Resource Summary Report

Generated by NIF on May 20, 2025

Computer Retrieval of Information on Scientific Projects Thesaurus

RRID:SCR_005301

Type: Tool

Proper Citation

Computer Retrieval of Information on Scientific Projects Thesaurus (RRID:SCR_005301)

Resource Information

URL: http://purl.bioontology.org/ontology/CRISP

Proper Citation: Computer Retrieval of Information on Scientific Projects Thesaurus

(RRID:SCR_005301)

Description: Ontology of Computer retrieval of Information on Scientific Projects (CRISP).

Abbreviations: CRISP

Resource Type: data or information resource, controlled vocabulary, ontology

Keywords: umls

Funding:

Resource Name: Computer Retrieval of Information on Scientific Projects Thesaurus

Resource ID: SCR_005301

Alternate IDs: nlx_157374

Record Creation Time: 20220129T080229+0000

Record Last Update: 20250519T203354+0000

Ratings and Alerts

No rating or validation information has been found for Computer Retrieval of Information on

Scientific Projects Thesaurus.

No alerts have been found for Computer Retrieval of Information on Scientific Projects Thesaurus.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 41 mentions in open access literature.

Listed below are recent publications. The full list is available at NIF.

Casaletto JA, et al. (2025) Analyzing the relationship between gene expression and phenotype in space-flown mice using a causal inference machine learning ensemble. Scientific reports, 15(1), 2363.

Emery JD, et al. (2023) The Colorectal cancer RISk Prediction (CRISP) trial: a randomised controlled trial of a decision support tool for risk-stratified colorectal cancer screening. The British journal of general practice: the journal of the Royal College of General Practitioners, 73(733), e556.

Poklukar K, et al. (2023) A meta-analysis of genetic and phenotypic diversity of European local pig breeds reveals genomic regions associated with breed differentiation for production traits. Genetics, selection, evolution: GSE, 55(1), 88.

Brewer MS, et al. (2023) Killer Knots: Molecular Evolution of Inhibitor Cystine Knot Toxins in Wandering Spiders (Araneae: Ctenidae). Toxins, 15(2).

Mathema VB, et al. (2022) CRISP: a deep learning architecture for GC?x?GC-TOFMS contour ROI identification, simulation and analysis in imaging metabolomics. Briefings in bioinformatics, 23(2).

Pechar GS, et al. (2022) Editing melon eIF4E associates with virus resistance and male sterility. Plant biotechnology journal, 20(10), 2006.

Chatterjee P, et al. (2021) The Effect of an Automated Mobile Patient Engagement Application on Emergency Department Revisits: Prospective Observational Study. JMIR formative research, 5(12), e17839.

Bovo S, et al. (2021) Describing variability in pig genes involved in coronavirus infections for a One Health perspective in conservation of animal genetic resources. Scientific reports, 11(1), 3359.

Lakhssassi N, et al. (2021) TILLING-by-Sequencing+ to Decipher Oil Biosynthesis Pathway

in Soybeans: A New and Effective Platform for High-Throughput Gene Functional Analysis. International journal of molecular sciences, 22(8).

Wei YB, et al. (2020) A functional variant in the serotonin receptor 7 gene (HTR7), rs7905446, is associated with good response to SSRIs in bipolar and unipolar depression. Molecular psychiatry, 25(6), 1312.

Gemperle J, et al. (2019) The interaction of p130Cas with PKN3 promotes malignant growth. Molecular oncology, 13(2), 264.

Moeck P, et al. (2018) Accurate lattice parameters from 2D-periodic images for subsequent Bravais lattice type assignments. Advanced structural and chemical imaging, 4(1), 5.

Walker JG, et al. (2018) The use of a risk assessment and decision support tool (CRISP) compared with usual care in general practice to increase risk-stratified colorectal cancer screening: study protocol for a randomised controlled trial. Trials, 19(1), 397.

Srivastava AK, et al. (2017) High-frequency torsional Alfvén waves as an energy source for coronal heating. Scientific reports, 7, 43147.

Doyle SR, et al. (2017) Genome-wide analysis of ivermectin response by Onchocerca volvulus reveals that genetic drift and soft selective sweeps contribute to loss of drug sensitivity. PLoS neglected tropical diseases, 11(7), e0005816.

Lodovicho ME, et al. (2017) Investigating possible biological targets of Bj-CRP, the first cysteine-rich secretory protein (CRISP) isolated from Bothrops jararaca snake venom. Toxicology letters, 265, 156.

Walker JG, et al. (2017) The CRISP colorectal cancer risk prediction tool: an exploratory study using simulated consultations in Australian primary care. BMC medical informatics and decision making, 17(1), 13.

Chen J, et al. (2016) Identifying Genetic Signatures of Natural Selection Using Pooled Population Sequencing in Picea abies. G3 (Bethesda, Md.), 6(7), 1979.

Pinos S, et al. (2016) Compartmentalization in PVC super-phylum: evolution and impact. Biology direct, 11, 38.

Anand S, et al. (2016) Next Generation Sequencing of Pooled Samples: Guideline for Variants' Filtering. Scientific reports, 6, 33735.