

Supplementary Materials: Therapeutic effect of endothelin-converting enzyme inhibitor on chronic kidney disease through the inhibition of endoplasmic reticulum stress and the NLRP3 inflammasome

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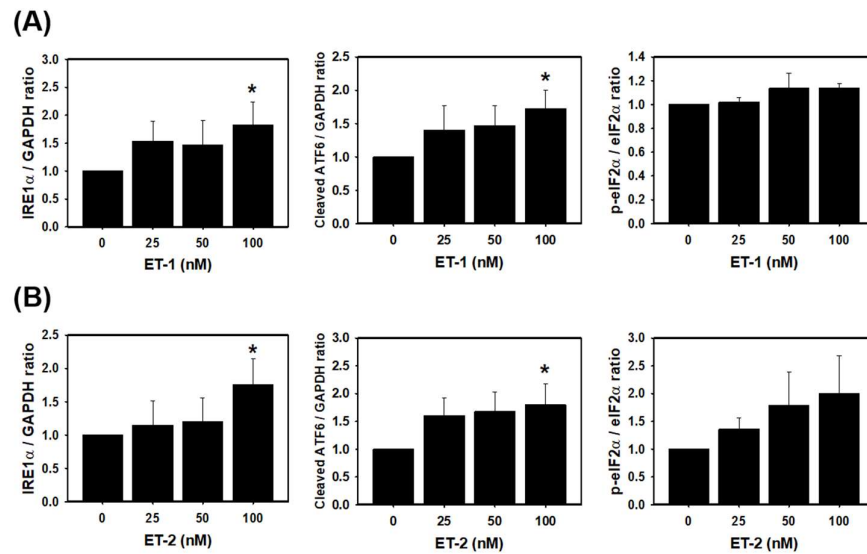


Figure S1. Effects of ET-1 (A) or ET-2 (B) exposure on the ER stress in human kidney cells. The ER stress-related proteins expression of histogram represent the average normalized densitometric values. GAPDH was used as the internal control. Data are presented as the means \pm standard deviation of three independent experiments. *P < 0.05 compared with the control.

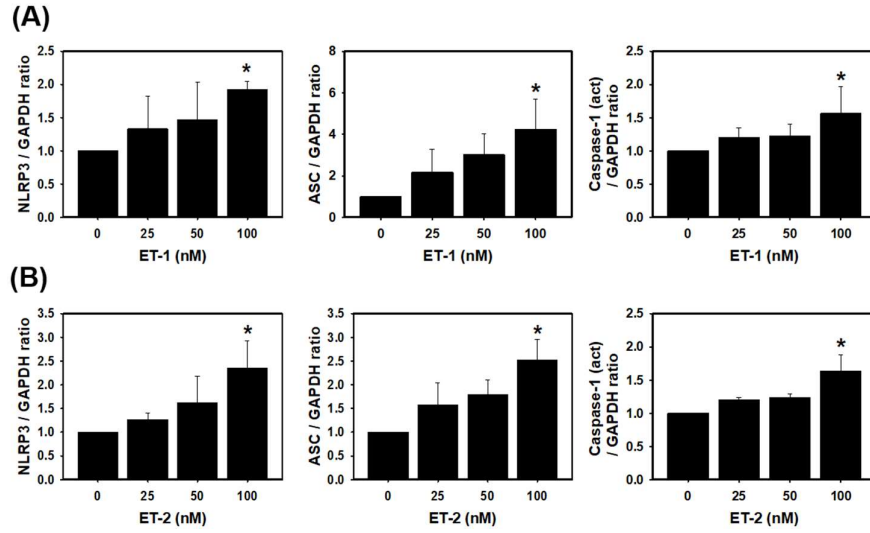


Figure S2. Effects of ET-1 (A) or ET-2 (B) exposure on the NLRP3 inflammasome in human kidney cells. The NLRP3 inflammasome-related proteins expression of histogram represent the average normalized densitometric values. GAPDH was used as the internal control. Data are presented as the means \pm standard deviation of three independent experiments. *P < 0.05 compared with the control.

