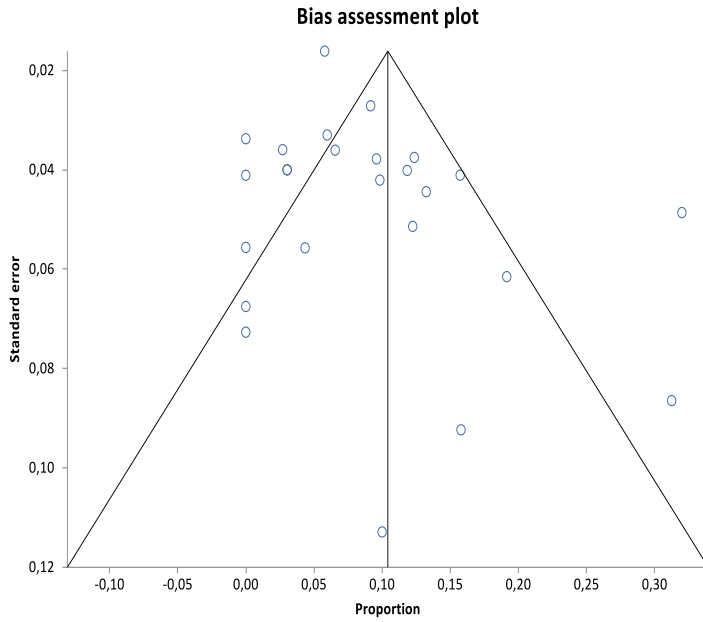**Supplementary Figure 1.** Funnel plots assessing publication bias, for studies reporting PIK3CA mutational status pooled discordance rates. A. Overall discordance rate; B. Overall discordance rate from PIK3CA-mutated to PIK3CA-wild-type; C. Overall discordance rate from PIK3CA-wild-type to PIK3CA-mutated
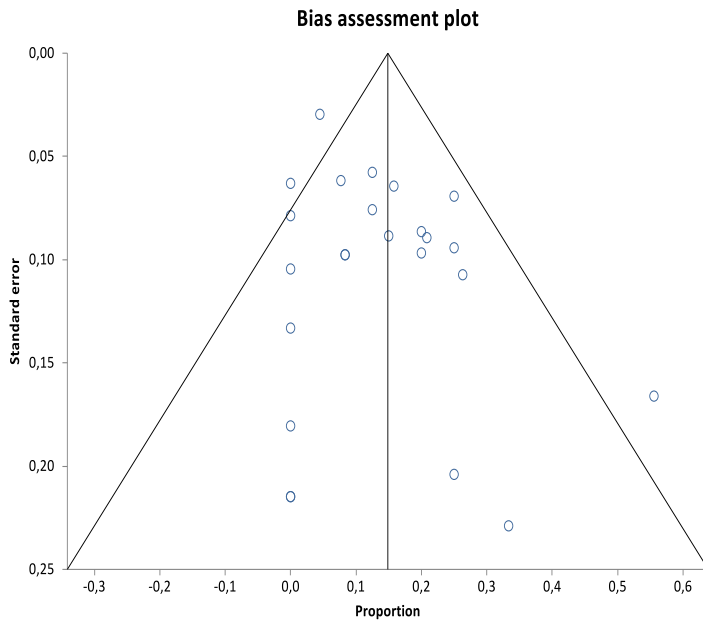
A

Begg-Mazumdar: Kendall's 0,217391 P = 0,1348

Egger: bias = 1,148938 (95% CI = -0,535455 to 2,833331) P = 0,1716

Harbord: bias = -0,790959 (92,5% CI = -2,818292 to 1,236375) P = 0,4742
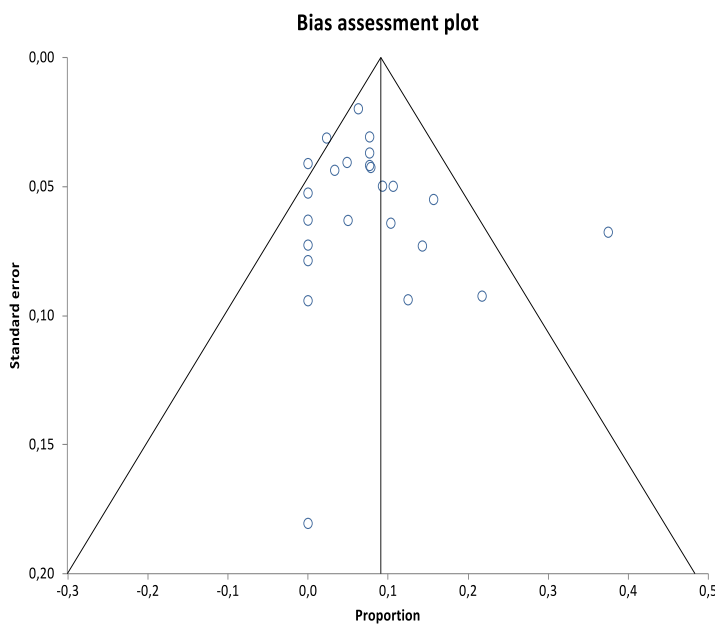
B

Begg-Mazumdar: Kendall's 0,094891 P = 0,5347

Egger: bias = 0,996692 (95% CI = -0,021999 to 2,015382) P = 0,0547

Harbord: bias = -0,013746 (92,5% CI = -1,235155 to 1,207664) P = 0,9834

C

Begg-Mazumdar: Kendall's 0,166667 P = 0,2675

Egger: bias = 0,618025 (95% CI = -0,707182 to 1,943233) P = 0,344

Harbord: bias = -0,646081 (92,5% CI = -2,493965 to 1,201803) P = 0,5202