

Source**(95% CI)****Sequencing = TPM**

Fumet.2, Lung, n = 41	-1.81 [-3.87; 0.25]
Riaz, Melanoma, n = 33	-1.29 [-2.92; 0.34]
Jung, Lung, n = 26	-1.14 [-3.35; 1.07]
Fumet.1, Lung, n = 39	-0.90 [-2.88; 1.08]
Mariathasan, Bladder, n = 133	-0.60 [-1.50; 0.30]
Miao.1, Kidney, n = 28	-0.53 [-2.65; 1.59]
Braun, Kidney, n = 139	-0.36 [-1.30; 0.58]
Van_Allen, Melanoma, n = 39	-0.28 [-2.12; 1.56]
Snyder, Ureteral, n = 22	0.10 [-2.31; 2.51]
Mariathasan, Kidney, n = 46	0.28 [-1.66; 2.22]
Total	-0.60 [-1.08; -0.12]
Heterogeneity: $\chi^2_9 = 3.82$ ($P = .92$), $I^2 = 0\%$ [0%; 62%]	

Sequencing = FPKM

Liu, Melanoma, n = 112	-0.84 [-1.80; 0.12]
Nathanson, Melanoma, n = 24	-0.83 [-2.46; 0.80]
Hugo, Melanoma, n = 27	0.33 [-1.67; 2.33]
Total	-0.67 [-1.43; 0.10]
Heterogeneity: $\chi^2_2 = 1.12$ ($P = .57$), $I^2 = 0\%$ [0%; 90%]	
Total	-0.62 [-1.02; -0.21]
Heterogeneity: $\chi^2_{12} = 4.96$ ($P = .96$), $I^2 = 0\%$ [0%; 57%]	
Test for overall effect: $z = -2.99$ ($P = .003$)	
Test for subgroup differences: $\chi^2_1 = 0.02$ ($P = .88$)	

