

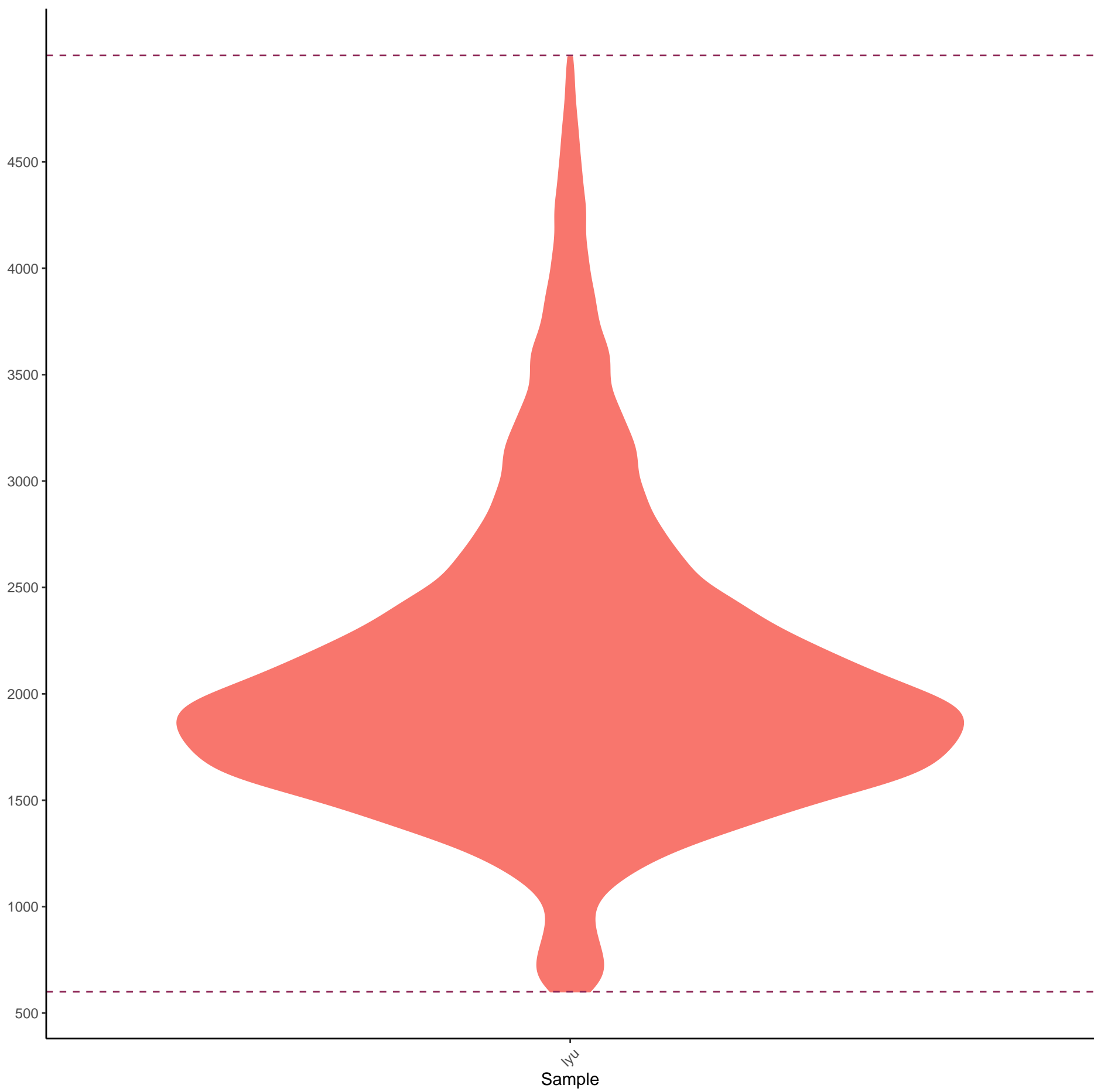
log10(number of transcripts)

4.4
4.2
4.0
3.8
3.6
3.4
3.2
3.0

lyu
Sample



number of detected genes



percentage of mitochondrial transcripts

