

RNA-seq data analysis

Fachprojekt Reproduzierbare Datenanalyse mit
Snakemake am Beispiel der Bioinformatik

Dr. Johannes Köster

Gruppe 1:

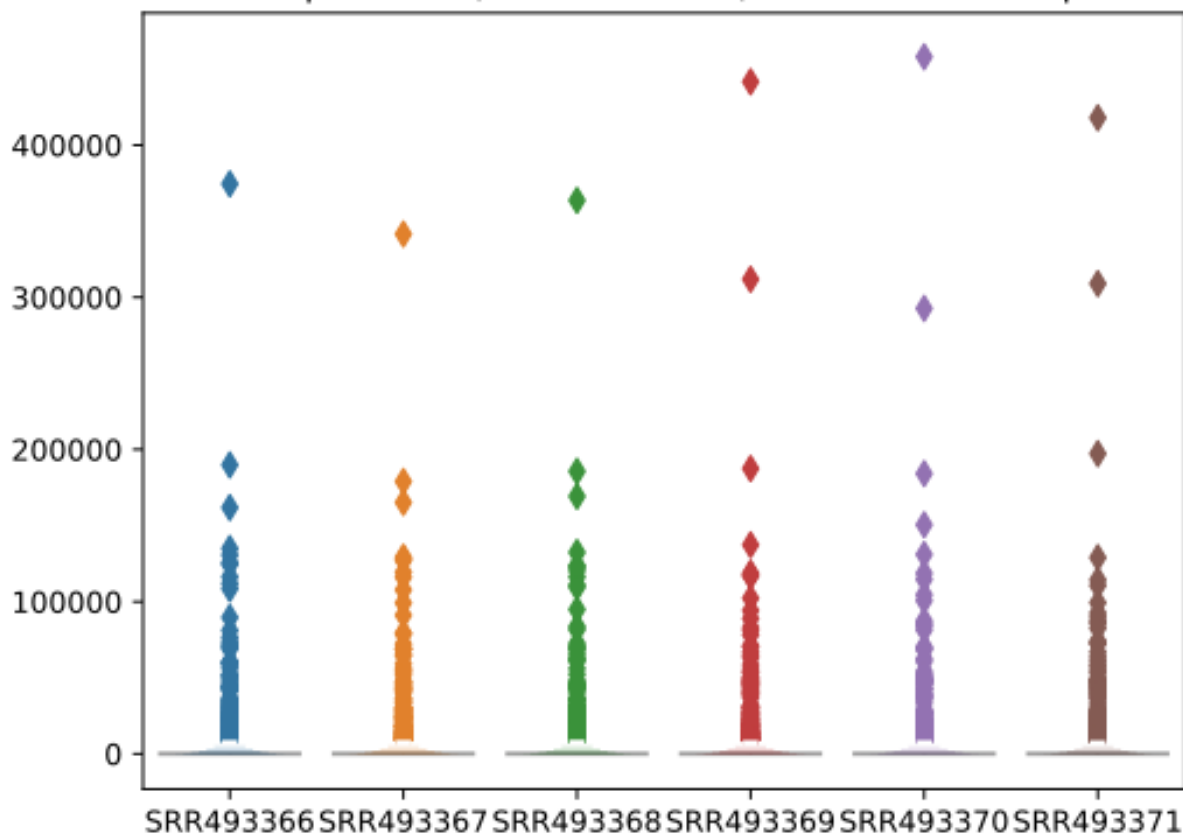
Jana Jansen

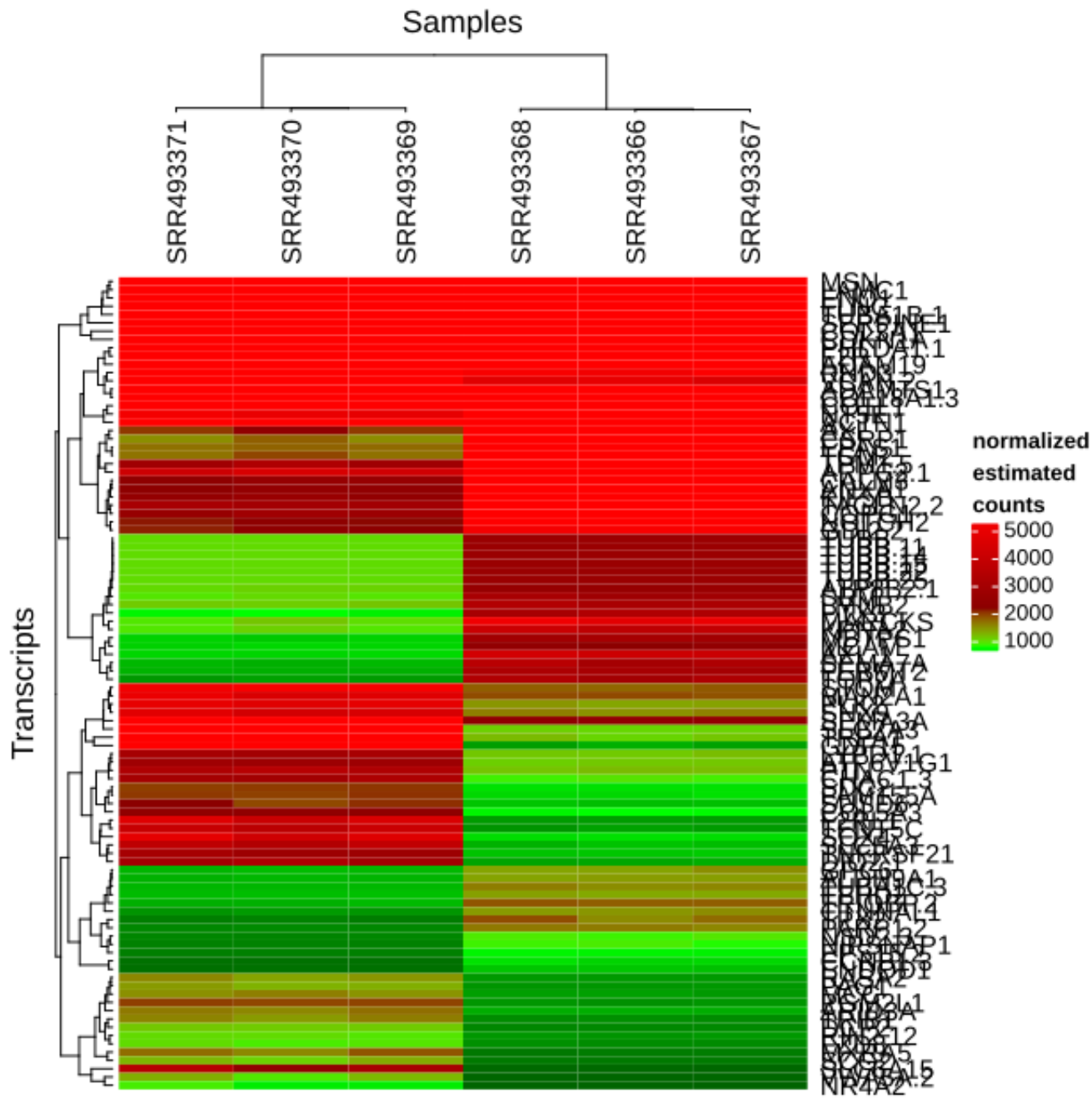
Ludmila Janzen

Sophie Sattler

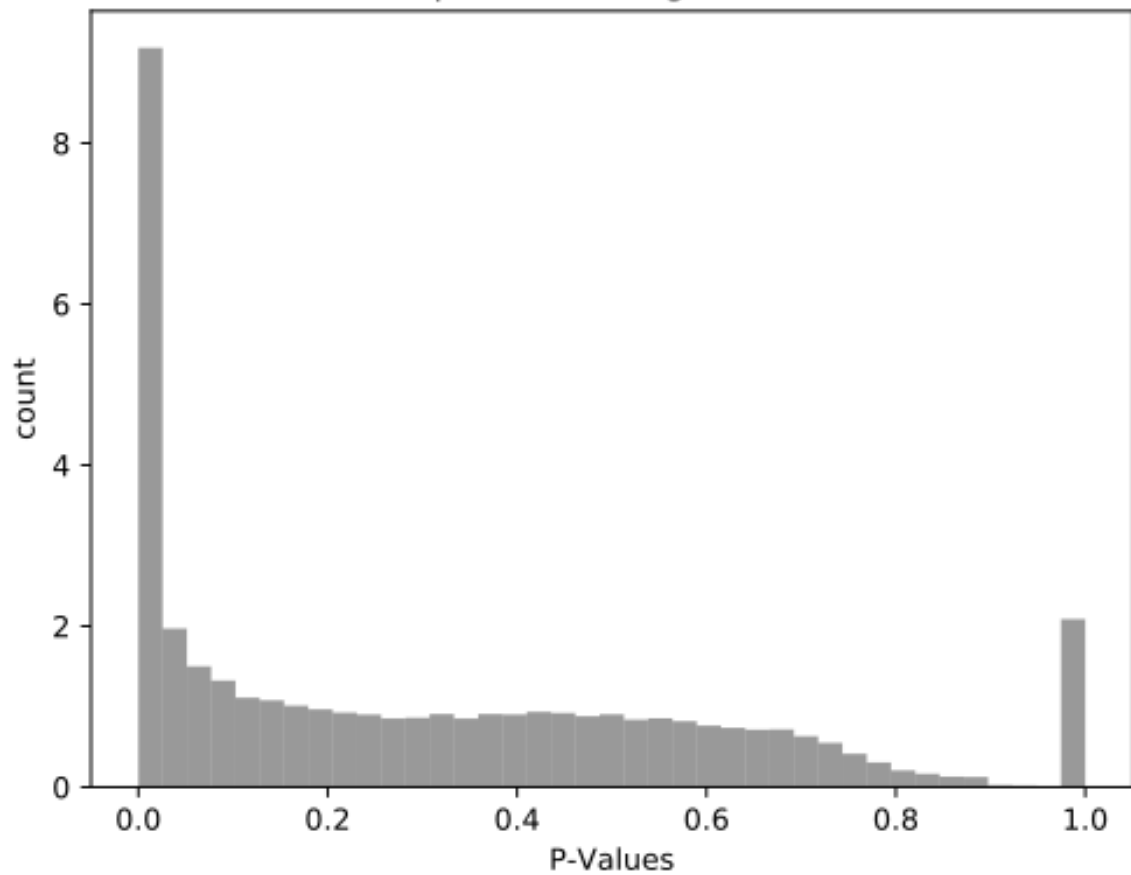
