

Supplementary Material

A Hypercaloric Diet Induces Early Podocyte Damage in Aged, Non-Diabetic Rats

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Supplementary Table 1. Gender differences in HFHCD-induced renal injury – Glomerular parameters.

HFHCD = high-fat-high-carbohydrate diet, n = number, SD = standard deviation,

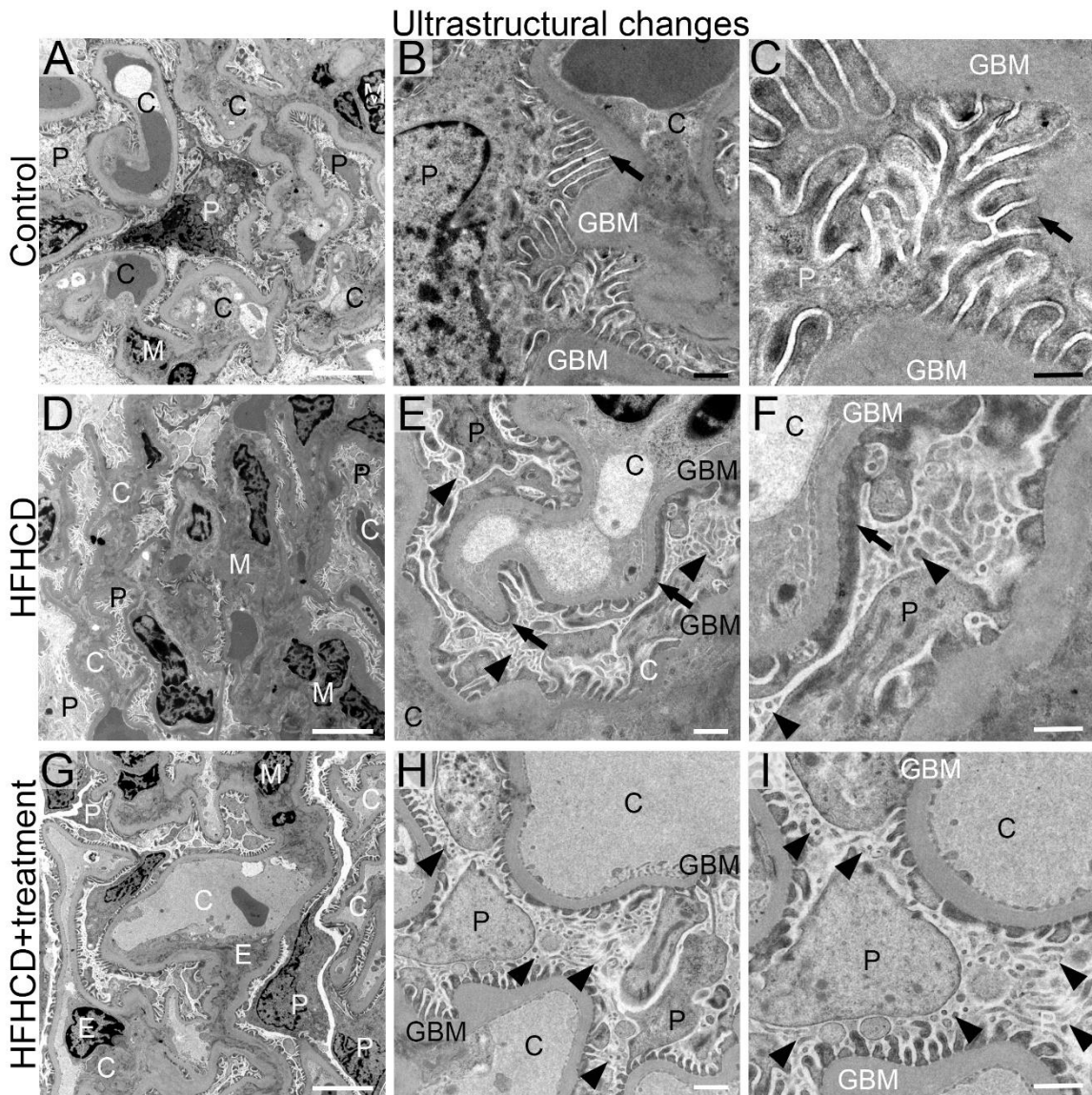
* p<0.05, ** p<0.01 versus HFHCD group.