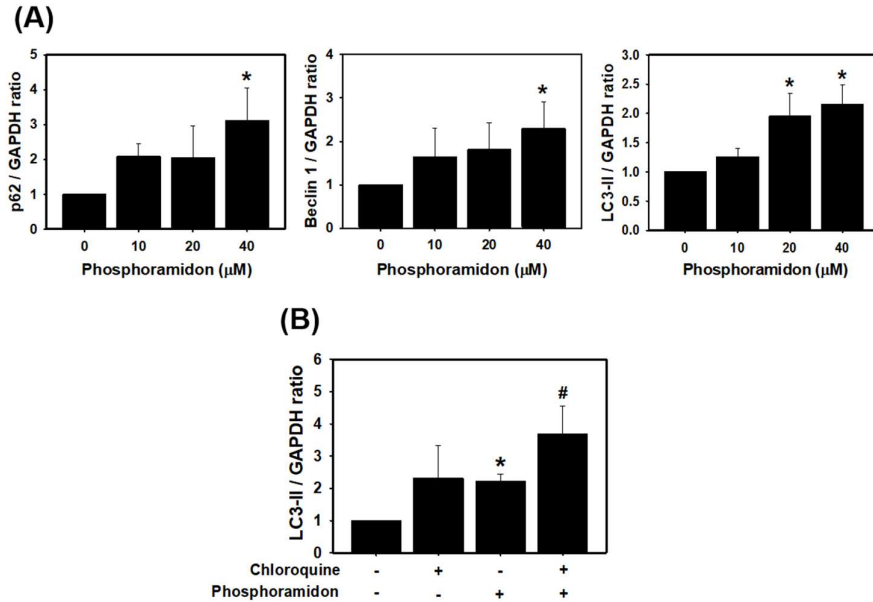


Figure S3. Effects of phosphoramidon treatment on autophagy in human kidney cells. (A) The autophagy-related proteins expression of histogram represent the average normalized densitometric values. *P < 0.05 compared with the control. (B) Quantification of LC3-II in HK-2 cells treated with chloroquine and phosphoramidon. The cells were pretreated with chloroquine

(5 μM) for 1 h and then treated with phosphoramidon (20 μM) for 24 h. GAPDH was used as the internal control. Data are presented as the means \pm standard deviation of three independent experiments. *P < 0.05 compared with the control. #P < 0.05 compared with the control. # p < 0.05 compared with the phosphoramidon.

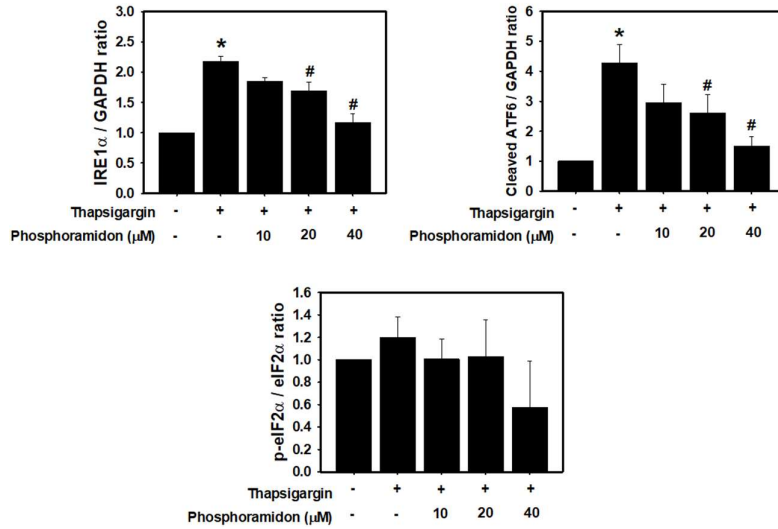


Figure S4. Effects of phosphoramidon treatment on the thapsigargin-induced ER stress in human kidney cells. The ER stress-related proteins expression of histogram represent the average normalized densitometric values. GAPDH was used as the internal control. Data are presented as the means \pm standard deviation of three independent experiments. *P < 0.05 compared with the control. # p < 0.05 compared with thapsigargin.

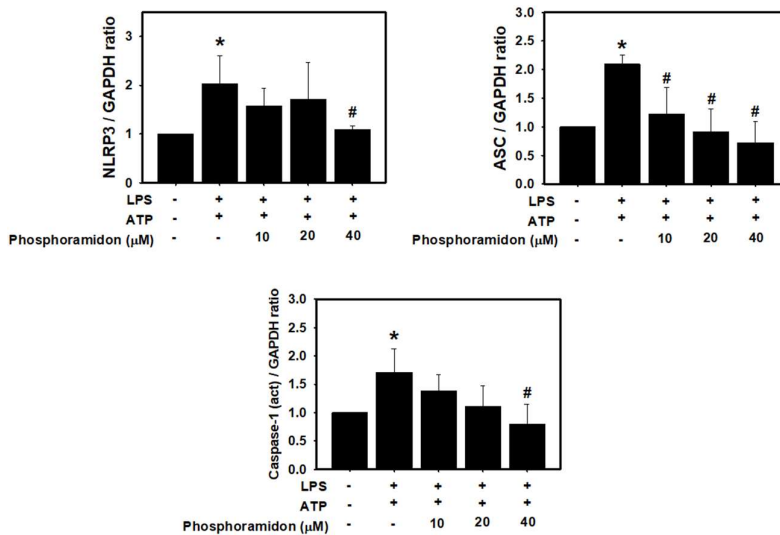


Figure S5. Effects of phosphoramidon treatment on the LPS+ATP-induced NLRP3 inflammasom in human kidney cells. The NLRP3 inflammasom-related proteins expression of histogram represent the average normalized densitometric values. GAPDH was used as the internal control. Data are presented as the means \pm standard deviation of three independent experiments. *P < 0.05 compared with the control. # p < 0.05 compared with the LPS+ATP group.

