

biotools_link	download_url	download_type
https://bio.tools/2020plus	https://github.com/KarchinLab/2020plus	Source code
https://bio.tools/3dnetmod	https://bitbucket.org/creminslab/3dnetmod_method_v1_0_10_06_17/src	Source code
https://bio.tools/4pipe4	https://github.com/StuntsPT4Pipe4/releases	Source package
https://bio.tools/aaindexextract	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/aaindexextract	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/abiview	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/abiview	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/acdc	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/acdc	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/acdc_estimation	https://github.com/mlux86/acdc	Source code
https://bio.tools/acdpretty	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/acdpretty	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/acitable	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/acitable	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/acdtrace	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/acdtrace	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/acdvalid	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/acdvalid	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/ActiveDriver	https://CRAN.R-project.org/package=ActiveDriver	Source package
https://bio.tools/ActiveDriver	https://github.com/reimandlab/ActiveDriver	Source code
https://bio.tools/adaptest	http://bioconductor.org/packages/release/bioc/vignettes/adaptest/inst/doc/differentialExpression_R	Source code
https://bio.tools/affyqcreport	http://bioconductor.org/packages/release/bioc/vignettes/affyQCReport/inst/doc/affyQCReport_R	Source code
https://bio.tools/agglolndel	https://bibiserv.cebitec.uni-bielefeld.de/applications/agglolndel/resources/downloads/agglolndel.jar	Source code
https://bio.tools/agglolndel	https://bibiserv.cebitec.uni-bielefeld.de/applications/agglolndel/resources/downloads/agglolndel.jar	Binaries
https://bio.tools/aglip	http://bioconductor.org/packages/release/bioc/vignettes/aglip/inst/doc/aglip_manual_R	Source code
https://bio.tools/agmicrorna	http://bioconductor.org/packages/release/bioc/vignettes/AgliMicroRna/inst/doc/AgliMicroRna_R	Source code
https://bio.tools/ahtpdb	http://crdd.osdd.net/raghava/ahtpdb/download.php	Biological data
https://bio.tools/alientrimmer	ftp://ftp.pasteur.fr/pub/gensoft/projects/AlienTrimmer/	Source code
https://bio.tools/alientrimmer	ftp://ftp.pasteur.fr/pub/gensoft/projects/AlienTrimmer/	Binaries
https://bio.tools/aligncopy	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/aligncopy	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/aligncopypair	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/aligncopypair	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/alignmentcomparator	http://bioinfweb.info/AlignmentComparator/Download	Binaries
https://bio.tools/allotpred	https://github.com/jgreener64/allotpred	Source code
https://bio.tools/alra	https://github.com/KlugerLab/ALRA	Source code
https://bio.tools/amatreader	https://github.com/Idkerlab/aMatReader	Source code
https://bio.tools/amen	https://sourceforge.net/projects/amen/	Source code
https://bio.tools/andi	https://github.com/EvoBioInf/andi/releases	Source code
https://bio.tools/angsd	https://github.com/ANGSD/angsd	Source code
https://bio.tools/angsd	https://github.com/ANGSD/angsd	Binaries
https://bio.tools/anota2seq	http://bioconductor.org/packages/release/bioc/vignettes/anota2seq/inst/doc/anota2seq_R	Source code
https://bio.tools/antigenic	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/antigenic	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/apache_laverna	http://laverna.incubator.apache.org/download/code/	Source code
https://bio.tools/apache_laverna	http://laverna.incubator.apache.org/download/	Binaries
https://bio.tools/apeglm	http://bioconductor.org/packages/release/bioc/vignettes/apeglm/inst/doc/apeglm.R	Source code
https://bio.tools/apt	https://github.com/ChangLab/ATACPrimerTool	Binaries
https://bio.tools/apt	https://github.com/ChangLab/ATACPrimerTool/blob/master/Running_APT_with_Galaxy.md	Tool wrapper (galaxy)
https://bio.tools/aquaria	https://aquaria.ws/help/Aquaria3D_QR_v1.pdf	Source code
https://bio.tools/aquaria	https://aquaria.ws/help/Aquaria3D_QR_v1.pdf	Binaries
https://bio.tools/arachne_neural	https://github.com/LeonidSavchenko/Arachne	Source code
https://bio.tools/arb	http://download.arb-home.de/release/latest/	Binaries
https://bio.tools/arem	https://github.com/jakebiesinger/AREM	Source code
https://bio.tools/arts_antibiotics	https://bitbucket.org/ziemertlab/artswebapp	Source code
https://bio.tools/arts_antibiotics	https://bitbucket.org/ziemertlab/arts	Source code
https://bio.tools/ascat	https://github.com/Crick-CancerGenomics/ascat	Source code
https://bio.tools/askomics	https://github.com/askomics/askomics/releases	Binary package
https://bio.tools/aspasia	https://www.york.ac.uk/computational-immunology/software/aspasia/	Binaries
https://bio.tools/assemblyget	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/assemblyget	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/assist	http://compbiotoolbox.fmach.it/assist/	Source code
https://bio.tools/assist	http://compbiotoolbox.fmach.it/assist/	Binaries
https://bio.tools/asialavista	http://sammeth.net/confluence/display/ASTA/2+-+Download	Binaries
https://bio.tools/atlas2	https://sourceforge.net/projects/atlas2/files/	Source code
https://bio.tools/atomicchargecalculator	http://webchem.ncbr.muni.cz/Platform/ChargeCalculator/DownloadService	Binaries
https://bio.tools/autohd	http://www.ms-utilis.org/autohd_download.html	Source code
https://bio.tools/avpdb	http://crdd.osdd.net/servers/avpdb/dd.php	Biological data
https://bio.tools/bacddiver	https://github.com/TIBHannover/BacDiverR	Source code
https://bio.tools/backtranambig	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/backtranambig	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/backtranseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/backtranseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/baldr	https://github.com/BosingerL/ab/BALDR	Source code
https://bio.tools/bamsi	http://www.1000genomes.org/	Biological data
https://bio.tools/bamsi	https://github.com/NGDSG/BAMS1	Source package
https://bio.tools/bamsi	https://github.com/NGDSG/BAMS1-API	API specification
https://bio.tools/banana	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/banana	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/banocc	http://bioconductor.org/packages/release/bioc/vignettes/banocc/inst/doc/banocc-vignette.R	Source code
https://bio.tools/barmap	https://github.com/tseemann/barmap	Source code
https://bio.tools/basecallqc	http://bioconductor.org/packages/release/bioc/vignettes/basecallQC/inst/doc/basecallQC.R	Source code
https://bio.tools/basic_assembly	http://ftic.uchicago.edu/~aakhan/BASIC/	Source package
https://bio.tools/basic_assembly	https://github.com/akds/BASIC	Source code
https://bio.tools/bat	http://www.bioinf.uni-leipzig.de/Software/BAT/download/	Source code
https://bio.tools/bbknn	https://github.com/Teichlab/bbknn	Source package
https://bio.tools/bcseq	http://bioconductor.org/packages/release/bioc/vignettes/bcSeq/inst/doc/bcSeq.R	Source code
https://bio.tools/bcseq_bias-corrected_sequencing_analysis	http://www-rcf.usc.edu/~liangche/software.html	Binary package

biotools_link	download_url	download_type
https://bio.tools/bdbm	https://github.com/sing-group/BDBM/	Source code
https://bio.tools/bdbm	http://www.sing-group.org/BDBM/download.html	Binaries
https://bio.tools/bepipred	http://www.cbs.dtu.dk/services/doc/bepipred-1.0.readme	Source code
https://bio.tools/bepipred	http://www.cbs.dtu.dk/services/doc/bepipred-1.0.readme	Binaries
https://bio.tools/betaware	http://betaware.biocomp.unibo.it/BetAware/default/software	Source code
https://bio.tools/betaware	http://betaware.biocomp.unibo.it/BetAware/default/software	Binaries
https://bio.tools/bicycle	https://www.sing-group.org/bicycle/download.html	Binary package
https://bio.tools/bids_apps	https://github.com/BIDS-Apps	Source code
https://bio.tools/bignasim	http://mmb.irbbarcelona.org/BigNASim/help.php?id=BNSdownload	Source code
https://bio.tools/bignasim	http://mmb.irbbarcelona.org/BigNASim/help.php?id=BNSdownload	Binaries
https://bio.tools/binat	http://dmb.iasi.cnr.it/Downloads/BINAT/binat-1.0-src.rar	Source package
https://bio.tools/binat	http://dmb.iasi.cnr.it/Downloads/BINAT/binat-1.0-desktop-client.rar	Source package
https://bio.tools/bio.tools	https://github.com/bio-tools/biotoolsRegistry	Source code
https://bio.tools/biobb	https://github.com/bioexcel	Source code
https://bio.tools/biobb	https://anaconda.org/bioconda/biobb_common	Source package
https://bio.tools/biobb	https://anaconda.org/bioconda/biobb_io	Source package
https://bio.tools/biobb	https://anaconda.org/bioconda/biobb_md	Source package
https://bio.tools/bioexcel_seqqc	https://github.com/bioexcel/BioExcel_SeqQC	Source code
https://bio.tools/biomaj	https://github.com/genouest/biomaj	Source package
https://bio.tools/biomaj	https://hub.docker.com/r/osalou/biomaj-docker/	Container file
https://bio.tools/bionetstat	http://bioconductor.org/packages/3.7/bioc/src/contrib/Archive/BioNetStat/	Source package
https://bio.tools/bionode	https://github.com/bionode/bionode	Source code
https://bio.tools/bioplex_2.0	http://bioplex.hms.harvard.edu/downloadInteractions.php	Biological data
https://bio.tools/bioplex_2.0	http://bioplex.hms.harvard.edu/downloadData.php	Biological data
https://bio.tools/biosed	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/biosed	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/bioservices	https://pypi.org/project/bioservices/	Source code
https://bio.tools/biosunms	https://sourceforge.net/projects/biosunms/files/latest/download	Binaries
https://bio.tools/biosyntax	https://github.com/bioSyntax/bioSyntax/releases	Source package
https://bio.tools/biq_analyzer	https://biq-analyzer.bioinf.mpi-inf.mpg.de/download.php	Binary package
https://bio.tools/bison	https://github.com/dpryan79/bison	Source code
https://bio.tools/blackfan_anemia	http://www.dbagene.unibo.it/docs/LOVD_manual_2.0.pdf	Source code
https://bio.tools/blackfan_anemia	http://www.dbagene.unibo.it/docs/LOVD_manual_2.0.pdf	Binaries
https://bio.tools/blast_viewer	https://github.com/pgdurand/BlastViewer/releases	Binary package
https://bio.tools/blast-filter-tool	https://github.com/pgdurand/BLAST-Filter-Tool	Source code
https://bio.tools/blast-filter-tool	https://github.com/pgdurand/BLAST-Filter-Tool/releases	Binaries
https://bio.tools/blma	http://bioconductor.org/packages/release/bioc/vignettes/BLMA/inst/doc/BLMA.pdf	Source code
https://bio.tools/bmge	ftp://ftp.pasteur.fr/pub/gensoft/projects/BMGE/	Source code
