

# RNA-seq data analysis

Fachprojekt Reproduzierbare Datenanalyse mit  
Snakemake am Beispiel der Bioinformatik

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Gruppe 1:

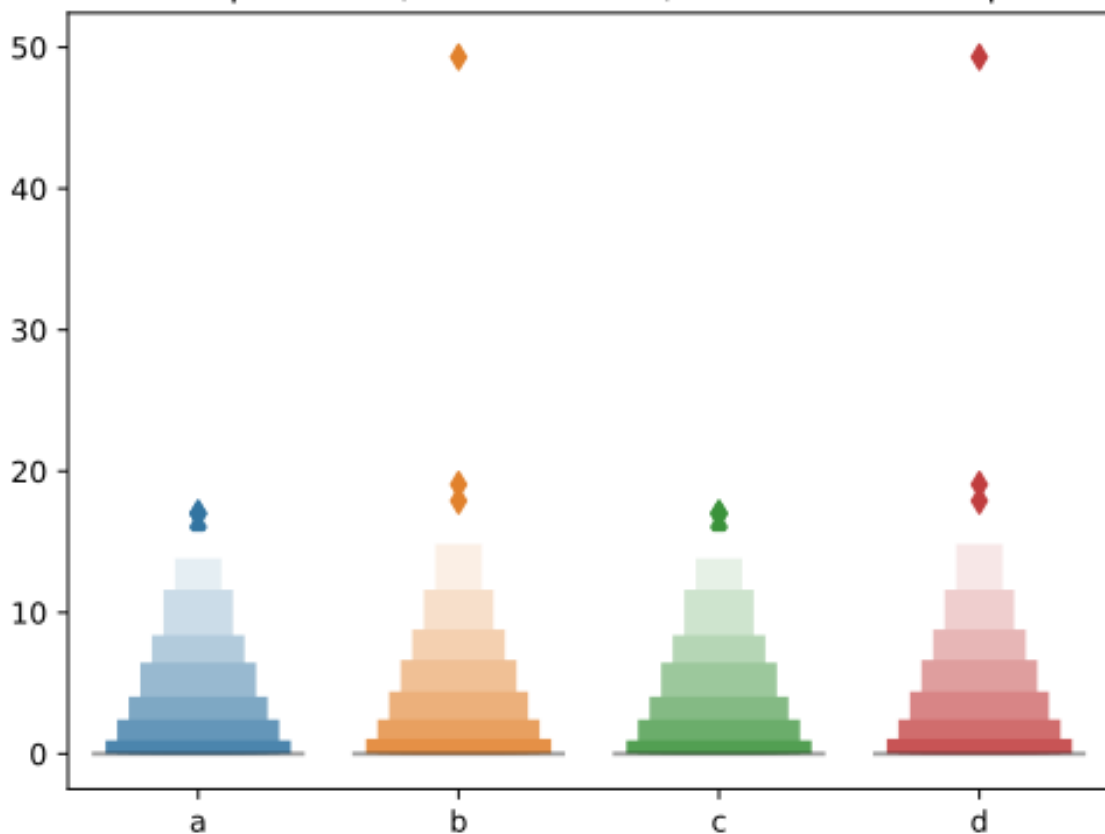
Jana Jansen

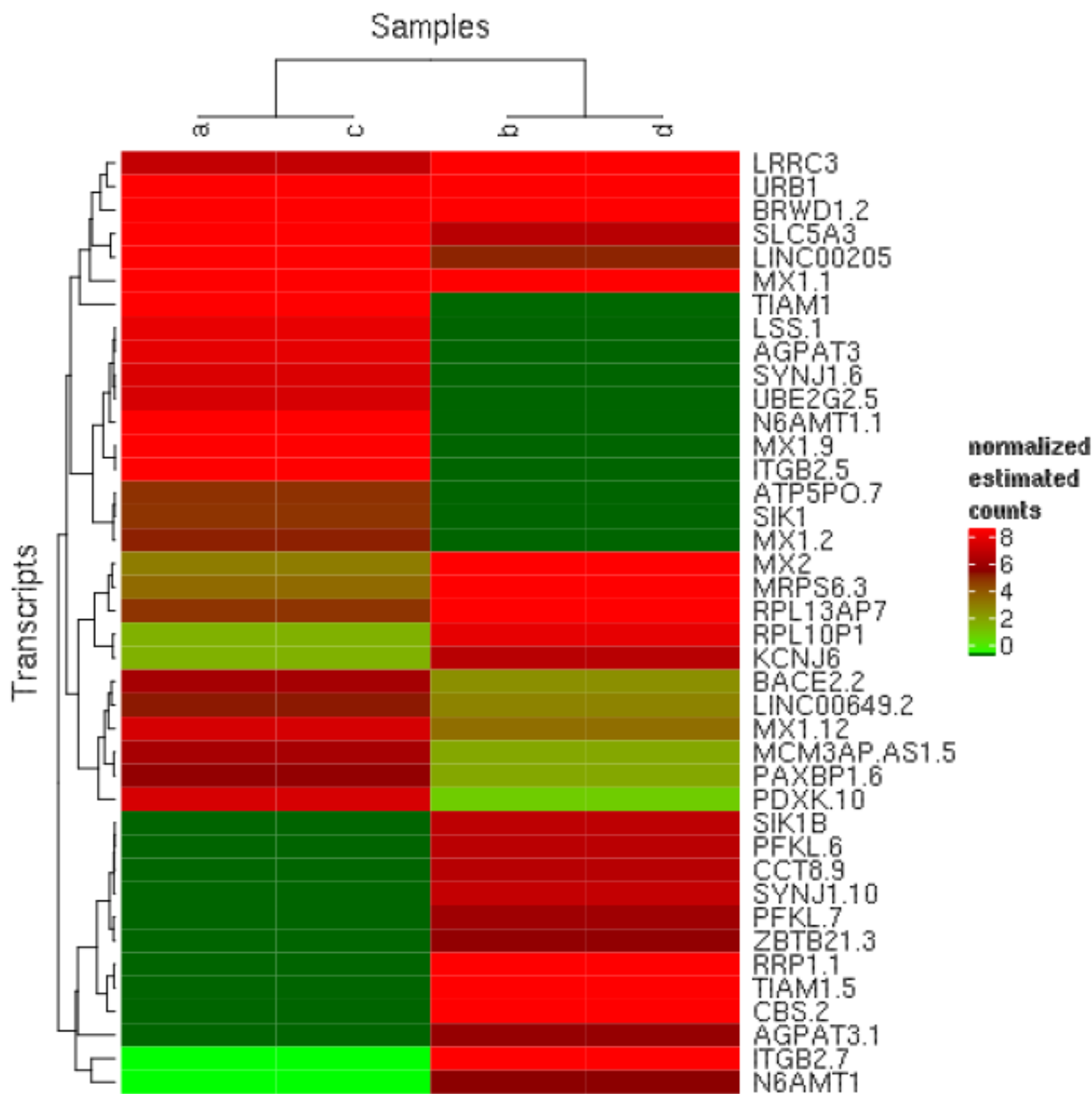
Ludmila Janzen

Sophie Sattler

