

## Multimedia Appendix 2. List of OS Projects Contacted for Survey

Project Name	Web Page
3D Slicer	<a href="http://www.slicer.org/">http://www.slicer.org/</a>
Apollo	<a href="http://www.fruitfly.org/annot/apollo/">http://www.fruitfly.org/annot/apollo/</a>
Biobuilder	<a href="http://www.biomedcentral.com/1471-2105/5/43">http://www.biomedcentral.com/1471-2105/5/43</a>
Bioconductor	<a href="http://www.bioconductor.org">http://www.bioconductor.org</a>
Biojava	<a href="http://www.biojava.org/">http://www.biojava.org/</a>
Biomail Scientific References Automation	<a href="http://biomail.sourceforge.net/biomail/index.html">http://biomail.sourceforge.net/biomail/index.html</a>
Bioperl	<a href="http://bioperl.org/">http://bioperl.org/</a>
Biophp	<a href="http://biophp.org">http://biophp.org</a>
Biopython	<a href="http://www.biopython.org/">http://www.biopython.org/</a>
Bioquery	<a href="http://www.bioquery.org/">http://www.bioquery.org/</a>
Biowarehouse	<a href="http://bioinformatics.ai.sri.com/biowarehouse/">http://bioinformatics.ai.sri.com/biowarehouse/</a>
Cd-Hit Sequence Clustering Software	<a href="http://bioinformatics.org/cd-hit/">http://bioinformatics.org/cd-hit/</a>
Chemistry Development Kit	<a href="http://almost.cubic.uni-koeln.de/cdk/">http://almost.cubic.uni-koeln.de/cdk/</a>
Coasim	<a href="http://www.daimi.au.dk/~mailund/CoaSim/">http://www.daimi.au.dk/~mailund/CoaSim/</a>
Cytoscape	<a href="http://www.cytoscape.org">http://www.cytoscape.org</a>
Das	<a href="http://biodas.org/">http://biodas.org/</a>
E-Cell System	<a href="http://sourceforge.net/projects/ecell/">http://sourceforge.net/projects/ecell/</a>
Emboss	<a href="http://emboss.sourceforge.net/">http://emboss.sourceforge.net/</a>
Ensemble	<a href="http://www.ensembl.org/info/software/versions.html">http://www.ensembl.org/info/software/versions.html</a>
Eviewbox Dicom Java Project	<a href="http://sourceforge.net/projects/eviewbox/">http://sourceforge.net/projects/eviewbox/</a>
Freemed Project	<a href="http://bioinformatics.org/project/?group_id=298">http://bioinformatics.org/project/?group_id=298</a>
Gchemical	<a href="http://www.bioinformatics.org/gchemical/">http://www.bioinformatics.org/gchemical/</a>
Gnumed	<a href="http://www.gnumed.org">http://www.gnumed.org</a>
Medical Dataserver	<a href="http://www.mii.ucla.edu/dataserver">http://www.mii.ucla.edu/dataserver</a>
Medical Image Analysis	<a href="http://sourceforge.net/projects/mia">http://sourceforge.net/projects/mia</a>
Moby	<a href="http://biomoby.open-bio.org/">http://biomoby.open-bio.org/</a>
Olduvai	<a href="http://sourceforge.net/projects/olduvai/">http://sourceforge.net/projects/olduvai/</a>
Openclinica	<a href="http://www.openclinica.org">http://www.openclinica.org</a>
Openemed	<a href="http://openemed.org/">http://openemed.org/</a>
Openemr	<a href="http://www.oemr.org/">http://www.oemr.org/</a>
Oscarmcmaster	<a href="http://sourceforge.net/projects/oscarcmcmaster/">http://sourceforge.net/projects/oscarcmcmaster/</a>
Probemaker	<a href="http://probemaker.sourceforge.net/">http://probemaker.sourceforge.net/</a>
Res Medicinae	<a href="http://resmedicinae.sourceforge.net/">http://resmedicinae.sourceforge.net/</a>
Rtools	<a href="http://www.bioconductor.org">http://www.bioconductor.org</a>
