

**Electronic supplementary information (ESI) for**  
**Ferulic acid supplementation alleviates hyperuricemia in high-fructose/fat diet-**  
**fed rats *via* promoting uric acid excretion and mediating the gut microbiota**

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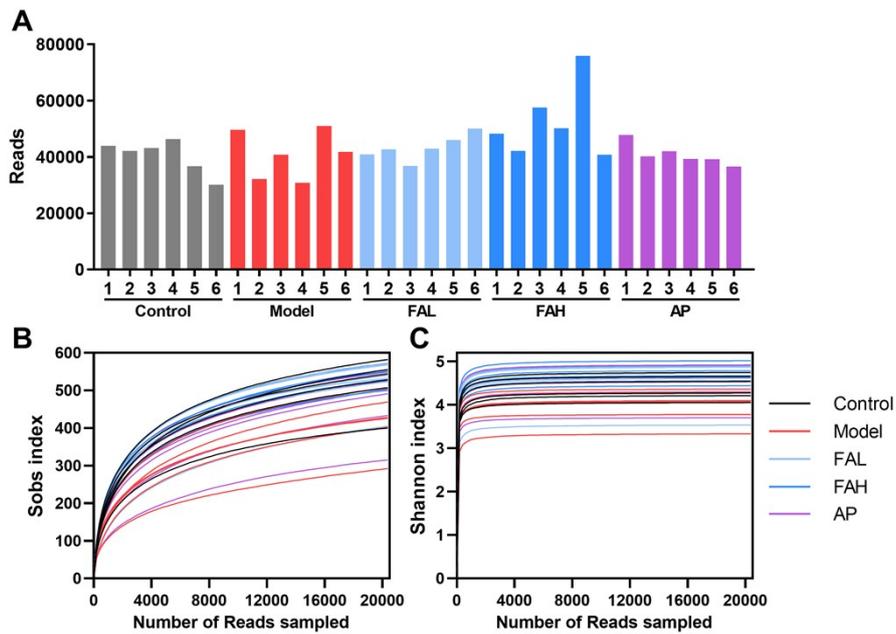
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## Figure



**Fig. S1** Diversity estimation of the gut microbiota in each sample. (A) The number of clean reads. Rarefaction curves based on Sobs (B) and Shannon (C) indices at the operational taxonomic unit level. Control: control group supplied with a standard diet; Model: model group supplied with a HFFD; FAL: low-dose FA group supplied with a HFFD containing 0.05% FA; FAH: high-dose FA group supplied with a HFFD containing 0.1% FA; AP: allopurinol group supplied with a HFFD containing 0.0145% allopurinol.

## Tables

**Table S1** Composition of diets in different groups

|                                  | Control | Model | FAL | FAH | AP    |
|----------------------------------|---------|-------|-----|-----|-------|
| corn starch (g/kg)               | 250     | -     |     |     |       |
| wheat middling (g/kg)            | 200     | -     |     |     |       |
| refined wheat flour (g/kg)       | 219     | 420   | 420 | 420 | 420   |
| soybean meal (g/kg)              | 200     | 80    | 80  | 80  | 80    |
| fructose (g/kg)                  | -       | 180   | 180 | 180 | 180   |
| lard (g/kg)                      | -       | 200   | 200 | 200 | 200   |
| chicken powder (g/kg)            | -       | 80    | 80  | 80  | 80    |
| sodium cholate (g/kg)            | -       | 2     | 2   | 2   | 2     |
| miscellaneous meal (g/kg)        | 45      | -     | -   | -   | -     |
| vegetable oil (g/kg)             | 20      | -     | -   | -   | -     |
| fish meal (g/kg)                 | 20      | -     | -   | -   | -     |
| calcium bicarbonate (g/kg)       | 10      | 10    | 10  | 10  | 10    |
| mountain flour (g/kg)            | 16      | 8     | 8   | 8   | 8     |
| vitamin & mineral mixture (g/kg) | 20      | 20    | 20  | 20  | 20    |
| ferulic acid (g/kg)              | -       | -     | 0.5 | 1   | -     |
| allopurinol (g/kg)               | -       | -     | -   | -   | 0.145 |

Control: control group supplied with a standard diet; Model: model group supplied with a HFFD; FAL: low-dose FA group supplied with a HFFD containing 0.05% FA; FAH: high-dose FA group supplied with a HFFD containing 0.1% FA; AP: allopurinol group supplied with a HFFD containing 0.0145% allopurinol.

