Supplementary Figure 2. Energy metabolism-related genes could distinguish glioma patients with different clinical and molecular features. (**A**) Consensus clustering CDF for k = 2 to k = 10. (**B**) Relative change in area under CDF curve for k = 2 to k = 10. (**C**) Consensus clustering matrix of 309 samples from CGGA dataset for k = 2. (**D**) Heat map of two clusters defined by the top 50 variable expression genes. (**E**) survival analysis of patients in cluster 1 and cluster 2.