

1 Additional file 15. Fig. S6. Comparisons of functional genes between CON and naïve asthma group. a. Functional genes of level 1 (human disease)
2 with significant differences between CON and untreated asthma group were showed (Kruskal-Wallis rank-sum test), corrections were made using
3 the False Discovery Rate multiple testing correction. Results were considered statistically significant for p-values ≤ 0.05 . b. Functional genes of
4 level 3 with significant differences between CON and untreated asthma group were showed (Kruskal-Wallis rank-sum test), corrections were made
5 using the False Discovery Rate multiple testing correction. Results were considered statistically significant for p-values ≤ 0.05 .