https://bio.tools/bmge	ftp://ftp.pasteur.fr/pub/gensoft/projects/BMGE/	Binaries
https://bio.tools/bnbc	http://bioconductor.org/packages/release/bioc/vignettes/bnbc/inst/doc/bnbc.R	Source code
https://bio.tools/bowhead	https://gitlab.com/engel/bowhead	Source code
https://bio.tools/bowtie2	https://gitlab.com/sibyllewohlgemuth/cwl_files/raw/master/bowtie2.cwl	CWL file
https://bio.tools/bowtie2	https://gitlab.com/sibyllewohlgemuth/cwl_files/raw/master/bowtie2_build.cwl	CWL file
https://bio.tools/bowtie2	https://docker-ui.genouest.org/app/#container/bioconda/bowtie2	Container file
https://bio.tools/bowtie2	https://anaconda.org/bioconda/bowtie2	Container file
https://bio.tools/breath_easy_ed	https://github.com/johnksander/BEEEDA_sample_data	Biological data
https://bio.tools/btwisted	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/btwisted	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/bufet	https://github.com/diwis/BUFET	Source code
https://bio.tools/bumhmm	http://bioconductor.org/packages/release/bioc/vignettes/BUMHMM/inst/doc/BUMHMM.pdf	Source code
https://bio.tools/bwa	https://gitlab.com/sibyllewohlgemuth/cwl_files/raw/master/bwa_index.cwl	CWL file
https://bio.tools/bwa	https://gitlab.com/sibyllewohlgemuth/cwl_files/raw/master/bwa_mem.cwl	CWL file
https://bio.tools/bwa	https://docker-ui.genouest.org/app/#container/bioconda/bwa	Container file
https://bio.tools/bwa	https://anaconda.org/bioconda/bwa	Container file
https://bio.tools/bwa-pssm	https://github.com/pkerpedjiev/bwa-pssm	Source code
https://bio.tools/bwa-pssm	https://github.com/pkerpedjiev/bwa-pssm	Binaries
https://bio.tools/cachedas	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/cachedas	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/cachedbfetch	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/cachedbfetch	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/cacheebeyesearch	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/cacheebeyesearch	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/cacheensembl	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/cacheensembl	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/cai	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/cai	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/caver_pymol_plugin	http://www.caver.cz/fil/download/manual/caver_plugin_userguide.pdf	Source code
https://bio.tools/caver_pymol_plugin	http://www.caver.cz/fil/download/manual/caver_plugin_userguide.pdf	Binaries
https://bio.tools/caw	https://github.com/SciLifeLab/Sarek/releases/tag/2.0.0	Binaries
https://bio.tools/cbaf	http://bioconductor.org/packages/release/bioc/vignettes/cbaf/inst/doc/cbaf.R	Source code
https://bio.tools/ccfindr	http://bioconductor.org/packages/release/bioc/vignettes/ccfindr/inst/doc/ccfindr.R	Source code
https://bio.tools/cdfread	http://sourceforge.net/p/cdfread/code/HEAD/tree/src/cdfread.c	Source code
https://bio.tools/cellalign	https://github.com/shenorrLab/cellAlign	Binary package
https://bio.tools/cellbaseR	http://bioconductor.org/packages/release/bioc/vignettes/cellbaseR/inst/doc/cellbaseR.R	Source code
https://bio.tools/cellscape	https://bioconductor.org/packages/release/bioc/vignettes/cellscape/inst/doc/cellscape_vignette.html	Command-line specification
https://bio.tools/cg-cat	http://bibiserv.cebitec.uni-bielefeld.de/cgcat?id=cgcat_download	Source code
https://bio.tools/cg-cat	http://bibiserv.cebitec.uni-bielefeld.de/cgcat?id=cgcat_download	Binaries
https://bio.tools/chaos	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/chaos	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/charge	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/charge	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/checktrans	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/checktrans	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/chimera_bioconductor	http://bioconductor.org/packages/release/bioc/vignettes/chimera/inst/doc/chimera.R	Source code
https://bio.tools/chimeraviz	http://bioconductor.org/packages/release/bioc/vignettes/chimeraviz/inst/doc/chimeraviz_vignette.R	Source code
https://bio.tools/chips	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/chips	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/chlorop	http://www.cbs.dtu.dk/services/doc/chlorop-1.1.readme	Source code

biotools_link	download_url	download_type
https://bio.tools/chlorop	http://www.cbs.dtu.dk/services/doc/chlorop-1.1.readme	Binaries
https://bio.tools/chromatin_dynamics	https://www.dropbox.com/s/w8e09wqs92l0g9r/ChromatineDynamicsNAflex_ON.qcow2?dl=1	VM image
https://bio.tools/chromatin_dynamics	http://inb.bsc.es/MuGimages/ChromatineDynamicsNAflex_ON.qcow2	VM image
https://bio.tools/chromswitch	http://bioconductor.org/packages/release/bioc/vignettes/chromswitch/inst/doc/chromswitch_intro.R	Source code
https://bio.tools/circle-seq	https://github.com/tsailabSJC/circleseq	Binaries
https://bio.tools/cirdna	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/cirdna	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/cistem	https://cistem.org/software	Binaries
https://bio.tools/clark	http://clark.cs.ucr.edu/Tool/	Source code
https://bio.tools/clark	http://clark.cs.ucr.edu/Tool/	Binaries
https://bio.tools/cloudaligner	https://github.com/tungnguyen/CloudAligner	Source code
https://bio.tools/clusterseq	http://bioconductor.org/packages/release/bioc/vignettes/clusterSeq/inst/doc/clusterSeq.R	Source code
https://bio.tools/clustvis	https://hub.docker.com/r/taunometsalu/clustvis/	Container file
https://bio.tools/cmd	https://eds.gene.le.ac.uk/docs/L_OVD_manual_2.0.pdf	Source code
https://bio.tools/cmd	https://eds.gene.le.ac.uk/docs/L_OVD_manual_2.0.pdf	Binaries
https://bio.tools/cnogpro	https://cran.r-project.org/web/packages/CNOGpro/index.html	Source code
https://bio.tools/codcmp	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/codcmp	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/codcopy	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/codcopy	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/coderet	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/coderet	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/cofactory	http://www.cbs.dtu.dk/services/doc/cofactory-1.0.readme	Source code
https://bio.tools/cofactory	http://www.cbs.dtu.dk/services/doc/cofactory-1.0.readme	Binaries
https://bio.tools/collect_mgf	http://www.ms-utis.org/collect_mgf.c	Source code
https://bio.tools/comet	https://sourceforge.net/projects/comet-ms/files/	Source code
https://bio.tools/comet	https://sourceforge.net/projects/comet-ms/files/	Binaries
https://bio.tools/comparems2	http://www.ms-utis.org/compareMS2.c	Source code
https://bio.tools/comparems2	http://www.ms-utis.org/compareMS2.html	Binaries
https://bio.tools/compartments	https://compartments.jensenlab.org/Downloads	Biological data
https://bio.tools/compil	https://github.com/robinparky/prolicidComPIL	Binaries
https://bio.tools/CompositeDriver	https://github.com/khuranalab/CompositeDriver	Source code
https://bio.tools/compseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/compseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/compass	http://mmbarcelona.org/~andrio/PyMDSSetup-v2.qcow2	VM image
https://bio.tools/compass	https://www.bsc.es/research-and-development/software-and-apps/software-list/comp-superscalar/downloads	Source package
https://bio.tools/comspari	http://www.biomechanic.org/comspari/#download	Source code
https://bio.tools/conet	http://psbweb05.psb.ugent.be/conet/documents/CoNetDoc.pdf	Source code
https://bio.tools/conet	http://psbweb05.psb.ugent.be/conet/documents/CoNetDoc.pdf	Binaries
https://bio.tools/cons	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/cons	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/consambig	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/consambig	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/contrail	https://github.com/ndimiduk/contrail-bio	Source code
https://bio.tools/convert-matrix	http://rsat.eu/	Source code
https://bio.tools/convert-matrix	http://rsat.eu/	Binaries
https://bio.tools/copasi	http://copasi.org/Download/	Binaries
https://bio.tools/copheism	https://bitbucket.org/izhbannikov/copheism/downloads/	Source package
https://bio.tools/coseq	http://bioconductor.org/packages/release/bioc/vignettes/coseq/inst/doc/coseq.R	Source code
https://bio.tools/cpgplot	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/cpgplot	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/cpgreport	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/cpgreport	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/cpmd	http://www.fz-juelich.de/ias/jsc/EN/Expertise/Support/Software/Chemistry/CPMD.html	VM image
https://bio.tools/crispcasfinder	https://crispcas.i2bc.paris-saclay.fr/Home/Download	Source package
https://bio.tools/cuneiform	http://www.cuneiform-lang.org/download	Source code
https://bio.tools/cuneiform	http://www.cuneiform-lang.org/download	Binaries
https://bio.tools/cusp	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/cusp	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/cutgextract	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/cutgextract	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/cutseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/cutseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/cydar	http://bioconductor.org/packages/release/bioc/html/cydar.html	Binaries
https://bio.tools/cytolib	http://bioconductor.org/packages/release/bioc/vignettes/cytolib/inst/doc/cytolib.R	Source code
https://bio.tools/dan	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/dan	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/dataanalysis2tpp	http://www.ms-utis.org/DataAnalysis2TPP.awk	Source code
https://bio.tools/dbiblast	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/dbiblast	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/dbifasta	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/dbifasta	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/dbiflat	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/dbiflat	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/dbigcg	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/dbigcg	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/dbitell	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/dbitell	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/dbxcompress	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/dbxcompress	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/dbxedam	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/dbxedam	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/dbxfasta	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/dbxfasta	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/dbxflat	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/dbxflat	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/dbxgcg	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/dbxgcg	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries

biotools_link	download_url	download_type
https://bio.tools/dbxobo	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/dbxobo	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/dbxreport	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/dbxreport	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/dbxresource	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/dbxresource	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/dbxstat	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/dbxstat	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/dbxtax	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/dbxtax	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/dbxuncompress	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/dbxuncompress	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/dcell	https://github.com/dekerlab/DCell/	Biological data
https://bio.tools/decontools	http://ncrr.pnl.gov/software/sourcecode_register.asp?id=18	Source code
https://bio.tools/deepsig	https://deepsig.biocomp.unibo.it/deepsig/default/software	Source code
https://bio.tools/defcom	https://bitbucket.org/bryancquach/defcom/downloads/	Source package
https://bio.tools/degapseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/degapseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/deltamass	https://github.com/chhh/deltamass	Source package
https://bio.tools/density	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/density	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/descseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/descseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/detectlr	https://sourceforge.net/projects/detectlr/	Source code
https://bio.tools/dexsi	https://github.com/DExSI/DExSI/releases/download/v1.11/DExSI_setup_exe	Binaries
https://bio.tools/diffseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/diffseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/diffusion	https://github.com/dekerlab/heat-diffusion	Source code
https://bio.tools/dig	http://www.ms-utlis.org/dig/dig.c	Source code
https://bio.tools/dig2	http://www.ms-utlis.org/dig2/dig2.c	Source code
https://bio.tools/digenome-seq	http://www.rgenome.net/digenome-js/standalone	Binaries
https://bio.tools/dinamo	https://github.com/bonsai-team/DiNAMO/releases	Binaries
https://bio.tools/discordant	http://bioconductor.org/packages/release/bioc/vignettes/discordant/inst/doc/Discordant_vignette.R	Source code
https://bio.tools/discotope	http://www.cbs.dtu.dk/services/doc/discotope-1.1_readme	Source code
https://bio.tools/discotope	http://www.cbs.dtu.dk/services/doc/discotope-1.1_readme	Binaries
https://bio.tools/diseases	https://diseases.jensenlab.org/Downloads	Biological data
https://bio.tools/disgenet	http://www.disgenet.org/web/DisGeNET/menu/downloads	Source code
https://bio.tools/disgenet	http://www.disgenet.org/web/DisGeNET/menu/downloads	Binaries
https://bio.tools/dismat	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/dismat	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/disvis	https://github.com/haddocking/disvis	Source code
https://bio.tools/dotmatcher	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/dotmatcher	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/dotpath	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/dotpath	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/dotup	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/dotup	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/dreg	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/dreg	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/drfindata	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/drfindata	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/drfindformat	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/drfindformat	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/drfindid	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/drfindid	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/drfindresource	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/drfindresource	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/drget	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/drget	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/drtext	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/drtext	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/e-Driver	https://github.com/eduardporta/e-Driver	Source code
https://bio.tools/eagle_gxe	https://github.com/davidaknowles/eagle	Command-line specification
https://bio.tools/ebprot	https://github.com/cssblab/EBprot	Source package
https://bio.tools/ec-clem	https://www.nature.com/articles/nmeth.4170	Source code
https://bio.tools/edamdef	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/edamdef	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/edamhasinput	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/edamhasinput	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/edamhasoutput	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/edamhasoutput	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/edamisformat	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/edamisformat	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/edamidsid	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/edamidsid	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/edamname	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/edamname	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/edialign	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/edialign	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/effectivedb	http://www.effectivedb.org/download	Source code
https://bio.tools/effectivedb	http://www.effectivedb.org/download	Binaries
https://bio.tools/effective3	http://www.effectivedb.org/method/effective3	Source code
https://bio.tools/effective3	http://www.effectivedb.org/method/effective3	Binaries
https://bio.tools/eggnog	http://eggnogdb.embl.de/#/app/downloads	Biological data
https://bio.tools/einverted	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/einverted	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/embedsom	https://github.com/exaexa/EmbedSOM.git	Source code
https://bio.tools/embosdata	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code

biotools_link	download_url	download_type
https://bio.tools/embosssdata	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/embosssupdate	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/embosssupdate	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/embossversion	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/embossversion	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/emmma	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/emmma	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/emrowse	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/emrowse	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/empusa	http://download.systemsbioology.nl/empusa/EmpusaCodeGen.jar	Binary package
https://bio.tools/enrichedheatmap	https://bioconductor.org/packages/release/bioc/html/EnrichedHeatmap.html	Source code
https://bio.tools/entret	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/entret	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/eosc_pipeline-reproducibility	https://github.com/CRG-CNAG/EOSC-Pilot	Source code
https://bio.tools/epestfind	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/epestfind	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/epidb	https://epidb.animalgenome.org/resem_expression.php	Biological data
https://bio.tools/epilog	http://epilog-tool.org/documentation	Biological data
https://bio.tools/epilog	http://epilog-tool.org/downloads	Source package
https://bio.tools/epinew	http://bioconductor.org/packages/release/bioc/vignettes/epiNEM/inst/doc/epiNEM.R	Source code
https://bio.tools/epimer3	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/epimer3	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/epimer32	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/epimer32	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/equicktandem	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/equicktandem	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/erma	http://bioconductor.org/packages/release/bioc/vignettes/erma/inst/doc/erma.R	Source code
https://bio.tools/esimsa	http://www.ms-utlis.org/esimsa.c	Source code
https://bio.tools/esimsa2d	http://www.ms-utlis.org/esimsa2D.c	Source code
https://bio.tools/esprit-forest	https://www.acsu.buffalo.edu/~vijunsun/lab/ESPRIT-Tree.html	Source code
https://bio.tools/est2genome	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/est2genome	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/etandem	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/etandem	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/eva_inbred	https://www.nordgen.org/en/farm-animals/resources/ocs/	Binaries
https://bio.tools/evmutation	https://github.com/debbiemarkslab/EVmutation	Source code
https://bio.tools/evmutation	https://marks.hms.harvard.edu/evmutation/human_proteins.html	Biological data
https://bio.tools/exibd	http://akeylab.princeton.edu/downloads.html	Source package
https://bio.tools/extractalign	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/extractalign	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/extractfeat	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/extractfeat	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/extractseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/extractseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/faas	http://faas.bme.unc.edu/sample/Paxillin_test.tif	Test data
https://bio.tools/fateid	https://github.com/dgrun/FateID	Binary package
https://bio.tools/fc_analysis	https://static-content.springer.com/esm/art%3A10.1186%2F12859-018-2265-4/MediaObjects/12859_2018_2265_MOESM1_ESM.exe	Source package
https://bio.tools/featcopy	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/featcopy	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/featmerge	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/featmerge	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/featreport	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/featreport	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/feattext	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/feattext	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/feelnc	https://github.com/tdemrien/FEELnc	Command-line specification
https://bio.tools/fgmp	https://github.com/stajichlab/FGMP/releases	Source code
