

Source	(95% CI)
Sequencing = TPM	
Riaz, Melanoma, n = 33	-0.03 [-1.46; 1.40]
Van_Allen, Melanoma, n = 39	0.50 [-1.87; 2.87]
Jerby_Arnon, Melanoma, n = 96	1.06 [0.08; 2.04]
Total	0.67 [-0.15; 1.49]
Heterogeneity: $\chi^2_2 = 1.55$ ($P = .46$), $I^2 = 0\%$ [0%; 90%]	

Sequencing = FPKM	
Hugo, Melanoma, n = 27	0.00 [-2.29; 2.29]
Liu, Melanoma, n = 112	0.98 [-0.04; 2.00]
Nathanson, Melanoma, n = 24	1.37 [-0.69; 3.43]
Total	0.91 [0.06; 1.76]
Heterogeneity: $\chi^2_2 = 0.81$ ($P = .67$), $I^2 = 0\%$ [0%; 90%]	
Total	0.79 [0.22; 1.36]
Heterogeneity: $\chi^2_5 = 2.51$ ($P = .78$), $I^2 = 0\%$ [0%; 75%]	
Test for overall effect: $z = 2.72$ ($P = .006$)	
Test for subgroup differences: $\chi^2_1 = 0.16$ ($P = .69$)	