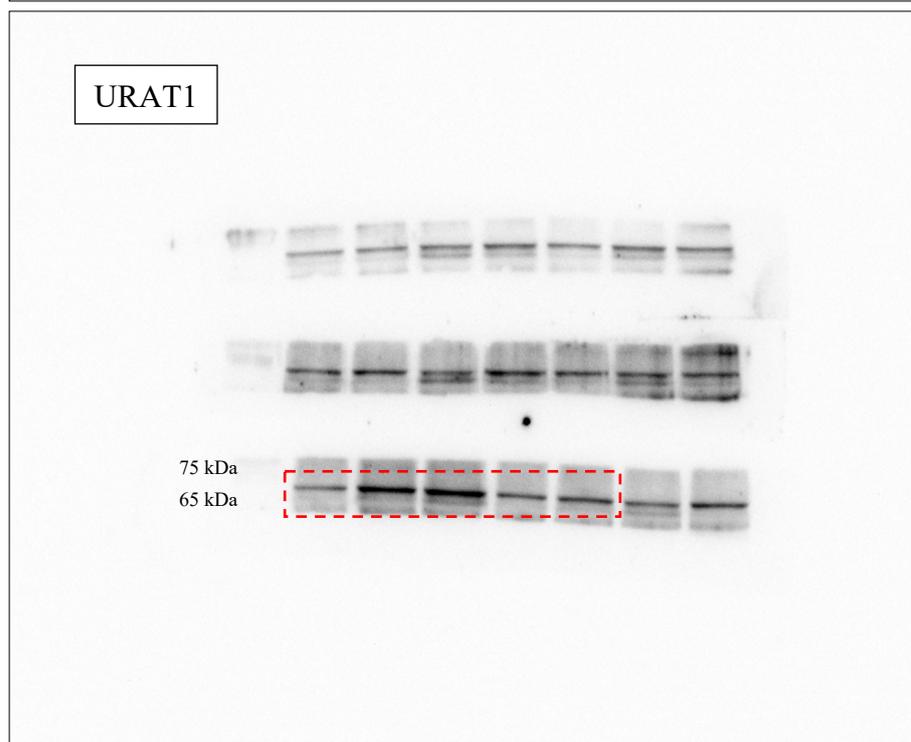
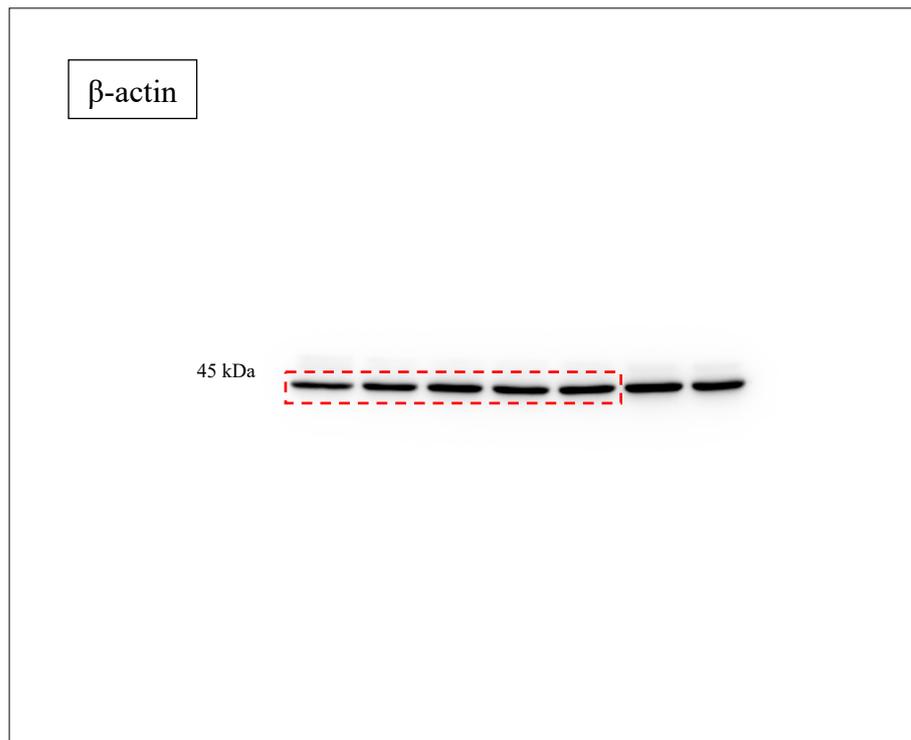
**Table S2** Sequences of the human-specific primers used for quantitative real-time polymerase chain reaction

