

Source	(95% CI)
Sequencing = TPM	
Mariathasan, Ureteral, n = 26	-0.88 [-3.09; 1.33]
Hwang, Lung, n = 21	-0.66 [-3.11; 1.79]
Braun, Kidney, n = 178	-0.42 [-1.54; 0.70]
Fumet.2, Lung, n = 43	-0.19 [-1.70; 1.32]
Mariathasan, Kidney, n = 67	-0.17 [-1.58; 1.24]
Van_Allen, Melanoma, n = 42	0.27 [-1.26; 1.80]
Riaz, Melanoma, n = 51	0.39 [-1.24; 2.02]
Snyder, Ureteral, n = 25	0.69 [-1.70; 3.08]
Mariathasan, Bladder, n = 194	0.80 [0.00; 1.60]
Mariathasan, Lymph_node, n = 26	1.62 [-0.87; 4.11]
Miao.1, Kidney, n = 33	3.35 [-0.06; 6.76]
Total	0.27 [-0.21; 0.74]
Heterogeneity: $\chi^2_{10} = 9.86$ ($P = .45$), $I^2 = 0\%$ [0%; 60%]	

Sequencing = FPKM	
Liu, Melanoma, n = 121	0.31 [-0.69; 1.31]
Hugo, Melanoma, n = 27	0.96 [-1.96; 3.88]
Nathanson, Melanoma, n = 24	4.20 [1.34; 7.06]
Total	1.57 [-0.72; 3.86]
Heterogeneity: $\chi^2_2 = 6.35$ ($P = .04$), $I^2 = 68\%$ [0%; >91%]	
Total	0.38 [-0.03; 0.78]
Heterogeneity: $\chi^2_{13} = 17.05$ ($P = .20$), $I^2 = 24\%$ [0%; 60%]	
Test for overall effect: $z = 1.81$ ($P = .07$)	
Test for subgroup differences: $\chi^2_1 = 1.19$ ($P = .28$)	

