

Supplementary Materials

Kaseda, S., et al.

Contents

Supplementary Method

Endothelin-1 concentration measurement

Supplementary Figures

Figure S1. UBE-1099 slightly reduced the body weight and increased the urine volume in Alport mice

Figure S2. UBE-1099 did not affect the heart rate and blood pressure in Alport mice

Figure S3. Transcriptome analysis reveals the comprehensive effects of UBE-1099 in the glomeruli of Alport mice

Figure S4. Dysregulated GO terms in the glomeruli of Alport mice

Figure S5. Up-regulated GO terms for Alport vehicle vs WT

Figure S6. Down-regulated GO terms for Alport Vehicle vs WT

Figure S7. Up-regulated GO terms for Alport UBE-1099 vs Alport Vehicle

Figure S8. Down-regulated GO terms for Alport UBE-1099 vs Alport Vehicle

Figure S9. UBE1099 altered genes in each condition

Figure S10. Expression level of cell specific markers in the glomerular cell

Figure S11. UBE-1099 did not affect the food intake and muscle weight in Alport mice and did not worsen the early renal pathology

Figure S12. UBE-1099 did not affect the endothelin expression

Figure S13. Full length blots for Figure 5A, B

Supplementary Method

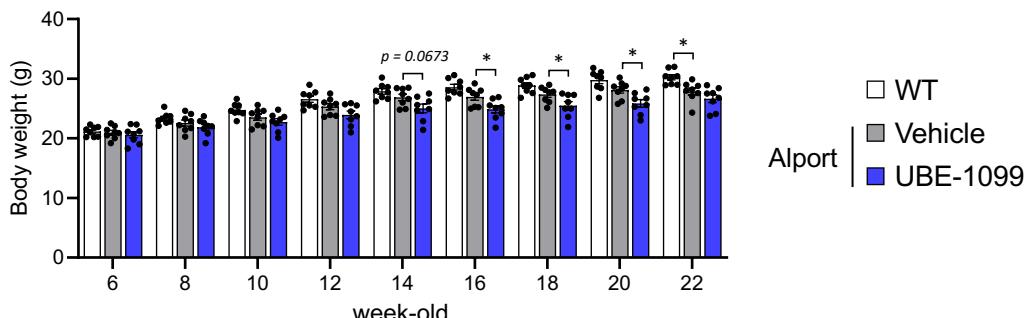
Endothelin-1 concentration measurement

Rat proximal tubular cells (NRK-52E cells) were treated with vehicle (dimethyl sulfoxide, 0.1% final concentration) or 10-100 nM of CDDO-Im or UBE-1099. After 24 h, cell culture media were collected and assessed for concentrations of secreted endothelin-1 (ET-1) using ELISA kit (R&D Systems, USA) according to the manufacturer's protocol. Cell viability was measured by Cell Titer-Glo Luminescent cell viability assay (Promega) according to the manufacturer's protocol.

Supplementary Figures

Figure S1

A



B

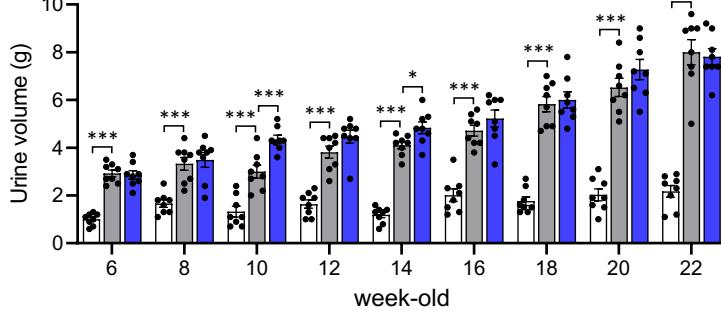
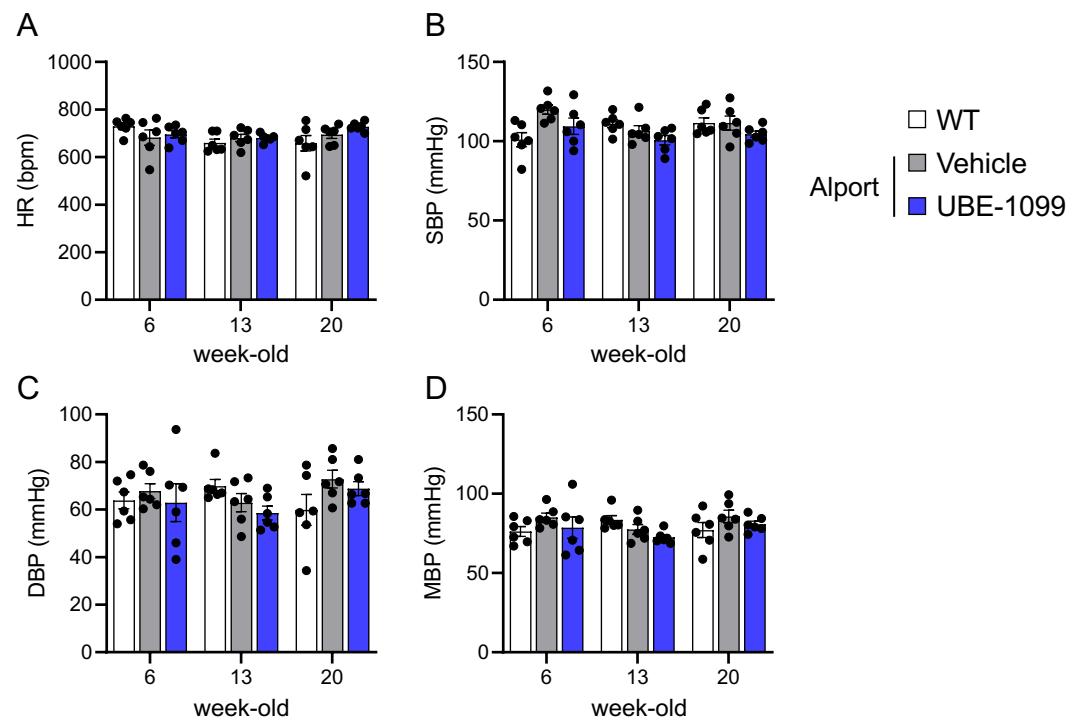


Figure S1. UBE-1099 slightly reduced the body weight and increased the urine volume in Alport mice

(A, B) Body weight and urine volume were measured every two weeks. Urine volume was measured using metabolic cages for 24h. Data are presented as mean \pm SE ($n = 8$ per group). P values were assessed by Dunnett's test. (* $p < 0.05$, *** $p < 0.001$).

Figure S2**Figure S2. UBE-1099 did not affect the heart rate and blood pressure in Alport mice**

(A-D) Heart rate (HR), Systolic blood pressure (SBR), Diastolic blood pressure (DBP) and Mean blood pressure MBP were measured by BP-98A-1 (Softron). Data are presented as mean \pm SE (n = 6 per group).