S2s	<a href="http://bioinformatics.org/S2S/">http://bioinformatics.org/S2S/</a>
Sashimi	<a href="http://sourceforge.net/projects/sashimi/">http://sourceforge.net/projects/sashimi/</a>
Sbeams	<a href="http://www.sbeams.org">http://www.sbeams.org</a>
Alkahest	<a href="http://www.alkahest.org">http://www.alkahest.org</a>

Array-A-Lizer	<a href="http://bioinformatics.org/cgi-bin/cvsweb.cgi/arrayalizer/">http://bioinformatics.org/cgi-bin/cvsweb.cgi/arrayalizer/</a>
Ball	<a href="http://bioinformatics.org/project/?group_id=506">http://bioinformatics.org/project/?group_id=506</a>
Biococoa Biolibrary For Obj-C	<a href="http://bioinformatics.org/project/?group_id=318">http://bioinformatics.org/project/?group_id=318</a>
Biomedical Informatics Research Network	<a href="http://www.nbirn.net">http://www.nbirn.net</a>
Biosignal Tools	<a href="http://sourceforge.net/projects/biosig">http://sourceforge.net/projects/biosig</a>
Brain Imaging Simulation System	<a href="http://sourceforge.net/projects/brainimage">http://sourceforge.net/projects/brainimage</a>
Cehr Net	<a href="http://sourceforge.net/projects/cehrnet/">http://sourceforge.net/projects/cehrnet/</a>
Clearhealth Practice Management System	<a href="http://sourceforge.net/projects/clearhealth/">http://sourceforge.net/projects/clearhealth/</a>
Common Open Source Medical Objects	<a href="http://sourceforge.net/projects/cosmos">http://sourceforge.net/projects/cosmos</a>
Compbio Perl Module Suite	<a href="http://sourceforge.net/projects/compbio/">http://sourceforge.net/projects/compbio/</a>
Delila-Genome Protein Binding Analysis	<a href="http://bioinformatics.org/project/?group_id=296">http://bioinformatics.org/project/?group_id=296</a>
Deltastat 2D Gel Analysis	<a href="http://bioinformatics.org/project/?group_id=371">http://bioinformatics.org/project/?group_id=371</a>
Dicom.Pm	<a href="http://dicomperl.sourceforge.net/">http://dicomperl.sourceforge.net/</a>
Dicomparser	<a href="http://sourceforge.net/projects/dicomparser/">http://sourceforge.net/projects/dicomparser/</a>
Disease Gene Profiler	<a href="http://sourceforge.net/projects/dgp/">http://sourceforge.net/projects/dgp/</a>
Eidors 3D	<a href="http://sourceforge.net/projects/eidors3d/">http://sourceforge.net/projects/eidors3d/</a>
Extractor For Esi Quadrupole ToF Tandem Ms Data Enabled For High Throughput Batch Processing	<a href="http://sourceforge.net/projects/protms/">http://sourceforge.net/projects/protms/</a>
Febrl	<a href="http://sourceforge.net/projects/febrl">http://sourceforge.net/projects/febrl</a>
Free Practice Management	<a href="http://sourceforge.net/projects/freepm/">http://sourceforge.net/projects/freepm/</a>
Free2bind	<a href="http://sourceforge.net/projects/free2bind/">http://sourceforge.net/projects/free2bind/</a>
Fs Kit Analysis Of Force Spectra	<a href="http://galaxie.cgb.ki.se/galaxieEST.html">http://galaxie.cgb.ki.se/galaxieEST.html</a>
Gata	<a href="http://sourceforge.net/projects/geckoe">http://sourceforge.net/projects/geckoe</a>
Gecko	<a href="http://www.cebitec.uni-bielefeld.de/groups/brf/software/gendb_info/license.html">http://www.cebitec.uni-bielefeld.de/groups/brf/software/gendb_info/license.html</a>
Gene Ontology Api For Java	<a href="http://combio.cs.brandeis.edu/GeneNotes/index.htm">http://combio.cs.brandeis.edu/GeneNotes/index.htm</a>
