

Resource Summary Report

Generated by RRID on Aug 5, 2026

Indiana University; Indiana; USA

RRID:SCR_011294

Type: Tool

Proper Citation

Indiana University; Indiana; USA (RRID:SCR_011294)

Resource Information

URL: <http://www.indiana.edu/>

Proper Citation: Indiana University; Indiana; USA (RRID:SCR_011294)

Description: Public research university in Indiana. Has 8 statewide campuses known for innovation, creativity, and academic freedom.

Abbreviations: IU

Synonyms: Indiana University

Resource Type: university

Resource Name: Indiana University; Indiana; USA

Resource ID: SCR_011294

Alternate IDs: grid.411377.7, ISNI:0000 0001 0790 959X, nlx_87622, Wikidata:Q1079140, Crossref funder ID:100010178

Alternate URLs: <https://ror.org/02k40bc56>

Record Creation Time: 20220129T080303+0000

Record Last Update: 20260801T190406+0000

Ratings and Alerts

No rating or validation information has been found for Indiana University; Indiana; USA.

No alerts have been found for Indiana University; Indiana; USA.

Data and Source Information

Usage and Citation Metrics

We found 14 mentions in open access literature.

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

Makki MS, et al. (2020) A Precision Medicine Approach Uncovers a Unique Signature of Neutrophils in Patients With Brushite Kidney Stones. *Kidney international reports*, 5(5), 663.

Pyle E, et al. (2018) Structural Lipids Enable the Formation of Functional Oligomers of the Eukaryotic Purine Symporter UapA. *Cell chemical biology*, 25(7), 840.

Bankers L, et al. (2017) De Novo Transcriptome Characterization of a Sterilizing Trematode Parasite (*Microphallus* sp.) from Two Species of New Zealand Snails. *G3 (Bethesda, Md.)*, 7(3), 871.

Pacholarz KJ, et al. (2017) Hybrid Mass Spectrometry Approaches to Determine How L-Histidine Feedback Regulates the Enzyme MtATP-Phosphoribosyltransferase. *Structure (London, England : 1993)*, 25(5), 730.

Yampolsky LY, et al. (2017) Net Evolutionary Loss of Residue Polarity in Drosophilid Protein Cores Indicates Ongoing Optimization of Amino Acid Composition. *Genome biology and evolution*, 9(10), 2879.

Dabrowski-Tumanski P, et al. (2017) LinkProt: a database collecting information about biological links. *Nucleic acids research*, 45(D1), D243.

Sluka JP, et al. (2016) A Liver-Centric Multiscale Modeling Framework for Xenobiotics. *PloS one*, 11(9), e0162428.

Townsend JT, et al. (2016) A Note on Drawing Conclusions in the Study of Visual Search and the Use of Slopes in Particular. *i-Perception*, 7(6), 2041669516674220.

Alberge F, et al. (2015) Dynamic subcellular localization of a respiratory complex controls bacterial respiration. *eLife*, 4.

Schwarz F, et al. (2015) Siglec receptors impact mammalian lifespan by modulating oxidative stress. *eLife*, 4.

Goenner HF, et al. (2014) On the History of Unified Field Theories. Part II. (ca. 1930-ca. 1965). *Living reviews in relativity*, 17(1), 5.

Snir S, et al. (2014) Universal pacemaker of genome evolution in animals and fungi and variation of evolutionary rates in diverse organisms. *Genome biology and evolution*, 6(6), 1268.

Cridland JM, et al. (2013) Abundance and distribution of transposable elements in two *Drosophila* QTL mapping resources. *Molecular biology and evolution*, 30(10), 2311.

Sporns O, et al. (2007) Identification and classification of hubs in brain networks. PloS one, 2(10), e1049.