

Source	(95% CI)
Sequencing = WES	
Rizvi.15, Lung, n = 34	-1.20 [-1.93; -0.47]
Snyder, Ureteral, n = 25	-0.62 [-1.38; 0.14]
Miao.2, Bladder, n = 27	-0.55 [-2.22; 1.12]
Jung, Lung, n = 58	-0.37 [-0.88; 0.14]
Liu, Melanoma, n = 144	-0.20 [-0.51; 0.11]
Miao.1, Kidney, n = 35	-0.20 [-1.77; 1.37]
Van_Allen, Melanoma, n = 112	-0.06 [-0.37; 0.25]
Braun, Kidney, n = 249	-0.02 [-0.26; 0.22]
Miao.2, Melanoma, n = 38	0.38 [-0.62; 1.38]
Miao.2, Lung, n = 34	2.56 [-0.83; 5.95]
Total	-0.22 [-0.44; 0.00]
Heterogeneity: $\chi^2_9 = 15.56$ ($P = .08$), $I^2 = 42\%$ [0%; 72%]	

Sequencing = TGS	
Samstein, Unknown, n = 34	-0.73 [-1.46; 0.00]
Rizvi.18, Lung, n = 29	-0.66 [-1.44; 0.12]
Samstein, Melanoma, n = 132	-0.59 [-0.98; -0.20]
Samstein, Esophagus, n = 21	-0.22 [-0.87; 0.43]
Samstein, HNC, n = 78	-0.21 [-0.60; 0.18]
Total	-0.43 [-0.67; -0.20]
Heterogeneity: $\chi^2_4 = 3.24$ ($P = .52$), $I^2 = 0\%$ [0%; 79%]	
Total	-0.30 [-0.47; -0.12]
Heterogeneity: $\chi^2_{14} = 22.79$ ($P = .06$), $I^2 = 39\%$ [0%; 67%]	
Test for overall effect: $z = -3.32$ ($P < .001$)	
Test for subgroup differences: $\chi^2_1 = 1.71$ ($P = .19$)	

