

<http://evolve.zoo.ox.ac.uk/Evolve/Software.html>

<http://virus.zoo.ox.ac.uk/rnavirusdb/>

<http://www.kuleuven.be/aidslab/phylogenybook/Software.html>

MasterPlex ReaderFit for ELISA values calculation

http://www.miraibio.com/masterplex-readerfit/curve-fitting-for-plate-readers.html?utm_source=Boomerang&utm_medium=Email&utm_term=MasterPlex+ReaderFit&utm_content=muhammad.munir@bvf.slu.se&utm_campaign=Curve+fitting+software+%24499+promotion

Image J for quantification of PCR gel or Westen blotting gel

<http://rsbweb.nih.gov/ij/docs/examples/index.html>

How:

http://www.ehow.com/how_5950558_quantify-electrophoresis-gels-images.html

or: <http://www.mun.ca/biology/dinnes/B2250/Lab2/ImageJ.htm>

Serial Clonar for Vector manipulation

http://serialbasics.free.fr/Serial_Cloner-Download.html

GeNorm: to normalize the real-time PCR

<http://medgen.ugent.be/~jvdesomp/genorm/index.php>

<http://genomebiology.com/2002/3/7/research/0034/>

<http://en.bio-soft.net/draw.html>

<http://en.bio-soft.net/dna.html>

<http://en.bio-soft.net/protein.html>

<http://en.bio-soft.net/primer.html>

<http://en.bio-soft.net/plasmid.html>

<http://en.bio-soft.net/tree.html>

<http://en.bio-soft.net/format.html>

<http://en.bio-soft.net/zhol.html>

<http://www.fidelitysystems.com/difficult.htm>

fasta to nexus for Mr. Bayes

<http://www.ub.edu/dnasp/>

Predict protein domain

http://smart.embl-heidelberg.de/help/smart_about.shtml

To calculate synonymous to non-synonymous substitution (Marti Cortey)

<http://www.hiv.lanl.gov/content/sequence/SNAP/SNAP.html>

Gel band intensity quantification:

http://imagejdocu.tudor.lu/doku.php?id=video:analysis:gel_quantification_analysis

Prediction of transcription binding sites:

http://alggen.lsi.upc.es/cgi-bin/promo_v3/promo/promoinit.cgi?dirDB=TF_8.3

Protein and protein interaction

<http://nar.oxfordjournals.org/content/32/21/6312.long>