

Resource Summary Report

Generated by T1D on Sep 5, 2026

DiProGB

RRID:SCR_005651

Type: Tool

Proper Citation

DiProGB (RRID:SCR_005651)

Resource Information

URL: <http://diprogb.fli-leibniz.de/>

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Description: Genome browser that encodes the genome sequence by physico-chemical dinucleotide properties such as stacking energy, melting temperature or twist angle. Analyses can be performed for the + and ?, as well as for the double strand.

Abbreviations: DiProGB

Synonyms: DiProGB - The Dinucleotide Properties Genome Browser, Dinucleotide Properties Genome Browser

Resource Type: software resource

Defining Citation: [PMID:19605418](#)

Keywords: genome, browser, bio.tools

Availability: Free, Freely available

Resource Name: DiProGB

Resource ID: SCR_005651

Alternate IDs: biotools:diprogb, OMICS_00880

Alternate URLs: <https://bio.tools/diprogb>

Record Creation Time: 20220129T080231+0000

Record Last Update: 20260903T234802+0000

Ratings and Alerts

No rating or validation information has been found for DiProGB.

No alerts have been found for DiProGB.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 4 mentions in open access literature.

Listed below are recent publications. The full list is available at [T1D](#).

Wang HT, et al. (2020) Identification of DNA N6-methyladenine sites by integration of sequence features. *Epigenetics & chromatin*, 13(1), 8.

Daga A, et al. (2015) Characterization of chromosomal translocation breakpoint sequences in solid tumours: "an in silico analysis". *The open medical informatics journal*, 9, 1.

Gupta Y, et al. (2014) ptRNAPred: computational identification and classification of post-transcriptional RNA. *Nucleic acids research*, 42(22), e167.

Maruyama H, et al. (2013) An alternative beads-on-a-string chromatin architecture in *Thermococcus kodakarensis*. *EMBO reports*, 14(8), 711.