

Source	(95% CI)
Sequencing = FPKM	
Hugo, Melanoma, n = 27	-1.08 [-2.12; -0.04]
Liu, Melanoma, n = 112	-0.03 [-0.40; 0.34]
Nathanson, Melanoma, n = 24	0.22 [-0.62; 1.06]
Total	-0.18 [-0.77; 0.40]
Heterogeneity: $\chi^2_2 = 4.11$ ($P = .13$), $I^2 = 51\%$ [0%; 86%]	

Sequencing = TPM	
Snyder, Ureteral, n = 22	-0.83 [-1.81; 0.15]
Jung, Lung, n = 26	-0.61 [-1.49; 0.27]
Mariathasan, Kidney, n = 46	-0.48 [-1.26; 0.30]
Van_Allen, Melanoma, n = 39	-0.44 [-1.15; 0.27]
Mariathasan, Bladder, n = 133	-0.40 [-0.77; -0.03]
Riaz, Melanoma, n = 33	-0.23 [-1.11; 0.65]
Braun, Kidney, n = 139	0.14 [-0.23; 0.51]
Miao.1, Kidney, n = 28	0.20 [-0.66; 1.06]
Fumet.2, Lung, n = 41	0.30 [-0.41; 1.01]
Fumet.1, Lung, n = 39	0.42 [-0.29; 1.13]
Total	-0.15 [-0.40; 0.10]
Heterogeneity: $\chi^2_9 = 13.01$ ($P = .16$), $I^2 = 31\%$ [0%; 67%]	
Total	-0.14 [-0.35; 0.06]
Heterogeneity: $\chi^2_{12} = 17.17$ ($P = .14$), $I^2 = 30\%$ [0%; 64%]	
Test for overall effect: $z = -1.36$ ($P = .18$)	
Test for subgroup differences: $\chi^2_1 = 0.01$ ($P = .92$)	