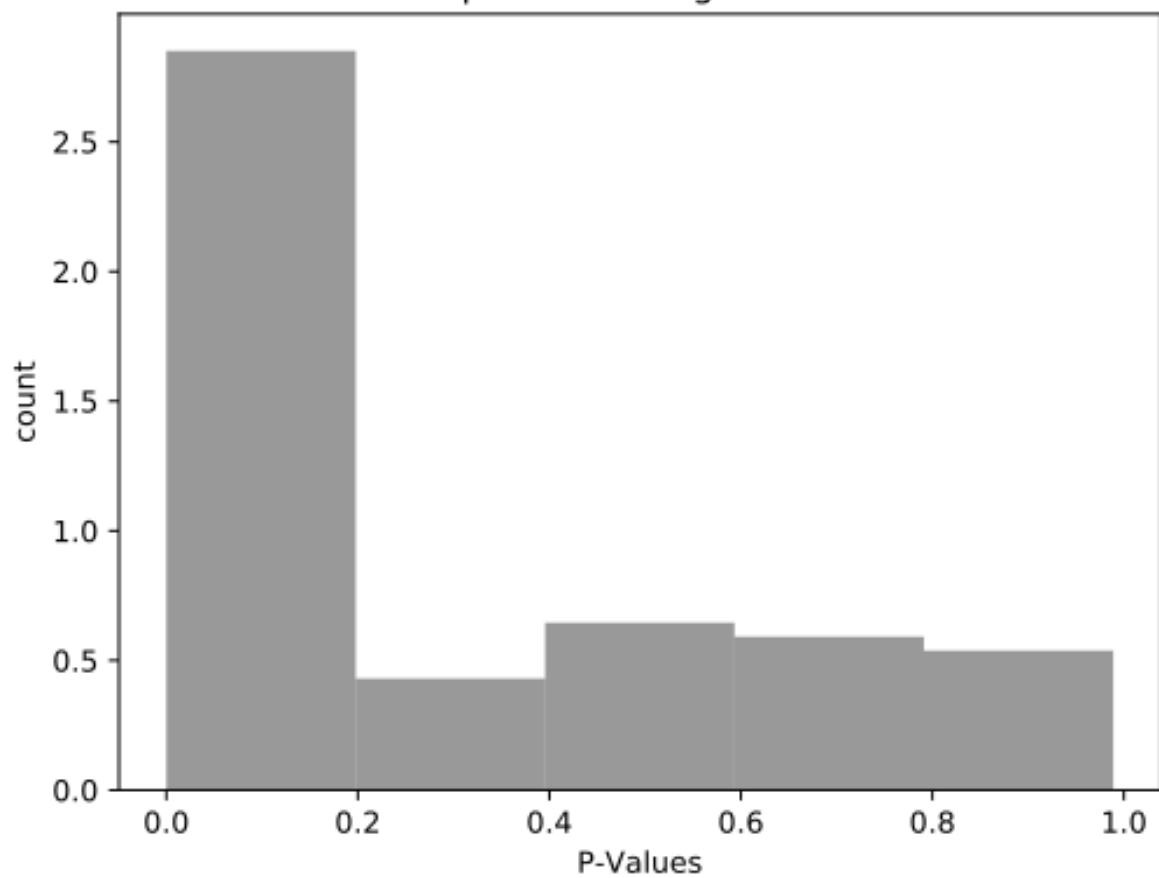
Antonie Vietor

Boxenplots der (normalisierten) Counts aller Samples

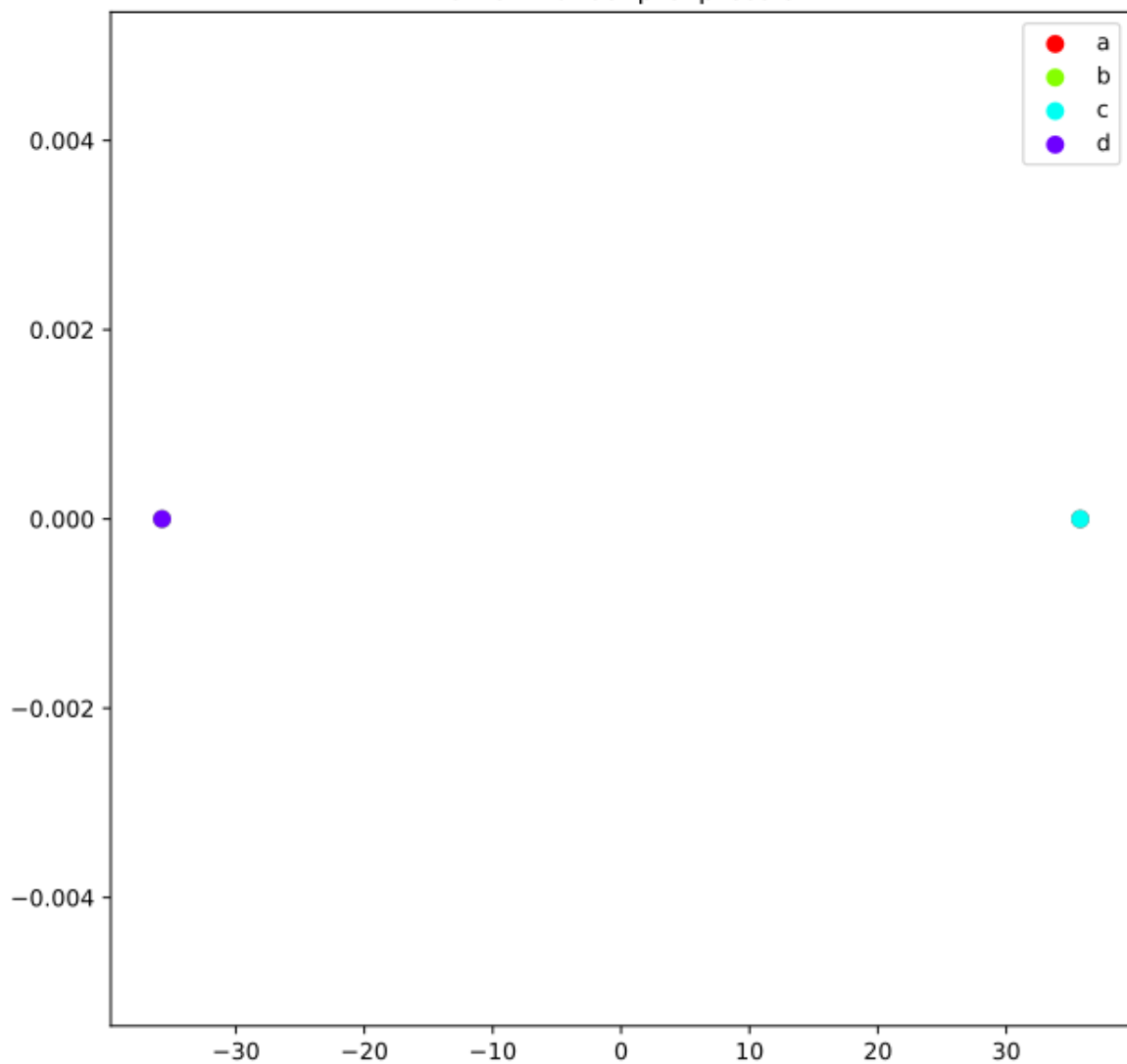




