Forest Plot

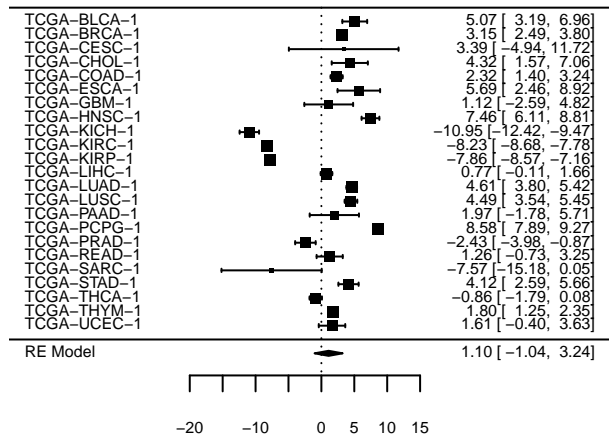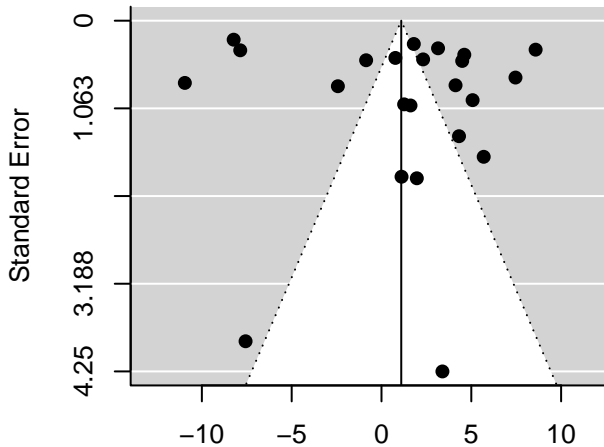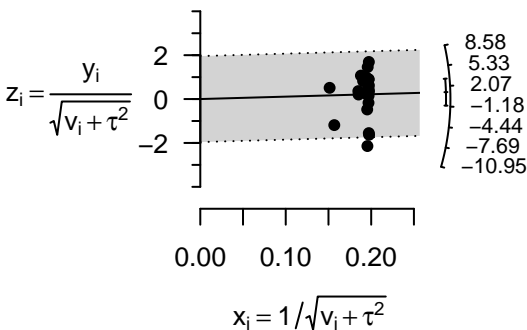| Study	SMD [95% CI]
TCGA-BLCA-1	5.07 [3.19, 6.96]
TCGA-BRCA-1	3.15 [2.49, 3.80]
TCGA-CESC-1	3.39 [-4.94, 11.72]
TCGA-CHOL-1	4.32 [1.57, 7.06]
TCGA-COAD-1	2.32 [1.40, 3.24]
TCGA-ESCA-1	5.69 [2.46, 8.92]
TCGA-GBM-1	1.12 [-2.59, 4.82]
TCGA-HNSC-1	7.46 [6.11, 8.81]
TCGA-KICH-1	-10.95 [-12.42, -9.47]
TCGA-KIRC-1	-8.23 [-8.68, -7.78]
TCGA-KIRP-1	-7.86 [-8.57, -7.16]
TCGA-LIHC-1	0.77 [-0.11, 1.66]
TCGA-LUAD-1	4.61 [3.80, 5.42]
TCGA-LUSC-1	4.49 [3.54, 5.45]
TCGA-PAAD-1	1.97 [-1.78, 5.71]
TCGA-PCPG-1	8.58 [7.89, 9.27]
TCGA-PRAD-1	-2.43 [-3.98, -0.87]
TCGA-READ-1	1.26 [-0.73, 3.25]
TCGA-SARC-1	-7.57 [-15.18, 0.05]
TCGA-STAD-1	4.12 [2.59, 5.66]
TCGA-THCA-1	-0.86 [-1.79, 0.08]
TCGA-THYM-1	1.80 [1.25, 2.35]
TCGA-UCEC-1	1.61 [-0.40, 3.63]
RE Model	1.10 [-1.04, 3.24]

Standardized Mean Difference

RE Model for All Studies (Q = 3147.33, df = 22, p = 0.00; I^2 = 99.0%)
RE Model for All Studies (beta = 1.10, se = 1.09, zval = 1.01; P = 3.13E−01)
LHX1

Funnel Plot

Standard Error

Standardized Mean Difference

Radial Plot

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

8.58
5.33
2.07
−1.18
−4.44
−7.69
−10.95

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

Standardized Residuals

Study