Figure S3

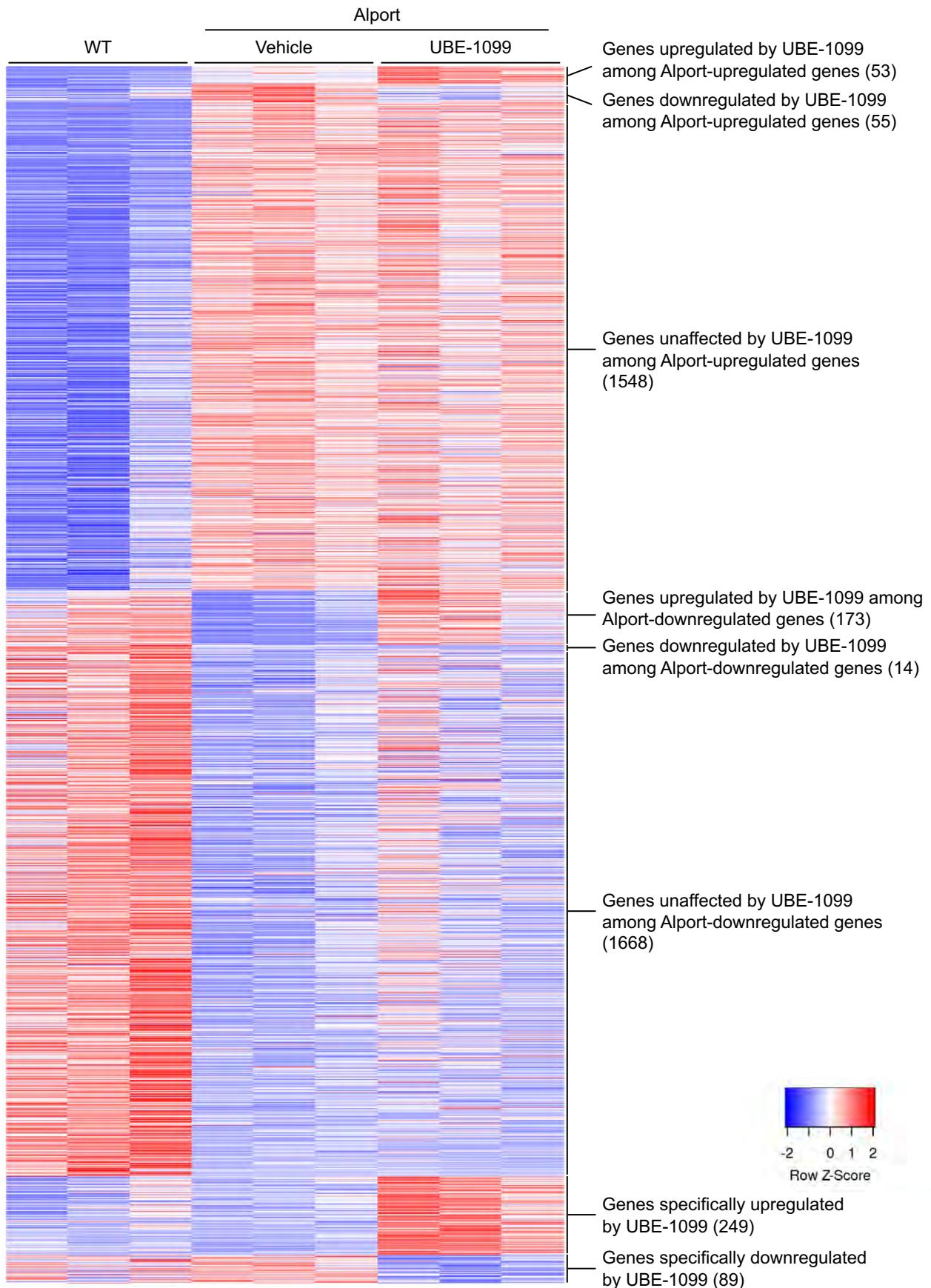


Figure S3. Transcriptome analysis reveals the comprehensive effects of UBE-1099 in the glomeruli of Alport mice

Heatmap shows the number of fluctuated genes in each condition (fold change > 1.2 or < -1.2 , $p < 0.05$).

Figure S4

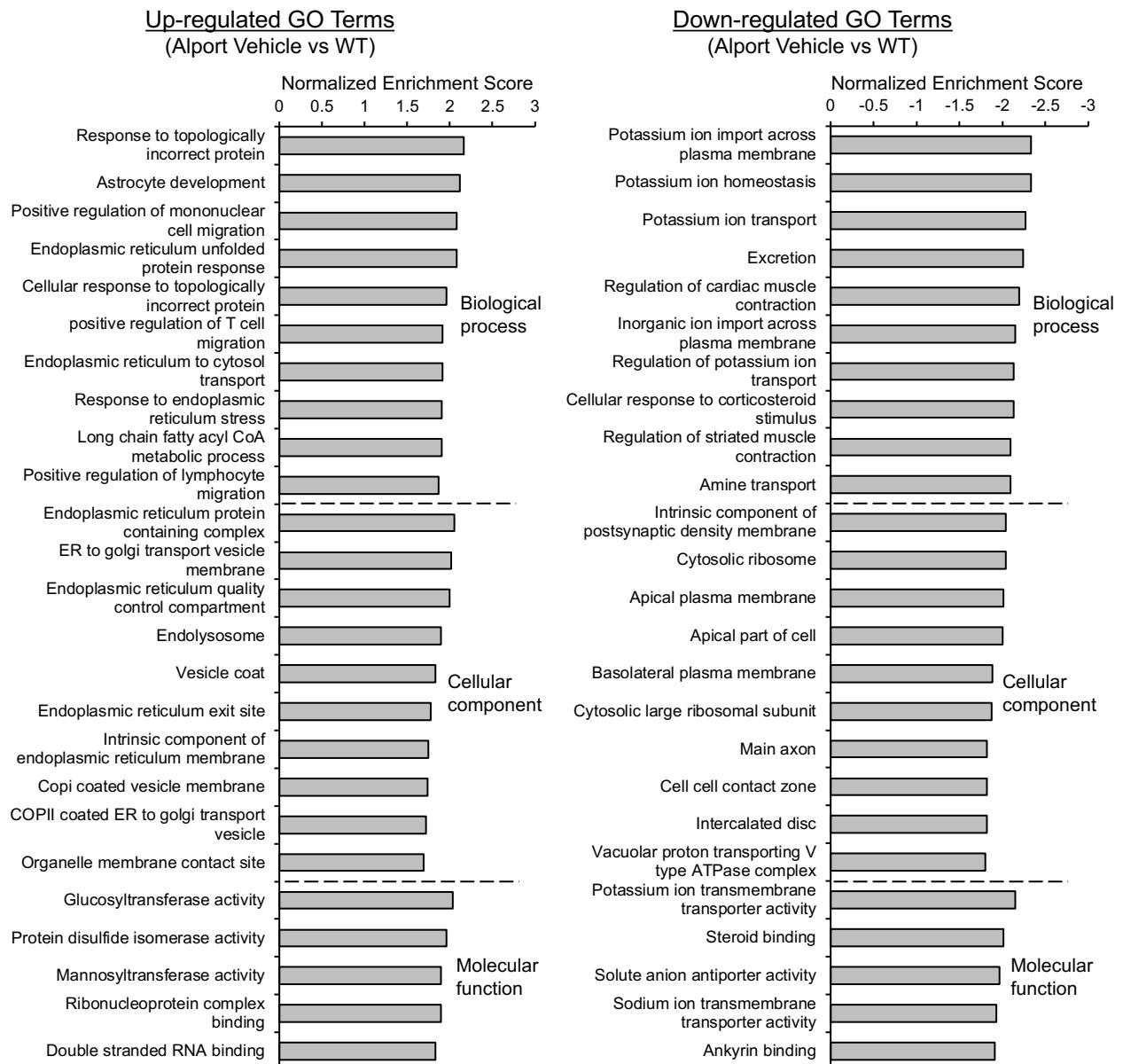
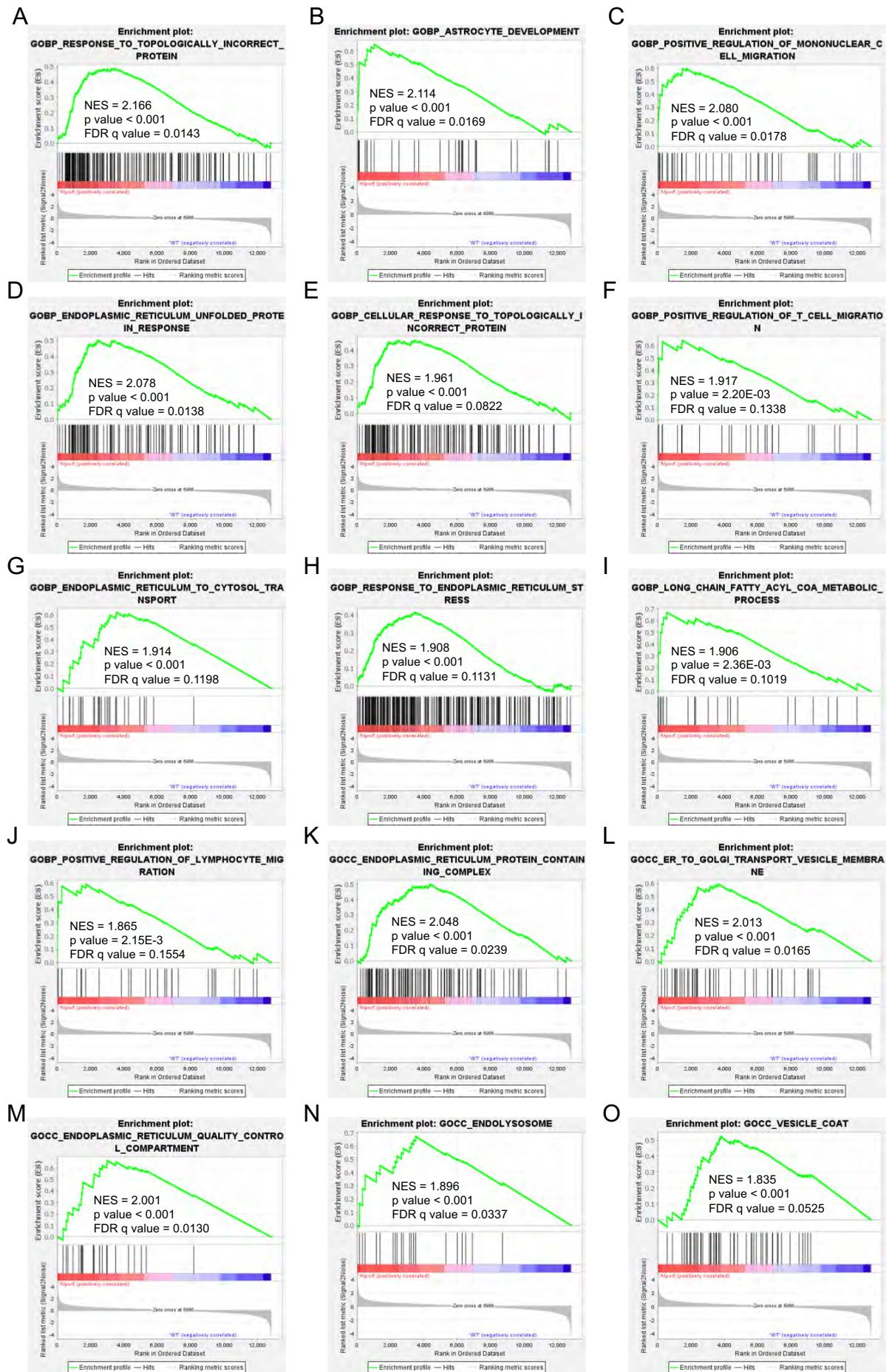