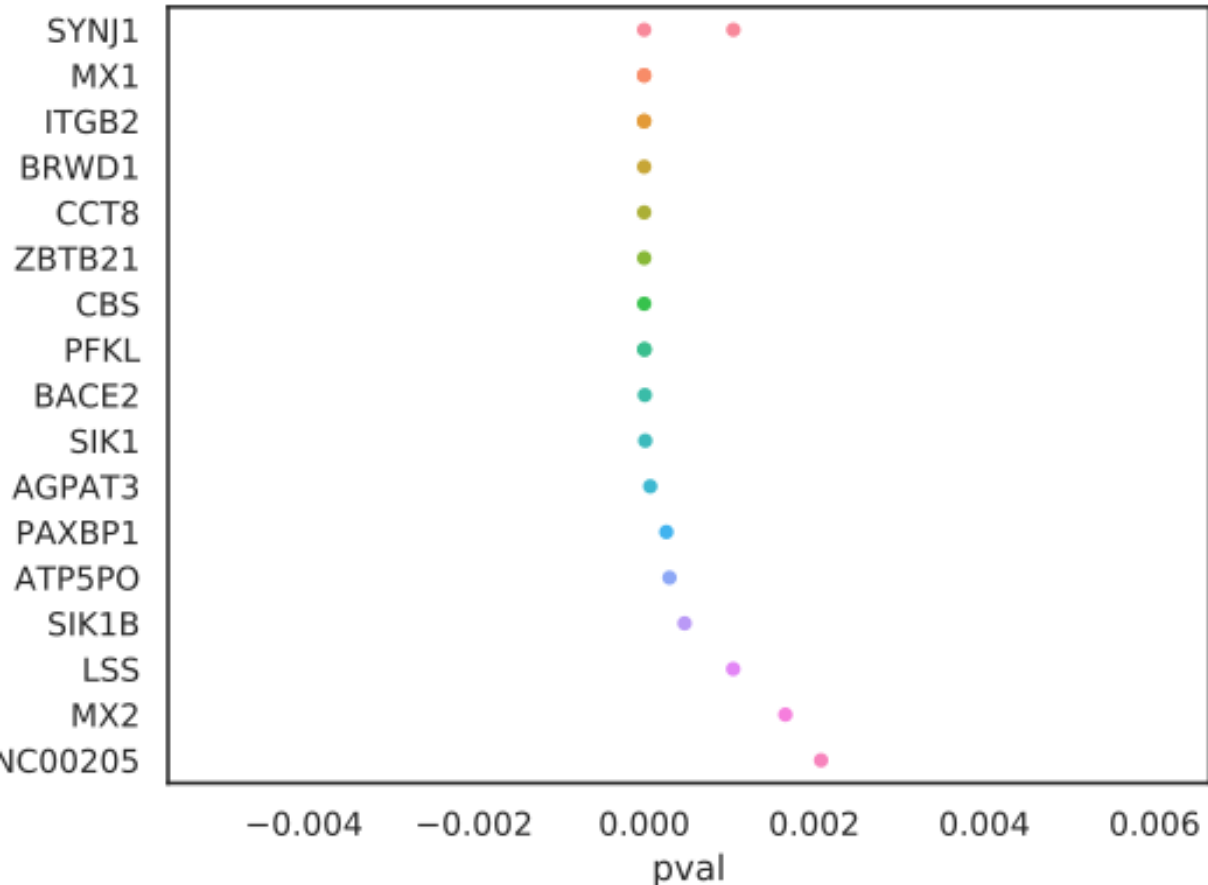
p-value Histogramm



PCA of Transcriptexpression



Stripchart der top20 differentiell exprimierten Gene



Volcano-Plot