https://bio.tools/fidpd	http://202.119.249.49/pubdata/fidpd.pl	Source code
https://bio.tools/fimtrack	https://www.uni-muenster.de/PRIA/en/FIM/download.shtml	Binary package
https://bio.tools/fimtrack	https://github.com/i-git/FIMTrack	Source code
https://bio.tools/findkm	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/findkm	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/flipper	http://bitlab-es.com/flipper/files/Flipper_supplementary_materialv3.pdf	Source code
https://bio.tools/flowsim	http://maide.org/~keili/biohaskell/flowsim/	Source code
https://bio.tools/fluvidencesynthesis	https://github.com/MJomaba/flu-evidence-synthesis	Source code
https://bio.tools/freak	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/freak	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/funcexplorer	https://gl.cs.ut.ee/bit/funcexplorer_public	Source code
https://bio.tools/funel	http://ico2s.org/data/code/funel.tar.xz	Source code
https://bio.tools/funel	http://ico2s.org/data/code/funel.tar.xz	Binaries
https://bio.tools/fuzznuc	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/fuzznuc	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/fuzzpro	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/fuzzpro	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/fuzztran	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/fuzztran	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/fuzzyclust	https://bitbucket.org/veitveit/fuzzyclust	Source code
https://bio.tools/gamier	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/gamier	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/gasoline	https://alpha.dmi.unict.it/~gmicale/gasoline/#download	Biological data
https://bio.tools/gatk-lodn	https://bitbucket.org/BBDA-UNIBO/wes-pipeline/wiki/Home	Source code
https://bio.tools/gcocoa	https://pypi.python.org/pypi/gprofiler-official	Binary package
https://bio.tools/gcocoa	https://cran.r-project.org/web/packages/gProfileR/index.html	Binary package
https://bio.tools/gcocoa	https://www.npmjs.com/package/biojs-vis-gprofiler	Binary package
https://bio.tools/gcocoa	https://toolshed.g2.bx.psu.edu/repository?repository_id=2d3d786121020d7a	Tool wrapper (galaxy)
https://bio.tools/gconvert	https://pypi.python.org/pypi/gprofiler-official	Binary package
https://bio.tools/gconvert	https://cran.r-project.org/web/packages/gProfileR/index.html	Binary package
https://bio.tools/gconvert	https://www.npmjs.com/package/biojs-vis-gprofiler	Binary package

biotools_link	download_url	download_type
https://bio.tools/gconvert	https://toolshed.g2.bx.psu.edu/repository?repository_id=2d3d786121020d7a	Tool wrapper (galaxy)
https://bio.tools/gecko	https://sourceforge.net/projects/gecko-aligner/	Source code
https://bio.tools/gecko-csb	http://bitlab-es.com/gecko/template.php#GECKO-CSB	Binaries
https://bio.tools/gecko-mgv	http://pistacho.ac.uma.es/help	Source code
https://bio.tools/gecko-mgv	http://pistacho.ac.uma.es/help	Binaries
https://bio.tools/geceee	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/geceee	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/gem3	https://github.com/smarco/gem3-mapper	Source code
https://bio.tools/gemma_algorithm	https://github.com/genetics-statistics/GEMMA/releases	Binary package
https://bio.tools/gemmapper	https://sourceforge.net/projects/gemlibrary/files/gem-library/Binary%20pre-release%203/	Binaries
https://bio.tools/gemoma	http://www.jstacs.de/download.php?which=GeMoMa	Source package
https://bio.tools/geneid	http://genome.crg.es/software/geneid/index.html	Binaries
https://bio.tools/genevalidator	https://github.com/wurmlab/genevalidator	Source code
https://bio.tools/gennon-h	http://genome.crg.es/phylo_mod/gennonhom/	Source code
https://bio.tools/genomemai	http://www.genomemai.org/DownloadAllExperimentsForm	Source code
https://bio.tools/genomemai	http://www.genomemai.org/DownloadAllExperimentsForm	Binaries
https://bio.tools/genonets	http://eu-genonets.uzh.ch/python_package	Source code
https://bio.tools/gepia	http://gepia.cancer-pku.cn/api.html	API specification
https://bio.tools/germonline	http://www.germonline.org/gol_4_userguide.pdf	Source code
https://bio.tools/germonline	http://www.germonline.org/gol_4_userguide.pdf	Binaries
https://bio.tools/getorf	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/getorf	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/gff2aplot	http://genome.crg.es/software/gfftools/GFF2PLOT.html#DOWNLOADING	Source code
https://bio.tools/ggost	https://pypi.python.org/pypi/gprofiler-official	Binary package
https://bio.tools/ggost	https://toolshed.g2.bx.psu.edu/repository?repository_id=2d3d786121020d7a	Tool wrapper (galaxy)
https://bio.tools/ggost	https://cran.r-project.org/web/packages/gProfileR/index.html	Binary package
https://bio.tools/ggost	https://www.npmjs.com/package/biojs-vis-gprofiler	Binary package
https://bio.tools/giant_2.0	http://giant-v2.princeton.edu/download/	Biological data
https://bio.tools/giant_2.0	http://giant-v2.princeton.edu/data/	Biological data
https://bio.tools/gibbscluster	http://www.cbs.dtu.dk/cgi-bin/sw_request?gibbscluster	Source package
https://bio.tools/giggle	https://github.com/ryanlayer/giggle	Binary package
https://bio.tools/goa_gubbar	http://www.ms-utills.org/goa_gubbar/goa_gubbar.c	Source code
https://bio.tools/godef	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/godef	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/goname	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/goname	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/goprofiles	http://bioconductor.org/packages/release/bioc/vignettes/goProfiles/inst/doc/goProfiles.R	Source code
https://bio.tools/gorth	https://pypi.python.org/pypi/gprofiler-official	Binary package
https://bio.tools/gorth	https://cran.r-project.org/web/packages/gProfileR/index.html	Binary package
https://bio.tools/gorth	https://www.npmjs.com/package/biojs-vis-gprofiler	Binary package
https://bio.tools/gorth	https://toolshed.g2.bx.psu.edu/repository?repository_id=2d3d786121020d7a	Tool wrapper (galaxy)
https://bio.tools/gosemsim	http://bioconductor.org/packages/release/bioc/vignettes/GOSemSim/inst/doc/GOSemSim.R	Source code
https://bio.tools/gprofile_r	https://pypi.python.org/pypi/gprofiler-official	Binary package
https://bio.tools/gprofile_r	https://CRAN.R-project.org/package=gProfileR	Binary package
https://bio.tools/gprofile_r	https://www.npmjs.com/package/biojs-vis-gprofiler	Binary package
https://bio.tools/gprofiler	https://toolshed.g2.bx.psu.edu/repository?repository_id=2d3d786121020d7a	Tool wrapper (galaxy)
https://bio.tools/gprofiler	https://pypi.python.org/pypi/gprofiler-official	Binary package
https://bio.tools/gpufit	https://toolshed.g2.bx.psu.edu/repository?repository_id=2d3d786121020d7a	Tool wrapper (galaxy)
https://bio.tools/gqlstats	https://github.com/gpufit/Gpufit	Binaries
https://bio.tools/grape_2.0	http://bioconductor.org/packages/release/bioc/vignettes/gQTLstats/inst/doc/gQTLstats.R	Source code
https://bio.tools/grape_2.0	https://github.com/grape-pipeline/grape	Source code
https://bio.tools/greatscanmultipatterns	https://absynth.issb.genopole.fr/Bioinformatics/tools/GREAT/help#docker_integrated	Source code
https://bio.tools/greatscanmultipatterns	https://absynth.issb.genopole.fr/Bioinformatics/tools/GREAT/help#docker_integrated	Binaries
https://bio.tools/greatscanpatterns	https://absynth.issb.genopole.fr/Bioinformatics/tools/GREAT/help#docker_patterns	Source code
https://bio.tools/greatscanpatterns	https://absynth.issb.genopole.fr/Bioinformatics/tools/GREAT/help#docker_patterns	Binaries
https://bio.tools/groc-svs	https://github.com/grocsvs/grocsvs	Command-line specification
https://bio.tools/gromacs	http://www.gromacs.org/Downloads	Source package
https://bio.tools/gsean	http://bioconductor.org/packages/release/bioc/vignettes/gsean/inst/doc/gsean.R	Source code
https://bio.tools/guides	https://github.com/sanjanalab/GUIDES	Binaries
https://bio.tools/guugle	https://bibiserv.cebitec.uni-bielefeld.de/guugle?id=guugle_view_download	Binaries
https://bio.tools/gwaspi	http://www.gwaspi.org/?page_id=182	Source code
https://bio.tools/gwaspi	http://www.gwaspi.org/?page_id=182	Binaries
https://bio.tools/gxa	https://www.ebi.ac.uk/gxa/download.html	Container file
https://bio.tools/hasappy	https://github.com/gdiminin/HasAPPy	Source code
https://bio.tools/helixturnhelix	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/helixturnhelix	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/hichipper	http://hichipper.readthedocs.io/en/latest/	Binary package
https://bio.tools/hichipper	http://hichipper.readthedocs.io/en/latest/	Source code
https://bio.tools/hicrep	http://bioconductor.org/packages/release/bioc/vignettes/hicrep/inst/doc/hicrep-vignette.R	Source code
https://bio.tools/hieranoid	http://hieranoid.sbc.su.se/download/H2/	Biological data
https://bio.tools/hieranoid	http://software.sbc.su.se/cgi-bin/request.cgi?project=hieranoid2	Source code
https://bio.tools/hilbertcurve	https://bioconductor.org/packages/release/bioc/html/EnrichedHeatmap.html	Source code
https://bio.tools/hipathia	http://bioconductor.org/packages/release/bioc/vignettes/hipathia/inst/doc/hipathia-vignette.R	Source code
https://bio.tools/hipstr	https://github.com/HipSTR-Tool/HipSTR	Source code
https://bio.tools/histocat	https://github.com/BodenmillerGroup/histoCAT/releases	Binaries
https://bio.tools/hmdbquery	http://bioconductor.org/packages/release/bioc/vignettes/hmdbQuery/inst/doc/hmdbQuery.R	Source code
https://bio.tools/hmoment	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/hmoment	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/htseqtools	http://bioconductor.org/packages/release/bioc/vignettes/htSeqTools/inst/doc/htSeqTools.R	Source code
https://bio.tools/hu_map	http://proteincomplexes.org/download	Biological data
https://bio.tools/hx-express	http://www.hxms.com/HXExpress/HX-Express%20Beta.xls	Source code
https://bio.tools/hyasp	http://biomaging.usc.edu/software.html#HYSF	Binary package
https://bio.tools/icnv	http://bioconductor.org/packages/release/bioc/vignettes/iCNV/inst/doc/iCNV-vignette.R	Source code
https://bio.tools/idconvert	http://proteowizard.sourceforge.net/downloads.shtml	Source code
https://bio.tools/idconvert	http://proteowizard.sourceforge.net/downloads.shtml	Binaries
https://bio.tools/idrem	https://github.com/phoenixding/idrem/tree/master/sourcecode/edu/cmu/cs/sb	Source code
https://bio.tools/iep	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/iep	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries

biotools_link	download_url	download_type
https://bio.tools/igdiscover	https://pypi.python.org/pypi/igdiscover	Source package
https://bio.tools/igvr	http://bioconductor.org/packages/3.7/bioc/src/contrib/Archive/igvr/	Source package
