Forest Plot

Study	SMD [95% CI]
TCGA–BLCA–1	2.24 [1.63, 2.86]
TCGA–BRCA–1	1.12 [1.01, 1.24]
TCGA–CESC–1	2.16 [1.74, 2.58]
TCGA–CHOL–1	1.46 [0.87, 2.04]
TCGA–COAD–1	1.45 [1.29, 1.60]
TCGA–ESCA–1	2.03 [1.48, 2.58]
TCGA–GBM–1	1.22 [0.84, 1.59]
TCGA–HNSC–1	2.18 [1.94, 2.41]
TCGA–KICH–1	−0.23 [−0.54, 0.09]
TCGA–KIRC–1	0.46 [0.31, 0.62]
TCGA–KIRP–1	1.32 [1.11, 1.54]
TCGA–LIHC–1	1.47 [1.22, 1.71]
TCGA–LUAD–1	1.62 [1.43, 1.81]
TCGA–LUSC–1	2.35 [2.15, 2.55]
TCGA–PAAD–1	−0.10 [−1.04, 0.84]
TCGA–PCPG–1	1.25 [0.87, 1.64]
TCGA–PRAD–1	0.47 [0.26, 0.69]
TCGA–READ–1	1.82 [1.60, 2.05]
TCGA–SARC–1	1.16 [0.84, 1.47]
TCGA–STAD–1	1.83 [1.60, 2.05]
TCGA–THCA–1	−0.43 [−0.59, −0.27]
TCGA–THYM–1	−0.27 [−1.13, 0.59]
TCGA–UCEC–1	2.51 [2.28, 2.74]
RE Model	1.28 [0.93, 1.64]

Standardized Mean Difference

Funnel Plot

Standardized Mean Difference
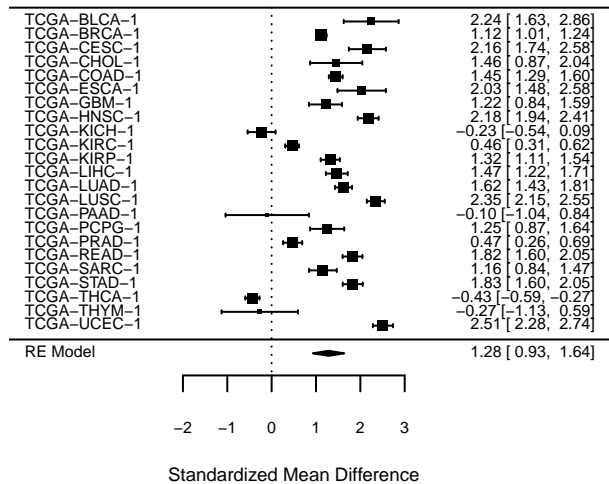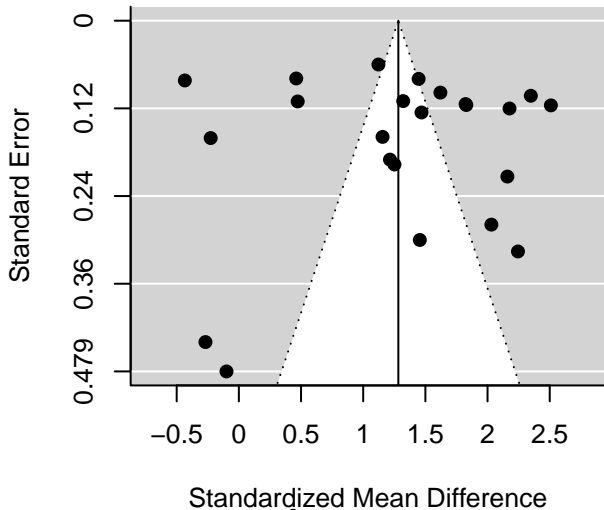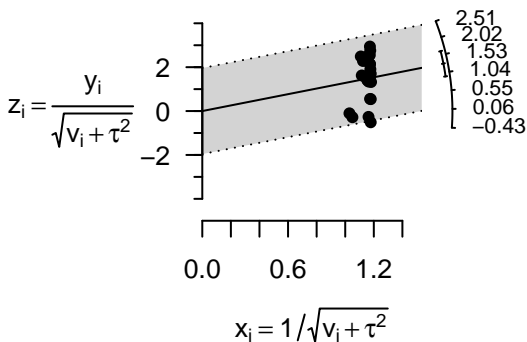
RE Model for All Studies (Q = 1084.83, df = 22, p = 1.67E−215; I^2 = 98.0%)
RE Model for All Studies (beta = 1.28, se = 0.18, zval = 7.09; P = 1.36E−12)
DNMT3B

Radial Plot

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

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

Standardized Residuals

Study