

Resource Summary Report

Generated by PRECISE-TBI on Sep 18, 2026

Purl

RRID:SCR_010238

Type: Tool

Proper Citation

Purl (RRID:SCR_010238)

Resource Information

URL: <http://purl.org/>

Proper Citation: Purl (RRID:SCR_010238)

Description: Web addresses that act as permanent identifiers in the face of a dynamic and changing Web infrastructure. Instead of resolving directly to Web resources, PURLs provide a level of indirection that allows the underlying Web addresses of resources to change over time without negatively affecting systems that depend on them. This capability provides continuity of references to network resources that may migrate from machine to machine for business, social or technical reasons.

Synonyms: Persistent Uniform Resource Locators

Resource Type: data or information resource, narrative resource, standard specification

Resource Name: Purl

Resource ID: SCR_010238

Alternate IDs: nlx_156858

Record Creation Time: 20220129T080257+0000

Record Last Update: 20260912T195716+0000

Ratings and Alerts

No rating or validation information has been found for Purl.

No alerts have been found for Purl.

Data and Source Information

Usage and Citation Metrics

We found 208 mentions in open access literature.

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

Han Y, et al. (2025) Antibiotic-associated changes in *Akkermansia muciniphila* alter its effects on host metabolic health. *Microbiome*, 13(1), 48.

Cox SJD, et al. (2021) Ten simple rules for making a vocabulary FAIR. *PLoS computational biology*, 17(6), e1009041.

Sharma N, et al. (2020) Role of allosteric switches and adaptor domains in long-distance cross-talk and transient tunnel formation. *Science advances*, 6(14), eaay7919.

Bamparopoulos G, et al. (2016) Towards exergaming commons: composing the exergame ontology for publishing open game data. *Journal of biomedical semantics*, 7, 4.

Rodríguez-Iglesias A, et al. (2016) Publishing FAIR Data: An Exemplar Methodology Utilizing PHI-Base. *Frontiers in plant science*, 7, 641.

Guralnick RP, et al. (2015) Community next steps for making globally unique identifiers work for biocollections data. *ZooKeys*(494), 133.

Zaher H, et al. (2014) Consuming viscous prey: a novel protein-secreting delivery system in neotropical snail-eating snakes. *BMC evolutionary biology*, 14, 58.

Gu W, et al. (2014) The impact of RNA structure on coding sequence evolution in both bacteria and eukaryotes. *BMC evolutionary biology*, 14, 87.

Sjökvist E, et al. (2014) Stereopsidales--a new order of mushroom-forming fungi. *PloS one*, 9(4), e95227.

Pozzi L, et al. (2014) A multilocus phylogeny reveals deep lineages within African galagids (Primates: Galagidae). *BMC evolutionary biology*, 14, 72.

Butler RJ, et al. (2014) New clade of enigmatic early archosaurs yields insights into early pseudosuchian phylogeny and the biogeography of the archosaur radiation. *BMC evolutionary biology*, 14, 128.

Xu L, et al. (2014) Complete genome sequence and comparative genomic analyses of the vancomycin-producing *Amycolatopsis orientalis*. *BMC genomics*, 15(1), 363.

Yonemaru J, et al. (2014) Genomic regions involved in yield potential detected by genome-wide association analysis in Japanese high-yielding rice cultivars. *BMC genomics*, 15(1), 346.

Klein A, et al. (2014) Benchmarking infrastructure for mutation text mining. *Journal of biomedical semantics*, 5(1), 11.

Lorén JG, et al. (2014) Molecular phylogenetics and temporal diversification in the genus *Aeromonas* based on the sequences of five housekeeping genes. PLoS one, 9(2), e88805.

Miyazawa H, et al. (2014) Molecular phylogeny of Myriapoda provides insights into evolutionary patterns of the mode in post-embryonic development. Scientific reports, 4, 4127.

Grant JR, et al. (2014) Building a phylogenomic pipeline for the eukaryotic tree of life - addressing deep phylogenies with genome-scale data. PLoS currents, 6.

Zhou CF, et al. (2014) A new specimen of *Manchurochelys manchoukuoensis* from the Early Cretaceous Jehol Biota of Chifeng, Inner Mongolia, China and the phylogeny of Cretaceous basal eucryptodiran turtles. BMC evolutionary biology, 14, 77.

Liu F, et al. (2014) Comparative genomic analysis of *Mycobacterium tuberculosis* clinical isolates. BMC genomics, 15(1), 469.

Yoder MJ, et al. (2014) Monograph of the Afrotropical species of *Scelio* Latreille (Hymenoptera, Platygasteridae), egg parasitoids of acridid grasshoppers (Orthoptera, Acrididae). ZooKeys(380), 1.