

	Source data	Website
Table 1, Figure 1 A-D	GEO, EGA, and TCGA	https://kmplot.com/analysis/
Citation: Lanczky A, Gyorffy B: Web-Based Survival Analysis Tool Tailored for Medical Research (KMplot): Development and Implementation, J Med Internet Res, 2021 Jul 26;23(7):e27633. doi: 10.2196/27633		
Figure 1 E-F	TCGA	http://www.proteinatlas.org
Citation: Uhlen M, Fagerberg L, Hallstrom BM, Lindskog C, Oksvold P, Mardinoglu A, et al. Proteomics. Tissue-based map of the human proteome. Science. 2015;347(6220):1260419.		
Figure 2,3	GEO, EGA, and TCGA	https://kmplot.com/analysis/
Citation: Lanczky A, Gyorffy B: Web-Based Survival Analysis Tool Tailored for Medical Research (KMplot): Development and Implementation, J Med Internet Res, 2021 Jul 26;23(7):e27633. doi: 10.2196/27633		
Figure 4	TCGA	http://bioinfo.life.hust.edu.cn/GSCA/#/immune
Citation: Liu CJ, Hu FF, Xia M, Han L, Zhang Q, Guo AY. (2018). GSCALite: A Web Server for Gene Set Cancer Analysis. Bioinformatics.		
Figure 5,6	TCGA	http://cis.hku.hk/TISIDB/index.php
Citation: Beibei Ru, Ching Ngar Wong, Yin Tong, Jia Yi Zhong, Sophia Shek Wa Zhong, Wai Chung Wu, Ka Chi Chu, Choi Yiu Wong, Chit Ying Lau, Ian Chen, Nam Wai Chan, Jiangwen Zhang. TISIDB: an integrated repository portal for tumor-immune system interactions. Bioinformatics. 2019; btz210.		
Figure 7	TCGA	http://cbioportal.org
Citation: Sumter TF, Xian L, Huso T, Koo M, Chang YT, Almasri TN, et al. The High Mobility Group A1 (HMGA1) Transcriptome in Cancer and Development. Current molecular medicine. 2016;16(4):353-93.		