Genetdb Regulatory Networks Platform	<a href="http://bioinformatics.org/project/?group_id=207">http://bioinformatics.org/project/?group_id=207</a>
Genquire Genome Browsing And Annotation	<a href="http://bioinformatics.org/project/?group_id=99">http://bioinformatics.org/project/?group_id=99</a>
Gentle	<a href="http://gentle.magnusmanske.de/">http://gentle.magnusmanske.de/</a>
G-Language Project	<a href="http://www.g-language.org/main.html">http://www.g-language.org/main.html</a>
Hdbstat	<a href="http://www.soph.uab.edu/ssg_content.asp?id=1289">http://www.soph.uab.edu/ssg_content.asp?id=1289</a>
Hospital Scenario Generator	<a href="http://sourceforge.net/projects/hsg/">http://sourceforge.net/projects/hsg/</a>
Imlib3d	<a href="http://imlib3d.sourceforge.net/">http://imlib3d.sourceforge.net/</a>
Internet Pathology Suite	<a href="http://sourceforge.net/projects/ipath/">http://sourceforge.net/projects/ipath/</a>
Isys	<a href="http://sourceforge.net/projects/isys2/">http://sourceforge.net/projects/isys2/</a>
Jaligner	<a href="http://jaligner.sourceforge.net/">http://jaligner.sourceforge.net/</a>

Jatlasview	<a href="http://www.biomedcentral.com/1471-2105/6/47/abstract">http://www.biomedcentral.com/1471-2105/6/47/abstract</a>
Jklinika	<a href="http://sourceforge.net/projects/jclinic/">http://sourceforge.net/projects/jclinic/</a>
Knowtator	<a href="http://bionlp.sourceforge.net/Knowtator/index.shtml">http://bionlp.sourceforge.net/Knowtator/index.shtml</a>
Librarian Virtual Library Of Pdfs	<a href="http://bioinformatics.org/project/?group_id=131">http://bioinformatics.org/project/?group_id=131</a>
Mccore	<a href="http://sourceforge.net/projects/mccore">http://sourceforge.net/projects/mccore</a>
Medical Words	<a href="http://sourceforge.net/projects/medicalwords/">http://sourceforge.net/projects/medicalwords/</a>
MedsIt	<a href="http://sourceforge.net/projects/medslt">http://sourceforge.net/projects/medslt</a>
Msatfinder	<a href="http://www.bioinf.ceh.ac.uk/msatfinder/">http://www.bioinf.ceh.ac.uk/msatfinder/</a>
Mugen	<a href="http://genome.jouy.inra.fr/MuGeN/">http://genome.jouy.inra.fr/MuGeN/</a>
Netepi	<a href="http://sourceforge.net/projects/netepi">http://sourceforge.net/projects/netepi</a>
Ondex Suite	<a href="http://ondex.sourceforge.net/">http://ondex.sourceforge.net/</a>
Open Infrastructure For Outcomes	<a href="http://www.txoutcome.org/">http://www.txoutcome.org/</a>
Openvista	<a href="http://www.pacifichui.org/OpenVista/">http://www.pacifichui.org/OpenVista/</a>
Osirix	<a href="http://sourceforge.net/projects/osirix/">http://sourceforge.net/projects/osirix/</a>
Paclims	<a href="https://sourceforge.net/projects/paclims/">https://sourceforge.net/projects/paclims/</a>
Paracelsus	<a href="http://sourceforge.net/projects/paracelsus">http://sourceforge.net/projects/paracelsus</a>
Patient Record Anonymization Program	<a href="http://sourceforge.net/project">http://sourceforge.net/project</a>
Pentaplot	<a href="http://sourceforge.net/projects/pentaplot">http://sourceforge.net/projects/pentaplot</a>
Phaccs	<a href="http://sourceforge.net/projects/phaccs/">http://sourceforge.net/projects/phaccs/</a>
Ping	<a href="http://sourceforge.net/projects/pinghealth/">http://sourceforge.net/projects/pinghealth/</a>
