

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

Fumet.2, Lung, n = 41	-1.78 [-3.64; 0.08]
Jung, Lung, n = 26	-1.18 [-3.06; 0.70]
Riaz, Melanoma, n = 33	-1.17 [-2.58; 0.24]
Miao.1, Kidney, n = 28	-0.86 [-2.80; 1.08]
Van_Allen, Melanoma, n = 39	-0.75 [-2.53; 1.03]
Mariathanan, Kidney, n = 46	-0.64 [-2.42; 1.14]
Mariathanan, Bladder, n = 133	-0.57 [-1.37; 0.23]
Braun, Kidney, n = 139	-0.40 [-1.24; 0.44]
Snyder, Ureteral, n = 22	0.09 [-1.71; 1.89]
Fumet.1, Lung, n = 39	0.93 [-0.81; 2.67]
Total	-0.58 [-1.00; -0.15]
Heterogeneity: $\chi^2_9 = 6.36$ ($P = .70$), $I^2 = 0\%$ [0%; 62%]	

Sequencing = FPKM

Liu, Melanoma, n = 112	-0.88 [-1.72; -0.04]
Nathanson, Melanoma, n = 24	-0.54 [-2.01; 0.93]
Hugo, Melanoma, n = 27	0.28 [-1.50; 2.06]
Total	-0.64 [-1.32; 0.04]
Heterogeneity: $\chi^2_2 = 1.35$ ($P = .51$), $I^2 = 0\%$ [0%; 90%]	
Total	-0.60 [-0.95; -0.24]
Heterogeneity: $\chi^2_{12} = 7.74$ ($P = .81$), $I^2 = 0\%$ [0%; 57%]	
Test for overall effect: $z = -3.25$ ($P = .001$)	
Test for subgroup differences: $\chi^2_1 = 0.02$ ($P = .88$)	