| Gene            | Sense | Sequence (5'-3')        | No.            |
|-----------------|-------|-------------------------|----------------|
| <i>SLC22A12</i> | F     | CGTGTACTGCCTGTTCCGCT    | NM_144585.4    |
|                 | R     | CCACTCCATCAGGAGAGTGCC   |                |
| <i>SLC2A9</i>   | F     | CCTCTACGGCTACAACCT      | NM_020041.3    |
|                 | R     | GATGGCGAATATGGACACA     |                |
| <i>TLR4</i>     | F     | GGACCTTTCCAGCAACAAGA    | NM_138554.5    |
|                 | R     | CAGGGACAGGTCTAAAGAGAGA  |                |
| <i>MYD88</i>    | F     | CCAGTTTGTGCAGGAGATGA    | NM_001172567.2 |
|                 | R     | GCTCACTAGCAATAGACCAGAC  |                |
| <i>IKBKB</i>    | F     | CCAGCCAAGAAGAGTGAAGAA   | NM_001556.3    |
|                 | R     | CCGTGAAACTCTGGTCTTGT    |                |
| <i>NFKBIA</i>   | F     | CTCCGAGACTTTCGAGGAAATAC | NM_020529.3    |
|                 | R     | CCATTGTAGTTGGTAGCCTTCA  |                |
| <i>NFKB1</i>    | F     | GAGACATCCTTCCGCAAACCT   | NM_003998.4    |
|                 | R     | GGTCCTTCCTGCCATAATC     |                |
| <i>GAPDH</i>    | F     | GATTCCACCCATGGCAAATTC   | NM_002046.7    |
|                 | R     | CTGGAAGATGGTGATGGGATT   |                |

