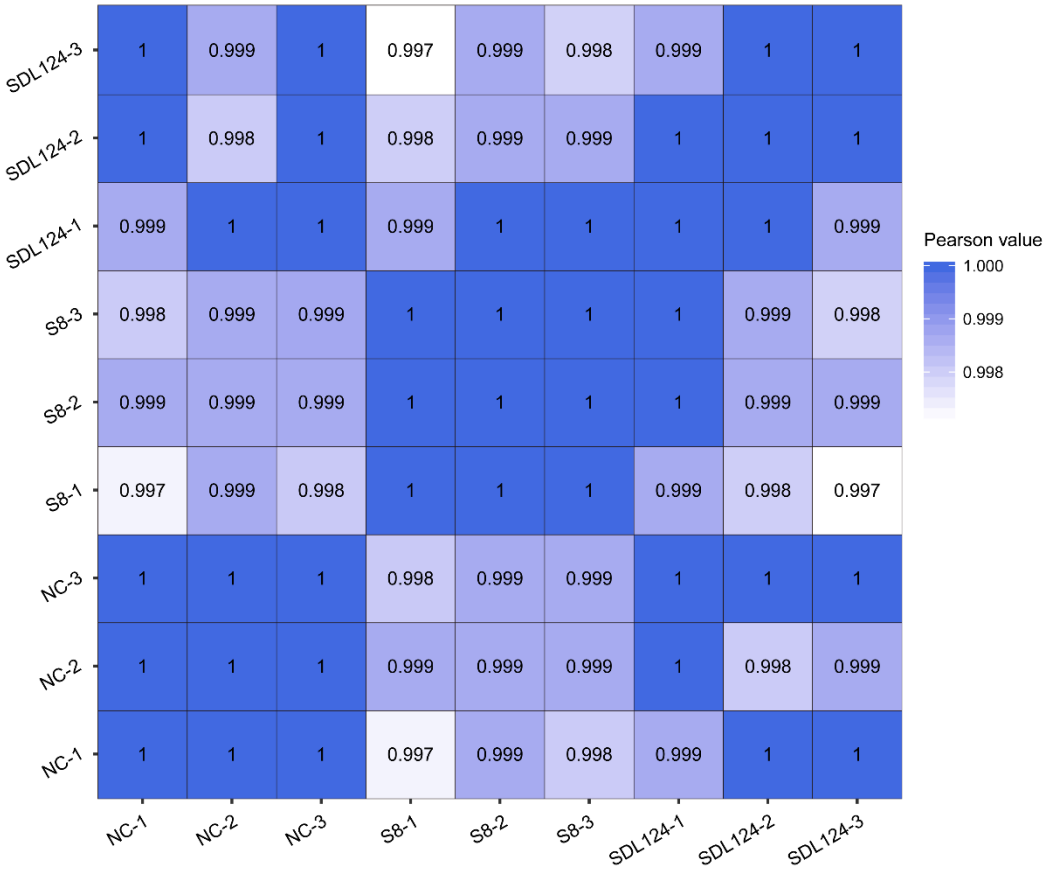


Figure S1. Heatmap of correlation between samples. The correlation between samples was calculated based on FPKM values of genes, and an inter-sample correlation heatmap was generated using *R*.