

# Links and references

## Literature

Benjamini and Speed, Nucleic Acids Research (2012): <http://nar.oxfordjournals.org/content/40/10/e72>

Diaz et al., Stat. Appl. Gen. Mol. Biol. (2012): <http://www.degruyter.com/view/j/sagmb.2012.11.issue-3/1544-6115.1750/1544-6115.1750.xml>

For more NGS-related literature, see our collection at the deepTools web server: <http://deeptools.ie-freiburg.mpg.de/u/fduendar/p/useful-bioinfo-literature>

## Additional bioinformatic tools

bowtie2: <http://bowtie-bio.sourceforge.net/bowtie2/manual.shtml>

cluster3: <http://bonsai.hgc.jp/~mdehoon/software/cluster/>

IGV (Integrative Genome Browser): <http://www.broadinstitute.org/igv/>

k-means clustering: [http://en.wikipedia.org/wiki/K-means\\_clustering](http://en.wikipedia.org/wiki/K-means_clustering)

R: <http://www.r-project.org/>

## File format information

SAM file specification: <http://samtools.sourceforge.net/SAMv1.pdf>

File formats explained at UCSC: <http://genome.ucsc.edu/FAQ/FAQformat.html#format1>