Forest Plot

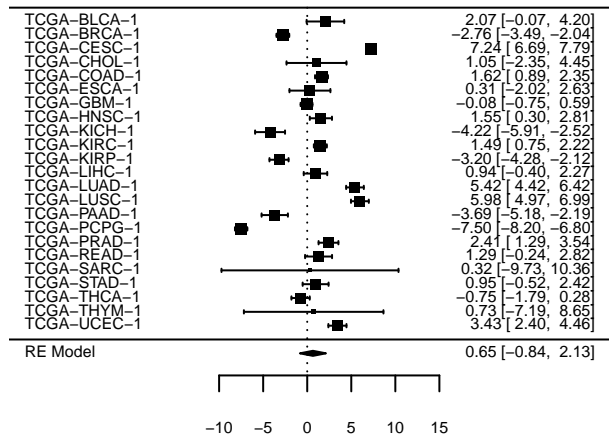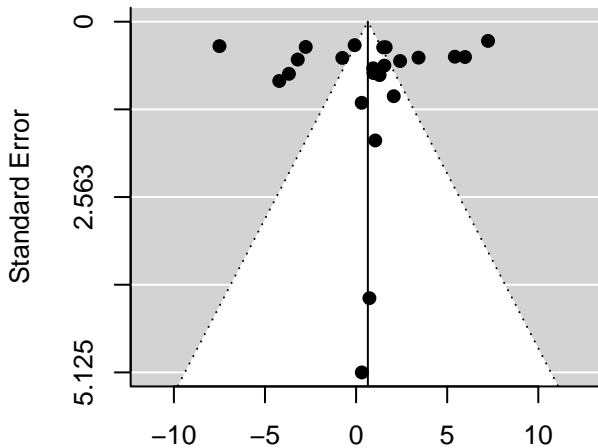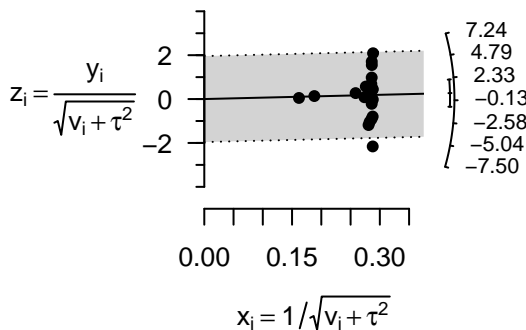| Study	SMD [95% CI]
TCGA–BLCA–1	2.07 [−0.07, 4.20]
TCGA–BRCA–1	−2.76 [−3.49, −2.04]
TCGA–CESC–1	7.24 [6.69, 7.79]
TCGA–CHOL–1	1.05 [−2.35, 4.45]
TCGA–COAD–1	1.62 [0.89, 2.35]
TCGA–ESCA–1	0.31 [−2.02, 2.63]
TCGA–GBM–1	−0.08 [−0.75, 0.59]
TCGA–HNSC–1	1.55 [0.30, 2.81]
TCGA–KICH–1	−4.22 [−5.91, −2.52]
TCGA–KIRC–1	1.49 [0.75, 2.22]
TCGA–KIRP–1	−3.20 [−4.28, −2.12]
TCGA–LIHC–1	0.94 [−0.40, 2.27]
TCGA–LUAD–1	5.42 [4.42, 6.42]
TCGA–LUSC–1	5.98 [4.97, 6.99]
TCGA–PAAD–1	−3.69 [−5.18, −2.19]
TCGA–PCPG–1	−7.50 [−8.20, −6.80]
TCGA–PRAD–1	2.41 [1.29, 3.54]
TCGA–READ–1	1.29 [−0.24, 2.82]
TCGA–SARC–1	0.32 [−9.73, 10.36]
TCGA–STAD–1	0.95 [−0.52, 2.42]
TCGA–THCA–1	−0.75 [−1.79, 0.28]
TCGA–THYM–1	0.73 [−7.19, 8.65]
TCGA–UCEC–1	3.43 [2.40, 4.46]
RE Model	0.65 [−0.84, 2.13]

Standardized Mean Difference

RE Model for All Studies (Q = 1523.83, df = 22, p = 2.34E−309; I^2 = 97.7%)
RE Model for All Studies (beta = 0.65, se = 0.76, zval = 0.85; P = 3.93E−01)
NR2E1

Funnel Plot

Standard Error

Standardized Mean Difference

Radial Plot

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

7.24
4.79
2.33
−0.13
−2.58
−5.04
−7.50

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

Standardized Residuals

Study