**Table S3** Sequences of the rat-specific primers used for quantitative real-time polymerase chain reaction

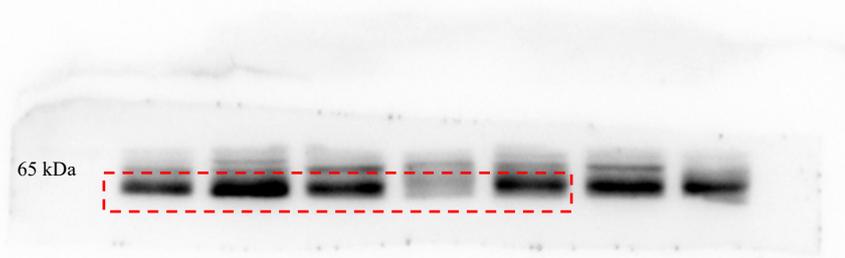
| Gene            | Sense | Sequence (5'-3')        | No.            |
|-----------------|-------|-------------------------|----------------|
| <i>Slc22a12</i> | F     | TCTTCTGGTGTCTGTGTCT     | NM_001034943.1 |
|                 | R     | GCTGGCTGTGTTTCATCAT     |                |
| <i>Slc2a9</i>   | F     | CTGAGAGTCCACGATACCT     | NM_001191551.1 |
|                 | R     | CGCATTGAGTCCACATAGT     |                |
| <i>Slc22a9</i>  | F     | AAGTATGAGGCTGGTGAGA     | NM_177481.1    |
|                 | R     | GTGGTCGTCAATGTAGGAG     |                |
| <i>Slc22a13</i> | F     | GAGGCTCACTACTGTTCAG     | NM_001126285.1 |
|                 | R     | GTGGCTCAGGATGTCTTC      |                |
| <i>Abcg2</i>    | F     | GGACAAGCATAGGAATGGA     | NM_181381.2    |
|                 | R     | CTGGTGAATGGAGAAGATGA    |                |
| <i>Slc22a6</i>  | F     | AATGTGGAATGGATGCCTAT    | NM_017224.2    |
|                 | R     | GAAGAACCAAGAGTAGATGAAG  |                |
| <i>Slc22a8</i>  | F     | TGAGCACCGTTATCTTGAAT    | NM_031332.1    |
|                 | R     | CACTGAGGAATGGCATAGG     |                |
| <i>Slc22a1</i>  | F     | CTGTGTGAACTTGGGCTTCT    | NM_012697.1    |
|                 | R     | GTCACCAAGAGACAGAGCTTAC  |                |
| <i>Slc22a2</i>  | F     | CTGGAGTGGCCTATGTGATTC   | NM_031584.2    |
|                 | R     | CCGGTATGCACCAGAAATAGAG  |                |
| <i>Slc22a4</i>  | F     | GAGGAGATTTGAAGAGGCAGAA  | NM_022270.2    |
|                 | R     | GCTCCAGAGGATCGAATATCAC  |                |
| <i>Slc22a5</i>  | F     | CCTCAGTCGCACCACATTTA    | NM_019269.1    |
|                 | R     | ATGGTCAGCCACAGGATTATG   |                |
| <i>Slc17a1</i>  | F     | GTGCATTCCCTGGAGTCTTGATA | NM_133554.2    |
|                 | R     | CATTGGGTATGCTCCAGATAGG  |                |
| <i>Tlr4</i>     | F     | CAGAGCCGTTGGTGTATCTT    | NM_019178.2    |
|                 | R     | AGCAAGGACTTCTCCACTTTC   |                |
| <i>Myd88</i>    | F     | GAAGCGACTGATCCCTATCAAG  | NM_198130.2    |
|                 | R     | TTGGTGCAAGGGTTGGTATAG   |                |
| <i>Ikkkb</i>    | F     | CCAAGAGACCAAAGGACAGAAG  | NM_053355.3    |
|                 | R     | GAGAGTAAATCACCCGCACTTT  |                |
| <i>Nfkbia</i>   | F     | AGTAACCTACCAGGGCTACTC   | NM_001105720.2 |
|                 | R     | ATAGCTCTCCTCATCCTCACTC  |                |
| <i>Nfkb1</i>    | F     | GGTTACGGGAGATGTGAAGATG  | NM_001276711.1 |
|                 | R     | GTGGATGATGGCTAAGTGTAGG  |                |
| <i>Gapdh</i>    | F     | GCTGCCTTCTCTTGTGAC      | NM_017008.4    |
|                 | R     | CTTGACTGTGCCGTTGAA      |                |

