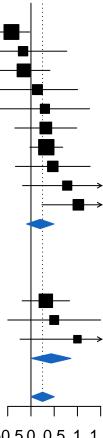
## Source (95% CI)

## Sequencing = TPM

Braun, Kidney, n = 178-0.42 [-0.81; -0.03] Mariathasan, Ureteral, n = 26 -0.17 [-1.11; 0.77] Mariathasan, Kidney, n = 67-0.16 [-0.73; 0.41] Miao.1, Kidney, n = 330.14 [-0.72; 1.00] Snyder, Ureteral, n = 250.30 [-0.66; 1.26] Riaz, Melanoma, n = 51 0.32 [-0.35; 0.99] Mariathasan, Bladder, n = 1940.33 [-0.02; 0.68] Fumet.2, Lung, n = 430.47 [-0.33; 1.27] Mariathasan, Lymph\_node,  $n = 26 \ 0.78 \ [-0.18; \ 1.74]$ Van\_Allen, Melanoma, n = 42 1.02 [ 0.24; 1.80] 0.20 [-0.09; 0.49] Total Heterogeneity:  $\chi_{Q}^{2} = 18.02 \ (P = .03), \ I^{2} = 50\% \ [0\%, 76\%]$ 

## Sequencing = FPKM



-0.5 0 0.5 1 1.5 logHR estimate