Antonie Vietor

Boxenplots der (normalisierten) Counts aller Samples

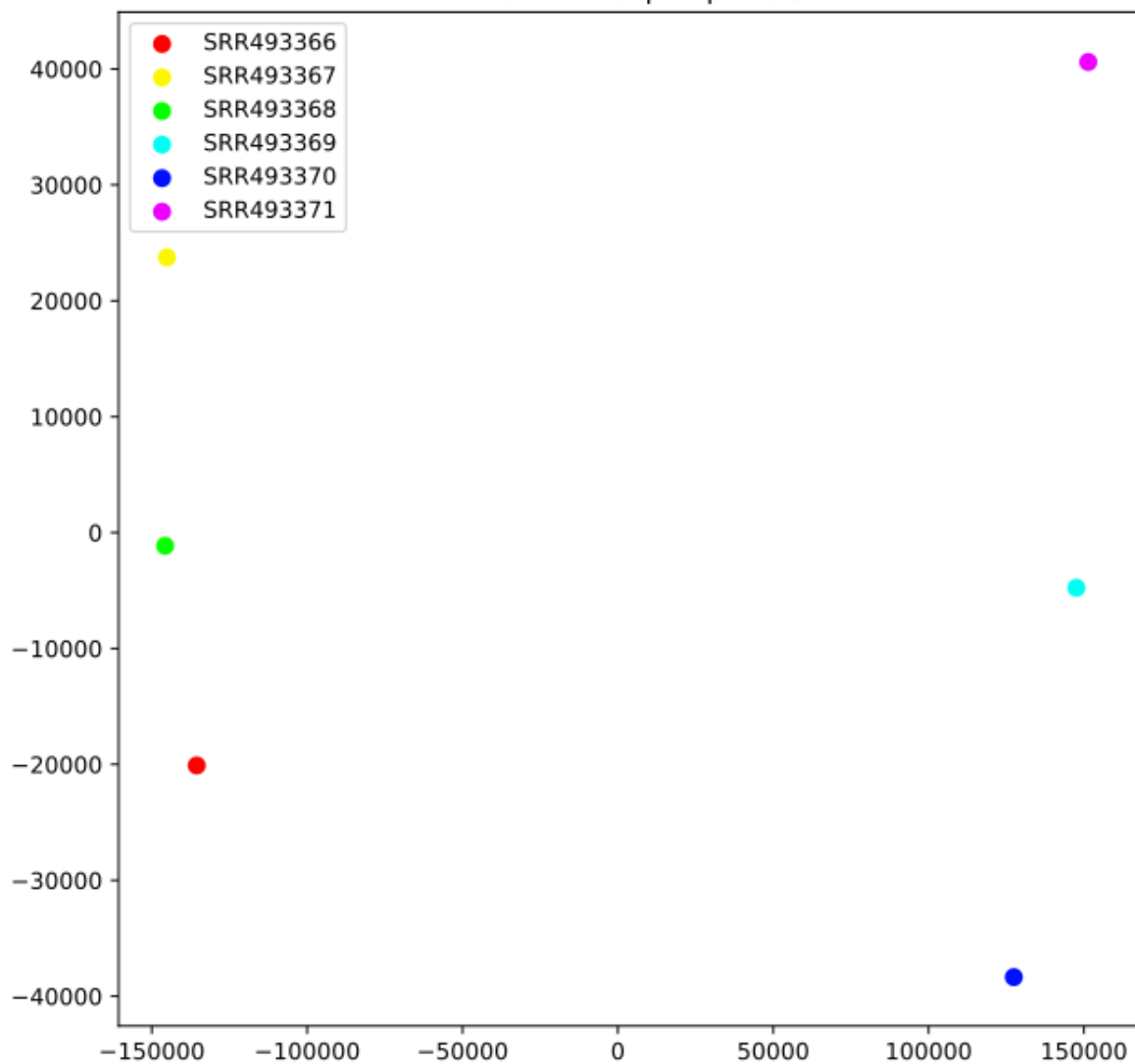