| Groups | Males (n=29) | | Females (n=28) | |
|--|-----------------|--------------|-------------------|--------------|
| | n | mean ± SD | n | mean ± SD |
| Tuft size (µm²) | | | | |
| Control | 7 | 10294 ± 1308 | 6 | 8585 ± 757 |
| HFHCD | 6 | 9351 ± 1269 | 8 | 7968 ± 632 |
| HFHCD+liraglutide | 8 | 9145 ± 562 | 4 | 7978 ± 85.3 |
| HFHCD+metformin | 8 | 9428 ± 391 | 7 | 8164 ± 573 |
| Glomerular podocin (% area) | | | | |
| Control | 7 | 30.4 ± 3.0 | 6 | 30.9 ± 3.9 |
| HFHCD | 6 | 28.3 ± 3.9 | 8 | 27.2 ± 3.6 |
| HFHCD+liraglutide | 8 | 28.4 ± 3.9 | 5 | 33.1 ± 4.9 |
| HFHCD+metformin | 8 | 25.9 ± 1.7 | 7 | 29.5 ± 3.2 |
| Glomerular nestin (% area) | | | | |
| Control | 7 | 9.1 ± 1.7 | 6 | 11.1 ± 2.1* |
| HFHCD | 6 | 8.0 ± 1.8 | 8 | 7.2 ± 1.8 |
| HFHCD+liraglutide | 8 | 7.2 ± 3.2 | 5 | 6.9 ± 1.9 |
| HFHCD+metformin | 8 | 8.2 ± 2.1 | 7 | 11.0 ± 2.1** |
| Glomerular desmin (% area) | | | | |
| Control | 7 | 5.5 ± 2.9 | 6 | 3.7 ± 1.0 |
| HFHCD | 6 | 4.3 ± 2.8 | 8 | 3.5 ± 1.8 |
| HFHCD+liraglutide | 8 | 7.0 ± 4.2 | 5 | 3.9 ± 2.2 |
| HFHCD+metformin | 8 | 6.0 ± 3.0 | 7 | 4.2 ± 3.0 |
| Glomerular CD44 (% no of glom.) | | | | |
| Control | 7 | 23.3 ± 3.3* | 6 | 18.7 ± 8.0 |
| HFHCD | 6 | 40.0 ± 10.1 | 8 | 19.0 ± 9.6 |
| HFHCD+liraglutide | 8 | 40.5 ± 15.1 | 5 | 17.8 ± 7.3 |
| HFHCD+metformin | 8 | 38.3 ± 11.4 | 7 | 11.9 ± 5.6 |
| Glomerular collagen IV (% area) | | | | |
| Control | 7 | 15.6 ± 3.6 | 6 | 16.3 ± 2.0 |
| HFHCD | 6 | 10.2 ± 5.0 | 8 | 15.2 ± 5.6 |
| HFHCD+liraglutide | 8 | 13.6 ± 1.8 | 5 | 18.3 ± 3.8 |
| HFHCD+metformin | 8 | 13.0 ± 3.7 | 7 | 15.3 ± 5.8 |

Supplementary Table 2. Gender differences in HFHCD-induced renal injury- Cortex and mRNA.

aSMA = alpha smooth muscle actin, HFHCD = high-fat-high-carbohydrate diet, n = number, SD = standard deviation, TI = tubulointerstitial * p<0.05, ** p<0.01 versus HFHCD group.

