



Fig. S2 t-distributed Stochastic Neighbor Embedding (t-SNE) visualization identifying 20 and 22 putative cell clusters in aerial and subterranean pegs, respectively. **A, B** t-SNE plot of nuclei according to transcriptomic profiles in aerial (A) and subterranean (B) pegs; **C, D** Two replicates in aerial (C) and subterranean (D) pegs; **E, F** The number of cells in each cell cluster in in aerial (E) and subterranean (F) pegs