## Raw images of Western blot



GLUT9

65 kDa



ABCG2

75 kDa

65 kDa



OAT3



OCT1



OCT2

65 kDa

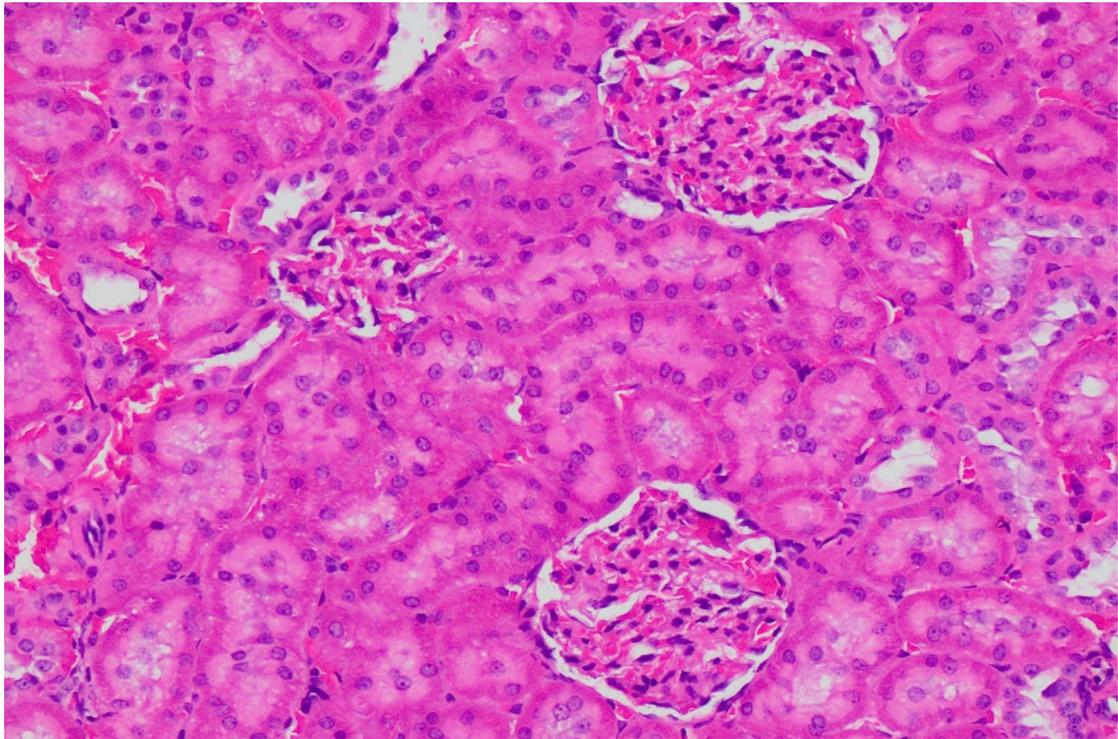


OCTN2