Figure S4. Dysregulated GO terms in the glomeruli of Alport mice

Gene Set Enrichment Analysis (GSEA) for Alport Vehicle vs WT

Figure S5



(Continue to the next page)

Figure S5

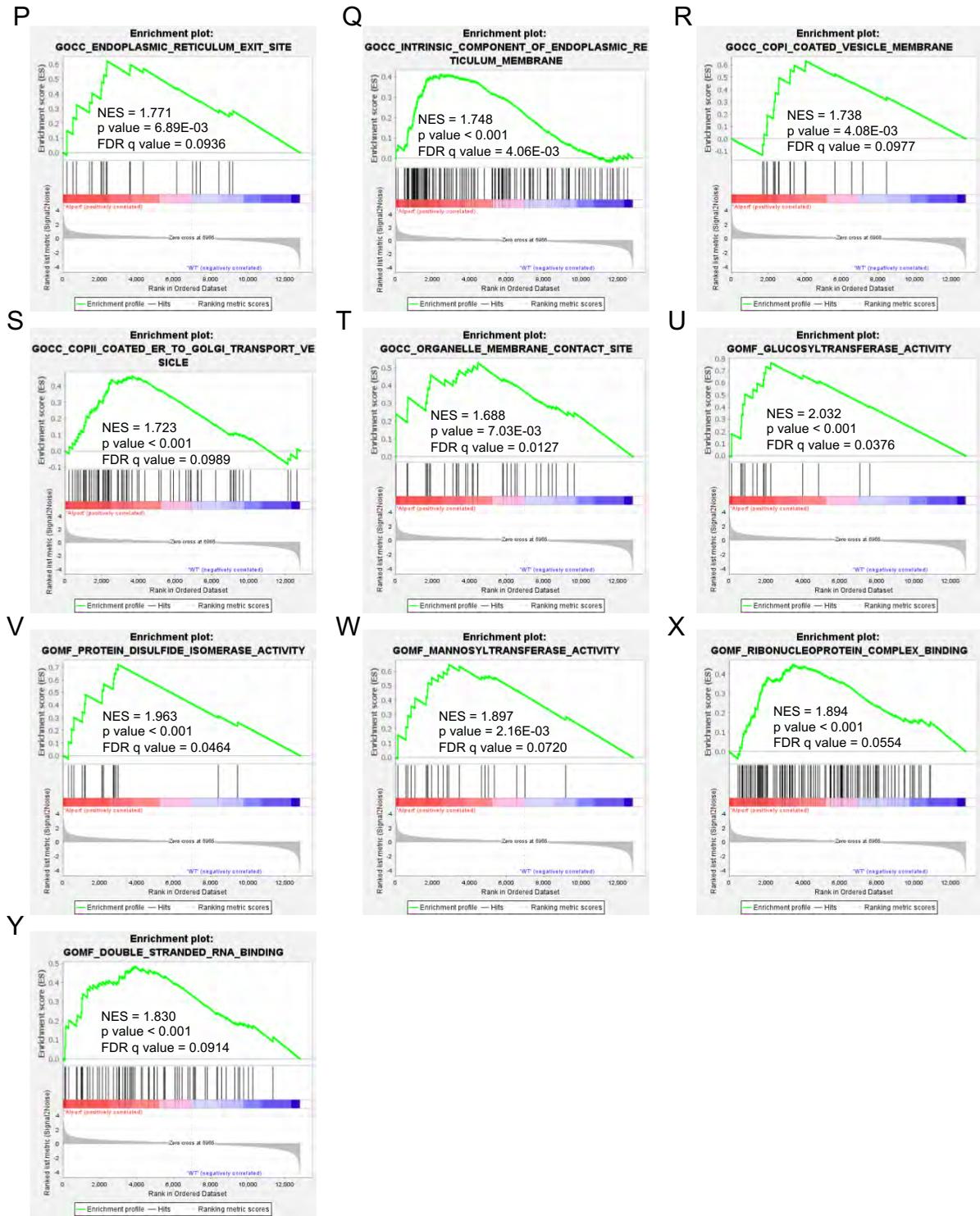
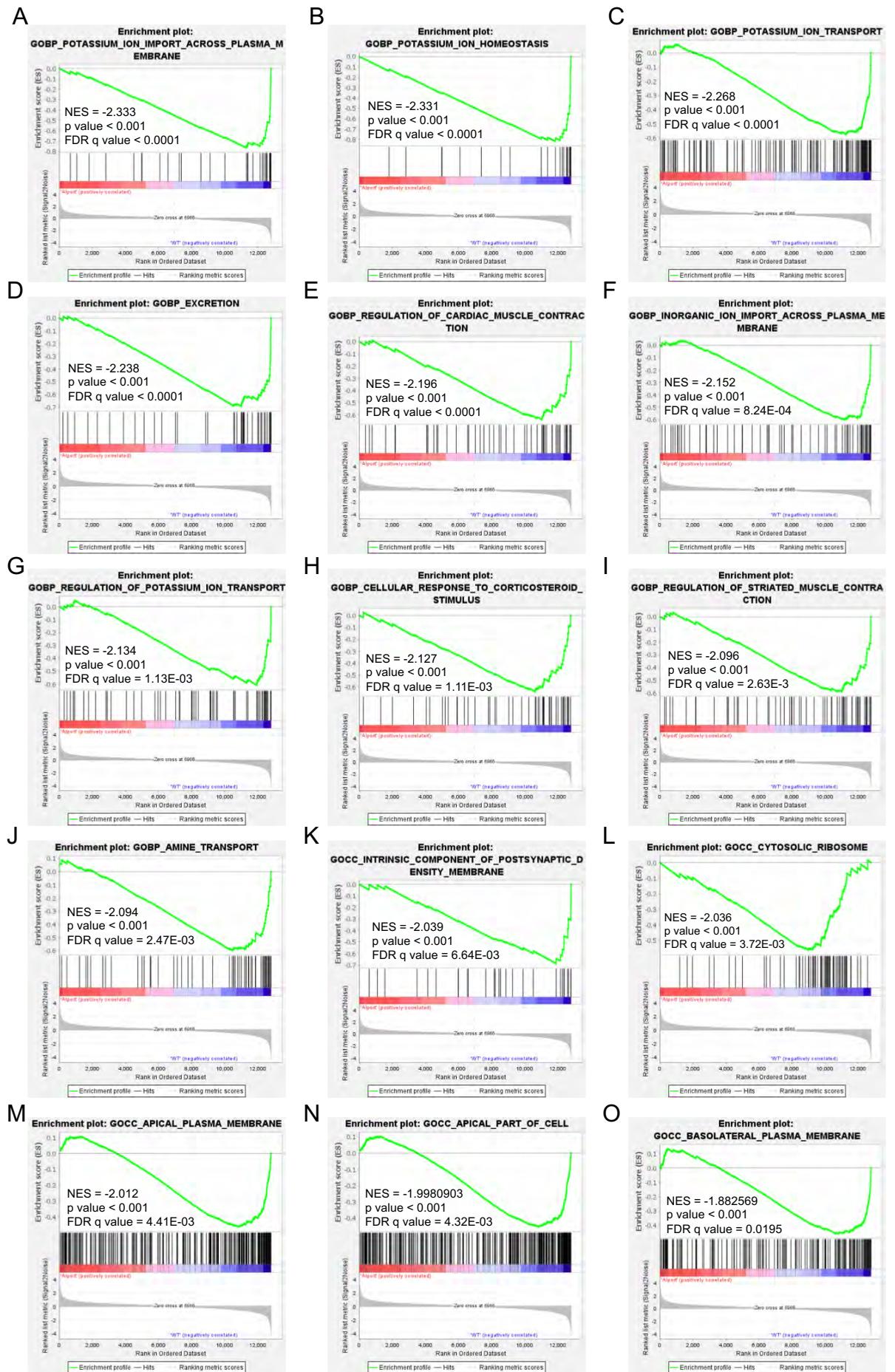


