This repository contains the readcounts for the paper ‘’Variable expression of eighteen common housekeeping genes in human non-cancerous kidney biopsies’’.

An overview over which data corresponds to which dataset I the paper can be found in Table 1 in the main paper.

All readcounts presented are normalized as described in the paper. Note that the readcounts from the fabry sequencing are normalized but are not batchcorrected. If for any reason the batchcorrected data is required, please contact the author directly.

On the fabry data;

For the validation of housekeeping genes we only used the Norwegian cohort and compared Fabry patients with 10 years of follow-up with normal controls. All comparisons were kept to the same compartment.

PT: Proximal tubule

DT: Distal Tubule

A: artery

G: glomeruli

0: biopsy from baseline

5: biopsy after 5 years of follow-up

10: biopsy from 10 years follow-up

NO: Norwegian cohort

CHRO: swiss cohort

N\_CTRL : normal control