

Network and co-expression analysis of airway smooth muscle cell transcriptome delineates potential gene signatures in asthma

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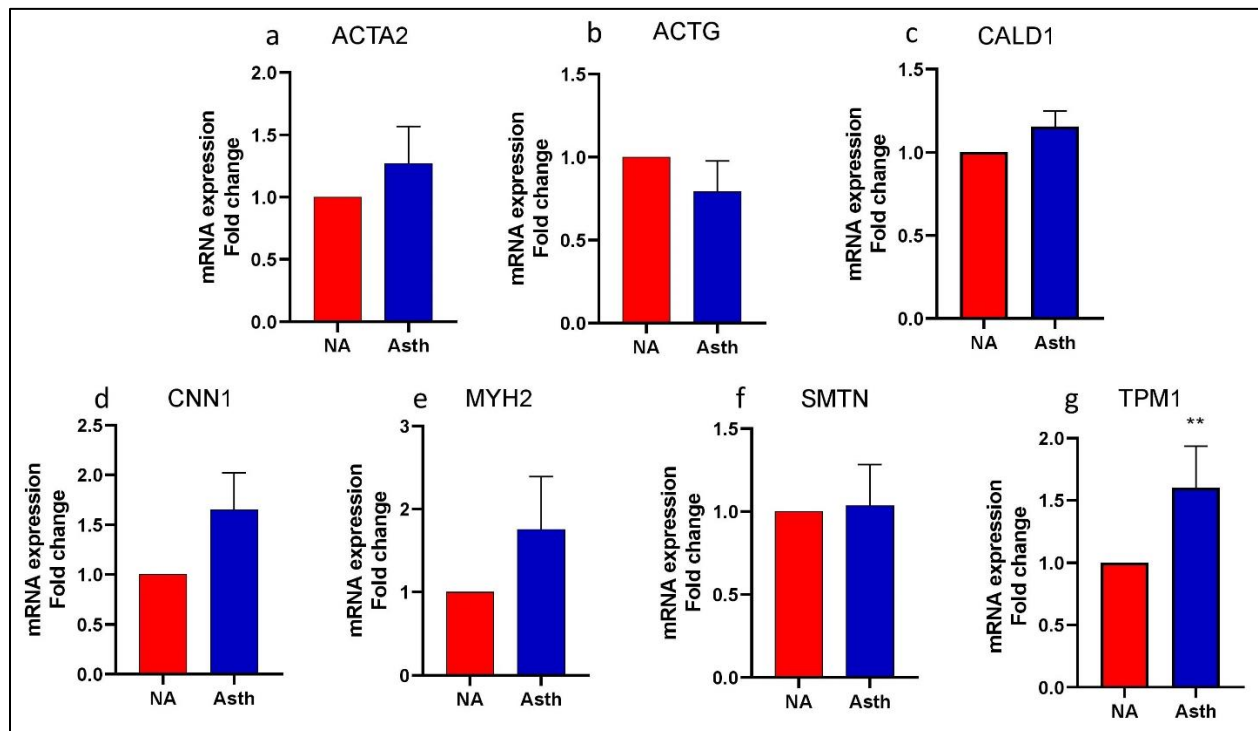
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SUPPLEMENTARY FIGURE



Supplementary Figure S1: Differential mRNA expression of smooth muscle markers in asthmatic and non-asthmatic human ASM samples. The fold change for mRNA was evaluated in both groups. The genes were validated in human ASM samples collected from non-asthmatic (NA) and asthmatic (Asth) patients (N = 5 in each group). Data represented as mean $\pm$ SEM in each group. **p-value<0.001.