Figure S5. Up-regulated GO terms for Alport vehicle vs WT

Gene Set Enrichment Analysis (GSEA) data for up-regulated GO terms in Figure S4. NES = Normalized Enrichment Score.

Figure S6



(Continue to the next page)

Figure S6

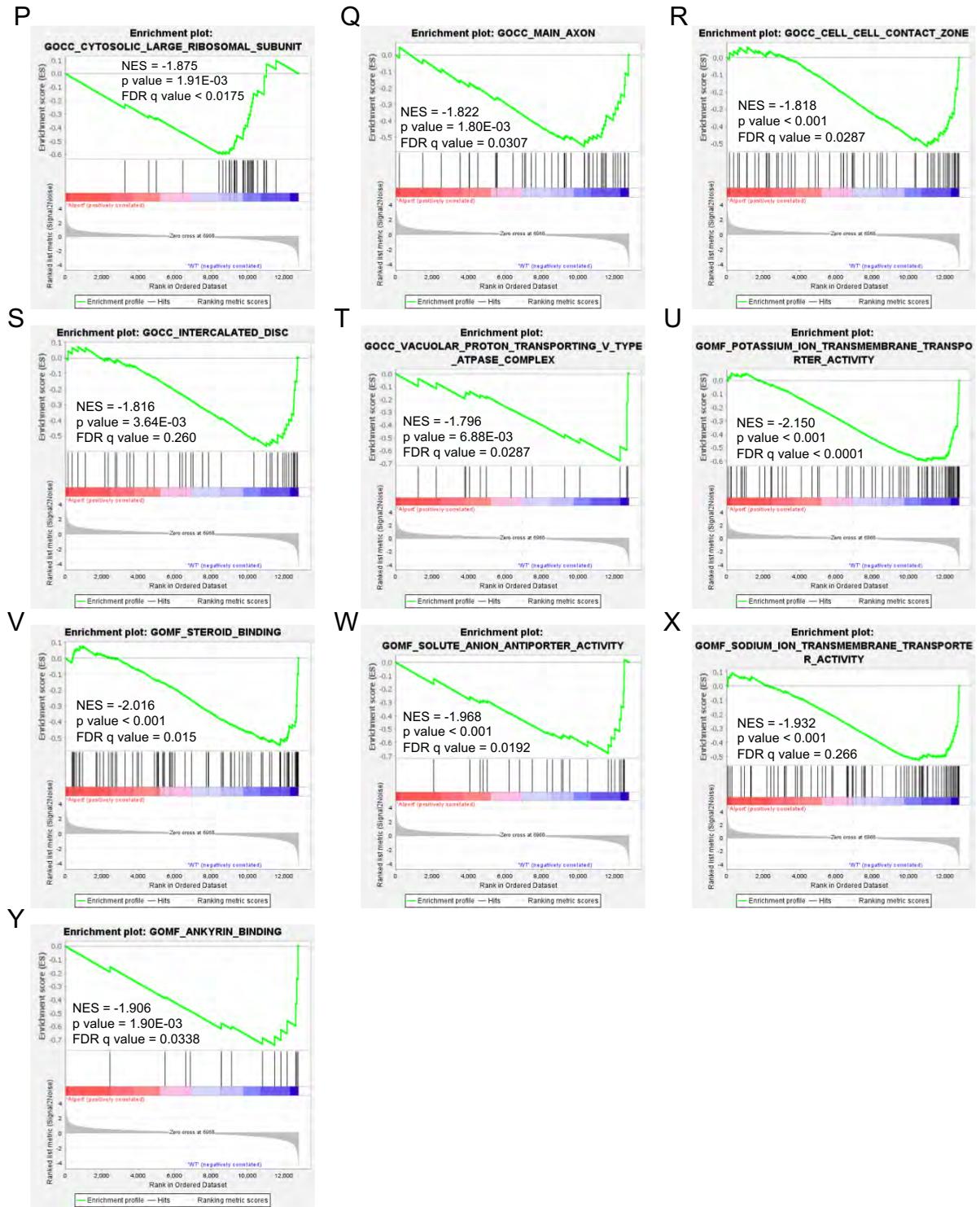
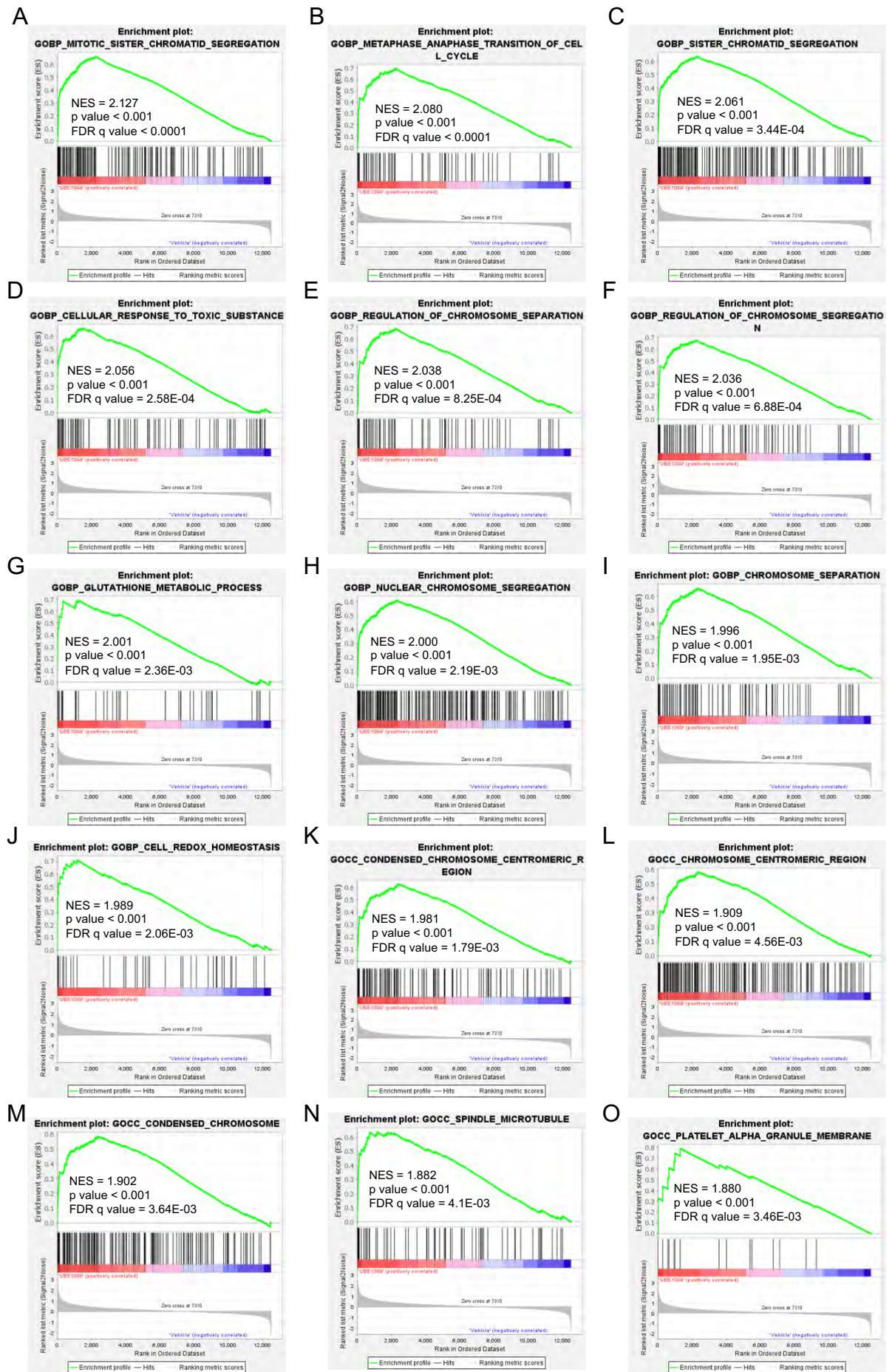


