

Supplementary Table 1. RNA-seq datasets used in this study.

Analyzed Data	Identifier	Reference
scRNA-seq of mouse tracheal epithelium: basal, club, ciliated, and other rare cells (7193 3' scRNA-seq cells + 301 full-length scRNA-seq cells)	GSE103354	1
scRNA-seq of mouse tracheal epithelium (7661 cells from uninjured tissue) & scRNA-seq of differentiated human bronchial epithelial cells (2970 cells)	GSE102580	2
bulk RNA-seq of sorted mouse AEP and AT2 cells (2 and 3 replicates per cell type)	GSE97055	3
scRNA-seq of sorted mouse AT1 cells (at P3, P15, P60 stages; 3149, 2940, and 1337 cells) and AT2 cells (P60; 2093 cells)	GSE106960	4
bulk RNA-seq of sorted mouse BASCs, AT2, and club cells (2 replicates per cell type)	GSE129440	5
scRNA-seq of mouse BASC, AT2, and club + ciliated cells (480 cells)	GSE118891	6
scRNA-seq of distal adult murine lung airway epithelium: H2-K1 ^{high} cells within club-like cells (3278 cells after filtering)	GSE129937	7

References

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