https://bio.tools/igvr	http://bioconductor.org/packages/release/bioc/vignettes/igvr/inst/doc/igvr.R	Source code
https://bio.tools/imas	http://bioconductor.org/packages/release/bioc/vignettes/IMAS/inst/doc/IMAS.pdf	Source code
https://bio.tools/impulse2	http://bioconductor.org/packages/release/bioc/vignettes/ImpulseDE2/inst/doc/ImpulseDE2_Tutorial.R	Source code
https://bio.tools/infoalign	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/infoalign	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/infoassembly	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/infoassembly	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/infobase	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/infobase	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/inforesidue	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/inforesidue	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/informed-proteomics	https://github.com/PNNL-Comp-Mass-Spec/Informed-Proteomics	Binaries
https://bio.tools/infoseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/infoseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/inserm-remap	http://tagc.univ-mrs.fr/remap/index.php?page=download	Source code
https://bio.tools/inserm-remap	http://tagc.univ-mrs.fr/remap/index.php?page=download	Binaries
https://bio.tools/intad	http://bioconductor.org/packages/release/bioc/vignettes/IntAD/inst/doc/IntAD.R	Source code
https://bio.tools/interactome_insider	http://interactomeinsider.vulab.org/downloads.html	Biological data
https://bio.tools/intermine	http://github.com/intermine/	Source code
https://bio.tools/intermine	http://github.com/intermine/	Binaries
https://bio.tools/intervene	https://bitbucket.org/CBGR/intervene/downloads/	Source code
https://bio.tools/ioniser	https://bioconductor.org/packages/release/bioc/html/IONiseR.html	Source code
https://bio.tools/ipsa	https://github.com/pervouchine/ipsa-full	Source code
https://bio.tools/irbis	https://github.com/pervouchine/irbis/	Source code
https://bio.tools/irrefr	https://cran.r-project.org/web/packages/IRefR/index.html	Source code
https://bio.tools/irfescape	http://irfindex.org/wiki/index.php?title=IRfEscape	Source code
https://bio.tools/isafe	https://github.com/alek0991/SAFE	Command-line specification
https://bio.tools/isee	http://bioconductor.org/packages/3.7/bioc/src/contrib/Archive/isee/	Source package
https://bio.tools/isma	https://sourceforge.net/projects/isma/files/latest/download	Command-line specification
https://bio.tools/isochore	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/isochore	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/isotop	http://www.ms-utilis.org/isotop.html	Source code
https://bio.tools/isotop_fs	http://www.ms-utilis.org/isotop_fs.html	Source code
https://bio.tools/isotope.py	http://www.ms-utilis.org/isotope.py	Source code
https://bio.tools/iterclust	http://bioconductor.org/packages/release/bioc/src/contrib/iterClust_1.2.0.tar.gz	Source package
https://bio.tools/jali	http://bibiserv.cebitec.uni-bielefeld.de/jali?id=jali_view_download	Source code
https://bio.tools/jali	http://bibiserv.cebitec.uni-bielefeld.de/resources/download/jali	Binaries
https://bio.tools/jasextract	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/jasextract	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/jaspscan	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/jaspscan	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/jmztab-m	https://github.com/lifs-tools/jmzTab-m/releases	Binary package
https://bio.tools/jmztab-m	https://github.com/lifs-tools/jmzTab-m/releases	Source package
https://bio.tools/jorca	http://bitlab-es.com/jorca/documents/JORCA-full.pdf	Source code
https://bio.tools/jphyloio	http://bioinfweb.info/JPhyloIO/Download	Binary package
https://bio.tools/jphyloio	http://bioinfweb.info/JPhyloIO/Download	Source package
https://bio.tools/jphyloio	http://bioinfweb.info/JPhyloIO/Download	API specification
https://bio.tools/kaiju	https://github.com/bioinformatics-centre/kaiju	Source code
https://bio.tools/khmer	https://github.com/dib-lab/khmer/releases	Source code
https://bio.tools/khmer	https://pypi.python.org/pypi/khmer	Binaries
https://bio.tools/knime	https://www.knime.org/knime-analytics-platform	Binaries
https://bio.tools/kplogo	http://kplogo.wi.mit.edu/manual.html?#install-kpl-ogo-locally	Source code
https://bio.tools/kraken	https://ccb.jhu.edu/software/kraken/	Binaries
https://bio.tools/kwip	https://github.com/kdmurray91/kWIP	Source code
https://bio.tools/la-images	https://github.com/sing-group/la-images	Source code
https://bio.tools/la-images	http://sing.ei.uvigo.es/la-images/download.html	Binaries
https://bio.tools/lailabs-qsm	https://bitbucket.org/jpk_bit_team/bioescorte-suggestion/src	Source code
https://bio.tools/ida	http://genome.tugraz.at/ida2/ida_download.shtml	Command-line specification
https://bio.tools/libfid	https://bibiserv.cebitec.uni-bielefeld.de/libfid?id=libfid_download	Binaries
https://bio.tools/libralign	http://bioinfweb.info/LibrAlign/Download	Binaries
https://bio.tools/libralign	http://bioinfweb.info/LibrAlign/Download	Source code
https://bio.tools/libralign	http://bioinfweb.info/LibrAlign/Download	API specification
https://bio.tools/limmarp	https://bitbucket.org/veilveil/limmarp	Source code
https://bio.tools/lindna	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/lindna	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/listor	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/listor	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/litemol	https://github.com/dsehna/LiteMol	Source code
https://bio.tools/livemore_metagenomics_analysis_toolkit	https://sourceforge.net/projects/lmat/	Source code
https://bio.tools/livemore_metagenomics_analysis_toolkit	https://sourceforge.net/projects/lmat/	Binaries
https://bio.tools/ljn	https://git.facom.ufms.br/bioinfo/LNJ/tree/master/sources	Source code
https://bio.tools/loctree3	https://roslab.org/owiki/index.php/Packages	Source code
https://bio.tools/loctree3	https://roslab.org/owiki/index.php/Packages	Binaries
https://bio.tools/lulefisk	http://sourceforge.net/projects/lulefiskxp	Source code
https://bio.tools/m-jolite	http://miolite2.iceht.forth.gr	Container file
https://bio.tools/macaron	https://github.com/wagasuddinkhan/MACARON-GenMed-LabEx.git	Source code
https://bio.tools/macsfinder	https://github.com/gem-pasteur/macsfinder	Source code
https://bio.tools/mageri	https://github.com/mikeshh/mageri	Source code
https://bio.tools/magpie	https://github.com/christibald/magpie	Source code
https://bio.tools/make_random	http://www.ms-utilis.org/make_random.c	Source code
https://bio.tools/makenucseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/makenucseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/makeprotseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/makeprotseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/mambo	https://github.com/danielriosgarza/MAMBO	Binaries
https://bio.tools/mapi	http://bitlab-es.com/mapi/documents/systemcalls.pdf	Source code

biotools_link	download_url	download_type
https://bio.tools/mapi	http://bitlab-es.com/mapi/repository	Binaries
https://bio.tools/mapscape	https://bioconductor.org/packages/release/bioc/vignettes/mapscape/inst/doc/mapscape_vignette.html	Command-line specification
https://bio.tools/marcat	https://s1.sfb.uit.no/public/mar/	Biological data
https://bio.tools/mardb	https://s1.sfb.uit.no/public/mar/	Biological data
https://bio.tools/marref	https://s1.sfb.uit.no/public/mar/	Biological data
https://bio.tools/marscan	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/marscan	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/maskambignuc	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/maskambignuc	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/maskambigprot	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/maskambigprot	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/maskfeat	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/maskfeat	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/maskseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/maskseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/mass-up	http://www.sing-group.org/mass-up/download	Source code
https://bio.tools/mass-up	http://www.sing-group.org/mass-up/download	Binaries
https://bio.tools/matcher	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/matcher	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/matrix-clustering	http://rsat.eu/	Source package
https://bio.tools/mcsm_na	http://biosig.unimelb.edu.au/mcsm_na/data	Biological data
https://bio.tools/mdanalysis	https://github.com/MDAnalysis/mdanalysis	Source package
https://bio.tools/mdanalysis	http://pypi.python.org/pypi/MDAnalysis	Source package
https://bio.tools/medecom	https://github.com/lutsik/MeDeCom	Source code
https://bio.tools/megamerger	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/megamerger	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/meneco	https://codeload.github.com/bioaspi/meneco/zip/meneco-1.5.2	Source code
https://bio.tools/mentalist	https://github.com/WGS-TB/MentalIST	Source code
https://bio.tools/mentalist	https://anaconda.org/bioconda/mentalist	Binary package
https://bio.tools/merger	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/merger	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/meta	http://genome.crg.es/software/meta/index.html#code	Source code
https://bio.tools/meta-gecko	http://bitlab-es.com/gecko/documents/METAGECKO-GuidedExercise.pdf	Source code
https://bio.tools/metaboanalyst	http://www.metaboanalyst.ca/faces/docs/Resources.xhtml	Source code
https://bio.tools/metacoder	https://github.com/grunwaldt/metacoder	Source code
https://bio.tools/metadyn_view	http://metadyn.vsch.cz	Source code
https://bio.tools/metadyn_view	http://metadyn.vsch.cz	Binaries
https://bio.tools/metagxovarian	http://bioconductor.org/packages/release/bioc/vignettes/MetaGxOvarian/inst/doc/MetaGxOvarian.R	Source code
https://bio.tools/metaphors	http://betaorthology.phylomedb.org/download	Biological data
https://bio.tools/metaproteomeanalyzer	https://code.google.com/p/meta-proteome-analyzer/source/checkout	Source code
https://bio.tools/metarep	https://github.com/jcvi/METAREP/releases/tag/1.4.0-beta	Source code
https://bio.tools/metaseq	https://www.bioconductor.org/packages/release/bioc/html/metaseqR.html	Source code
https://bio.tools/micca	https://github.com/compmetagen/micca/releases	Source code
https://bio.tools/micropan	https://cran.r-project.org/web/packages/micropan/index.html	Source code
https://bio.tools/miga-taxa	https://github.com/bio-miga/miga.git	API specification
https://bio.tools/miga-taxa	https://github.com/bio-miga/miga.git	Command-line specification
https://bio.tools/mindboggle	http://mindboggle.readthedocs.io/en/latest/	Binaries