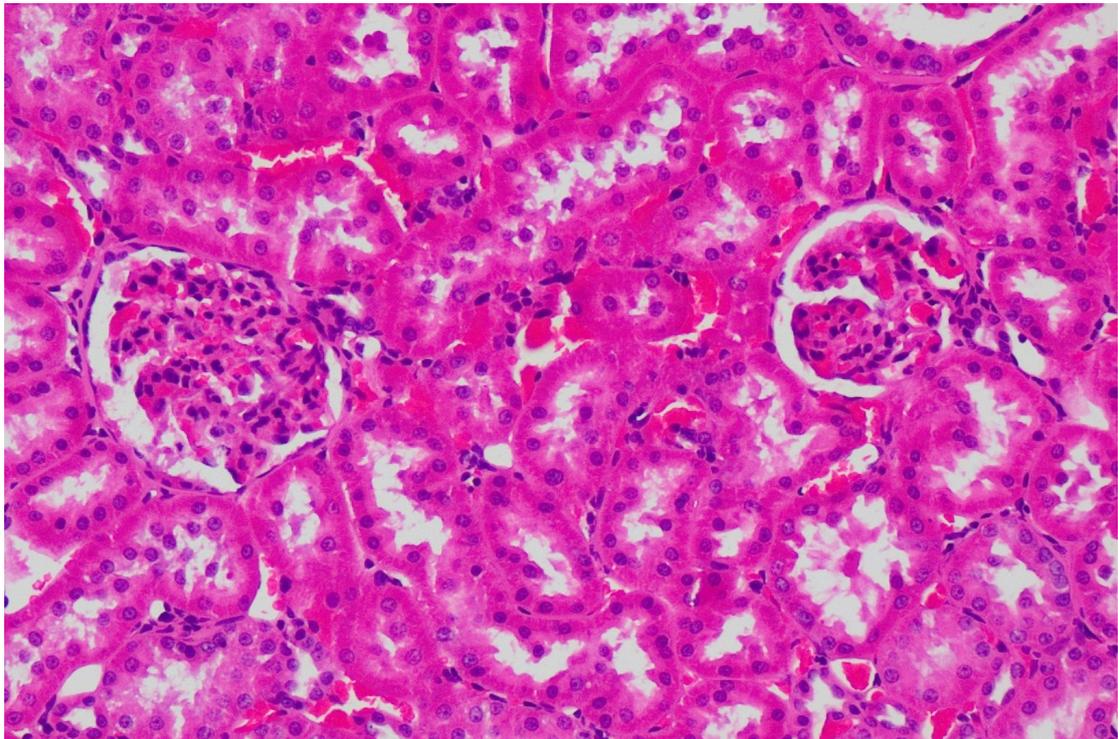
75 kDa



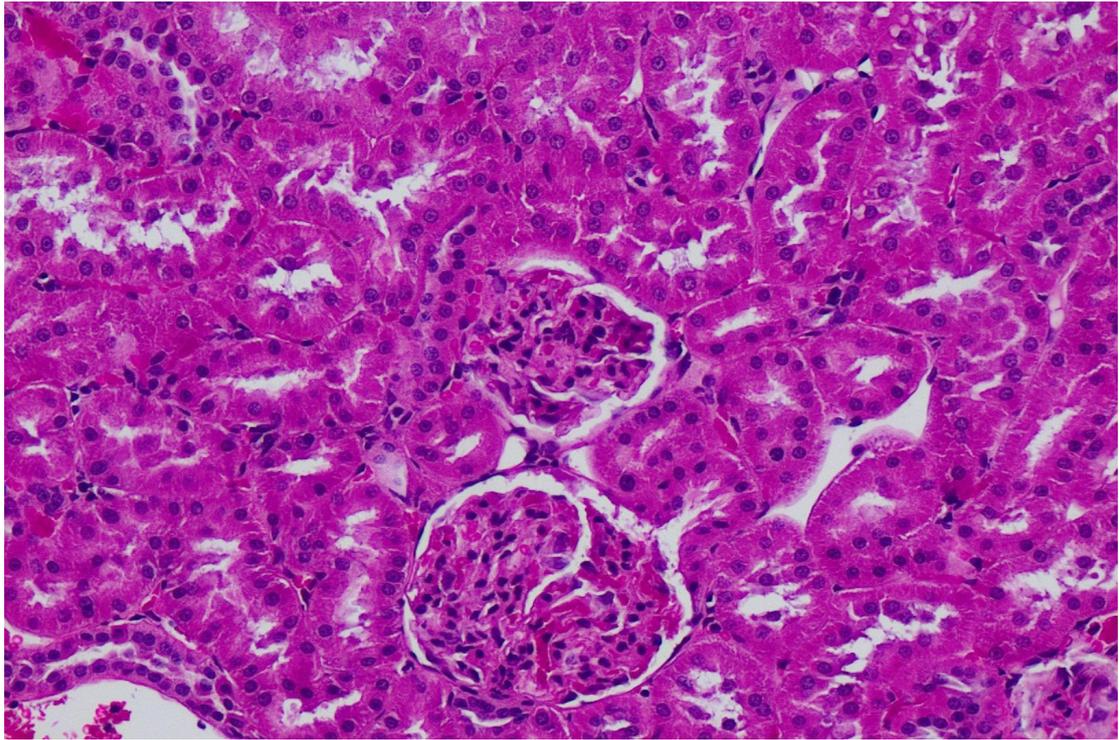
**Raw images of H&E staining micrographs in kidney tissues**



