

Source	(95% CI)
Sequencing = TPM	
Mariathanas, Ureteral, n = 26	-0.67 [-1.47; 0.13]
Miao.1, Kidney, n = 33	-0.29 [-1.00; 0.42]
Mariathanas, Kidney, n = 67	-0.24 [-0.69; 0.21]
Fumet.2, Lung, n = 43	-0.04 [-0.69; 0.61]
Snyder, Ureteral, n = 25	-0.02 [-0.78; 0.74]
Mariathanas, Bladder, n = 194	-0.01 [-0.30; 0.28]
Van_Allen, Melanoma, n = 42	0.04 [-0.59; 0.67]
Mariathanas, Lymph_node, n = 26	0.08 [-0.84; 1.00]
Riaz, Melanoma, n = 51	0.40 [-0.11; 0.91]
Braun, Kidney, n = 178	0.58 [0.21; 0.95]
Total	0.04 [-0.19; 0.27]
Heterogeneity: $\chi^2_9 = 15.35$ ($P = .08$), $I^2 = 41\%$ [0%; 72%]	

Sequencing = FPKM	
Liu, Melanoma, n = 121	0.12 [-0.27; 0.51]
Nathanson, Melanoma, n = 24	0.12 [-0.70; 0.94]
Hugo, Melanoma, n = 27	0.39 [-0.73; 1.51]
Total	0.14 [-0.19; 0.48]
Heterogeneity: $\chi^2_2 = 0.2$ ($P = .90$), $I^2 = 0\%$ [0%; 90%]	
Total	0.07 [-0.12; 0.26]
Heterogeneity: $\chi^2_{12} = 15.69$ ($P = .21$), $I^2 = 24\%$ [0%; 60%]	
Test for overall effect: $z = 0.73$ ($P = .46$)	
Test for subgroup differences: $\chi^2_1 = 0.25$ ($P = .62$)	

