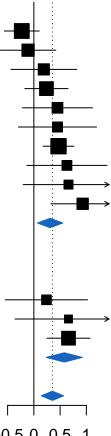
Source (95% CI)

Sequencing = TPM

Braun, Kidney, n = 178-0.23 [-0.56; 0.10] Riaz, Melanoma, n = 51-0.11 [-0.64; 0.42] 0.19 [-0.44; 0.82] Fumet.2, Lung, n = 43Mariathasan, Kidney, n = 670.24 [-0.17; 0.65] Mariathasan, Lymph_node, $n = 26 \ 0.45 \ [-0.22; 1.12]$ Miao.1, Kidney, n = 330.45 [-0.29; 1.19] Mariathasan, Bladder, n = 1940.47 [0.18; 0.76] Mariathasan, Ureteral, n = 260.63 [-0.13; 1.39] Snyder, Ureteral, n = 250.66 [-0.20; 1.52] Van_Allen, Melanoma, n = 42 0.93 [0.32; 1.54] 0.31 [0.07; 0.55] Total Heterogeneity: $\chi_9^2 = 19.16 \ (P = .02), \ I^2 = 53\% \ [4\%; 77\%]$

Sequencing = FPKM



-0.5 0 0.5 1 logHR estimate