Portablecbl	<a href="http://sourceforge.net/projects/cbl">http://sourceforge.net/projects/cbl</a>
Prehospital Medical Information System	<a href="http://sourceforge.net/projects/premis/">http://sourceforge.net/projects/premis/</a>
Probepicker	<a href="http://sourceforge.net/projects/probepicker/">http://sourceforge.net/projects/probepicker/</a>
Protms	<a href="http://www.protein-ms.de">http://www.protein-ms.de</a>
Pydaylight	<a href="http://www.dalkescientific.com/PyDaylight/">http://www.dalkescientific.com/PyDaylight/</a>
Real Time Medical Consultation System	<a href="http://sourceforge.net/projects/realmedic">http://sourceforge.net/projects/realmedic</a>
Semantic Moby	<a href="http://biomoby.open-bio.org/index.php/semantic-moby/">http://biomoby.open-bio.org/index.php/semantic-moby/</a>
Sequence Manipulation Suite 1	<a href="http://bioinformatics.org/project/?group_id=339">http://bioinformatics.org/project/?group_id=339</a>
Sight	<a href="http://jsight.sourceforge.net/index_SF.htm">http://jsight.sourceforge.net/index_SF.htm</a>
Simpacs	<a href="http://www.medical.toshiba.com/clinical/radiology/simpacs.htm">http://www.medical.toshiba.com/clinical/radiology/simpacs.htm</a>
Smith-Waterman On Greentea	<a href="http://bioinformatics.org/project/?group_id=134">http://bioinformatics.org/project/?group_id=134</a>
Sqlcare	<a href="http://sourceforge.net/projects/sqlcare/">http://sourceforge.net/projects/sqlcare/</a>
Tassel	<a href="http://sourceforge.net/projects/tassel/">http://sourceforge.net/projects/tassel/</a>
The Pogo Group	<a href="http://www.bioinformatics.org/pogo/">http://www.bioinformatics.org/pogo/</a>
Tls Motion Determination	<a href="http://pymmlib.sourceforge.net/">http://pymmlib.sourceforge.net/</a>
W-Curve Genomic String Visualization	<a href="http://www.bioinformatics.org/wcurve/">http://www.bioinformatics.org/wcurve/</a>
Xml4mat Xml/Matlab Format Converter	<a href="http://bioinformatics.org/project/?group_id=172">http://bioinformatics.org/project/?group_id=172</a>

3dspI	<a href="http://prisma.insa-lyon.fr/~kfrugier/index.php?page=3dspI">http://prisma.insa-lyon.fr/~kfrugier/index.php?page=3dspI</a>
A Taxonomic Search Engine: Federating Taxonomic Databases Using Web Services	<a href="http://www.biomedcentral.com/1471-2105/6/48">http://www.biomedcentral.com/1471-2105/6/48</a>
A Two-Way Interface Between Limited Systems Biology Markup Language And R	<a href="http://www.biomedcentral.com/1471-2105/5/190">http://www.biomedcentral.com/1471-2105/5/190</a>
Abc	<a href="http://mendel.stanford.edu/sidowlab/downloads/ABC_GERP/abcgerp.html">http://mendel.stanford.edu/sidowlab/downloads/ABC_GERP/abcgerp.html</a>
Amc Post Processing Package	<a href="http://idoimaging.com/cgi-bin/imaging/program.pl?ident=254">http://idoimaging.com/cgi-bin/imaging/program.pl?ident=254</a>
Amide	<a href="http://amide.sourceforge.net/">http://amide.sourceforge.net/</a>
An Edit Script For Taxonomic Classifications	<a href="http://darwin.zoology.gla.ac.uk/~rpage/forest/">http://darwin.zoology.gla.ac.uk/~rpage/forest/</a>
An Imaging System For Standardized Quantitative Analysis Of C. Elegans Behavior	<a href="http://www.pubmedcentral.nih.gov/articlerender.fcgi?tool=pubmed&amp;pubmedid=15331023">http://www.pubmedcentral.nih.gov/articlerender.fcgi?tool=pubmed&amp;pubmedid=15331023</a>
