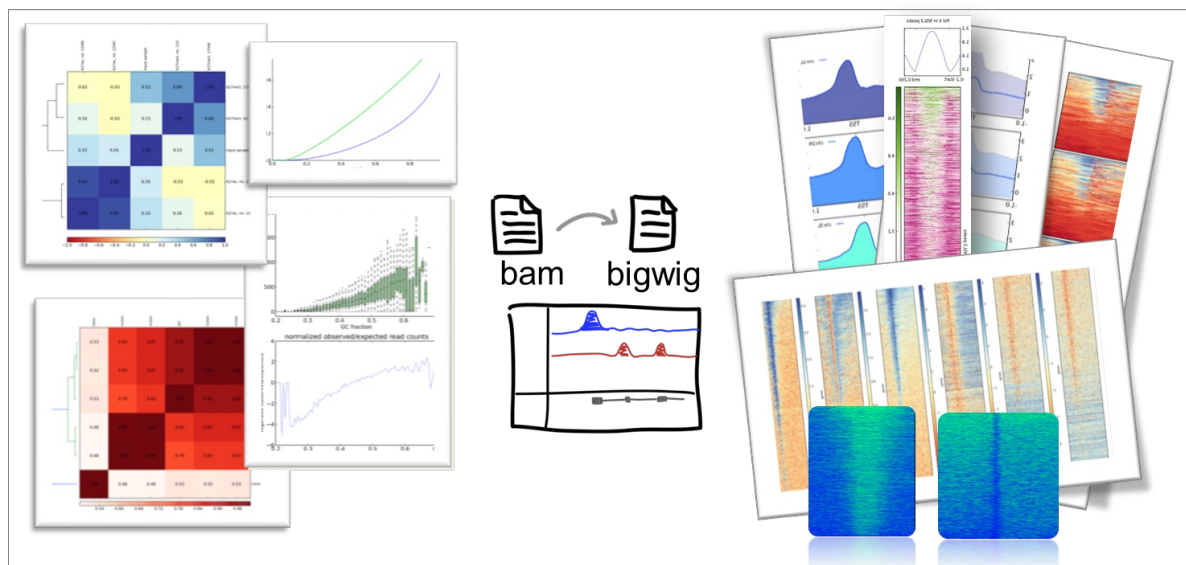


deepTools: a flexible platform for exploring deep-sequencing data

MANUAL

1. Why we built deepTools
2. How we use deepTools
3. What deepTools can do
4. Tool details
 - Quality controls of aligned reads
 - *bamCorrelate*
 - *computeGCbias*
 - *bamFingerprint*
 - Normalization and bigWig generation
 - *correctGCbias*
 - *bamCoverage*
 - *bamCompare*
 - Visualization: heatmaps and summary plots
5. Glossary: Abbreviations and file formats



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Web server (incl. sample data): deepTools.ie-freiburg.mpg.de

Code: github.com/fidelfram/deepTools