

Supplemental Table 2. NETs alter epithelial cell junction RNAs. List of functionally enriched RNAs for cellular components of anchored junctions and tight junctions found by RNA sequencing of HBE exposed to 5µg/ml NETs compared to PBS for 18h. Data analyzed using ToppGene (experiments=3, HBE donors=3, NET donors=3).

Cell Junction Organization	
Symbol	Fold Change
<i>ACTB</i>	2.89996469
<i>KIFC3</i>	2.944804134
<i>CLDN4</i>	3.207103512
<i>TGFB1</i>	3.726483192
<i>MAP4K4</i>	3.876084105
<i>LAMB3</i>	4.014288083
<i>TLR2</i>	4.487771729
<i>ESAM</i>	4.649265569
<i>EPHB3</i>	4.978859764
<i>DKK1</i>	6.847217229
<i>LAMC2</i>	8.371985395
<i>TIAM1</i>	9.666262173
<i>ARHGAP22</i>	-5.618308079
<i>CAV1</i>	-5.043682033
<i>WNT4</i>	-4.864411143
<i>CLDN8</i>	-4.838474606
<i>SNAI2</i>	-4.715354949
<i>PIK3R1</i>	-3.973412632
<i>MTSS1</i>	-3.870409255
<i>WNT3A</i>	-3.594393098
<i>WNT5A</i>	-3.247994985