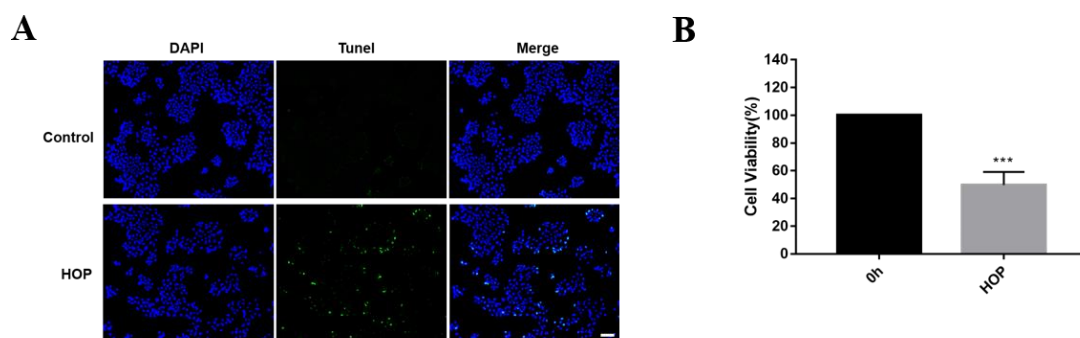


**Table1. Sequences of human primers used for RT-PCR**

Gene	Forward sequence(5'-3')	Reverse sequence(5'-3')
GAPDH	ATGTTTCGTCATGGGTGTGA A	GGTGCTAAGCAGTTGGTGG T
NLRP3	CGTGAGTCCCATTAAAGATG GAGT	CCCGACAGTGGATATAGAAC AGA
IL17A	TGTCACTGCTACTGCTGCT GAG	GGTGAGGTGGATCGGTTGT AGT
TNF- $\alpha$	CTCAGCAAGGACAGCAGA GGAC	TGGAGCCGTGGGTCAGTAT GT
Beclin1	TGTGTTGCTGCTCCATGCT CTG	GCCACTGCCTCCTGTGTCTT CA
P62	GAGCCTCATCTCCTCGGTG T	TTCTCAAGCCCCATGTTGCA C

**Table2. Sequences of mouse primers used for RT-PCR**

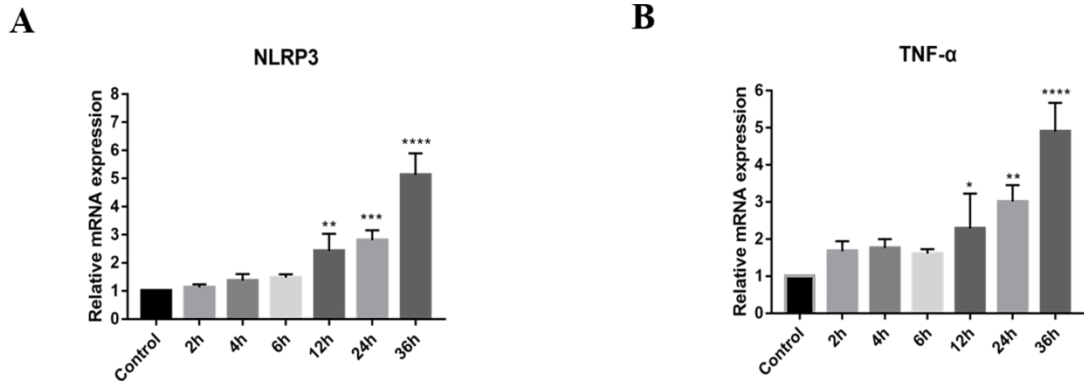
Gene	Forward sequence(5'-3')	Reverse sequence(5'-3')
GAPDH	TGTCCGTCGTGGATCTGA C	CCTGCTTCACCACCTTCTT G
ATG5	TGTGCTTCGAGATGTGTG GTT	GTCAAATAGCTGACTCTTG GCAA
Beclin1	TAAGGCGTCCAGCAGCA CCAT	AGAGACACCATCCTGGCGA GTT
P62	CCAGCACAGGCACAGAA GACAA	CCCACCGACTCCAAGGCTA TCT

