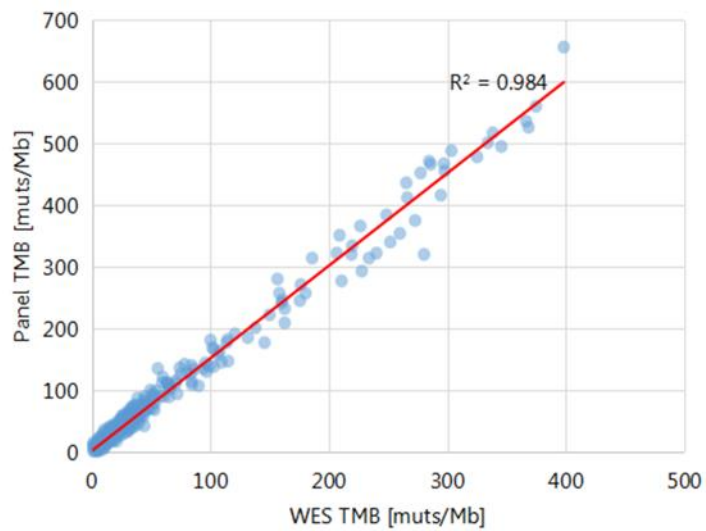


**A**



**B**

**S2 Fig.** The correlation of genepanels including 295-gene panel (A) and 1021-gene panel (B) in evaluating TMB when compared to TMB derived from WES, respectively. TMB, tumor mutational burden; WES, whole-exome sequencing.