https://bio.tools/milotip	https://github.com/sonneysa/MiloTIP	Source code
https://bio.tools/mixomics	https://cran.r-project.org/web/packages/mixOmics/	Binaries
https://bio.tools/mkesa	https://biserv.cebitec.uni-bielefeld.de/mkesa?id=mkesa_download	Binaries
https://bio.tools/mlibrary	http://www.sing-group.org/MLibrary/index.php?option=com_content&view=section&layout=blog&id=8&Itemid=58	Binaries
https://bio.tools/mmass	http://www.mmass.org/download/	Source code
https://bio.tools/mmeta	http://genome.crg.es/software/mmeta/index.html#code	Source code
https://bio.tools/mmp	https://s1.sfb.uit.no/public/mar/	Biological data
https://bio.tools/mobidb-lite	http://protein.bio.unipd.it/download/	Binary package
https://bio.tools/mocapy_plus_plus	http://sourceforge.net/projects/mocapy/files/latest/download?source=files	Source code
https://bio.tools/mocapy_plus_plus	http://sourceforge.net/projects/mocapy/files/latest/download?source=files	Binaries
https://bio.tools/mole	https://lcc.ncbr.muni.cz/whitezone/development/mole/web/download.php	Binaries
https://bio.tools/mole_pymol_plugin	http://webchem.ncbr.muni.cz/Platform/AppsBin/Mole/2.13.9.6/Mole2_manual.pdf	Source code
https://bio.tools/mole_pymol_plugin	http://mole.chemi.muni.cz	Binaries
https://bio.tools/molecular_weight_calculator	http://ncr.pnl.gov/software/software_register.asp?id=2	Source code
https://bio.tools/monocle	https://github.com/cole-trapnell-lab/monocle-release	Binary package
https://bio.tools/morpheus	http://www.chem.wisc.edu/~coon/Downloads/Morpheus/	Biological data
https://bio.tools/motiflab	http://www.motiflab.org/index.php?page=download	Binaries
https://bio.tools/motincor2	https://www.nature.com/articles/nmeth.4193#s1	Source code
https://bio.tools/motivevalidator	http://webchem.ncbr.muni.cz/Platform/MotiveValidator/DownloadService	Binaries
https://bio.tools/motus	http://www.bork.embl.de/software/mOTU/download.html	Source code
https://bio.tools/motus	http://www.bork.embl.de/software/mOTU/download.html	Binaries
https://bio.tools/mowsew	http://chirimoyo.ac.uma.es/gnv5/documents/homology_manual.pdf	Source code
https://bio.tools/mowsew	http://chirimoyo.ac.uma.es/gnv5/documents/homology_manual.pdf	Binaries
https://bio.tools/mrbayes	https://github.com/NBISweden/MrBayes/releases/tag/v3.2.6	Source code
https://bio.tools/mrbayes	https://github.com/NBISweden/MrBayes/releases/tag/v3.2.6	Binaries
https://bio.tools/msa	https://github.com/wilzbach/msa	Source code
https://bio.tools/msa	https://cdn.bio.sh/	Binaries
https://bio.tools/msaccess	http://proteowizard.sourceforge.net/downloads.shtml	Source code
https://bio.tools/msalign	http://www.ms-utils.org/msalign/msalign.c	Source code
https://bio.tools/msalign2	http://www.ms-utils.org/msalign/msalign.c	Source code
https://bio.tools/msviewer	https://github.com/wilzbach/msa	Source code
https://bio.tools/msviewer	https://cdn.bio.sh/	Binaries
https://bio.tools/msbar	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/msbar	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/msfilerader_python_bindings	https://github.com/frallain/MSFileReader_Python_bindings/blob/master/MSFileReader.py	Source code
https://bio.tools/msfragger	http://www.nesvilab.org/software	Source code
https://bio.tools/msgfdb2pepxml	http://ms-utils.org/msgfdb2pepxml/msgfdb2pepxml.py	Source code
https://bio.tools/mspolygraph	http://omics.pnl.gov/software/MSPolygraph.php	Source code
https://bio.tools/msqrob	https://github.com/ludgergoemmel/MSqRob/	Source package
https://bio.tools/msstats	https://github.com/MeenaChoi/MSstats	Source code
https://bio.tools/msviz	https://github.com/vitalit-sib/msviz-docker	Container file

biotools_link	download_url	download_type
https://bio.tools/ontotext	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/ontotext	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/open_phacts_api	http://data.openphacts.org/	Source code
https://bio.tools/open_phacts_api	https://github.com/openphacts/ops-docker	Container file
https://bio.tools/open_phacts_explorer	http://github.com/openphacts/explorer2	Source code
https://bio.tools/openmm	https://github.com/pandegroup/openmm	Source code
https://bio.tools/openms	https://github.com/OpenMS/OpenMS	Source code
https://bio.tools/openms	http://sourceforge.net/projects/open-ms/files/OpenMS/OpenMS-2.0/	Binaries
https://bio.tools/openprimer	http://bioconductor.org/packages/release/bioc/vignettes/openPrimerR/inst/doc/openPrimerR_vignette.R	Source code
https://bio.tools/orange	https://orange.biolab.si/download/	Source code
https://bio.tools/orange_network	http://orange.biolab.si/download/	Source code
https://bio.tools/organism_dplyr	http://bioconductor.org/packages/release/bioc/vignettes/Organism.dplyr/inst/doc/Organism.dplyr.R	Source code
https://bio.tools/organisms	https://organisms.jensenlab.org/Downloads	Biological data
https://bio.tools/oswitch	https://github.com/wurmlab/oswitch	Source code
https://bio.tools/oswitch	https://github.com/Homebrew/homebrew-science/blob/master/oswitch.rb	Source package
https://bio.tools/oswitch	https://launchpad.net/~nebc/+archive/ubuntu/bio-linux	Binary package
https://bio.tools/out2summary	http://sashimi.cvs.sourceforge.net/sashimi/out2summary	Source code
https://bio.tools/overlap	http://genome.crg.cat/software/overlap/64bits/overlap	Binaries
https://bio.tools/palindrome	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/palindrome	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/pandrugs	https://github.com/sing-group/pandrugs-docker	Container file
https://bio.tools/pasteseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/pasteseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/patmatdb	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/patmatdb	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/patmatmotifs	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/patmatmotifs	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/patternlab	http://pcarvalho.com/patternlab/downloads/windows/patternlab/	Source code
https://bio.tools/patternquery	http://webchem.ncbr.muni.cz/Platform/PatternQuery/DownloadService	Binaries
https://bio.tools/pcasuite	http://mmb.pcb.ub.es/software/pcasuite/	Binaries
https://bio.tools/pcsf	https://github.com/IOR-Bioinformatics/PCSF	Binaries
https://bio.tools/pdb_redo	http://www.cmbi.ru.nl/pdb_redo/downloads.html	Source code
https://bio.tools/pdb_redo	http://www.cmbi.ru.nl/pdb_redo/downloads.html	Binaries
https://bio.tools/peakzilla	https://github.com/steinmann/peakzilla	Source code
https://bio.tools/pecan_peptide-centric_analysis	https://bitbucket.org/maccoss/pecan/src	Source code
https://bio.tools/peffect	https://roslab.org/owiki/index.php/Packages	Source code
https://bio.tools/peffect	https://roslab.org/owiki/index.php/Packages	Binaries
https://bio.tools/pepcoll	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/pepcoll	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/pepdigest	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/pepdigest	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/pepinfo	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/pepinfo	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/pepnet	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/pepnet	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/pepstats	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/pepstats	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/pepwheel	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/pepwheel	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/pepwindow	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/pepwindow	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/pepwindowall	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/pepwindowall	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/pgp	http://edal.ipk-gatersleben.de/download.html	Source code
https://bio.tools/pgp	http://edal.ipk-gatersleben.de/download.html	Binaries
https://bio.tools/phmmer	http://hmmer.org/download.html	Source code
https://bio.tools/pia	https://github.com/mpc-bioinformatics/pia/releases	Binaries
https://bio.tools/pickle	http://www.pickle.gr/Downloads	Biological data
https://bio.tools/pickle	http://www.pickle.gr/Downloads	Ontology
https://bio.tools/picky	https://github.com/TheJacksonLaboratory/Picky	Binary package
https://bio.tools/piped	https://github.com/jmeyerucsd/PIQEDia/releases/tag/v0.1.2	Source code
https://bio.tools/piqmie	https://github.com/arnikz/PIQMie	Source code
https://bio.tools/plexy	http://www.bioinf.uni-leipzig.de/Software/PLEXY/plexy.pl	Binaries
https://bio.tools/plotcon	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/plotcon	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/plotorf	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/plotorf	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/plyranges	http://bioconductor.org/packages/3.7/bioc/src/contrib/Archive/plyranges/	Source package
https://bio.tools/pmx	https://github.com/dseeliger/pmx	Source package
https://bio.tools/polydot	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/polydot	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/poppunk	https://github.com/johnlees/PopPUNK	Source code
https://bio.tools/possumsearch	https://bibiserv.cebitec.uni-bielefeld.de/possumsearch?id=possumsearch_download	Source code
https://bio.tools/possumsearch	https://bibiserv.cebitec.uni-bielefeld.de/resources/download/possumsearch/	Binaries
https://bio.tools/possumsearch2	https://bibiserv.cebitec.uni-bielefeld.de/possumsearch2?id=possumsearch_download	Source code
https://bio.tools/possumsearch2	http://bibiserv.cebitec.uni-bielefeld.de/resources/download/possumsearch	Binaries
https://bio.tools/powerfit	https://github.com/haddocking/powerfit	Source code
https://bio.tools/ppinfer	http://bioconductor.org/packages/release/bioc/vignettes/PPInfer/inst/doc/PPInfer.R	Source code
https://bio.tools/precog	http://precog.lundberg.gu.se/PrecogInstall/	Binaries
https://bio.tools/predictnls	https://roslab.org/owiki/index.php/Packages	Source code
https://bio.tools/predictnls	https://roslab.org/owiki/index.php/Packages	Binaries
https://bio.tools/predictsnp	http://il47.sci.muni.cz:6212/predictsnp/USER_GUIDE.pdf	Source code
https://bio.tools/predictsnp	http://il47.sci.muni.cz:6212/predictsnp/USER_GUIDE.pdf	Binaries
https://bio.tools/predictsnp2	http://loschmidt.chemi.muni.cz/predictsnp2/docs/userguide.pdf	Source code
https://bio.tools/predictsnp2	http://loschmidt.chemi.muni.cz/predictsnp2/docs/userguide.pdf	Binaries
https://bio.tools/predrem	http://server.cs.ucf.edu/predrem/download.jsp	Biological data
https://bio.tools/preg	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/preg	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries

biotools_link	download_url	download_type