**Figure S1. Hyperosmolarity induced cellular apoptosis and death in HCECs. (A)**

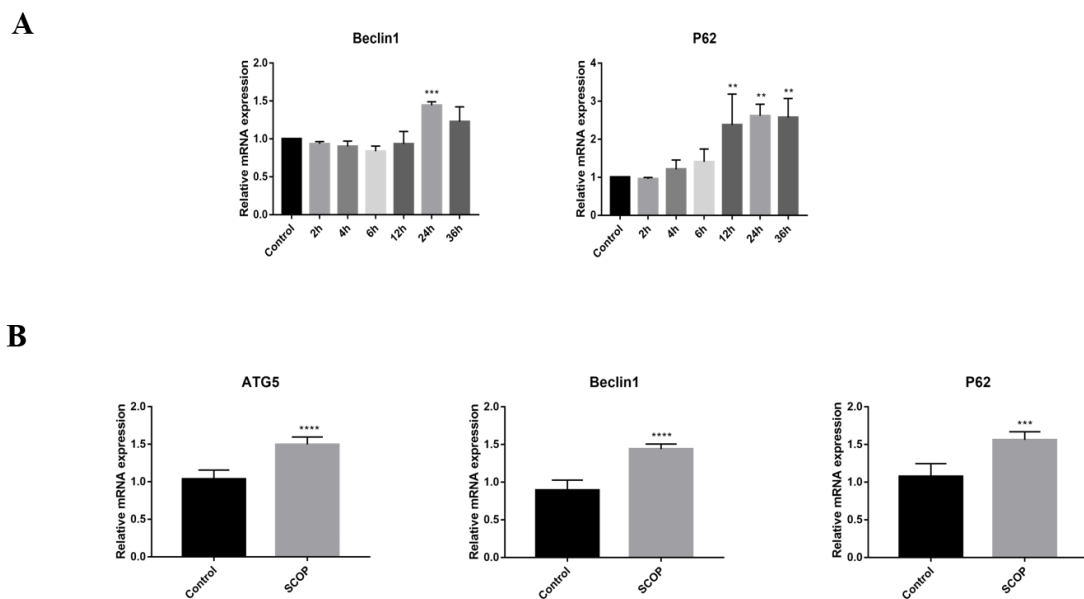
Tunel staining images in HCECs exposed to normal or hyperosmotic medium.

Bar=50 $\mu$ m. **(B)** Changes to cell viability of HCECs by CCK8 assay. The results are

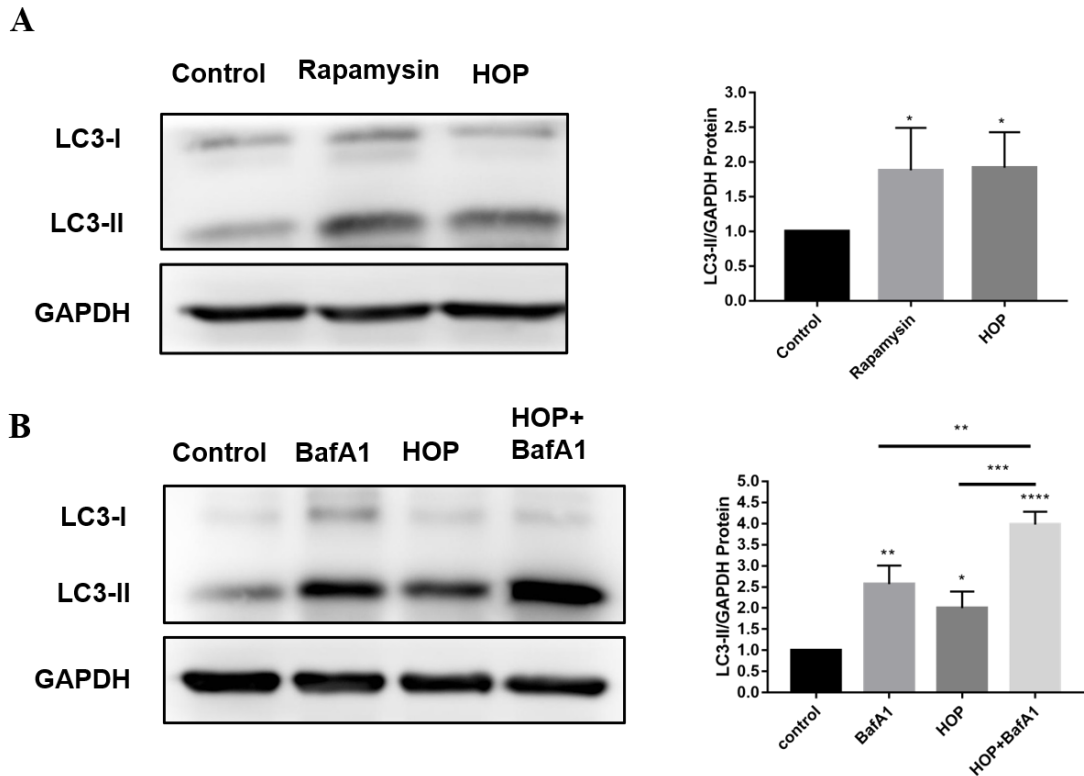
shown as mean  $\pm$  SD of three independent experiments. \*\*\*P < 0.001.



**Figure S2. Hyperosmolarity stimulated inflammatory factors in HCECs. (A)** Gene expression results of NLRP3. **(B)** Gene expression of TNF $\alpha$ . The results are shown as mean  $\pm$  SD of three independent experiments. \*P < 0.05, \*\* P < 0.01, \*\*\*P < 0.001, \*\*\*\*P < 0.0001.