Anovarray	<a href="http://www.biomedcentral.com/1471-2105/6/150">http://www.biomedcentral.com/1471-2105/6/150</a>
Arb	<a href="http://www.arb-home.de/">http://www.arb-home.de/</a>
Arraycghbase	<a href="http://medgen.ugent.be/arrayCGHbase/">http://medgen.ugent.be/arrayCGHbase/</a>
Arrayd	<a href="http://www.igib.res.in/scientists/arrayd/arrayd.html">http://www.igib.res.in/scientists/arrayd/arrayd.html</a>
Assessing Local Structural Perturbations In Proteins	<a href="http://www.biomedcentral.com/1471-2105/6/226">http://www.biomedcentral.com/1471-2105/6/226</a>
Autofact	<a href="http://megasun.bch.umontreal.ca/Software/AutoFACT.htm">http://megasun.bch.umontreal.ca/Software/AutoFACT.htm</a>
Autoprime	<a href="http://www.autoprime.de/">http://www.autoprime.de/</a>
Biofloweb	<a href="http://urgi.infobiogen.fr/BioFloWeb">http://urgi.infobiogen.fr/BioFloWeb</a>
Biomap	<a href="http://proteinontology.info/">http://proteinontology.info/</a>
Bionets	<a href="http://x.amath.unc.edu:16080/BioNetS">http://x.amath.unc.edu:16080/BioNetS</a>
Biotext	<a href="http://biotext.berkeley.edu/">http://biotext.berkeley.edu/</a>
Blast Score Ration Analysis	<a href="http://www.microbialgenomics.org/BSR/">http://www.microbialgenomics.org/BSR/</a>
Blox	<a href="http://gecco.org.chemie.uni-frankfurt.de/h-blox/hblox.html">http://gecco.org.chemie.uni-frankfurt.de/h-blox/hblox.html</a>
Caadapter	<a href="http://ncicb.nci.nih.gov/download/">http://ncicb.nci.nih.gov/download/</a>
Caryoscope	<a href="http://caryoscope.stanford.edu/">http://caryoscope.stanford.edu/</a>
Cghpro	<a href="http://www.molgen.mpg.de/~abt_rop/molecular_cytogenetics/ArrayCGH/CGHPRO/">http://www.molgen.mpg.de/~abt_rop/molecular_cytogenetics/ArrayCGH/CGHPRO/</a>
Cgmim	<a href="http://www.biomedcentral.com/1471-2105/6/78/abstract">http://www.biomedcentral.com/1471-2105/6/78/abstract</a>
Codaln	<a href="http://www.bioinf.uni-leipzig.de/Software/codaln/">http://www.bioinf.uni-leipzig.de/Software/codaln/</a>
Dishevelled	<a href="http://www.pubmedcentral.nih.gov/articlerender.fcgi?artid=533865">http://www.pubmedcentral.nih.gov/articlerender.fcgi?artid=533865</a>
Efficient Decoding Algorithms For Generalized Hidden Markov Model	<a href="http://www.pubmedcentral.nih.gov/articlerender.fcgi?tool=pubmed&amp;pubmedid=15667658#N0x865898">http://www.pubmedcentral.nih.gov/articlerender.fcgi?tool=pubmed&amp;pubmedid=15667658#N0x865898</a>

Gene Finders	0.0x86eb7b8
Emboss-Gui	<a href="http://bioinfo.pbi.nrc.ca/~lukem/EMBOSS/">http://bioinfo.pbi.nrc.ca/~lukem/EMBOSS/</a>
Emerencia	<a href="http://emerencia.math.chalmers.se/">http://emerencia.math.chalmers.se/</a>
Estima	<a href="http://titan.biotec.uiuc.edu/ESTIMA/">http://titan.biotec.uiuc.edu/ESTIMA/</a>
Expander	<a href="http://www.cs.tau.ac.il/~rshamir/expander/">http://www.cs.tau.ac.il/~rshamir/expander/</a>
Fact	<a href="http://www.factweb.de/">http://www.factweb.de/</a>
Fieldtrip	<a href="http://www2.ru.nl/fcdonders/fieldtrip/download.html">http://www2.ru.nl/fcdonders/fieldtrip/download.html</a>
Forest	<a href="http://darwin.zoology.gla.ac.uk/~rpage/forest/">http://darwin.zoology.gla.ac.uk/~rpage/forest/</a>
Galaxieest	<a href="http://www.pdbj.org/GASH/">http://www.pdbj.org/GASH/</a>
Gash	<a href="http://bioinformatics.rcsi.ie/~redwards/gasp/">http://bioinformatics.rcsi.ie/~redwards/gasp/</a>
