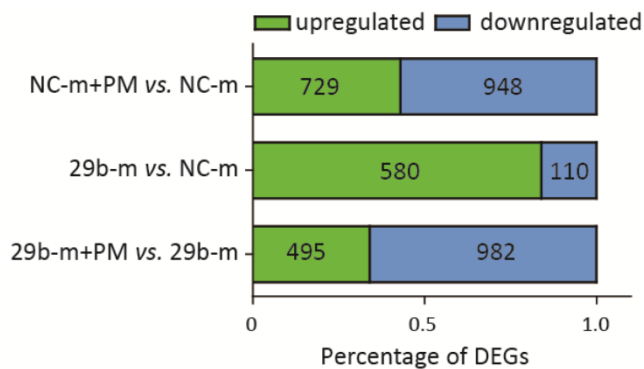
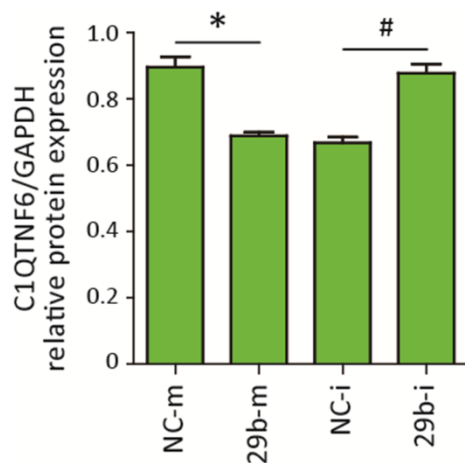


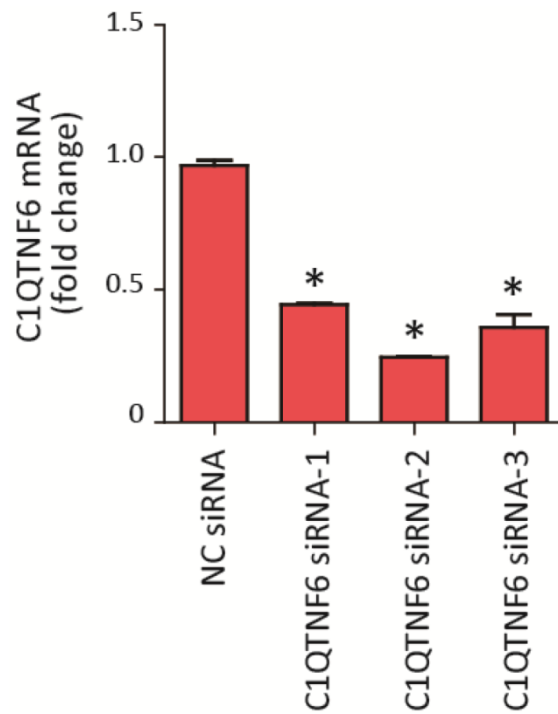
SUPPLEMENTARY FIGURE



Supplementary Figure 1. The number of differentially expressed genes was identified in NC-m vs. 29b-m group, NC-m + PM vs. 29b-m + PM group, and NC-m vs. NC-m + PM group.



Supplementary Figure 2. The optical densities of protein bands for Figure 4F. Values were shown as mean ± SEM; *, P<0.05, compared with the NC-m group; #, P<0.05, compared with the NC-i group; n=3.



Supplementary Figure 3. The optimum C1QTNF6 siRNA was identified according to the inhibition efficiency of three designed siRNAs using RT-PCR. The C1QTNF6 siRNAs were transfected into HBECs and the optimum C1QTNF6 siRNA was selected using real-time PCR. Values were shown as mean \pm SEM; *, $P < 0.05$, compared with the NC siRNA group; $n = 3$.