https://bio.tools/prettyplot	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/prettyplot	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/prettyseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/prettyseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/price-orf	https://www.nature.com/articles/nmeth.4631#methods	Source code
https://bio.tools/price-orf	https://github.com/erhard-lab/gedi	Source code
https://bio.tools/primersearch	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/primersearch	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/printextract	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/printextract	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/procon	http://www.ruhr-uni-bochum.de/mpc/software/ProCon/index.html.en	Source code
https://bio.tools/procon	http://www.ruhr-uni-bochum.de/mpc/software/ProCon/index.html.en	Binaries
https://bio.tools/prodisis	https://roslab.org/owiki/index.php/Packages	Source code
https://bio.tools/prodisis	https://roslab.org/owiki/index.php/Packages	Binaries
https://bio.tools/profsis	https://roslab.org/owiki/index.php/Packages	Source code
https://bio.tools/profsis	https://roslab.org/owiki/index.php/Packages	Binaries
https://bio.tools/profit	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/profit	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/proftmb	https://roslab.org/owiki/index.php/Packages	Source code
https://bio.tools/proftmb	https://roslab.org/owiki/index.php/Packages	Binaries
https://bio.tools/project	http://genome.crg.cat/software/project/64bits/project	Binaries
https://bio.tools/proline	http://proline.profpoteomics.fr/download/	Binaries
https://bio.tools/proline	http://proline.profpoteomics.fr/download/	Test data
https://bio.tools/prophet	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/prophet	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/props	http://bioconductor.org/packages/release/bioc/vignettes/PROPS/inst/doc/props.R	Source code
https://bio.tools/prosextract	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/prosextract	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/proteinhistorian	http://ighthouse.ucsf.edu/ProteinHistorian/downloads.html	Source code
https://bio.tools/proteinhistorian	http://ighthouse.ucsf.edu/ProteinHistorian/downloads.html	Biological data
https://bio.tools/proteome-pi	http://soelectricpointdb.org/download.html	Biological data
https://bio.tools/proteomedigestsim	http://www.ncbi.nlm.nih.gov/pubmed/22324799	Source code
https://bio.tools/proteometools	http://www.proteometools.org/index.php?id=51	Biological data
https://bio.tools/proteosafe	https://web.archive.org/web/20070706084404/http://peptide.ucsd.edu/ftprepos.html	Source code
https://bio.tools/proxli	https://github.com/yeastro/proxli-web-app	Source package
https://bio.tools/pscan	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/pscan	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/psichomics	https://bioconductor.org/packages/psichomics	Source package
https://bio.tools/psichomics	https://github.com/nuno-agostinho/psichomics	Source code
https://bio.tools/psiphi	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/psiphi	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/psaslib	http://mosaic.mpi-cbg.de/?q=downloads&stochastic_chemical_net	Source code
https://bio.tools/pview	http://compbio.cs.princeton.edu/pview/	Source code
https://bio.tools/pydock	https://life.bsc.es/pid/pydock/get_pydock.html	Binaries
https://bio.tools/pyldm	https://github.com/gadorhjac/PyLDM	Source code
https://bio.tools/pymdsetup	https://www.dropbox.com/s/uxgcm4rkm554d28/PyMDSetup-v2.qcow?dl=1	VM image
https://bio.tools/pymdsetup	https://github.com/bioexcel/pymdsetup	Source code
https://bio.tools/pymdsetup	http://mmb.irbbarcelona.org/~andrio/PyMDSetup-v2.qcow2	VM image
https://bio.tools/pyms	http://code.google.com/p/pyms/	Source code
https://bio.tools/q4sm	https://github.com/PNNL-Comp-Mass-Spec/Q4SRM	Source package
https://bio.tools/qctool	http://www.well.ox.ac.uk/~gav/qctool/documentation/download.html	Binary package
https://bio.tools/qmmm	http://www.ks.uiuc.edu/Research/qmmm/	Binary package
https://bio.tools/qsbio	http://www.weizmann.ac.il/Structural_Biology/faculty_pages/ELevy/downloads/QSbio.html	Biological data
https://bio.tools/quant	http://sourceforge.net/project/showfiles.php?group_id=109078&package_id=236902	Source code
https://bio.tools/raven_toolbox	https://github.com/SysBioChalmers/RAVEN/releases	Binaries
https://bio.tools/raven_toolbox	https://github.com/SysBioChalmers/RAVEN	Source code
https://bio.tools/raw2mzdb	https://github.com/mzdb/pwiz-mzdb/releases	Binaries
https://bio.tools/raw2mzdb	https://github.com/mzdb/pwiz-mzdb	Source code
https://bio.tools/rawtools	https://github.com/kevinkovalchik/RawTools	Source code
https://bio.tools/ray	http://sebhml.github.io/Ray.web/download.html	Source code
https://bio.tools/ray	http://sebhml.github.io/Ray.web/download.html	Binaries
https://bio.tools/rcy3	http://bioconductor.org/packages/3.7/bioc/src/contrib/Archive/RCy3/	Source package
https://bio.tools/reactome_graph_database	https://reactome.org/dev/graph-database	Binary package
https://bio.tools/readtools	https://github.com/magicDGS/ReadTools/releases	Binaries
https://bio.tools/readxplorer	http://readxplorer.org	Binaries
https://bio.tools/rebaseextract	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/rebaseextract	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/recal2	http://www.ms-utls.org/recalibrate_using_MSMS.c	Source code
https://bio.tools/recoder	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/recoder	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/redata	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/redata	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/refseqget	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/refseqget	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/relna	https://github.com/ashishbaghudana/mthesis-ashish	Source code
https://bio.tools/relna	https://github.com/ashishbaghudana/mthesis-ashish	Binaries
https://bio.tools/remap	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/remap	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/remaps	https://confluence.csc.fi/display/REMS/Installation	Source code
https://bio.tools/repeat_explorer	https://bitbucket.org/petrmovak/replex_jarean	Source code
https://bio.tools/repeat_explorer	https://bitbucket.org/petrmovak/replex_jarean	Binaries
https://bio.tools/reputer	http://bibiserv.cibitec.uni-bielefeld.de/reputer?id=reputer_view_download	Source code
https://bio.tools/reputer	http://bibiserv.cibitec.uni-bielefeld.de/reputer?id=reputer_view_download	Binaries
https://bio.tools/restover	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/restover	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/restrict	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/restrict	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/reverseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code

biotools_link	download_url	download_type
https://bio.tools/revseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/rfam	ftp://ftp.ebi.ac.uk/pub/databases/Rfam/CURRENT	Biological data
https://bio.tools/rightfield	http://www.rightfield.org.uk/download	Source code
https://bio.tools/rightfield	http://www.rightfield.org.uk/download	Binaries
https://bio.tools/rnacentral	ftp://ftp.ebi.ac.uk/pub/databases/RNACentral/current_release	Biological data
https://bio.tools/rnadecay	http://bioconductor.org/packages/3.7/bioc/src/contrib/Archive/RNAdelay/	Source package
https://bio.tools/rmahybrid	http://bibiserv.cebitec.uni-bielefeld.de/mahybrid?id=mahybrid_view_download	Source code
https://bio.tools/rmahybrid	http://bibiserv.cebitec.uni-bielefeld.de/resources/download/mahybrid	Binaries
https://bio.tools/rmbeads	http://bioconductor.org/packages/release/bioc/html/RnBeads.html	Binaries
https://bio.tools/rmbeads	https://quay.io/repository/biocontainers/bioconductor-rmbeads	Container file
https://bio.tools/rosettaes	https://www.rosettacommons.org/docs/latest/release-notes	API specification
https://bio.tools/rsd	https://github.com/todddeluca/reciprocal_smallest_distance	Source code
https://bio.tools/rt	http://www.ms-utlis.org/rt4.c	Source code
https://bio.tools/rtxi	http://rtxi.org/install/	Binaries
https://bio.tools/rtxi	https://github.com/rtxi	Source code
https://bio.tools/rxnorm_api	https://www.nlm.nih.gov/research/umls/rxnorm/docs/rxnormfiles.html	Biological data
https://bio.tools/s2p	http://sing-group.org/s2p/download.html	Binaries
https://bio.tools/s2p	https://github.com/sing-group/s2p	Source code
https://bio.tools/salmon	https://github.com/COMBINE-lab/salmon	Source code
https://bio.tools/salmonella_crispr_typing	https://github.com/C3BI-pasteur-fr/Salmonella-CRISPR-Typing/tree/master/galaxy	Tool wrapper (galaxy)
https://bio.tools/sampleclassifier	http://bioconductor.org/packages/release/bioc/vignettes/sampleClassifier/inst/doc/sampleClassifier.R	Source code
https://bio.tools/sapp	http://download.systemsbio.org/sapp/	Binaries
https://bio.tools/sarek	https://github.com/SciLifeLab/Sarek/releases	Source code
https://bio.tools/scale4c	http://bioconductor.org/packages/release/bioc/vignettes/Scale4C/inst/doc/vignette.R	Source code
https://bio.tools/scaller	https://github.com/biosinodx/SCcaller/	Source code
https://bio.tools/scenic	https://github.com/aertslab/SCENIC	Binaries
https://bio.tools/scipion	https://github.com/I2PC/scipion/wiki/How-to-Install	Source code
https://bio.tools/scipion	https://github.com/I2PC/scipion/wiki/How-to-Install	Binaries
https://bio.tools/scmap	http://bioconductor.org/packages/release/bioc/html/scmap.html	Source package
https://bio.tools/scnorm	https://github.com/rhondabacher/SCnorm	Source code
https://bio.tools/scope-scrna	https://github.com/aertslab/SCope	Source code
https://bio.tools/scry	https://pypi.python.org/pypi/scry	Source code
https://bio.tools/seda	http://www.sing-group.org/seda/download.html	Binaries
https://bio.tools/seealso	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/seealso	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/seek_for_science	http://docs.seek4science.org/tech/install.html	Source code
https://bio.tools/segway	https://bitbucket.org/hoffmanlab/segway/src	Source code
https://bio.tools/selenoprofiles3	http://genome.crg.es/~mmaniotti/selenoprofiles_3_installation.tar	Source code
https://bio.tools/selenoprofiles3	http://genome.crg.es/~mmaniotti/selenoprofiles_manual.pdf	Binaries
https://bio.tools/seqan	http://packages.seqan.de/	Source code
https://bio.tools/seqan	http://packages.seqan.de/	Binaries
https://bio.tools/seqbench	http://www.icbi.at/svnseqbench/seqbench/trunk/	Source code
https://bio.tools/seqcluster	http://github.com/lpantano/seqcluster#cite	Source code
https://bio.tools/seqcluster	http://github.com/bioconda	Binaries