Gasp	<a href="http://gata.sourceforge.net/">http://gata.sourceforge.net/</a>
Gendb	<a href="http://bioinformatics.org/project/?group_id=474">http://bioinformatics.org/project/?group_id=474</a>
Genenotes	<a href="http://combio.cs.brandeis.edu/GeneNotes/index.htm">http://combio.cs.brandeis.edu/GeneNotes/index.htm</a>
Geneorder Coregenes	<a href="http://binf.gmu.edu/genometools.html">http://binf.gmu.edu/genometools.html</a>
Generalizations Of Markov Model To Characterize Biological Sequences	<a href="http://www.biomedcentral.com/1471-2105/6/219">http://www.biomedcentral.com/1471-2105/6/219</a>
Generating Quantitative Models Describing The Sequence Specificity Of Biological Processes With The Stabilized Matrix Method	<a href="http://www.mhc-pathway.net/smm">http://www.mhc-pathway.net/smm</a>
Genexplorer	<a href="http://www.biomedcentral.com/1471-2105/5/141">http://www.biomedcentral.com/1471-2105/5/141</a>
Genezilla	<a href="http://bioinformatics.org/project/?group_id=470">http://bioinformatics.org/project/?group_id=470</a>
Genomeviz	<a href="http://www.biomedcentral.com/1471-2105/5/198">http://www.biomedcentral.com/1471-2105/5/198</a>
Genome-Wide Prediction Display And Refinement Of Binding Sites With Information Theory-Based Models	<a href="http://www.biomedcentral.com/1471-2105/4/38">http://www.biomedcentral.com/1471-2105/4/38</a>
Genomic Multiple Sequence Alignments: Refinement Using A Genetic Algorithm	<a href="http://www.biomedcentral.com/1471-2105/6/200">http://www.biomedcentral.com/1471-2105/6/200</a>
Go-Perl Amigo	<a href="http://search.cpan.org/~cmungall/go-perl/">http://search.cpan.org/~cmungall/go-perl/</a>
Graphical Oncology Diagnostic System	<a href="http://www.cs.umb.edu/gods/">http://www.cs.umb.edu/gods/</a>
Gridstuffer	<a href="http://cmgm.stanford.edu/~cparnot/xgrid-stanford/index.html">http://cmgm.stanford.edu/~cparnot/xgrid-stanford/index.html</a>
Grock	<a href="http://egee-na4.ct.infn.it/biomed/GROCK.html">http://egee-na4.ct.infn.it/biomed/GROCK.html</a>
High Througput Snp Detection Pipeline	<a href="http://research.imb.uq.edu.au/seqdoc/">http://research.imb.uq.edu.au/seqdoc/</a>
High-Throughput Gominer	<a href="http://discover.nci.nih.gov/gominer/htgm.jsp">http://discover.nci.nih.gov/gominer/htgm.jsp</a>
Html4blast	<a href="http://bioperl.org/wiki/Module:Bio::Tools::Run::Pis eApplication::html4blast">http://bioperl.org/wiki/Module:Bio::Tools::Run::Pis eApplication::html4blast</a>
Htsnper1.0	<a href="http://bmc.ub.uni-potsdam.de/1471-2105-6-38/">http://bmc.ub.uni-potsdam.de/1471-2105-6-38/</a>
Identitag	<a href="http://pbil.univ-lyon1.fr/software/identitag/">http://pbil.univ-lyon1.fr/software/identitag/</a>

Integrated Web Service For Improving Alignment Quality Based On Segments Comparison	<a href="http://www.biomedcentral.com/1471-2105/5/98">http://www.biomedcentral.com/1471-2105/5/98</a>
Ipath	<a href="http://ipath.ch/team/brauchli">http://ipath.ch/team/brauchli</a>
Kclms	<a href="http://puggy.symonds.net/~hareesh/kidwai/">http://puggy.symonds.net/~hareesh/kidwai/</a>
Ldmas	<a href="http://molpath.his.path.cam.ac.uk/bioinformatics/LDMAS.shtml">http://molpath.his.path.cam.ac.uk/bioinformatics/LDMAS.shtml</a>
Libcov	<a href="http://users.cs.dal.ca/~cblouin/libcov/">http://users.cs.dal.ca/~cblouin/libcov/</a>
