Supplementary figure 1. Heat map and hierarchical tree comparing gene expression alterations of the UC-ulcerative colitis group and NC-normal controls.

The red and blue colors represent increased and decreased gene expression compared with NC, respectively. Each line represents an individual sample according to its group, and each row represents the changes of a given transcript. The transcript used for the rows are the 1050 altered transcripts in the CLDP group. Taking together, each cell represents a change of a single transcript in a single sample. The direction and the magnitude of the changes are coded by color.