## **Resource Summary Report**

Generated by dkNET on May 20, 2025

# 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

**Funding:** 

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: 20250519T203654+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

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We found 14 mentions in open access literature.

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

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 Enzyzme MtATP-Phosphoribosyltransferase. Structure (London, England: 1993), 25(5), 730.

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

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.

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.