

Deprecated Apps Documentation

DEPRECATED APPS

The apps in this list have been deprecated and are no longer available in the DE. If you need an app reinstated or have questions about a suitable replacement app, please contact support@cyverse.org.

- 1000 Bulls GATK Genotyping (archived)
- 1000 Bulls GATK Genotyping 1.00 (archived)
- ABySS Short Read Assembler (archived)
- AdjustPvalue 0.0.1 (archived)
- Agbase GOanna 1.02 (archived)
- Agbase GOanna 2.03 (archived)
- AgMIP QuadUI (archived)
- Autodock Vina (archived)
- BEDTools annotateBed (archived)
- BEDTools bedToBam (archived)
- BEDTools closestBed (archived)
- BEDTools clusterBed (archived)
- BEDTools complementBed (archived)
- BEDTools flankBed (archived)
- BEDTools getOverlap (archived)
- BEDTools groupBy (archived)
- BEDTools mergeBed (archived)
- BEDTools multiBamCov (archived)
- BEDTools multiIntersectBed (archived)
- BEDTools nucBed (archived)
- BEDTools shuffleBed (archived)
- BEDTools sortBed (archived)
- BEDTools subtractBed (archived)
- BEDTools windowBed (archived)
- Bismark 0.7.4 (archived)
- Bismark Genome Preparation 0.7.4 (archived)
- Bismark Methylation Extractor 0.7.4 (archived)
- Blastp-2.2.26 (archived)
- BLASTX Contig Annotation (Stage 2) (archived)
- Blat_the_most_basic_version (archived)
- BWA 0.5.9 (archived)
- BWA 0.59 (Paired-End Illumina Reads) (archived)
- cd-hit 4.6.1 (archived)
- CD-HIT-EST (archived)
- CD-HIT-EST-2D-ISU-GROUP-D (archived)
- Change Case (archived)
- CloudSTRUCTURE (archived)
- CONTRAST (archived)
- Convert standard files for AntEpiSeeker (CvtAntEpiSeeker) (archived)
- Convert standard files for Fast-LMM (CvtFastlmm) (archived)
- Convert standard files to Random Jungle format (CvtRandomJungle) (archived)
- Copy of Agbase GOanna 1.02 (archived)
- Copy of Infernal Stage 4 (cmsearch) (archived)
- Create BAM from SAM file (archived)
- CuffCompare (archived)
- Cuffdiff2a (archived)
- cufflinks2gff3 (archived)
- DACE (archived)
- demultiplex samples (archived)
- de novo construction genetic map (archived)
- Detect text file types (archived)
- Disparity through Time (archived)
- DNAML (archived)
- EBI Interproscan (Web Service) (archived)
- EBI Interproscan 1.0.1 (Web Service) (archived)
- eFP Batch Retrieve (archived)

- [EMBOSS backtranseq \(archived\)](#)
- [EMBOSS Biosed \(archived\)](#)
- [EMBOSS Charge \(archived\)](#)
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- [Infernal Stage 1 \(cmbuild\) \(archived\)](#)
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- [Infernal Stage 3 \(cmcalibrate\)](#) (archived)
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