

Übersicht zu wichtigen Datenbanken und Programmen und deren allgemeinen Verwendung

Alignment/Stammbäume

CLUSTALW/Clustal Omega <https://www.ebi.ac.uk/Tools/msa/clustalo/>
MUSCLE <https://www.ebi.ac.uk/Tools/msa/muscle/>
PHYLP <https://evolution.genetics.washington.edu/phylip.html>
Datensätze zu biologische Größen/Biotechnologie/synthetische Biologie
BioNumbers <https://bionumbers.hms.harvard.edu>
BioBricks <https://biobricks.org/>
GoSynthetic <https://gosyn.bioapps.biozentrum.uni-wuerzburg.de/index.php>
Dotplot
Dotter <https://sonnhammer.sbc.su.se/Dotter.html>
GEPARD <https://mips.gsf.de/services/analysis/gepard>
JDotter <https://athena.bioc.uvic.ca/virology-ca-tools/jdotter/>

Funktionsdatenbanken

Functional Glycomics <https://www.functionalglycomics.org/>; <https://ncfg.hms.harvard.edu/>
Gene Ontology <https://www.geneontology.org>

Gehirn-Baupläne

Blue Brain Projekt (EU) <https://bluebrain.epfl.ch/>
Brain Activity Atlas <https://www.brainactivityatlas.org/>
Brain Activity Projekt (USA) <https://www.braininitiative.nih.gov/>
Connectome-Projekt <https://www.openconnectomeproject.org>
Mouse Brain Connectivity Atlas <https://mouse.brain-map.org/static/atlas>
Neuroaktivitätsdetektion <https://www.ncbi.nlm.nih.gov/pubmed/23537512>
Temporal lobe <https://www.temporal-lobe.com/background/connectome>
Virtual Insect Brain Lab <https://www.neurofly.de/>
WormWiring <https://wormwiring.org/>
Wurmatlas <https://www.wormatlas.org/>

Genomannotation/Genannotation/Sequenzanalyse/Online-Bibliotheken/Experimentelle Datensätze

BLAST <https://blast.ncbi.nlm.nih.gov/Blast.cgi>
GenScan <https://genes.mit.edu/GENSCAN.html>
RepeatMasker <https://www.repeatmasker.org/>
ENCODE <https://www.encodeproject.org>
Ensembl https://www.ensembl.org/Homo_sapiens/Info/Index
GATK Workshop <https://software.broadinstitute.org/gatk/guide/article?id=7869#1.3>
Genomic Science Program <https://genomics.energy.gov>
Human Genome Project https://web.ornl.gov/sci/techresources/Human_Genome/index.shtml
UCSC <https://genome.ucsc.edu/>
DDBJ (DNA Data Bank of Japan) <https://www.ddbj.nig.ac.jp/>
EBI <https://www.ebi.ac.uk/services>
iGEM Parts https://igem.org/Main_Page
MEDLINE/NCBI/PubMed <https://www.ncbi.nlm.nih.gov/pubmed/>
NIH <https://www.genome.gov>
OMIM <https://www.omim.org/>
Swiss Bioinformatics Institute <https://www.sib.swiss/>
WebDirectory <https://www.biologydir.com/over-population/p1.html>
Computational Population Biology <https://compbio.cs.uic.edu/>
GENEVESTIGATOR <https://genevestigator.com/gv/>
GEO <https://www.ncbi.nlm.nih.gov/geo/>

Grafikprogramme, Modellierung und Netzwerkanalyse

CellDesigner <https://www.celldesigner.org/>
CellNetAnalyzer <https://www2.mpi-magdeburg.mpg.de/projects/cna/cna.html>
Cytoscape <https://www.cytoscape.org/>
COBRA <https://opencobra.github.io/>
COPASI <https://copasi.org/>
Flux balance analysis <https://systemsbiology.ucsd.edu/Downloads/FluxBalanceAnalysis>
Jimena https://www.bioinfo.biozentrum.uni-wuerzburg.de/computing/jimena_c/
MATLAB <https://de.mathworks.com/products/matlab.html>

Metatool <https://pinguin.biologie.uni-jena.de/bioinformatik/networks/metatool/>
Odefy <https://www.helmholtz-muenchen.de/icb/software/odefy/index.html>
PLAS <https://enzymology.fc.ul.pt/software/plas/>
PottersWheel <https://www.potterswheel.de/Pages/>
SQUAD <https://www.vital-it.ch/software/SQUAD>
YANA/YANAsquare <https://www.bioinfo.biozentrum.uni-wuerzburg.de/computing/yanasquare/>

