



Figure S1. Validation of S3-enriched genes by quantitative RT-PCR. (A) TSNE plots showing the expression pattern of 12 selected genes. (B) Validation of selected genes by using qRT-PCR. Data obtained from qRT-PCR as well as bulk RNA-seq were displayed as mean \pm SEM (standard error of mean). According to the t-test, $p < 0.05$ is true for all genes. $n = 3$.