



Supplementary Fig S1. Scatter plot using FPKM values of AF1q and ICAM-1. Publicly available 1,222 RNA-Seq data of 1,092 breast cancer cases from TCGA database are used. When FPKM of AF1q is high, that of ICAM-1 is low, supporting that overexpression of AF1q attenuates ICAM-1 expression.