Interaktionsdatenbank, Medikamenten-Interaktionsdatenbank

catRAPID https://s.tartagliolab.com/page/catrapid_group
HPRD <https://hprd.org/>
iHOP <https://www.ihop-net.org/UniPub/iHOP/>
KEGG <https://www.genome.jp/kegg/>
NPInter <https://www.bioinfo.org/NPInter/>
PlateletWeb <https://plateletweb.bioapps.biozentrum.uni-wuerzburg.de/plateletweb.php>
Roche Pathways <https://biochemical-pathways.com/#/map/1>
STRING <https://string-db.org>
DrumPID <https://drumpid.bioapps.biozentrum.uni-wuerzburg.de/compounds/index.php>
STITCH <https://stitch.embl.de/>
EcoCyc <https://ecocyc.org/>

Lokalisations-/Motiv-Vorhersage

LocP <https://ekhidna2.biocenter.helsinki.fi/LOCP/>
LocSigDB <https://genome.unmc.edu/LocSigDB/>
nucloc <https://www.nucloc.org/>
NucPred <https://www.sbc.su.se/~maccallr/nucpred/>
SignalP <https://www.cbs.dtu.dk/services/SignalP/>
TMHMM <https://www.cbs.dtu.dk/services/TMHMM/>
Functional Glycomics <https://www.functionalglycomics.org/>
ELM <https://elm.eu.org/>

Programmiersprachen

Biojava <https://biojava.org/>
BioPerl <https://bioperl.org/>
C++ <https://www.cplusplus.com/>
Java <https://www.oracle.com/technetwork/java/index.html>
Perl <https://www.perl.org/>
Python <https://www.python.org/>
R <https://cran.r-project.org/>
Bioconductor <https://www.bioconductor.org/>

Promotoranalyse

ALGGEN PROMO
https://algggen.lsi.upc.es/cgi-bin/promo_v3/promo/promoinit.cgi?dirDB=TF_8.3
Genomatix <https://www.genomatix.de/>
JASPAR https://jaspar.genereg.net/cgi-bin/jaspar_db.pl
MotifMap <https://motifmap.igb.uci.edu/>
TESS <https://www.cbil.upenn.edu/tess/>
TRANSFAC <https://www.gene-regulation.com/pub/databases.html>

Proteinanalyse

AnDom
https://andom.bioapps.biozentrum.uni-wuerzburg.de/index_new.html
CATH <https://www.cathdb.info/>
Conserved Domains <https://www.ncbi.nlm.nih.gov/Structure/cdd/wrpsb.cgi>
ExpASy <https://www.expasy.org>
InterPro <https://www.ebi.ac.uk/interpro/>
MODELLER <https://salilab.org/modeller/tutorial/>
PDB <https://www.rcsb.org/pdb/home/home.do>
Pfam <https://pfam.xfam.org/>
ProDom <https://prodom.prabi.fr/prodom/current/html/home.php>
PRODORIC <https://prodoric.tu-bs.de/>
PROSITE <https://prosite.expasy.org>
PyMOL <https://www.pymol.org/>
QUARK <https://zhanglab.ccmb.med.umich.edu/QUARK/>
Ramachandran-Plot <https://mordred.bioc.cam.ac.uk/~rapper/rampage.php>
RasMol <https://www.openrasmol.org/>

SCOP (old) <https://scop.mrc-lmb.cam.ac.uk/scop/>
SCOP updated <https://scop.berkeley.edu/>
SMART <https://smart.embl-heidelberg.de/>
SWISS-MODEL <https://swissmodel.expasy.org>
UniProt/Swiss-Prot <https://www.uniprot.org/>

RNA-Analyse

ITS2 <https://its2.bioapps.biozentrum.uni-wuerzburg.de/>
LNCipedia <https://www.lncipedia.org/>
mfold Webserver <https://unafold.rna.albany.edu/?q=mfold/RNA-Folding-Form>
microRNA.org/miRanda <https://www.microrna.org/microrna/home.do>
miRBase <https://www.mirbase.org/>
regRNA <https://regrna2.mbc.nctu.edu.tw/>
Rfam <https://rfam.xfam.org/>
Riboswitch-Finder <https://riboswitch.bioapps.biozentrum.uni-wuerzburg.de/>
RNAAnalyzer <https://rnaanalyzer.bioapps.biozentrum.uni-wuerzburg.de/>
RNAfold Webserver <https://rna.tbi.univie.ac.at/cgi-bin/RNAWebSuite/RNAfold.cgi>
TargetScan <https://www.targetscan.org>
tRNAscan <https://lowelab.ucsc.edu/tRNAscan-SE/>
Vienna Package <https://www.tbi.univie.ac.at/RNA/>