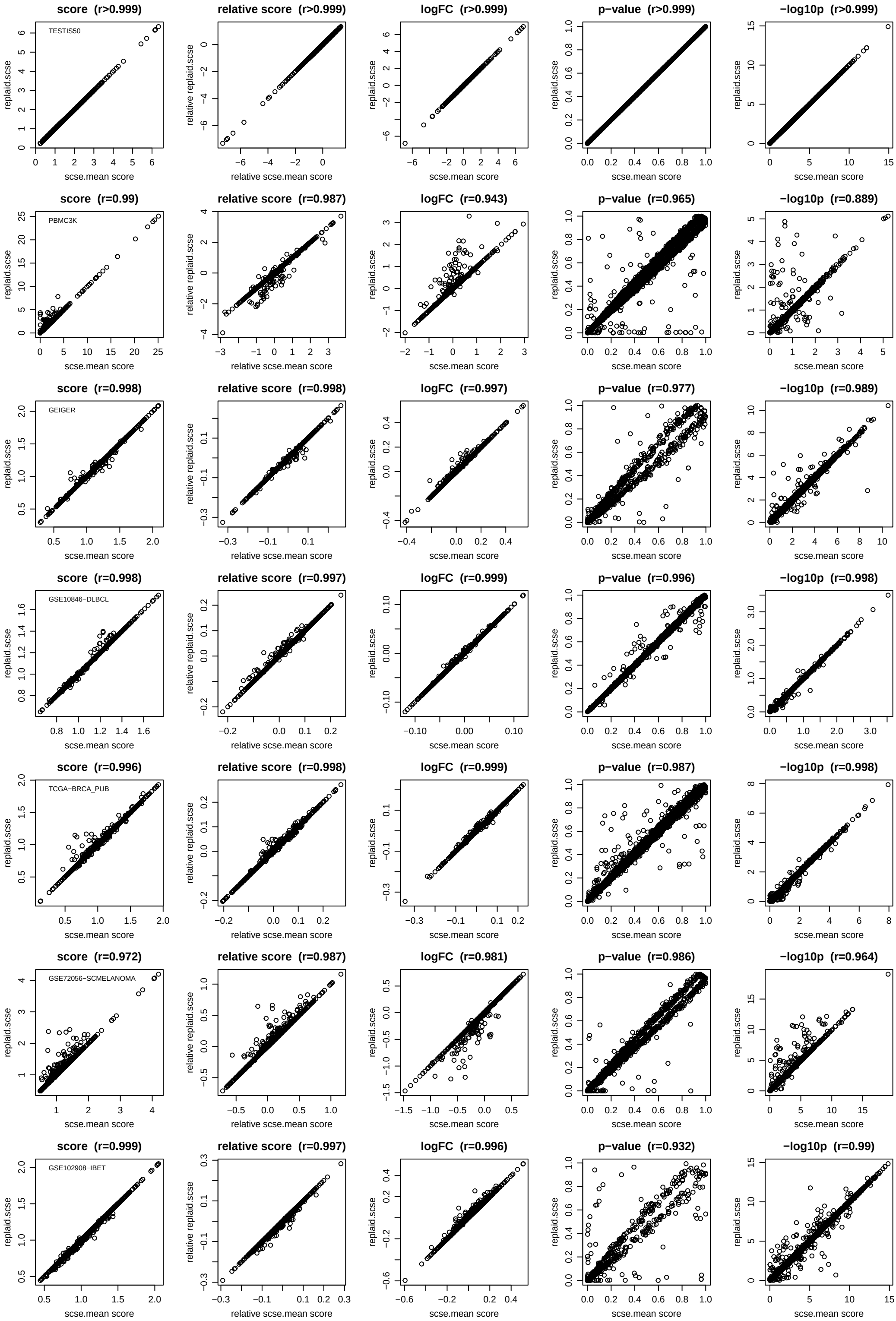
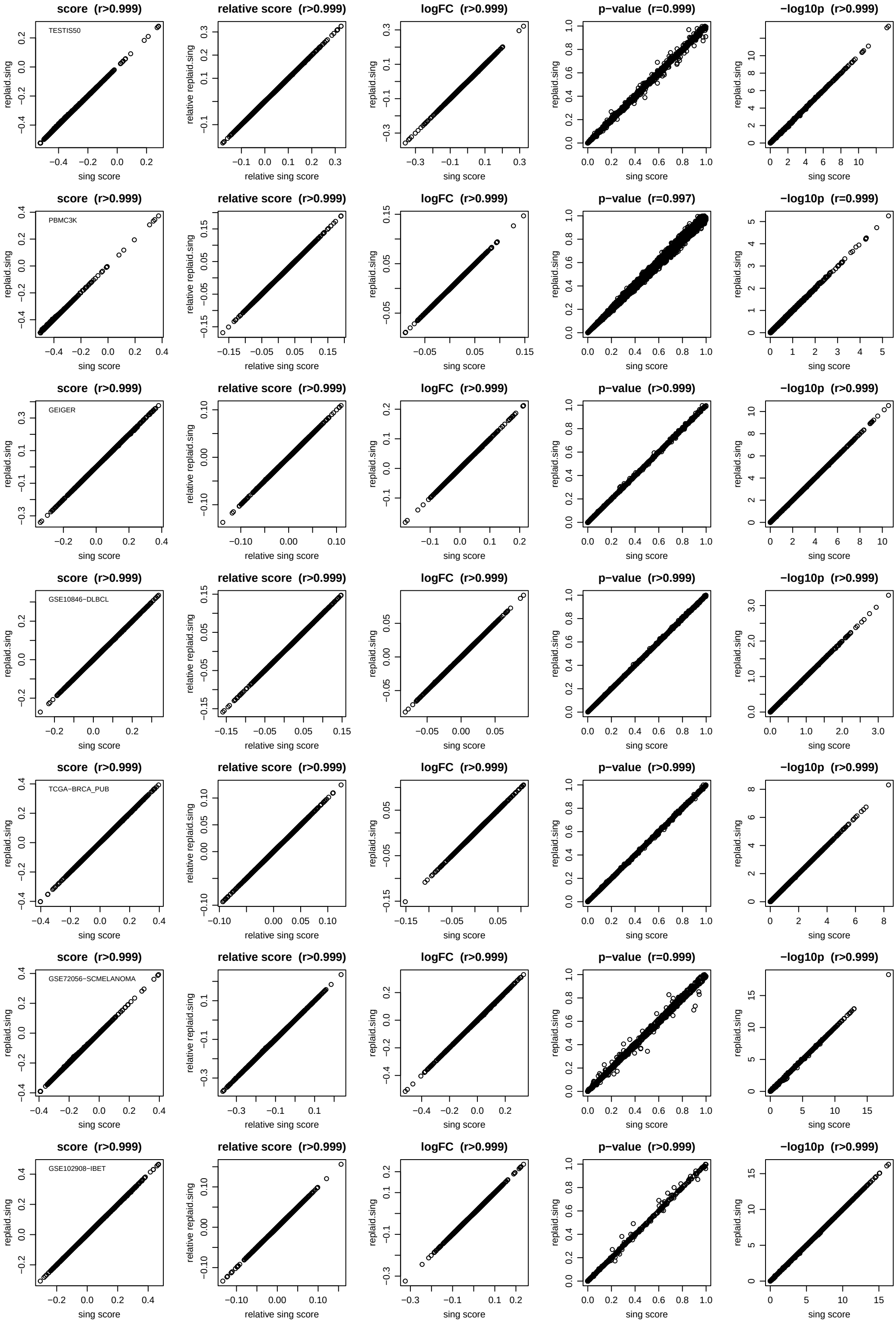
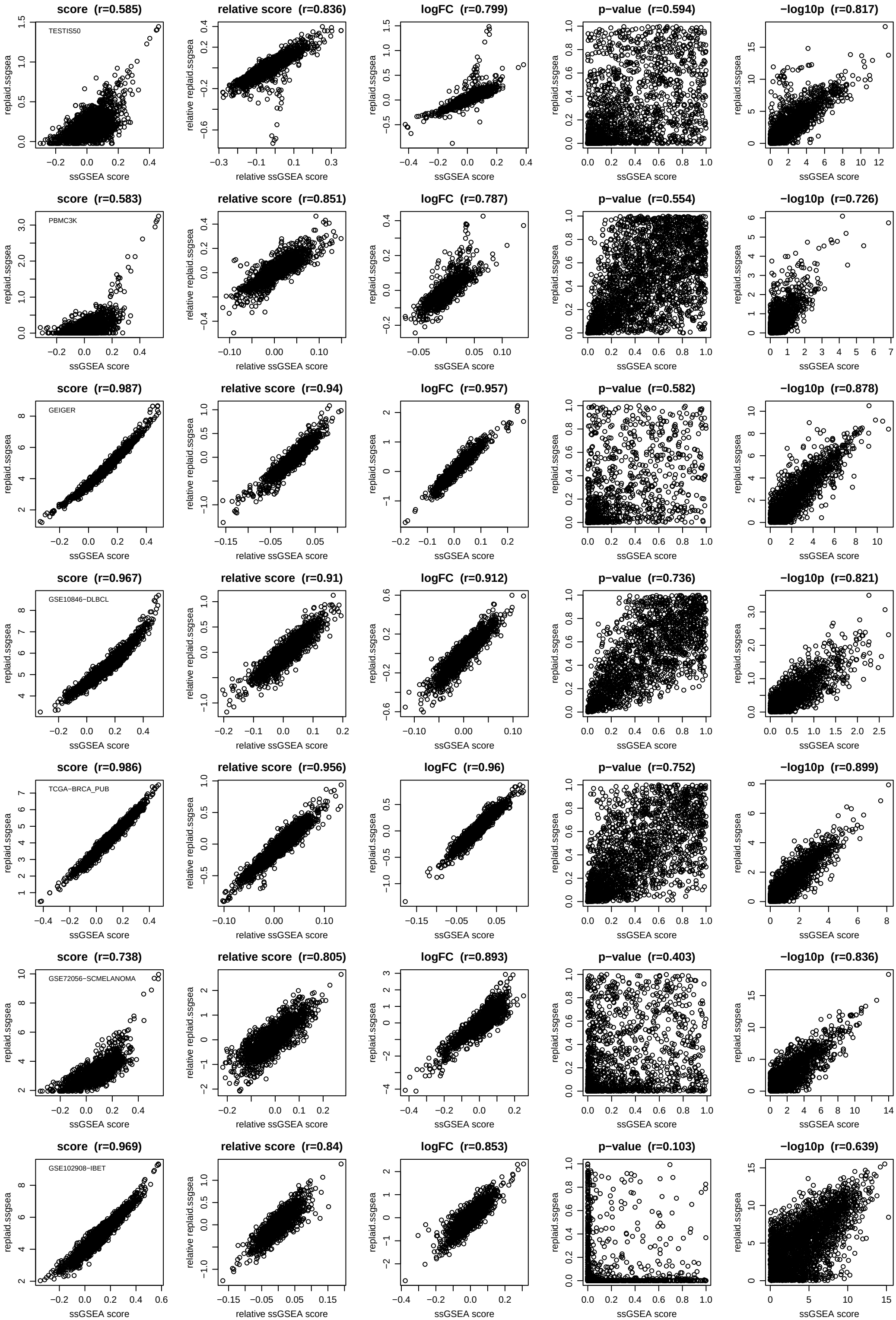

score (r>0.999)
relative score (r>0.999)
logFC (r>0.999)
p−value (r>0.999)
−log10p (r>0.999)
TESTIS50
score (r>0.999)
relative score (r=0.984)
logFC (r=0.974)
p−value (r=0.913)
−log10p (r=0.898)
PBMC3K
score (r>0.999)
relative score (r>0.999)
logFC (r>0.999)
p−value (r=0.987)
−log10p (r=0.997)
GEIGER
score (r=0.986)
relative score (r=0.995)
logFC (r=0.978)
p−value (r=0.986)
−log10p (r=0.986)
GSE10846−DLBCL
score (r=0.998)
relative score (r=0.999)
logFC (r>0.999)
p−value (r=0.988)
−log10p (r=0.993)
TCGA−BRCA_PUB
score (r=0.989)
relative score (r=0.986)
logFC (r=0.964)