Mars	<a href="http://www.biomedcentral.com/1471-2105/6/101">http://www.biomedcentral.com/1471-2105/6/101</a>
Maxd	<a href="http://bioinf.man.ac.uk/microarray/maxd/">http://bioinf.man.ac.uk/microarray/maxd/</a>
Mbetoolbox	<a href="http://www.biomedcentral.com/1471-2105/6/64#IDA0LZDJ">http://www.biomedcentral.com/1471-2105/6/64#IDA0LZDJ</a>
Mcgh	<a href="http://folk.uio.no/junbaiw/mcgh/">http://folk.uio.no/junbaiw/mcgh/</a>
Mevislab	<a href="http://www.mevislab.de/">http://www.mevislab.de/</a>
Modbiosql	<a href="http://mbiosql.biomembrane.hu/mbs_docs.htm">http://mbiosql.biomembrane.hu/mbs_docs.htm</a>
Moltalk	<a href="http://www.moltalk.org/">http://www.moltalk.org/</a>
Mouseatlas Software	<a href="http://www.loni.ucla.edu/MAP/">http://www.loni.ucla.edu/MAP/</a>
Mprime	<a href="http://kbrin.a-bldg.louisville.edu/Tools/MPrime/">http://kbrin.a-bldg.louisville.edu/Tools/MPrime/</a>
Multiclass Discoverer	<a href="http://www.thep.lu.se/~markus/software/classdiscoverer/">http://www.thep.lu.se/~markus/software/classdiscoverer/</a>
Muscle	<a href="http://www.drive5.com/muscle">http://www.drive5.com/muscle</a>
Opensputnik	<a href="http://www.opensputnik.org">http://www.opensputnik.org</a>
Paircomp Familyrelationsii And Cartwheel	<a href="http://family.caltech.edu/">http://family.caltech.edu/</a>
Partigene	<a href="http://www.nematodes.org/PartiGene/">http://www.nematodes.org/PartiGene/</a>
Pegasys	<a href="http://bioinformatics.ubc.ca/pegasys/">http://bioinformatics.ubc.ca/pegasys/</a>
Permol	<a href="http://www.biologie.uni-regensburg.de/Biophysik/Kalbitzer/permol/permol.html">http://www.biologie.uni-regensburg.de/Biophysik/Kalbitzer/permol/permol.html</a>
Phenotype And Disease Ontologies	<a href="http://diseaseontology.sourceforge.net">http://diseaseontology.sourceforge.net</a>
Phylogenetic Reconstruction Of Ancestral Character States For Gene Expression And Mrna Splicing Data	<a href="http://rossnes.org/phyrex/">http://rossnes.org/phyrex/</a>
Predicting Functional Sites With An Automated Algorithm Suitable For Heterogeneous Datasets	<a href="http://www.pmap.csupomona.edu/MINER/dl.html">http://www.pmap.csupomona.edu/MINER/dl.html</a>
Primermatch	<a href="http://bioinformatics.org/project/?group_id=415">http://bioinformatics.org/project/?group_id=415</a>
Progenexpress	<a href="http://progenexpress.sourceforge.net/">http://progenexpress.sourceforge.net/</a>
Promi	<a href="http://promi.mpimp-golm.mpg.de/home.shtml">http://promi.mpimp-golm.mpg.de/home.shtml</a>
Prot4est	<a href="http://bioinformatics.org/cgi-bin/cvsweb.cgi/PartiGene/prot4EST/">http://bioinformatics.org/cgi-bin/cvsweb.cgi/PartiGene/prot4EST/</a>
Psi-Blast-Iss	<a href="http://www.ibt.lt/bioinformatics/iss/">http://www.ibt.lt/bioinformatics/iss/</a>
Pyevolve	<a href="http://cbis.anu.edu.au/software.html#PyEvolve">http://cbis.anu.edu.au/software.html#PyEvolve</a>
Scompy	part of biopython
Seqdoc	<a href="http://research.imb.uq.edu.au/seqdoc/">http://research.imb.uq.edu.au/seqdoc/</a>

Seqx	<a href="http://janbiro.com/Downloads.html">http://janbiro.com/Downloads.html</a>
Simprot	<a href="http://www.uhnres.utoronto.ca/labs/tillier/software.htm#3">http://www.uhnres.utoronto.ca/labs/tillier/software.htm#3</a>
Spcr	<a href="http://moleco.sjtu.edu.cn/SPCR">http://moleco.sjtu.edu.cn/SPCR</a>