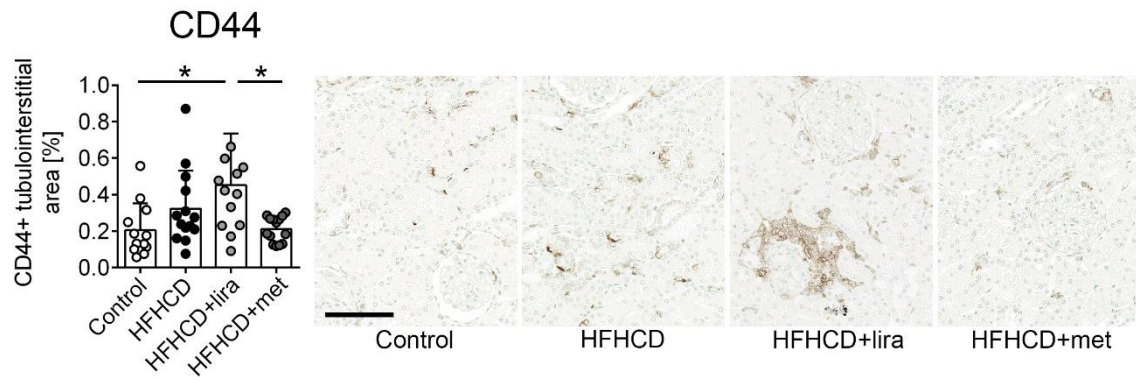
| Groups | Males (n=29) | | Females (n=28) | |
|---|-----------------|--------------|-------------------|-------------|
| | n | mean ± SD | n | mean ± SD |
| Cortex lipocalin 2 (no of tubules) | | | | |
| Control | 7 | 1.8 ± 0.6 | 6 | 0.6 ± 0.4 |
| HFHCD | 5 | 2.0 ± 0.3 | 8 | 1.0 ± 0.5 |
| HFHCD+liraglutide | 8 | 1.4 ± 0.8 | 5 | 1.1 ± 0.5 |
| HFHCD+metformin | 7 | 2.2 ± 0.7 | 6 | 0.8 ± 0.4 |
| Cortex TI injury (% area) | | | | |
| Control | 7 | 25.5 ± 20.1 | 6 | 14.4 ± 9.8 |
| HFHCD | 6 | 29.9 ± 21.3 | 8 | 10.6 ± 8.4 |
| HFHCD+liraglutide | 8 | 37.4 ± 15.4 | 5 | 22.5 ± 16.1 |
| HFHCD+metformin | 8 | 27.2 ± 21.3 | 7 | 18.7 ± 15.8 |
| Cortex CD68+ cells (no of cells) | | | | |
| Control | 7 | 20.0 ± 7.4 | 6 | 11.0 ± 3.6 |
| HFHCD | 6 | 32.5 ± 18.9 | 8 | 9.7 ± 4.2 |
| HFHCD+liraglutide | 8 | 33.8 ± 10.0 | 5 | 16.0 ± 4.9 |
| HFHCD+metformin | 8 | 28.2 ± 13.2 | 7 | 11.8 ± 5.3 |
| Cortex CD44 (% area) | | | | |
| Control | 6 | 0.1 ± 0.1 | 6 | 0.3 ± 0.2 |
| HFHCD | 6 | 0.5 ± 0.2 | 8 | 0.2 ± 0.1 |
| HFHCD+liraglutide | 8 | 0.5 ± 0.3 | 5 | 0.4 ± 0.2 |
| HFHCD+metformin | 7 | 0.3 ± 0.0 | 7 | 0.2 ± 0.1 |
| Cortex collagen I (% area) | | | | |
| Control | 7 | 4.4 ± 1.2 | 6 | 4.3 ± 1.3 |
| HFHCD | 6 | 5.2 ± 1.5 | 8 | 4.0 ± 1.3 |
| HFHCD+liraglutide | 8 | 5.8 ± 1.6 | 5 | 4.2 ± 1.0 |
| HFHCD+metformin | 8 | 5.0 ± 1.7 | 7 | 3.6 ± 0.8 |
| Cortex collagen IV (% area) | | | | |
| Control | 7 | 20.7 ± 2.7 | 6 | 20.3 ± 2.1 |
| HFHCD | 5 | 19.0 ± 2.1 | 8 | 19.9 ± 4.5 |
| HFHCD+liraglutide | 8 | 23.8 ± 2.2** | 5 | 22.6 ± 3.0 |
| HFHCD+metformin | 8 | 20.7 ± 3.1 | 7 | 18.9 ± 5.0 |
| Cortex aSMA (%area) | | | | |
| Control | 7 | 0.8 ± 0.4 | 6 | 0.9 ± 0.2 |
| HFHCD | 6 | 0.8 ± 0.2 | 8 | 0.8 ± 0.3 |
| HFHCD+liraglutide | 8 | 1.0 ± 0.3 | 5 | 1.3 ± 0.5* |
| HFHCD+metformin | 8 | 0.5 ± 0.2 | 7 | 0.8 ± 0.2 |
| Ccl2 mRNA expression (CCL2) | | | | |
| Control | 7 | 1.0 ± 0.6 | 6 | 1.1 ± 0.6 |
| HFHCD | 6 | 1.1 ± 1.0 | 8 | 1.1 ± 0.8 |
| HFHCD+liraglutide | 8 | 1.0 ± 0.6 | 5 | 0.5 ± 0.2 |
| HFHCD+metformin | 8 | 0.8 ± 0.3 | 7 | 0.8 ± 0.2 |
| Il1b mRNA expression (IL1-β) | | | | |
| Control | 7 | 1.0 ± 0.7 | 6 | 1.0 ± 1.3 |
| HFHCD | 6 | 1.5 ± 1.0 | 8 | 0.8 ± 0.4 |
| HFHCD+liraglutide | 8 | 2.2 ± 1.5 | 5 | 0.7 ± 0.2 |
| HFHCD+metformin | 8 | 0.9 ± 0.7 | 7 | 0.6 ± 0.4 |



Supplementary Figure 1. Ultrastructural changes in podocytes after HFHCD in female rats.

(A-C) Representative pictures from glomeruli of the control group. The glomerular structure was inconspicuous: only a few mesangial cells, open capillaries, fenestrated thin endothelium, thin basement membranes, podocytes with fine foot processes (arrow). (D-F) In the HFHCD group mesangial cells expanded in some glomeruli. Podocytes showed signs of stress: frequently, microvilli protruded into the urinary space (arrowheads) and very rare foot process effacement (arrow). (G-I) Both treatment groups, liraglutide and metformin, showed similar signs of podocyte stress. Microvilli formation (arrowheads) was observed in several glomeruli.

HFHCD = high-fat-high-carbohydrate diet; C = capillary, E = endothelial cell, GBM = glomerular basement membrane, M = mesangial cell, P = podocyte, US = urinary space, arrow = podocyte foot process, arrowhead = microvilli; scale bars = 5000, 1000 and 500 nm, respectively.



Supplementary Figure 2. Tubulointerstitial CD44 expression.

CD44 positive area of kidney cortex was quantified in immunohistochemical stainings. The HFHCD+liraglutide treated rats showed significantly higher expression compared to the control and HFHCD+metformin group.

HFHCD = high-fat-high-carbohydrate diet, lira = liraglutide, met = metformin; shown are individual animals and mean \pm SD; * $p < 0.05$.