Figure S6. Down-regulated GO terms for Alport Vehicle vs WT

Gene Set Enrichment Analysis (GSEA) data for up-regulated GO terms in Figure S4. NES = Normalized Enrichment Score.

Figure S7



(Continue to the next page)

Figure S7

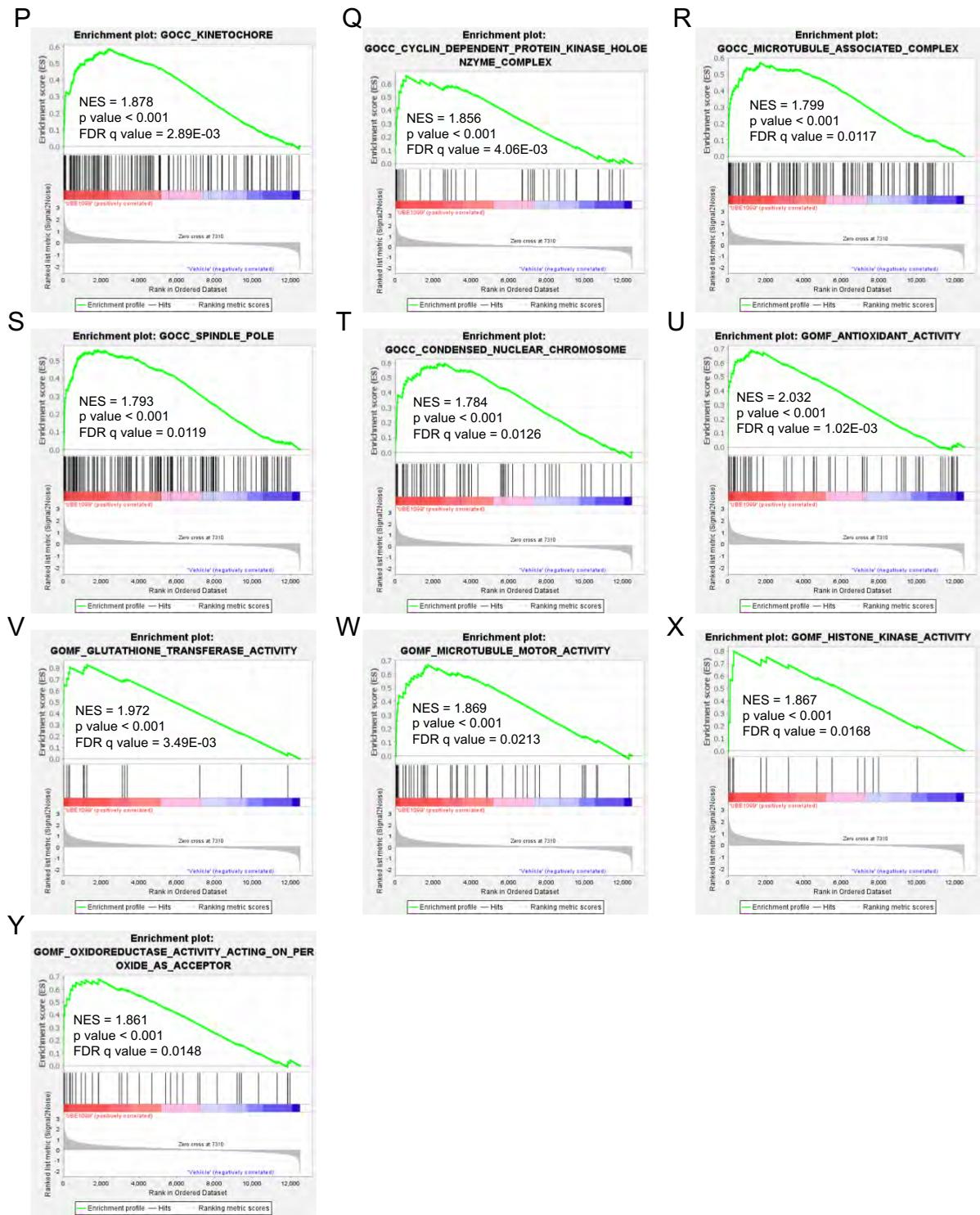
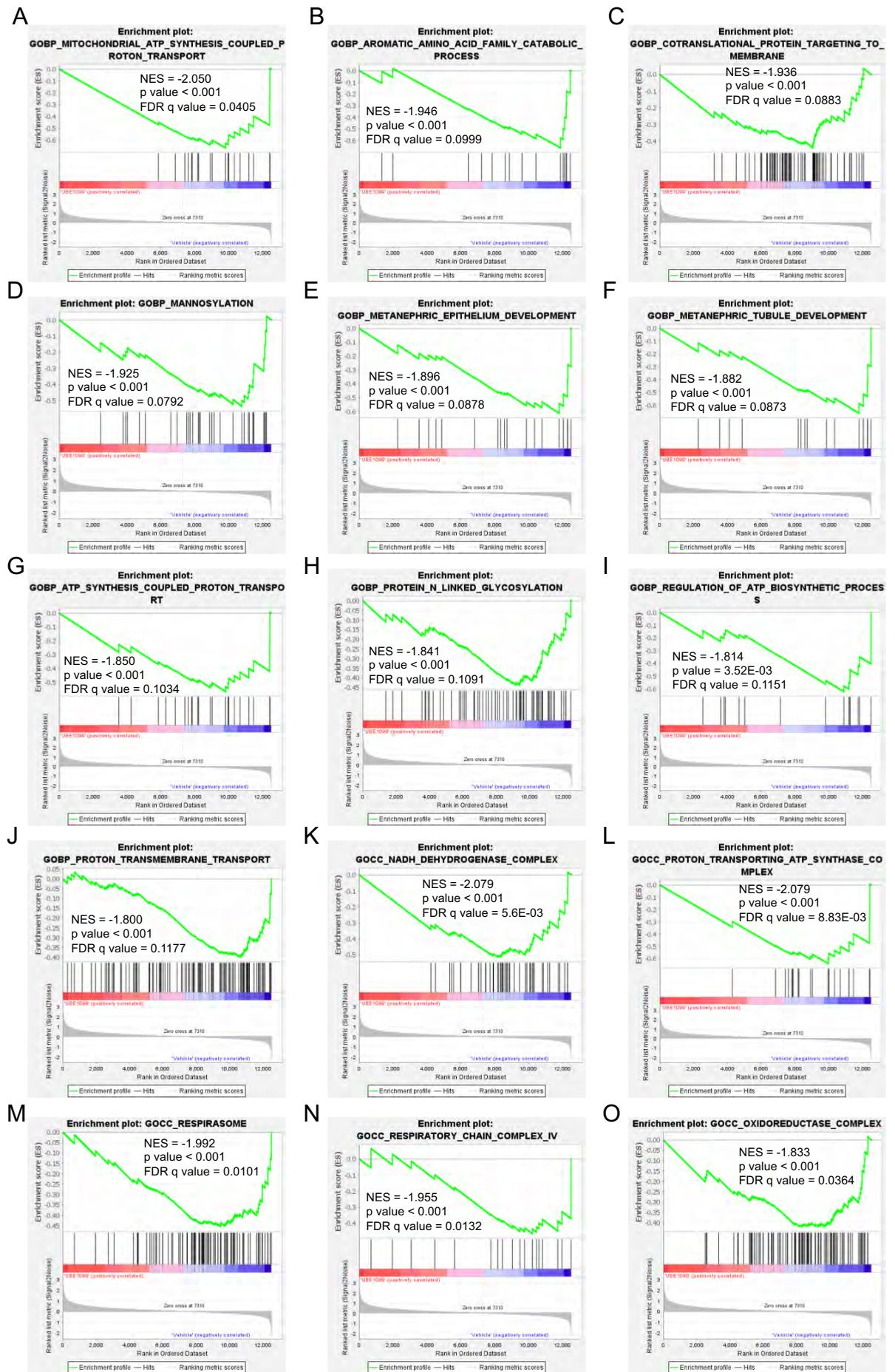