p−value (r=0.963)
−log10p (r=0.961)
GSE72056−SCMELANOMA
score (r>0.999)
relative score (r>0.999)
logFC (r=0.999)
p−value (r=0.956)
−log10p (r=0.991)
GSE102908−IBET
replaid.scse
relative replaid.scse
scse score
relative scse score


score (r>0.999)
relative score (r>0.999)
logFC (r>0.999)
p−value (r>0.999)
−log10p (r>0.999)
TESTIS50
score (r=0.99)
relative score (r=0.987)
logFC (r=0.943)
p−value (r=0.965)
−log10p (r=0.889)
PBMC3K
score (r=0.998)
relative score (r=0.998)
logFC (r=0.997)
p−value (r=0.977)
−log10p (r=0.989)
GEIGER
score (r=0.998)
relative score (r=0.997)
logFC (r=0.999)
p−value (r=0.996)
−log10p (r=0.998)
GSE10846−DLBCL
score (r=0.996)
relative score (r=0.998)
logFC (r=0.999)
p−value (r=0.987)
−log10p (r=0.998)
TCGA−BRCA_PUB
score (r=0.972)
relative score (r=0.987)
logFC (r=0.981)
p−value (r=0.986)
−log10p (r=0.964)
GSE72056−SCMELANOMA
score (r=0.999)
relative score (r=0.997)
logFC (r=0.996)
p−value (r=0.932)
−log10p (r=0.99)
GSE102908−IBET
replaid.scse
relative replaid.scse
scse.mean score
relative scse.mean score


score (r>0.999)
relative score (r>0.999)
logFC (r>0.999)
p−value (r=0.999)
−log10p (r>0.999)
TESTIS50
PBMC3K
GEIGER
GSE10846−DLBCL
TCGA−BRCA_PUB
GSE72056−SCMELANOMA
GSE102908−IBET
replaid.sing
relative replaid.sing
sing score
relative sing score


score (r=0.585)
relative score (r=0.836)
logFC (r=0.799)
p−value (r=0.594)
−log10p (r=0.817)
TESTIS50
score (r=0.583)
relative score (r=0.851)
logFC (r=0.787)
p−value (r=0.554)
−log10p (r=0.726)
PBMC3K
score (r=0.987)
relative score (r=0.94)
logFC (r=0.957)
p−value (r=0.582)
−log10p (r=0.878)
GEIGER
score (r=0.967)
relative score (r=0.91)
logFC (r=0.912)
p−value (r=0.736)
−log10p (r=0.821)
GSE10846−DLBCL
score (r=0.986)
relative score (r=0.956)
logFC (r=0.96)
p−value (r=0.752)
−log10p (r=0.899)
TCGA−BRCA_PUB
score (r=0.738)
relative score (r=0.805)
logFC (r=0.893)
p−value (r=0.403)
−log10p (r=0.836)
GSE72056−SCMELANOMA
score (r=0.969)
relative score (r=0.84)
logFC (r=0.853)
p−value (r=0.103)
−log10p (r=0.639)
GSE102908−IBET
replaid.ssgsea
relative replaid.ssgsea
ssGSEA score
relative ssGSEA score