

Figure S1 Flowchart of this study.

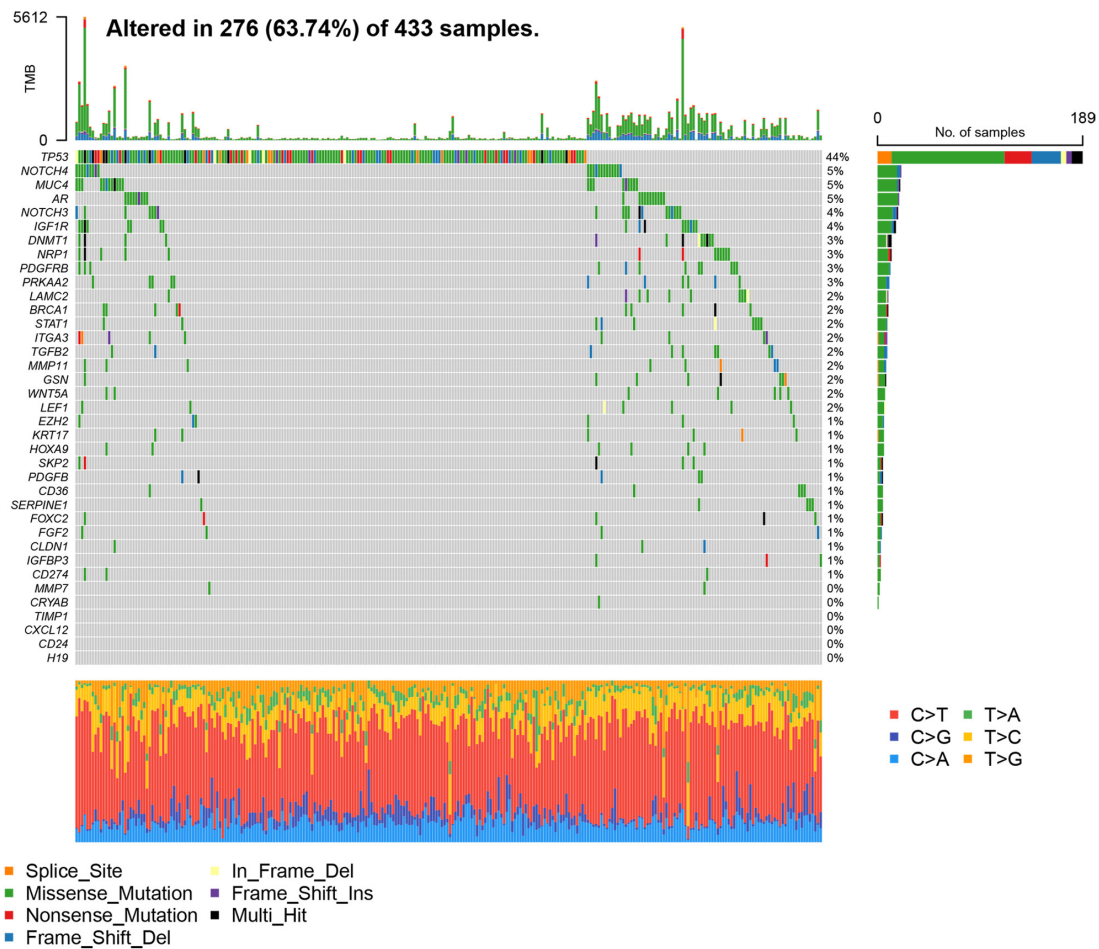


Figure S2 Mutational frequency analysis of AERGs in gastric cancer.

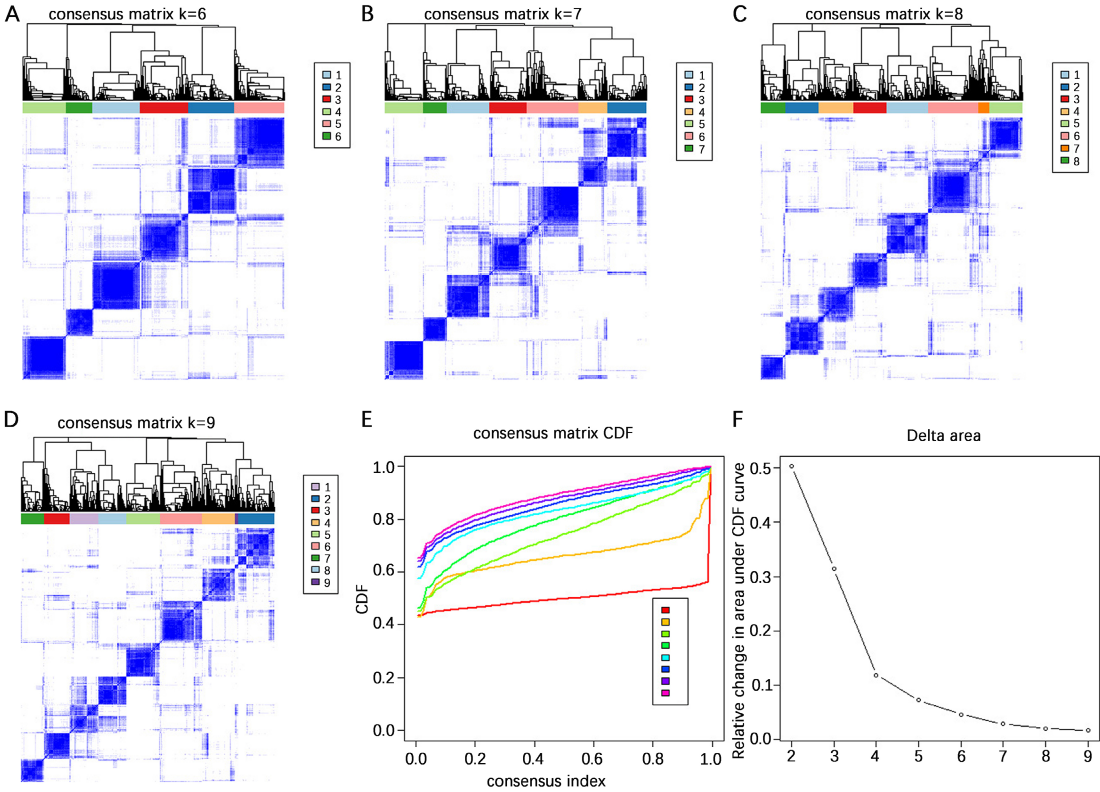


Figure S3 Consensus unsupervised clustering analysis of 37 AERGs in gastric cancer.

(A-D) Consensus matrix heatmaps of 37 AERGs ($k = 6-9$). (E) The CDF from $k = 2$ to 9. (F) Relative variation of the area under the CDF region from $k = 2$ to 9. CDF, cumulative distribution function.