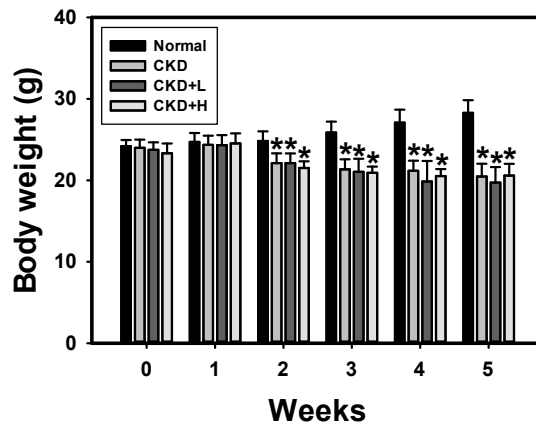
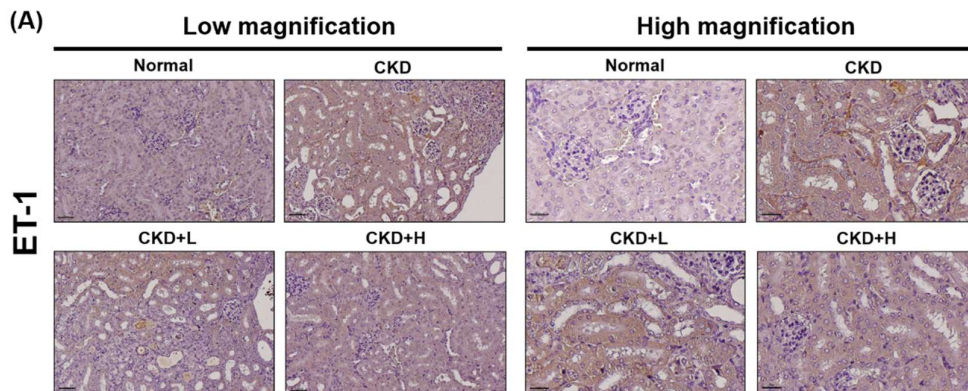


Figure S6. Measurement of body weights of C57BL/6 mice in various groups (5 mice per group). * p < 0.05 compared with the normal group. Data are presented as the means \pm standard deviation.



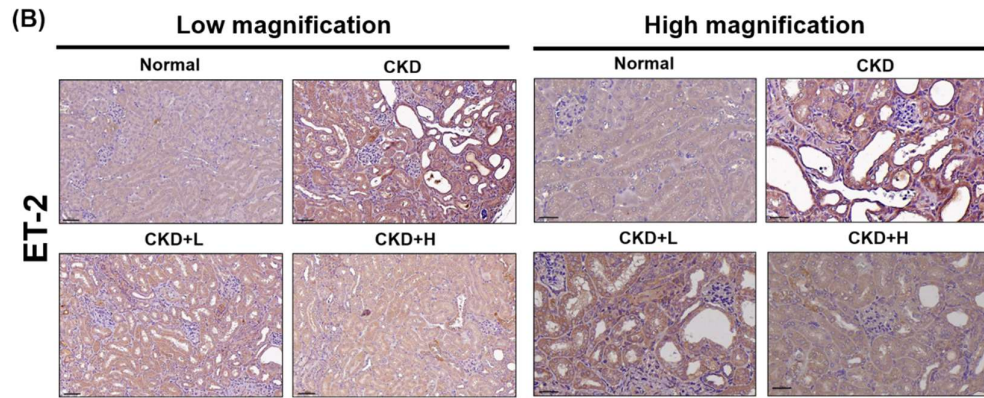


Figure S7. The ET-1 and ET-2 expression of kidneys after phosphoramidon treatment in the adenine diet-induced CKD model. IHC was used to determine the expression levels of ET-1 (A) and ET-2 (B) in kidney tissues. The scale bar of low magnification=60 μm . The scale bar of high magnification=30 μm .

