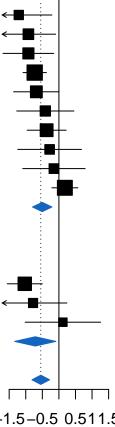
Source (95% CI)

Sequencing = TPM

Mariathasan, Lymph_node, n = 26 -1.21 [-2.21; -0.21]Fumet.2, Lung, n = 43-0.92 [-1.74; -0.10] -0.92 [-1.68; -0.16] Van Allen, Melanoma, n = 42 Mariathasan, Bladder, n = 194-0.73 [-1.08; -0.38] Riaz, Melanoma, n = 51-0.68 [-1.37; 0.01] Miao.1, Kidney, n = 33-0.41 [-1.27; 0.45] Mariathasan, Kidney, n = 67-0.37 [-0.96; 0.22] Snyder, Ureteral, n = 25-0.28 [-1.24; 0.68] Mariathasan, Ureteral, n = 26-0.15 [-1.09; 0.79] Braun, Kidney, n = 1780.18 [-0.21; 0.57] -0.50 [-0.80; -0.21] Total Heterogeneity: $\chi_{0}^{2} = 18.29 \ (P = .03), \ I^{2} = 51\% \ [0\%; 76\%]$

Sequencing = FPKM



-1.5-0.5 0.511.5 logHR estimate