

Resource Summary Report

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[tbl2asn](#)

RRID:SCR_016636

Type: Tool

Proper Citation

tbl2asn (RRID:SCR_016636)

Resource Information

URL: <https://www.ncbi.nlm.nih.gov/genbank/tbl2asn2/>

Proper Citation: tbl2asn (RRID:SCR_016636)

Description: Software tool as a command-line program that automates the creation of sequence records for submission to GenBank. Records need no additional manual editing before submission.

Resource Type: software resource, software application, data processing software

Keywords: command, line, program, automate, creation, sequence, record, submit, data, GenBank

Availability: Free, Freely available

Resource Name: tbl2asn

Resource ID: SCR_016636

Record Creation Time: 20220129T080331+0000

Record Last Update: 20260804T043631+0000

Ratings and Alerts

No rating or validation information has been found for tbl2asn.

No alerts have been found for tbl2asn.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 8 mentions in open access literature.

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

González-Vázquez R, et al. (2022) Genomic and Biochemical Characterization of *Bifidobacterium pseudocatenulatum* JCLA3 Isolated from Human Intestine. *Microorganisms*, 10(11).

Wang ZF, et al. (2021) The complete chloroplast genome of *Aganope dinghuensis* (Fabaceae). *Mitochondrial DNA. Part B, Resources*, 6(2), 460.

Pons J, et al. (2021) Aln2tbl: building a mitochondrial features table from a assembly alignment in fasta format. *Mitochondrial DNA. Part B, Resources*, 6(9), 2732.

Pace J, et al. (2020) PuMA: A papillomavirus genome annotation tool. *Virus evolution*, 6(2), veaa068.

Tisza MJ, et al. (2020) Discovery of several thousand highly diverse circular DNA viruses. *eLife*, 9.

Stroehlein AJ, et al. (2019) High-quality *Schistosoma haematobium* genome achieved by single-molecule and long-range sequencing. *GigaScience*, 8(9).

An K, et al. (2018) Spinal PKC activation - Induced neuronal HMGB1 translocation contributes to hyperalgesia in a bone cancer pain model in rats. *Experimental neurology*, 303, 80.

Raymond O, et al. (2018) The *Rosa* genome provides new insights into the domestication of modern roses. *Nature genetics*, 50(6), 772.