

Resource Summary Report

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CHAoS

RRID:SCR_005174

Type: Tool

Proper Citation

CHAoS (RRID:SCR_005174)

Resource Information

URL: <http://www.well.ox.ac.uk/~kgaulton/chaos.shtml>

Proper Citation: CHAoS (RRID:SCR_005174)

Description: A Perl-based system for annotation of variants identified in high-throughput sequencing experiments. Functionality includes annotation of variants with information relating to population genetics, known transcripts, positional records, and sequence motif-based prediction. In addition, annotated variants can be summarized and extracted to facilitate downstream analysis. There is also basic support for gene-based biological annotation, and eventually will include tools for variant and genotype analysis and visualization.

Abbreviations: CHAoS

Synonyms: chaos - Annotation analysis and visualization of variants from high-throughput sequencing experiments

Resource Type: software resource

Keywords: annotation, analysis, visualization, variant, high-throughput sequencing, perl, population genetic, transcript, positional record, sequence, motif, genotype

Availability: GNU General Public License, v2

Resource Name: CHAoS

Resource ID: SCR_005174

Alternate IDs: OMICS_00170

Record Creation Time: 20220129T080228+0000

Record Last Update: 20260912T195622+0000

Ratings and Alerts

No rating or validation information has been found for CHAoS.

No alerts have been found for CHAoS.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 29 mentions in open access literature.

Listed below are recent publications. The full list is available at [RRID](#) .

Rogers HF, et al. (2026) How do core surface flow models vary when inverted from IGRF-14 candidate field models? Earth, planets, and space : EPS, 78(1), 39.

Zhao Z, et al. (2025) Large-vocabulary segmentation for medical images with text prompts. NPJ digital medicine, 8(1), 566.

Tian R, et al. (2025) Integrating cross-sample and cross-modal data for spatial transcriptomics and metabolomics with SpatialMETA. Nature communications, 16(1), 8855.

Patel N, et al. (2024) Training Robust T1-Weighted Magnetic Resonance Imaging Liver Segmentation Models Using Ensembles of Datasets with Different Contrast Protocols and Liver Disease Etiologies. Research square.

von Stumm S, et al. (2024) Adolescents' Perceptions of Household Chaos Predict Their Adult Mental Health: A Twin-Difference Longitudinal Cohort Study. Psychological science, 35(7), 736.

Yoon S, et al. (2024) Accelerated muscle mass estimation from CT images through transfer learning. BMC medical imaging, 24(1), 271.

Moraitopoulou G, et al. (2024) No association between alexithymia and emotion recognition or theory of mind in a sample of adolescents enhanced for autistic traits. Autism : the international journal of research and practice, 28(8), 2066.

Wang T, et al. (2024) Weakly supervised chest X-ray abnormality localization with non-linear modulation and foreground control. Scientific reports, 14(1), 29181.

Patel N, et al. (2024) Training robust T1-weighted magnetic resonance imaging liver segmentation models using ensembles of datasets with different contrast protocols and liver disease etiologies. Scientific reports, 14(1), 20988.

Koster N, et al. (2022) Understanding personality pathology in a clinical sample of youth: study protocol for the longitudinal research project 'APOLO'. BMJ open, 12(6), e054485.

Li CF, et al. (2022) MultiR-Net: A Novel Joint Learning Network for COVID-19 segmentation and classification. *Computers in biology and medicine*, 144, 105340.

Mascheroni E, et al. (2022) The role of experience in parenting beliefs of British and Italian women during pregnancy. *Infant mental health journal*, 43(6), 835.

Camhi SM, et al. (2021) Socioecological Factors Associated with an Urban Exercise Prescription Program for Under-Resourced Women: A Mixed Methods Community-Engaged Research Project. *International journal of environmental research and public health*, 18(16).

Dubois-Comtois K, et al. (2021) Child Psychological Functioning During the COVID-19 Lockdown: An Ecological, Family-Centered Approach. *Journal of developmental and behavioral pediatrics : JDBP*, 42(7), 532.

Woolhead VL, et al. (2021) Breed predispositions, clinical findings, and prognostic factors for death in dogs with nonregenerative immune-mediated anemia. *Journal of veterinary internal medicine*, 35(1), 252.

Kloenne M, et al. (2020) Domain-specific cues improve robustness of deep learning-based segmentation of CT volumes. *Scientific reports*, 10(1), 10712.

Rayce SB, et al. (2020) Concordance within parent couples' perception of parental stress symptoms among parents to 1-18-year-olds with physical or mental health problems. *PloS one*, 15(12), e0244212.

Sandholm N, et al. (2018) Confirmation of GLRA3 as a susceptibility locus for albuminuria in Finnish patients with type 1 diabetes. *Scientific reports*, 8(1), 12408.

Flannick J, et al. (2017) Sequence data and association statistics from 12,940 type 2 diabetes cases and controls. *Scientific data*, 4, 170179.

Fuchsberger C, et al. (2016) The genetic architecture of type 2 diabetes. *Nature*, 536(7614), 41.