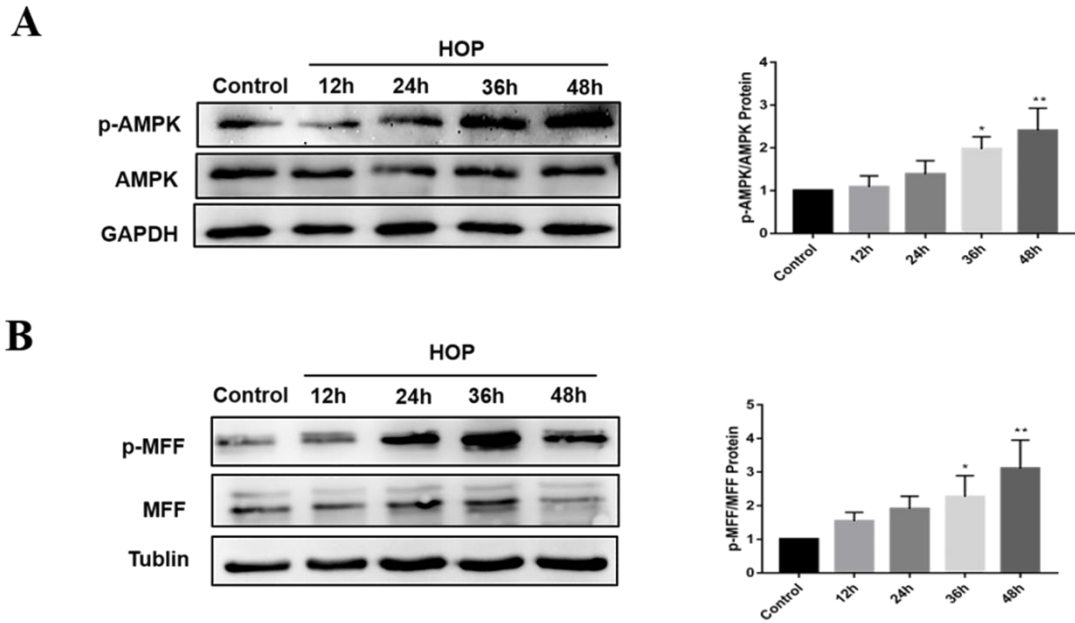


**Figure S3. Expression of autophagy genes in dry eye. (A)** RT-PCR results showing Beclin1 and P62 expression in untreated and HOP-stressed HCECs for indicated times (N=3). **(B)** Gene expression of ATG5, Beclin1 and P62 in normal and dry eye mice (N=5). The results are shown as mean  $\pm$  SD. \*\* P < 0.01, \*\*\*P < 0.001, \*\*\*\*P < 0.0001.



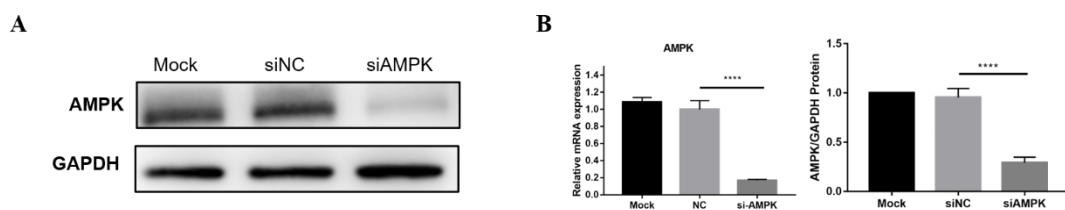
**Figure S4. Hyperosmolarity induced increased autophagy flux in HCECs. (A)**

Western blot assays were performed to detect LC3 expression in HCECs exposed to Rapamycin (1  $\mu$ M) or HOP for 24 hours. **(B)** HCECs were exposed to HOP for 24 hours and then treated or not treated with bafilomycin A1 (100 nM, 4 hours). Changes to LC-3-II expression were detected by Western blot. The results are shown as mean  $\pm$  SD of three or four independent experiments. \* $P < 0.05$ , \*\*  $P < 0.01$ , \*\*\*  $P < 0.001$ , \*\*\*\*  $P < 0.0001$ .



**Figure S5. The AMPK/MFF/DRP1 pathway was activated in HOP-stressed HCECs.**

(A) Western Blot detected AMPK and p-AMPK expression in untreated and HOP-stressed HCECs for indicated times. (B) Western Blot results showing MFF and p-MFF expression in untreated and HOP-stressed HCECs for indicated times. The results are shown as mean  $\pm$  SD of three independent experiments. \* $P < 0.05$ , \*\*  $P < 0.01$ .



**Figure S6. Validation of AMPK interference efficiency. (A)** Western Blot showed AMPK expression after AMPK-specific siRNAs transfection in HCECs. (B) Quantitative

analysis of the AMPK gene and protein expression after siAMPK transfection in HCECs. The results are shown as mean  $\pm$  SD of three independent experiments. \*\*\*\*

$P < 0.0001$ .