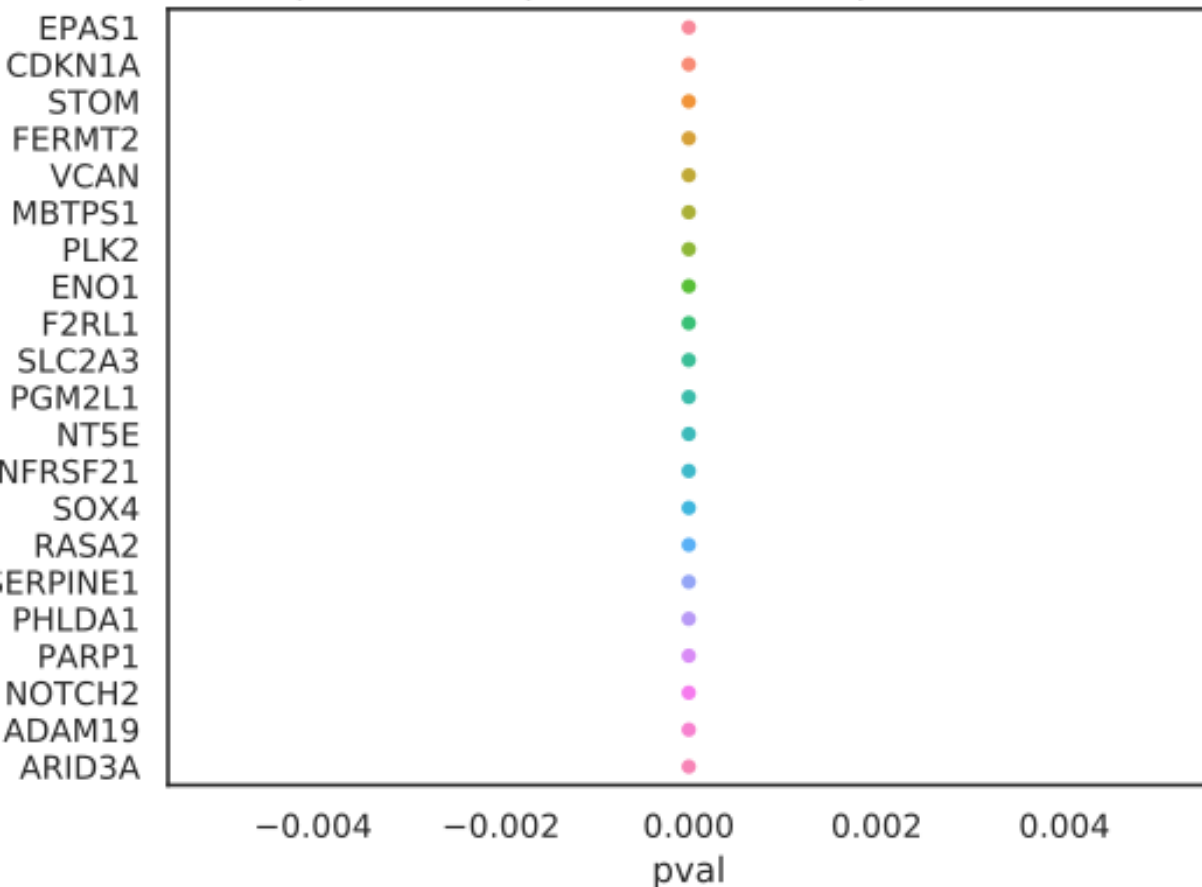
p-value Histogramm



PCA of Transcriptexpression



Stripchart der top20 differentiell exprimierten Gene



Volcano-Plot

