Resource Summary Report

Generated by dkNET on May 18, 2025

Kidney Tissue Atlas

RRID:SCR_021626

Type: Tool

Proper Citation

Kidney Tissue Atlas (RRID:SCR_021626)

Resource Information

URL: https://atlas.kpmp.org/

Proper Citation: Kidney Tissue Atlas (RRID:SCR_021626)

Description: Atlas is set of interactive tools built to promote retrieval, exploration, discovery, and analysis of Kidney Precision Medicine Project data by greater research community. Datasets available in repository are combination of raw and processed data from KPMP participant biopsies and reference tissue samples.

Resource Type: data or information resource, atlas

Keywords: Interactive tools, retrieval, exploration, discovery, analysis, Kidney Precision

Medicine Project data

Funding:

Availability: Free, Freely available

Resource Name: Kidney Tissue Atlas

Resource ID: SCR_021626

Record Creation Time: 20220129T080356+0000

Record Last Update: 20250516T054240+0000

Ratings and Alerts

No rating or validation information has been found for Kidney Tissue Atlas.

No alerts have been found for Kidney Tissue Atlas.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 13 mentions in open access literature.

Listed below are recent publications. The full list is available at dkNET.

Bueckle A, et al. (2025) Construction, Deployment, and Usage of the Human Reference Atlas Knowledge Graph for Linked Open Data. bioRxiv: the preprint server for biology.

Kösters P, et al. (2024) Adhesion G Protein-Coupled Receptor Gpr126 (Adgrg6) Expression Profiling in Diseased Mouse, Rat, and Human Kidneys. Cells, 13(10).

Zhang M, et al. (2024) Co-regulation and synteny of GFM2 and NSA2 links ribosomal function in mitochondria and the cytosol with chronic kidney disease. Molecular medicine (Cambridge, Mass.), 30(1), 176.

Sun Y, et al. (2024) The identification of key molecules and pathways in the crosstalk of calcium oxalate-treated TCMK-1 cells and macrophage via exosomes. Scientific reports, 14(1), 20949.

Hu H, et al. (2024) RBBP6-Mediated ERR? Degradation Contributes to Mitochondrial Injury in Renal Tubular Cells in Diabetic Kidney Disease. Advanced science (Weinheim, Baden-Wurttemberg, Germany), 11(46), e2405153.

Riedhammer KM, et al. (2024) Implication of transcription factor FOXD2 dysfunction in syndromic congenital anomalies of the kidney and urinary tract (CAKUT). Kidney international, 105(4), 844.

Wang J, et al. (2023) Dimension-agnostic and granularity-based spatially variable gene identification using BSP. Nature communications, 14(1), 7367.

Barry A, et al. (2023) Multi-population genome-wide association study implicates immune and non-immune factors in pediatric steroid-sensitive nephrotic syndrome. Nature communications, 14(1), 2481.

Sinning J, et al. (2023) The aging kidney is characterized by tubuloinflammaging, a phenotype associated with MHC-II gene expression. Frontiers in immunology, 14, 1222339.

Wang J, et al. (2023) Dimension-agnostic and granularity-based spatially variable gene

identification. bioRxiv: the preprint server for biology.

Ouyang JF, et al. (2023) Systems level identification of a matrisome-associated macrophage polarisation state in multi-organ fibrosis. eLife, 12.

Hansen J, et al. (2022) A reference tissue atlas for the human kidney. Science advances, 8(23), eabn4965.

Hinze C, et al. (2022) Single-cell transcriptomics reveals common epithelial response patterns in human acute kidney injury. Genome medicine, 14(1), 103.