

Source	(95% CI)
Sequencing = TPM	
Mariathasan, Ureteral, n = 26	-0.84 [-2.41; 0.73]
Mariathasan, Kidney, n = 67	-0.30 [-1.18; 0.58]
Snyder, Ureteral, n = 25	-0.27 [-1.62; 1.08]
Fumet.2, Lung, n = 43	-0.22 [-1.36; 0.92]
Miao.1, Kidney, n = 33	0.36 [-0.91; 1.63]
Braun, Kidney, n = 178	0.37 [-0.24; 0.98]
Riaz, Melanoma, n = 51	0.38 [-0.35; 1.11]
Mariathasan, Lymph_node, n = 26	0.46 [-0.72; 1.64]
Van_Allen, Melanoma, n = 42	0.58 [-0.65; 1.81]
Mariathasan, Bladder, n = 194	0.62 [0.15; 1.09]
Total	0.31 [0.04; 0.58]
Heterogeneity: $\chi^2_9 = 7.44$ ($P = .59$), $I^2 = 0\%$ [0%; 62%]	

Sequencing = FPKM	
Hugo, Melanoma, n = 27	0.52 [-1.30; 2.34]
Nathanson, Melanoma, n = 24	1.04 [0.02; 2.06]
Liu, Melanoma, n = 121	1.16 [0.43; 1.89]
Total	1.06 [0.50; 1.62]
Heterogeneity: $\chi^2_2 = 0.41$ ($P = .81$), $I^2 = 0\%$ [0%; 90%]	
Total	0.43 [0.16; 0.70]
Heterogeneity: $\chi^2_{12} = 13.44$ ($P = .34$), $I^2 = 11\%$ [0%; 50%]	
Test for overall effect: $z = 3.16$ ($P = .002$)	
Test for subgroup differences: $\chi^2_1 = 5.63$ ($P = .02$)	