https://bio.tools/seqcount	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/seqcount	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/seqmatchall	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/seqmatchall	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/seqret	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/seqret	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/seqretsetall	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/seqretsetall	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/seqretsplit	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/seqretsplit	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/sequenceserver	https://github.com/wurmlab/sequenceserver	Source code
https://bio.tools/sequenceserver	https://github.com/wurmlab/sequenceserver/blob/master/docker/Dockerfile	Container file
https://bio.tools/seqxref	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/seqxref	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/seqxrefget	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/seqxrefget	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/serverell	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/serverell	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/showalign	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/showalign	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/showdb	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/showdb	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/showfeat	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/showfeat	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/showorf	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/showorf	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/showpep	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/showpep	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/showseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/showseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/showserver	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/showserver	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/shuffleseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/shuffleseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/sider	http://sideeffects.embl.de/download/	Biological data
https://bio.tools/sigcleave	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/sigcleave	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/signalalign	https://github.com/ArtRand/signalAlign	Source code
https://bio.tools/signalp	http://www.cbs.dtu.dk/cgi-bin/sw_request?signalp	Source code
https://bio.tools/signalp	http://www.cbs.dtu.dk/cgi-bin/sw_request?signalp	Binaries
https://bio.tools/silent	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/silent	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/silva	https://www.arb-silva.de/download/archive/	Source code
https://bio.tools/silva	https://www.arb-silva.de/download/archive/	Binaries

biotools_link	download_url	download_type
https://bio.tools/simka	https://gatb.inria.fr/software/simka/	Source code
https://bio.tools/sina	https://github.com/epruesse/SINA/releases	Source code
https://bio.tools/sima	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/sima	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/sixpack	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/sixpack	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/sizeseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/sizeseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/sjcount	https://github.com/pervouchine/sjcount-full	Source code
https://bio.tools/sjcount	https://github.com/pervouchine/sjcount-full	Binaries
https://bio.tools/skipredundant	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/skipredundant	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/skipseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/skipseq	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/skylive	http://proteowizard.svn.sourceforge.net/viewvc/proteowizard/trunk/pwiz/pwiz_tools	Source code
https://bio.tools/sleuth	https://pachterlab.github.io/sleuth/download	Binaries
https://bio.tools/slimsuite	http://slimsuite.blogspot.com/p/downloads.html	Source code
https://bio.tools/snap2	https://github.com/Rostlab/SNAP2	Source code
https://bio.tools/snap2	https://github.com/Rostlab/SNAP2	Binaries
https://bio.tools/sniffles	https://github.com/fritzsedlazeck/Sniffles	Binary package
https://bio.tools/snooper	https://sourceforge.net/projects/snooper/files/	Source code
https://bio.tools/snp2fbs	ftp://ccg.vital-it.ch/snp2fbs/	Biological data
https://bio.tools/snpitty	https://bitbucket.org/ccbc/snpitty	Source code
https://bio.tools/snpitty	https://hub.docker.com/r/ccbc/	Binaries
https://bio.tools/snplice	https://code.google.com/p/snplice/	Binary package
https://bio.tools/sourcer	https://cran.r-project.org/web/packages/sourceR/index.html	Source code
https://bio.tools/spartaabc	http://spartaabc.tau.ac.il/webserver/source.php	Source code
https://bio.tools/spatialde	https://github.com/Teichlab/SpatialDE	Command-line specification
https://bio.tools/speaq	https://cran.r-project.org/web/packages/speaq/index.html	Binaries
https://bio.tools/speaq	https://github.com/beirmaert/speaq	Source code
https://bio.tools/speci	http://vm-lux.embl.de/~mende/specl/downloads.html	Source code
https://bio.tools/specter	https://github.com/rpeckner-broad/Specter	Binary package
https://bio.tools/splitsource	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/splitsource	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/splitter	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/splitter	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/sqlseeker	https://github.com/jmonlong/sQTLseeker	Source code
https://bio.tools/sqlseeker	https://github.com/jmonlong/sQTLseeker	Binaries
https://bio.tools/ssaha2	ftp://ftp.sanger.ac.uk/pub/resources/software/ssaha2/	Source package
https://bio.tools/stretcher	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/stretcher	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/stringapp	http://apps.cytoscape.org/apps/stringapp	Binaries
https://bio.tools/structure_threader	https://github.com/StunTSPT/Structure_threader/releases/latest	Source package
https://bio.tools/stssearch	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/stssearch	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/supermatcher	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/supermatcher	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/superquantnode	https://github.com/caetera/SuperQuantNode	Binaries
https://bio.tools/sv-callers	https://github.com/GooglingTheCancerGenome/sv-callers	Tool wrapper (other)
https://bio.tools/swicz	http://proteom.biomed.cas.cz	Source code
https://bio.tools/swicz	http://proteom.biomed.cas.cz	Binaries
https://bio.tools/swift	https://bibiserv.cebitec.uni-bielefeld.de/swift?id=swift_view_download	Binaries
https://bio.tools/swipe	https://github.com/torognes/swipe	Source code
https://bio.tools/swisstree	ftp://ftp.lausanne.isb-sib.ch/databases/SwissTree/	Biological data
https://bio.tools/syco	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/syco	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/syconn	https://github.com/StructuralNeurobiologyLab/SyConn	Source code
https://bio.tools/symcurv	http://genome.crg.es/SymCurv/documentation.html	Source code
https://bio.tools/symcurv	http://genome.crg.es/SymCurv/documentation.html	Binaries
https://bio.tools/tadbit	https://github.com/3DGenomes/tadbit	Source code
https://bio.tools/tam_2.0	http://www.scse.hebut.edu.cn/tam2/Public/static/data/tam_v2.rar	Source code
https://bio.tools/taxget	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/taxget	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/taxgetdown	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/taxgetdown	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/taxgetrank	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/taxgetrank	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/taxgetspecies	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/taxgetspecies	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/taxgetup	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/taxgetup	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/tcocode	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/tcocode	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/tensor_classification	https://github.com/laurafroelich/tensor_classification/blob/master/code/simulation/simulate_data.m	Test data
https://bio.tools/textget	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/textget	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/textsearch	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/textsearch	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/tfextract	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/tfextract	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/tfthaz	http://bioconductor.org/packages/release/bioc/vignettes/TFHAZ/inst/doc/TFHAZ_R	Source code
https://bio.tools/tfthaz	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/tfthaz	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/tfscan	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/tfscan	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/tfutils	http://bioconductor.org/packages/release/bioc/vignettes/TFutils/inst/doc/TFutils.R	Source code
https://bio.tools/tgnet	https://github.com/ksanao/TGNet	Source code
https://bio.tools/the_flux_capacitor	http://sammet.net/confluence/display/FLUX/2+-+Download	Source code

biotools_link	download_url	download_type
https://bio.tools/xina	https://www.bioconductor.org/packages/release/bioc/html/XINA.html	Binary package
https://bio.tools/xmlget	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/xmlget	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/xmltext	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/xmltext	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/xtalkdb	http://www.xtalkdb.org/download	Biological data
https://bio.tools/xtandem	ftp://ftp.thegpm.org/projects/tandem/	Source code
https://bio.tools/xtractor	http://www.ms-utils.org/Xtractor/Xtractor.c	Source code
https://bio.tools/yank	http://emboss.open-bio.org/html/adm/ch01s01.html	Source code
https://bio.tools/yank	http://emboss.open-bio.org/html/adm/ch01s01.html	Binaries
https://bio.tools/yara	https://docker-ui.genouest.org/app/#/container/bioconda/yara	Container file
https://bio.tools/yara	https://gitlab.com/sibyllewohlgemuth/cwl_files/raw/master/yara_indexer.cwl	CWL file
https://bio.tools/yara	https://gitlab.com/sibyllewohlgemuth/cwl_files/raw/master/yara_mapper.cwl	CWL file
https://bio.tools/yara	https://anaconda.org/bioconda/yara	Container file