

Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 8, 2026

[surankco](#)

RRID:SCR_024355

Type: Tool

Proper Citation

surankco (RRID:SCR_024355)

Resource Information

URL: <https://sourceforge.net/projects/surankco/>

Proper Citation: surankco (RRID:SCR_024355)

Description: Machine learning based software to score and rank contigs from de novo assemblies of next generation sequencing data. It trains with alignments of contigs with known reference genomes and predicts scores and ranking for contigs which have no related reference genome yet.

Resource Type: simulation software, software application, software resource

Defining Citation: [PMID:26224355](#)

Keywords: score and rank contigs, de novo assemblies of next generation sequencing data,

Availability: Free, Available for download, Freely available,

Resource Name: surankco

Resource ID: SCR_024355

Alternate IDs: OMICS_09480

Alternate URLs: <https://sources.debian.org/src/surankco/>

License: BSD License

Record Creation Time: 20230830T050217+0000

Record Last Update: 20260905T133030+0000

Ratings and Alerts

No rating or validation information has been found for surankco.

No alerts have been found for surankco.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Borda V, et al. (2024) Genetics of Latin American Diversity Project: Insights into population genetics and association studies in admixed groups in the Americas. Cell genomics, 4(11), 100692.