Figure S1. Heat map of gene expression and hierarchical cluster diagram showing the performance of 7 candidate genes and the 7-gene panel for clustering of 65 healthy controls and 102 SIL patients. (A) 7 separate candidate genes and (B) the 7-gene panel. The figure was generated using the 'Heatmap' function in R with default settings. SIL, squamous intraepithelial lesion.