Figure S1 cGAS-STING pathway expression in normal tissue compared to cancer tissue in cancer patients: (A) STING; (B) cGAS; (C) IFI16. Gene expression analysis was measured with ddPCR and normalized to ACTB. ddPCR, droplet digital polymerase chain reaction; ACTB, actin beta.
Figure S2 Kaplan-Meier survival analysis for stage I–III lung adenocarcinoma patients. The Kaplan-Meier plots were generated using gene expression data from microarray analysis and clinical data available at kmplot.com for (A) \textit{IFI16} (n=530), (B) \textit{IRF3} (n=370).