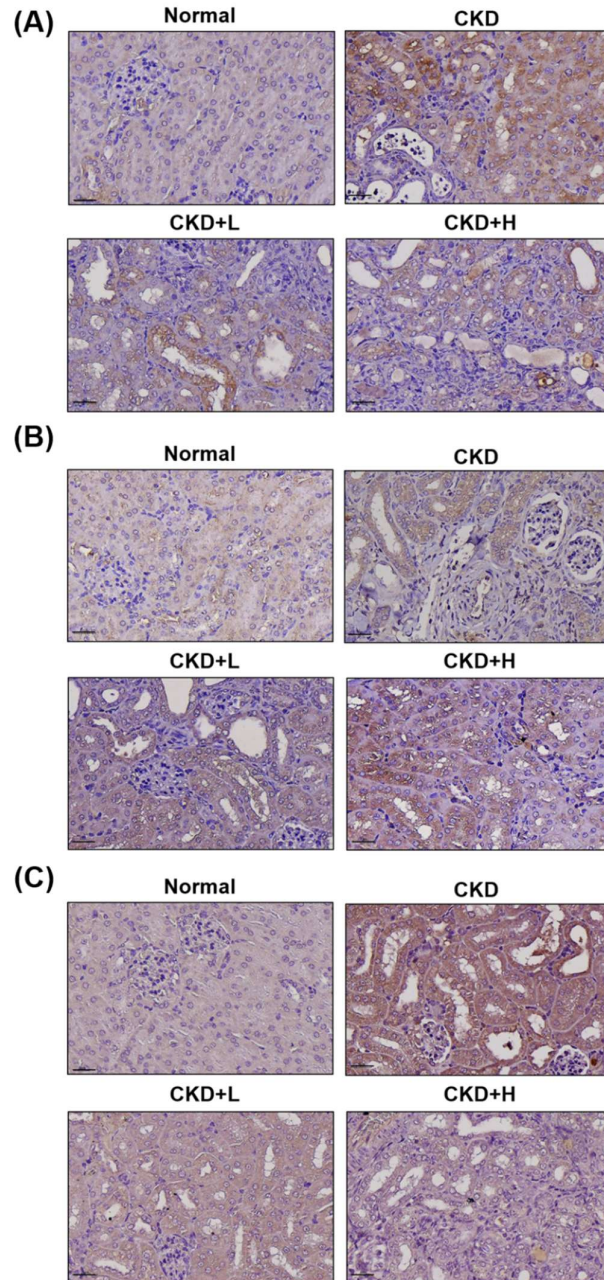


Figure S8. The high magnification of IHC images. Phosphoramidon regulates ER stress, autophagy and the NLRP3 inflammasome in the adenine diet-induced CKD model. The protein expression of IRE1 α (A), LC3 (B) and NLRP3 (C) in kidney sections in the indicated groups. Scale bar=30 μ m.

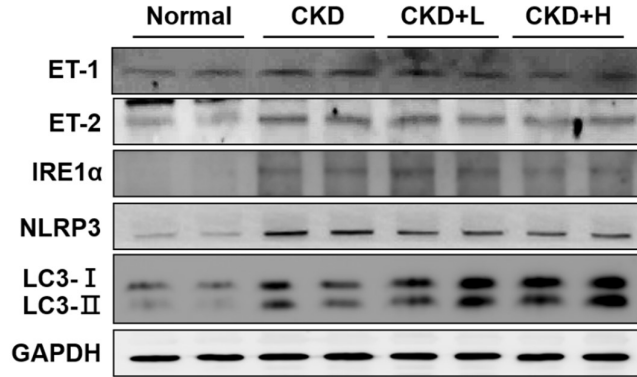


Figure S9. Western blot analysis of ET-1, ET-2, IRE1 α , LC3 and NLRP3 in kidney tissue sections in the indicated groups.

Table S1. The quantification of immunohistochemical (IHC) and masson staining.

Item / Group	Normal	CKD	CKD+L	CKD+H
IRE1 α area (%)	1.11 \pm 1.20	7.39 \pm 1.37*	6.25 \pm 1.72	3.25 \pm 1.91#
LC3 area (%)	2.10 \pm 1.62	5.56 \pm 1.66*	7.23 \pm 1.47#	8.90 \pm 1.35#
NLRP3 area (%)	1.91 \pm 1.06	26.19 \pm 5.87*	14.48 \pm 5.86#	10.75 \pm 2.97#
ET-1 area (%)	3.87 \pm 1.39	18.75 \pm 5.34*	15.96 \pm 5.18	9.10 \pm 3.57#
ET-2 area (%)	3.50 \pm 1.61	35.64 \pm 4.09*	21.99 \pm 3.19#	10.08 \pm 2.33#
Masson staining positive area (%)	0.2 \pm 0.13	5.8 \pm 1.86*	4.21 \pm 1.19#	2.02 \pm 0.61#

*p<0.05, CKD compared with normal.

#p<0.05, CKD+L or CKD+H compared with CKD.