Squid – A Simple Bioinformatics Grid	<a href="http://www.biomedcentral.com/1471-2105/6/197#B13">http://www.biomedcentral.com/1471-2105/6/197#B13</a>
Ss-Wrapper	<a href="ftp://ftp.genome.uab.edu">ftp://ftp.genome.uab.edu</a>
Stam	The R Project for Statistical Computing
Stanford Microarray Database	<a href="http://genome-www5.stanford.edu/">http://genome-www5.stanford.edu/</a>
Sting Millennium Suite	<a href="http://mirrors.rcsb.org/SMS/index_m_mirror.html">http://mirrors.rcsb.org/SMS/index_m_mirror.html</a>
Storing Linking And Mining Microarray Databases Using Srs	<a href="http://www.erasmusmc.nl/gatcplatform">http://www.erasmusmc.nl/gatcplatform</a>
Superficial	<a href="http://bioinformatics.charite.de/superficial/">http://bioinformatics.charite.de/superficial/</a>
T.I.M.S	<a href="http://bioinformatics.org/macrosHack/prog_list.html">http://bioinformatics.org/macrosHack/prog_list.html</a>
Taverna	<a href="http://www.mygrid.org.uk/index.php?module=page-master&amp;PAGE_user_op=view_page&amp;PAGE_id=44&amp;MMN_position=53:51:52">http://www.mygrid.org.uk/index.php?module=page-master&amp;PAGE_user_op=view_page&amp;PAGE_id=44&amp;MMN_position=53:51:52</a>
Tetra	<a href="http://www.megx.net/tetra">http://www.megx.net/tetra</a>
Text2knowledge Text Mining Tools	<a href="http://bioinformatics.org/textknowledge/">http://bioinformatics.org/textknowledge/</a>
The Cartwheel Bioinformatics Project	<a href="http://cartwheel.caltech.edu/">http://cartwheel.caltech.edu/</a>
The Indeval	<a href="http://www.sci.muni.cz/botany/elzdroje/indeval/">http://www.sci.muni.cz/botany/elzdroje/indeval/</a>
Tmb-Hunt	<a href="http://www.biomedcentral.com/1471-2105/6/56">http://www.biomedcentral.com/1471-2105/6/56</a>
Tools Enabling The Elucidation Of Molecular Pathways Active In Human Disease: Application To Hepatitis C Virus Infection	<a href="http://labs.systemsbiology.net/galitski/hepc/if_and_ct.html">http://labs.systemsbiology.net/galitski/hepc/if_and_ct.html</a>
Tools For Loading Medline Into A Local Relational Database	<a href="http://biotext.berkeley.edu/software">http://biotext.berkeley.edu/software</a>
Toophpix	<a href="http://truth.positive-internet.com/~mpreston/tp/tp0412intro.html">http://truth.positive-internet.com/~mpreston/tp/tp0412intro.html</a>
Transalign	<a href="http://www.tierzucht.tum.de/Bininda-Emonds/">http://www.tierzucht.tum.de/Bininda-Emonds/</a>
Trans-Proteomic Pipeline	<a href="http://tools.proteomecenter.org/TPP.php">http://tools.proteomecenter.org/TPP.php</a>
Vestige	<a href="http://cbis.anu.edu.au/software.html">http://cbis.anu.edu.au/software.html</a>
Vienna Rna Package	<a href="http://www.biomedcentral.com/1471-2105/6/89#IDAF2KLM">http://www.biomedcentral.com/1471-2105/6/89#IDAF2KLM</a>
Viral Bioinformatics Resource Center	<a href="http://athena.bioc.uvic.ca/">http://athena.bioc.uvic.ca/</a>
Visualization And Analysis Of Microarray And Gene Ontology Data With Treemaps	<a href="http://www.cs.umd.edu/hcil/treemap/">http://www.cs.umd.edu/hcil/treemap/</a>
Visualization Of Comparative Genomic Analyses By Blast Score Ratio	<a href="http://www.microbialgenomics.org/BSR/">http://www.microbialgenomics.org/BSR/</a>
Visualization-Based Discovery And	<a href="http://function.princeton.edu/ChARMView/charm_">http://function.princeton.edu/ChARMView/charm_</a>