Figure S7. Up-regulated GO terms for Alport UBE-1099 vs Alport Vehicle

Gene Set Enrichment Analysis (GSEA) data for up-regulated GO terms in Figure 6B. NES = Normalized Enrichment Score.

Figure S8



(Continue to the next page)

Figure S8

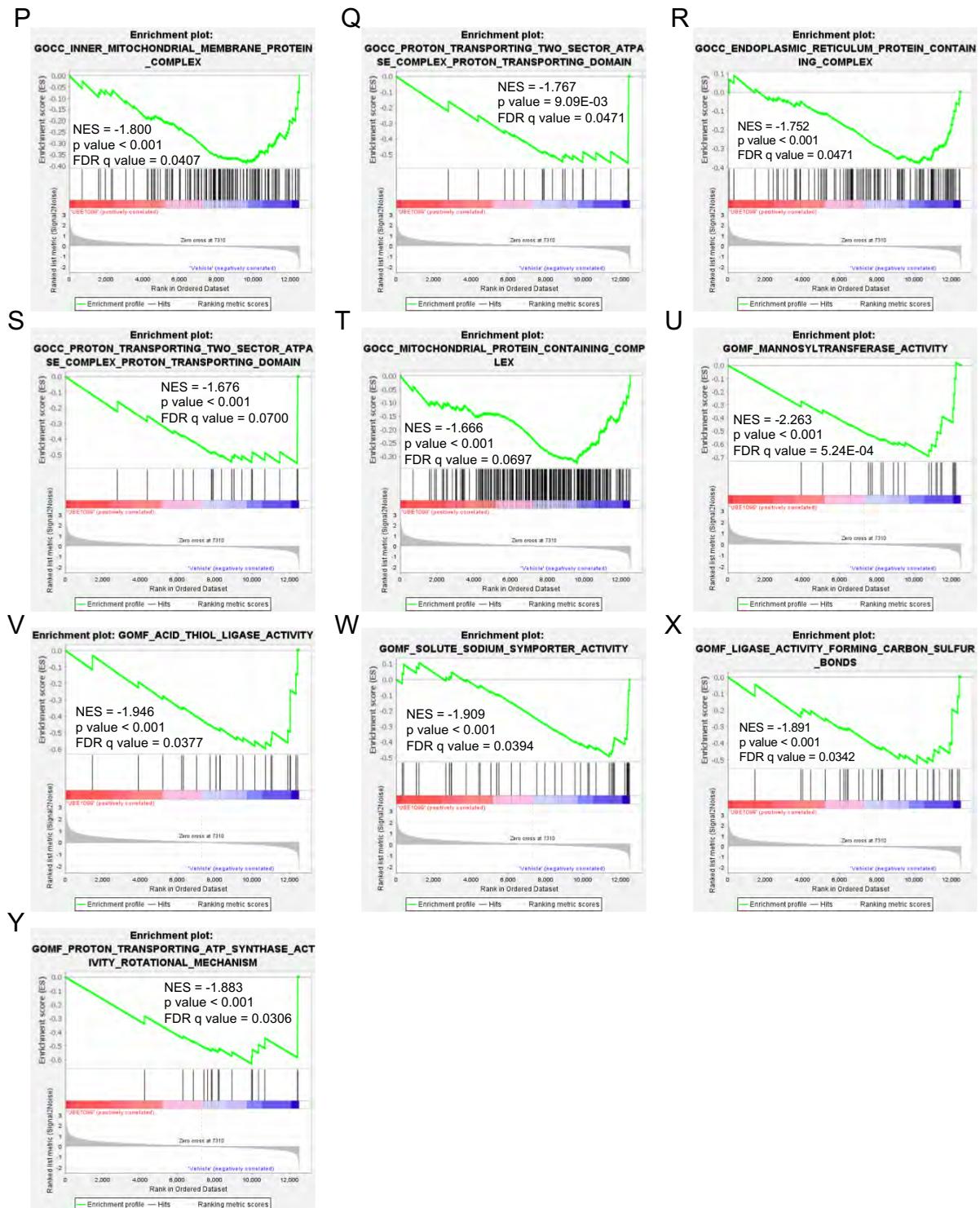
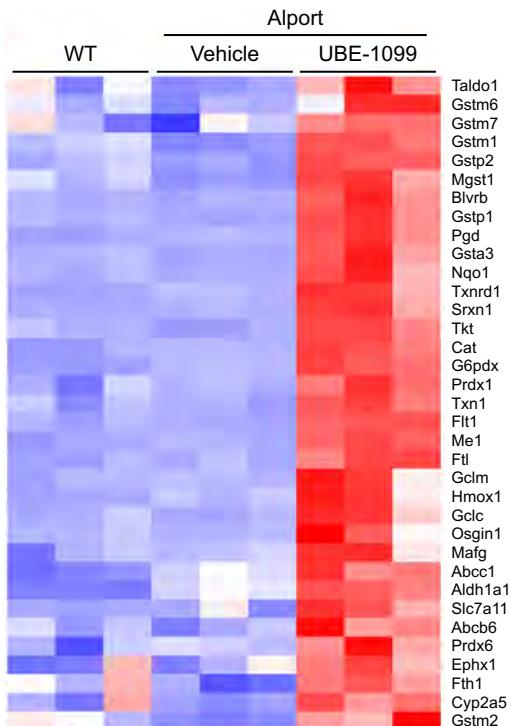


Figure S8. Down-regulated GO terms for Alport UBE-1099 vs Alport Vehicle

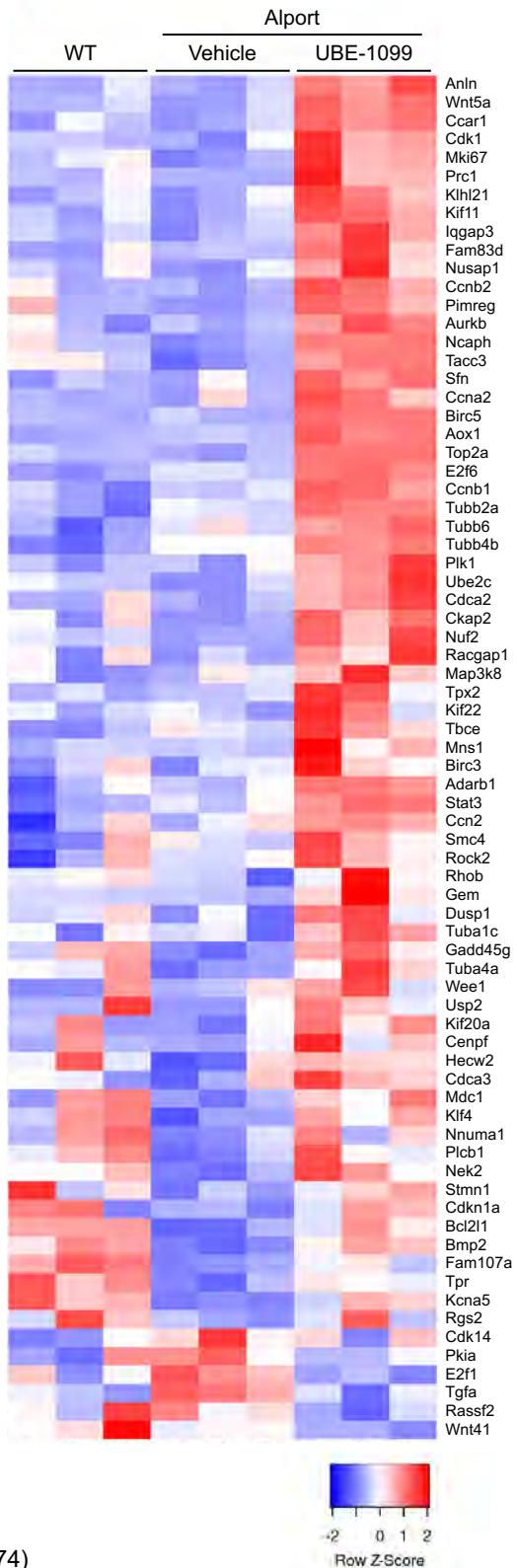
Gene Set Enrichment Analysis (GSEA) data for down-regulated GO terms in Figure 6B. NES = Normalized Enrichment Score.

