# Forest Plot



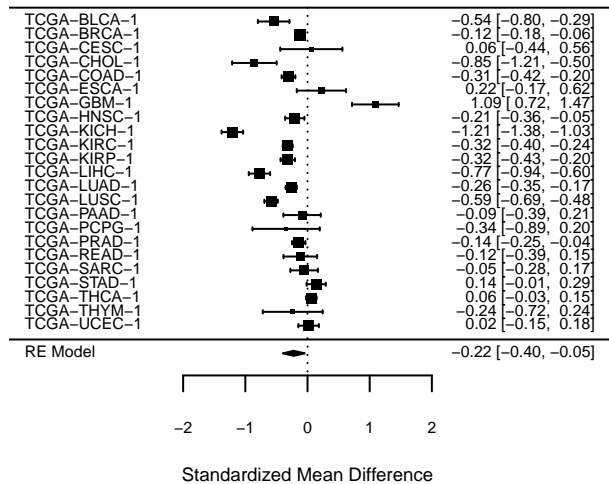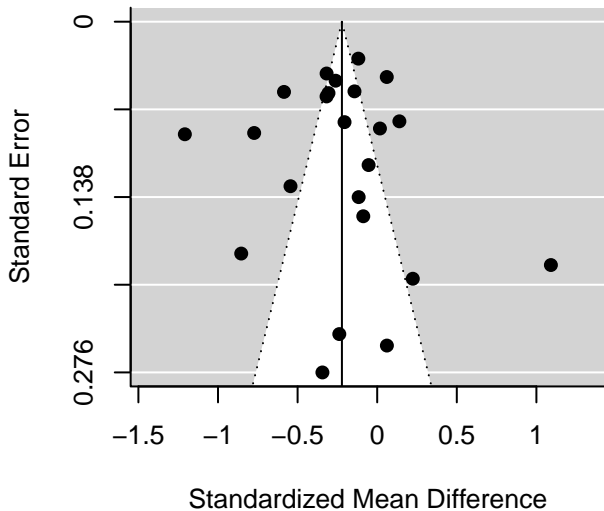| Study | SMD [95% CI] |
|---|---|
| TCGA−BLCA−1 | −0.54 [−0.80, −0.29] |
| TCGA−BRCA−1 | −0.12 [−0.18, −0.06] |
| TCGA−CESC−1 | 0.06 [−0.44, 0.56] |
| TCGA−CHOL−1 | −0.85 [−1.21, −0.50] |
| TCGA−COAD−1 | −0.31 [−0.42, −0.20] |
| TCGA−ESCA−1 | 0.22 [−0.17, 0.62] |
| TCGA−GBM−1 | 1.09 [0.72, 1.47] |
| TCGA−HNSC−1 | −0.21 [−0.36, −0.05] |
| TCGA−KICH−1 | −1.21 [−1.38, −1.03] |
| TCGA−KIRC−1 | −0.32 [−0.40, −0.24] |
| TCGA−KIRP−1 | −0.32 [−0.43, −0.20] |
| TCGA−LIHC−1 | −0.77 [−0.94, −0.60] |
| TCGA−LUAD−1 | −0.26 [−0.35, −0.17] |
| TCGA−LUSC−1 | −0.59 [−0.69, −0.48] |
| TCGA−PAAD−1 | −0.09 [−0.39, 0.21] |
| TCGA−PCPG−1 | −0.34 [−0.89, 0.20] |
| TCGA−PRAD−1 | −0.14 [−0.25, −0.04] |
| TCGA−READ−1 | −0.12 [−0.39, 0.15] |
| TCGA−SARC−1 | −0.05 [−0.28, 0.17] |
| TCGA−STAD−1 | 0.14 [−0.01, 0.29] |
| TCGA−THCA−1 | 0.06 [−0.03, 0.15] |
| TCGA−THYM−1 | −0.24 [−0.72, 0.24] |
| TCGA−UCEC−1 | 0.02 [−0.15, 0.18] |
| RE Model | −0.22 [−0.40, −0.05] |

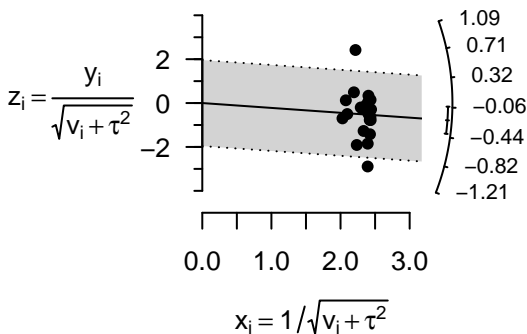Standardized Mean Difference

# Funnel Plot



Standardized Mean Difference

RE Model for All Studies (Q = 379.18, df = 22, p = 8.02E−67; $I^2$ = 97.2%)
RE Model for All Studies (beta = −0.22, se = 0.09, zval = −2.48; P = 1.32E−02)
STAT3.

# Radial Plot



$$z_i = \frac{y_i}{\sqrt{v_i + \tau^2}}$$

1.09
0.71
0.32
−0.06
−0.44
−0.82
−1.21

$$x_i = 1/\sqrt{v_i + \tau^2}$$

# Standardized Residuals



Study