

Source	(95% CI)
Sequencing = TPM	
Jung, Lung, n = 26	-2.41 [-5.39; 0.57]
Snyder, Ureteral, n = 22	-0.65 [-2.96; 1.66]
Mariathasan, Bladder, n = 133	-0.37 [-1.31; 0.57]
Braun, Kidney, n = 139	-0.35 [-1.31; 0.61]
Fumet.2, Lung, n = 41	0.33 [-1.14; 1.80]
Van_Allen, Melanoma, n = 39	1.09 [-1.22; 3.40]
Mariathasan, Kidney, n = 46	1.19 [-1.65; 4.03]
Total	-0.21 [-0.76; 0.34]
Heterogeneity: $\chi^2_6 = 5.09$ ($P = .53$), $I^2 = 0\%$ [0%; 71%]	

Sequencing = FPKM	
Nathanson, Melanoma, n = 24	-1.19 [-3.37; 0.99]
Hugo, Melanoma, n = 27	-0.84 [-3.00; 1.32]
Liu, Melanoma, n = 112	-0.26 [-1.34; 0.82]
Total	-0.51 [-1.39; 0.37]
Heterogeneity: $\chi^2_2 = 0.67$ ($P = .71$), $I^2 = 0\%$ [0%; 90%]	
Total	-0.29 [-0.76; 0.17]
Heterogeneity: $\chi^2_9 = 6.08$ ($P = .73$), $I^2 = 0\%$ [0%; 62%]	
Test for overall effect: $z = -1.23$ ($P = .22$)	
Test for subgroup differences: $\chi^2_1 = 0.32$ ($P = .57$)	