Figure S9

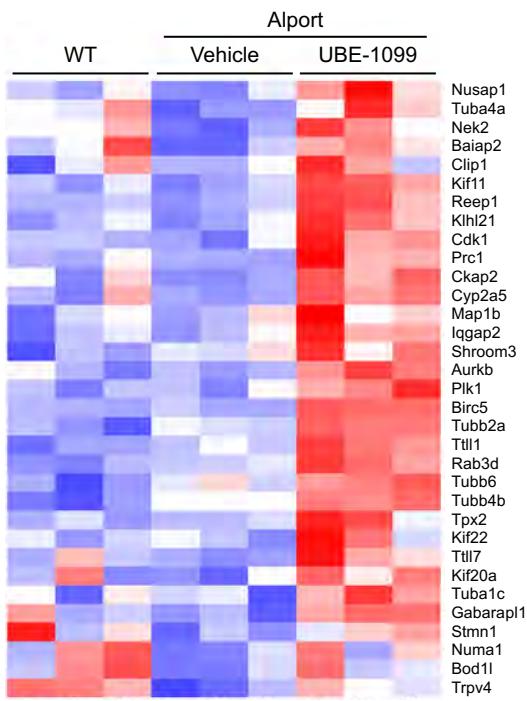
A Nrf2 target genes



B Cell cycle (GO:0007049)



C Microtubule (GO:0005874)



D Mitochondrial protein containing complex (GO:0005874)

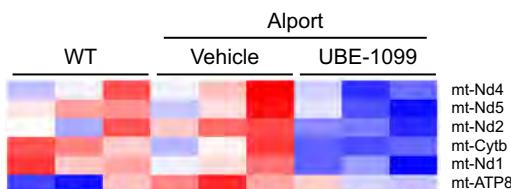
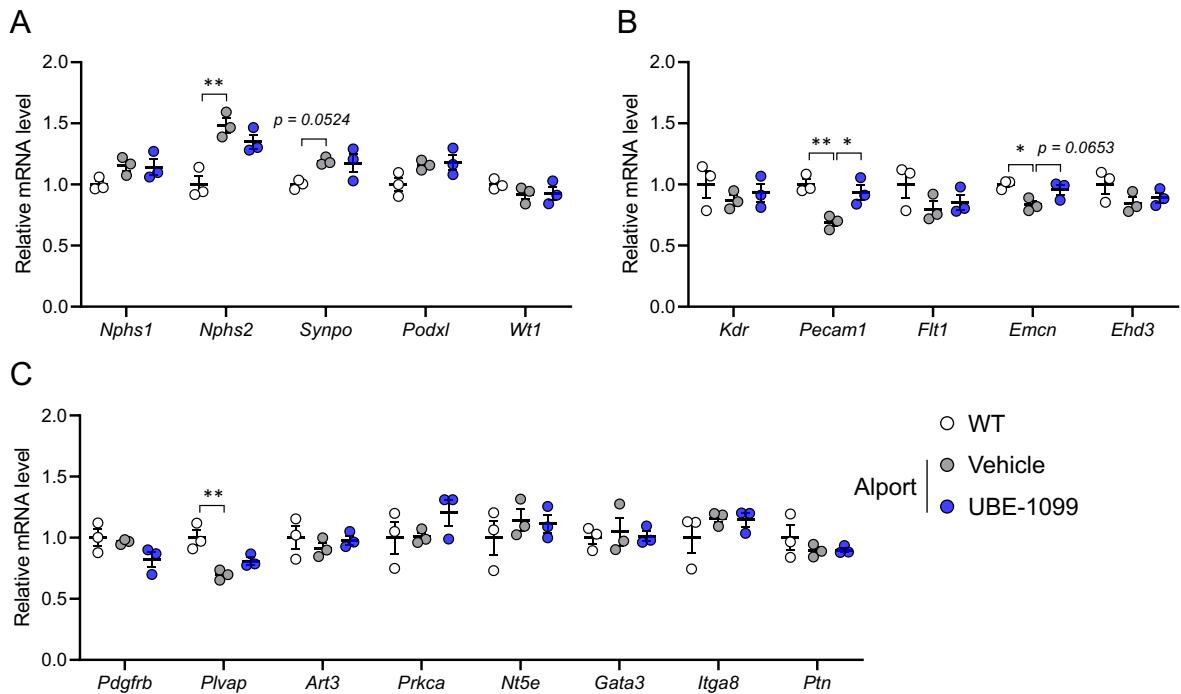


Figure S9. UBE1099 altered genes in each condition

Heat map shows the altered genes of Nrf2 target (A) and indicated GO terms (B-D) in UBE1099 group compared with Alport vehicle (fold change > 1.2 or < -1.2, p <0.05).

Figure S10**Figure S10. Expression level of cell specific markers in the glomerular cell**

Relative expression levels for cell specific markers for podocyte (A), endothelial cell (B) and mesangial cell (C) were measured by RNA-seq data (TPM). Data are presented as mean \pm SE (n = 3 per group). P values were assessed by Dunnett's test. (*p<0.05, **p<0.01).

