A comparison of protein expression levels across different treatments and conditions is shown in panel A. Panels B and C illustrate the fold change of PD-L1 mRNA and the percentage of PD-L1+ cells, respectively, with statistical significance indicated by p-values. Panel D presents a dose-response analysis with Rad001 at 0.25 μM, showing a notable increase in PD-L1+ cells with statistical significance. Panel E depicts protein expression analysis under NT and S6(-) conditions, with corresponding images of cell viability and apoptosis analysis.