## Forest Plot



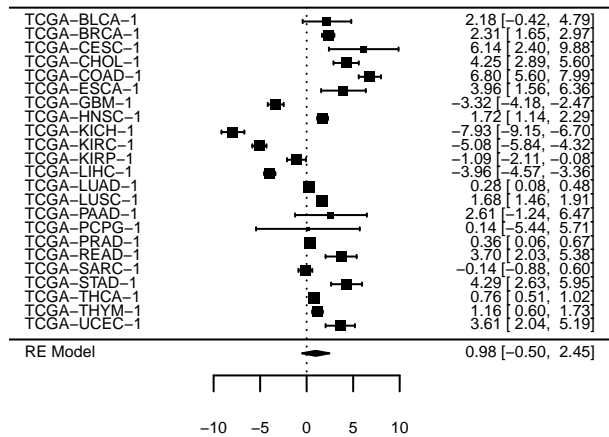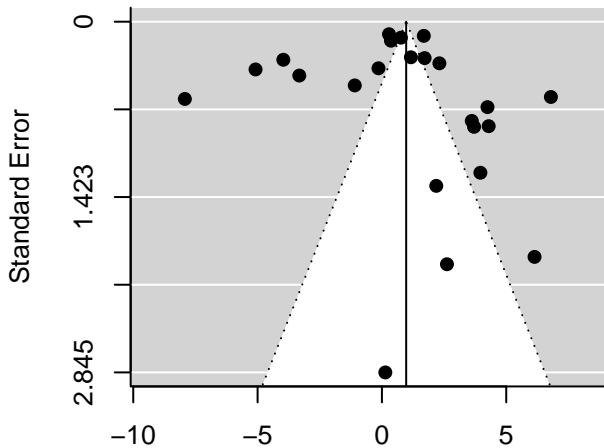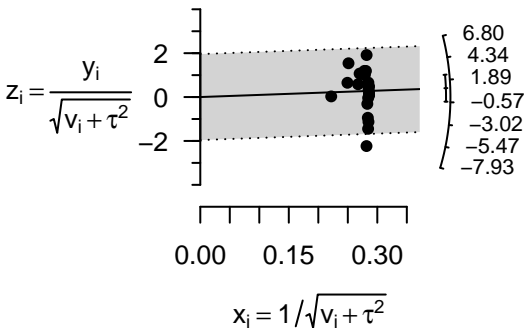| Study | SMD [95% CI] |
|---|---|
| TCGA−BLCA−1 | 2.18 [−0.42, 4.79] |
| TCGA−BRCA−1 | 2.31 [1.65, 2.97] |
| TCGA−CESC−1 | 6.14 [2.40, 9.88] |
| TCGA−CHOL−1 | 4.25 [2.89, 5.60] |
| TCGA−COAD−1 | 6.80 [5.60, 7.99] |
| TCGA−ESCA−1 | 3.96 [1.56, 6.36] |
| TCGA−GBM−1 | −3.32 [−4.18, −2.47] |
| TCGA−HNSC−1 | 1.72 [1.14, 2.29] |
| TCGA−KICH−1 | −7.93 [−9.15, −6.70] |
| TCGA−KIRC−1 | −5.08 [−5.84, −4.32] |
| TCGA−KIRP−1 | −1.09 [−2.11, −0.08] |
| TCGA−LIHC−1 | −3.96 [−4.57, −3.36] |
| TCGA−LUAD−1 | 0.28 [0.08, 0.48] |
| TCGA−LUSC−1 | 1.68 [1.46, 1.91] |
| TCGA−PAAD−1 | 2.61 [−1.24, 6.47] |
| TCGA−PCPG−1 | 0.14 [−5.44, 5.71] |
| TCGA−PRAD−1 | 0.36 [0.06, 0.67] |
| TCGA−READ−1 | 3.70 [2.03, 5.38] |
| TCGA−SARC−1 | −0.14 [−0.88, 0.60] |
| TCGA−STAD−1 | 4.29 [2.63, 5.95] |
| TCGA−THCA−1 | 0.76 [0.51, 1.02] |
| TCGA−THYM−1 | 1.16 [0.60, 1.73] |
| TCGA−UCEC−1 | 3.61 [2.04, 5.19] |
| RE Model | 0.98 [−0.50, 2.45] |

## Funnel Plot



RE Model for All Studies (Q = 1056.21, df = 22, p = 2.10E−209; $I^2$ = 99.4%)
RE Model for All Studies (beta = 0.98, se = 0.75, zval = 1.30, P = 1.99E−01)
WNT7B

Standardized Mean Difference

## Radial Plot



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

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

6.80
4.34
1.89
−0.57
−3.02
−5.47
−7.93

## Standardized Residuals



Study