Figure S11

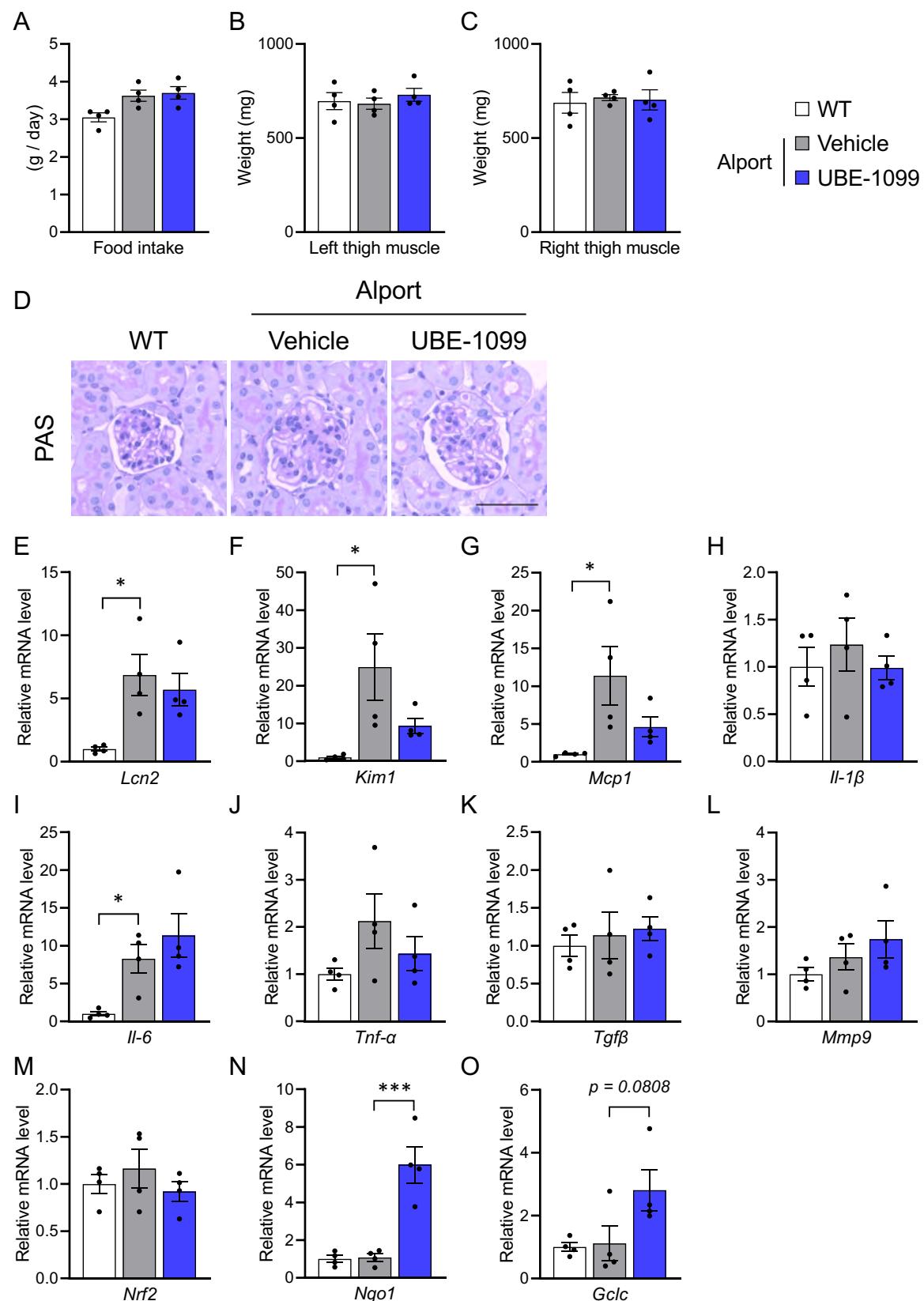
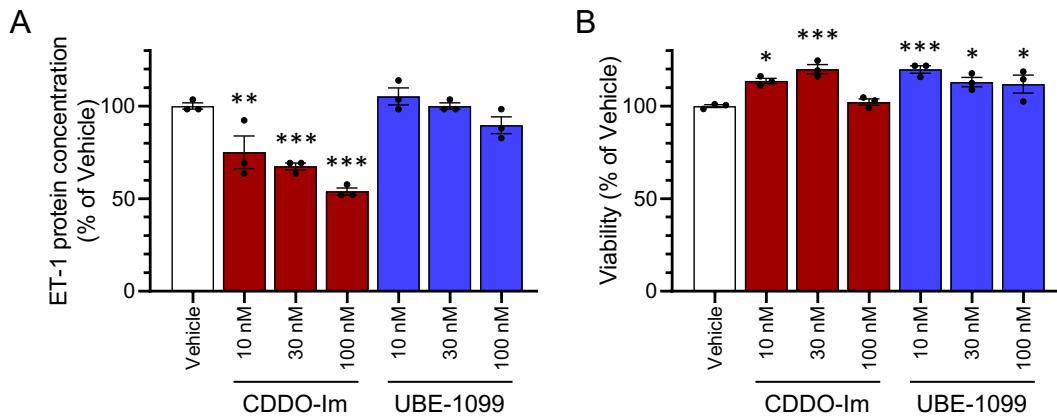
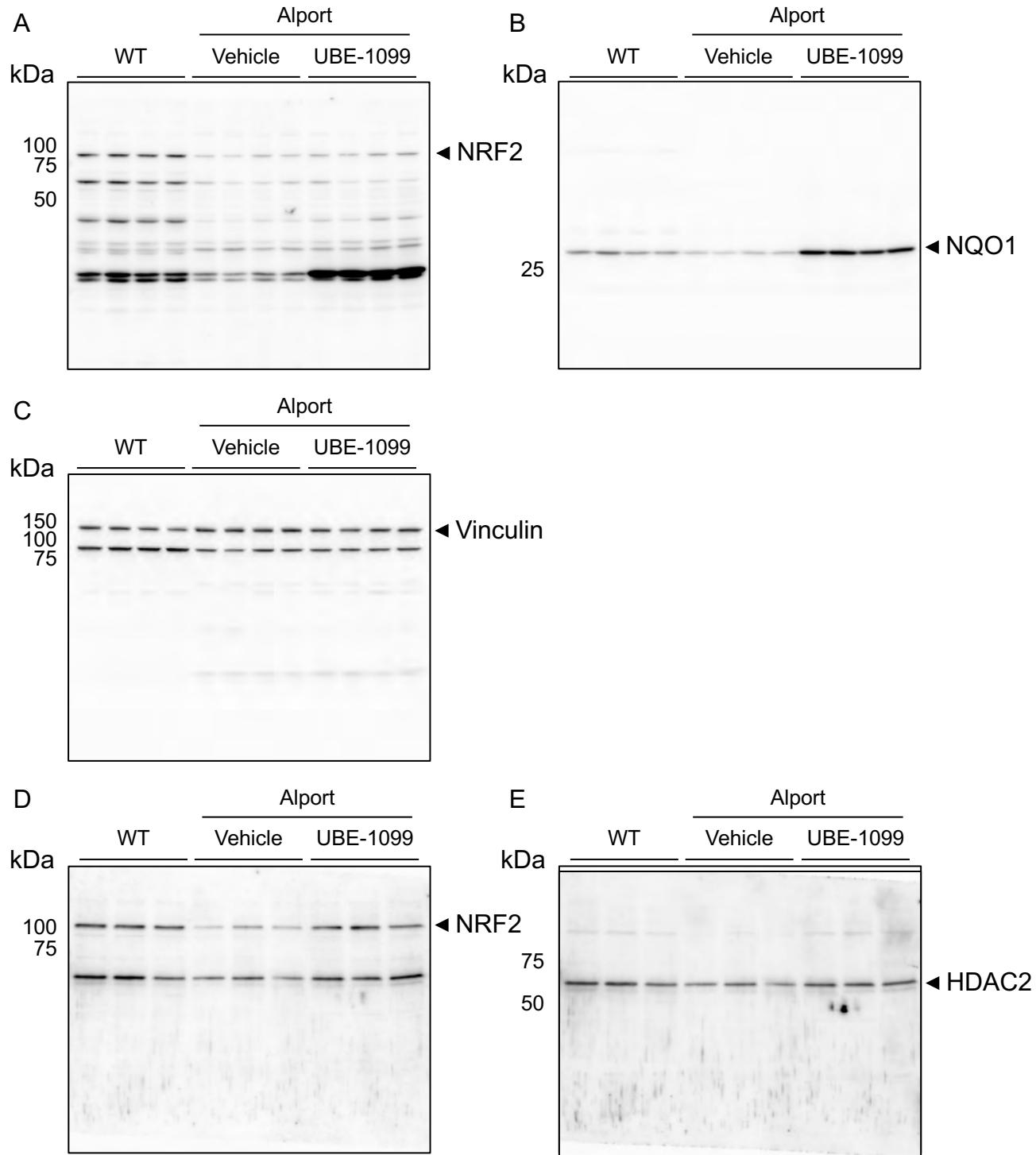


Figure S11. UBE-1099 did not affect the food intake and muscle weight in Alport mice and did not worsen the early renal pathology

(A-C) Food intake and thigh muscle weight were measured at 10 weeks old. (D) Renal sections of 10-week-old wild-type and Alport mice were analyzed by PAS staining. Representative images are shown. Scale bars = 50 μ m. (E-O) Total RNA was isolated from renal tissues of 10-week-old wild-type and Alport mice. The level of the indicated mRNA was measured and normalized to the level of Gapdh mRNA (internal control). Data are presented as mean \pm SE (n = 4 per group). P values were assessed by Dunnett's test. (*p<0.05, ***p<0.001).

Figure S12**Figure S12. UBE-1099 did not affect the endothelin expression**

(A) Endothelin-1 (ET-1) protein concentration (as % vehicle control) secreted into media from rat proximal tubule cells treated with vehicle, CDDO-Im or UBE-1099 (10-100 nM). (B) Cell viability of vehicle-, CDDO-Im- or UBE-1099-treated cells was measured by Cell Titer-Glo Luminescent cell viability assay. Data are presented as mean \pm SE (n = 3 per group). P values were assessed by Dunnett's test. (*p<0.05, ***p<0.01, ****p<0.001 vs Vehicle).

Figure S13**Figure S13. Full length blots for Figure 5A, B**

The full-length blots for Figure 5A, B with the indicated antibodies. Vinculin and HDAC2 were used as loading control. Samples were derived from the same experiment, and gels/blots were processed in parallel.