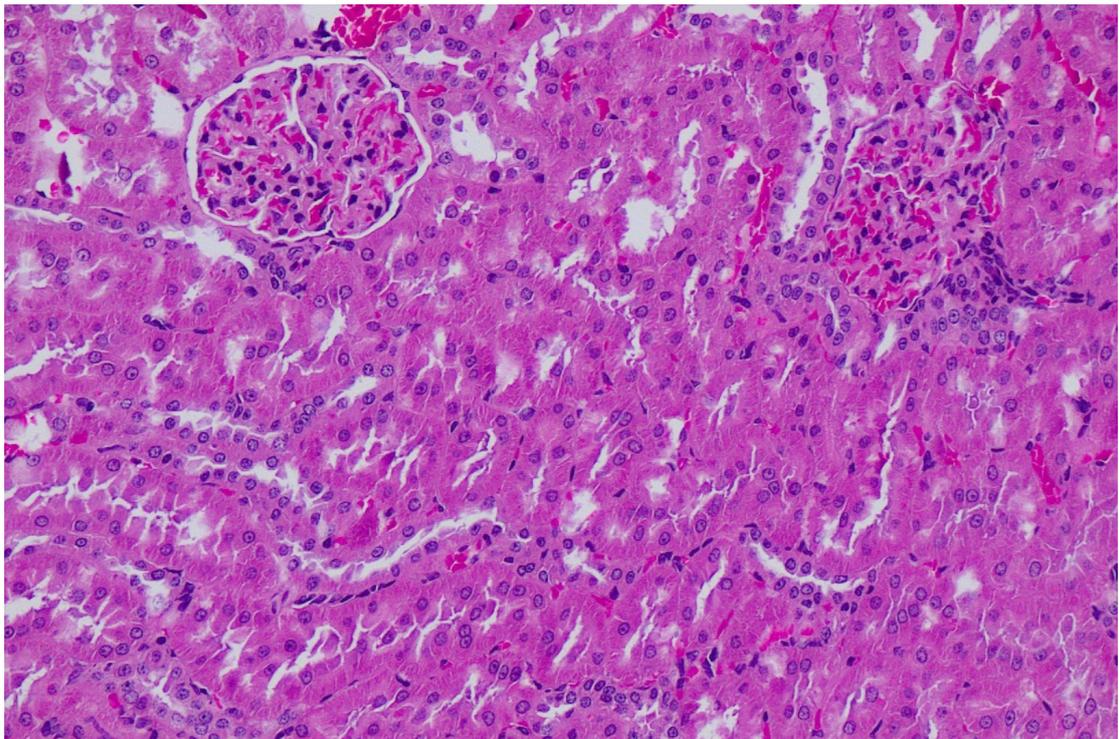
**Control**



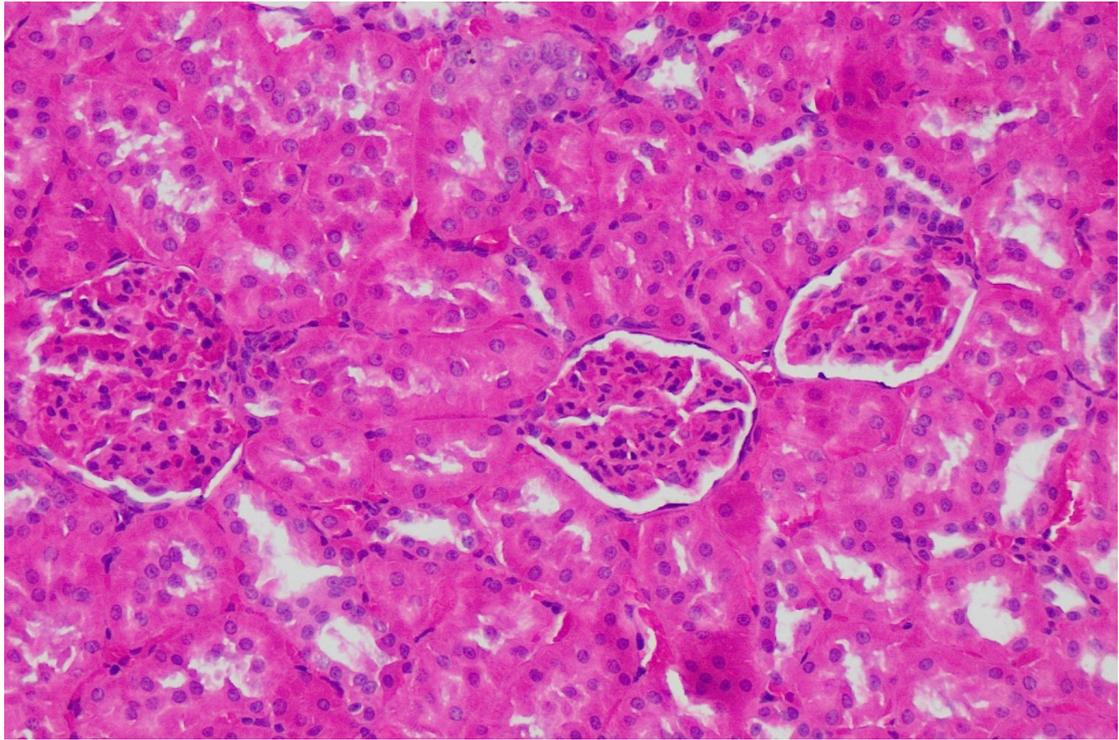
**Model**



**FAL**



**FAH**



AP