**BitSeq**

**RRID:** SCR_009904  
**Type:** Tool

**Proper Citation:**

BitSeq (RRID:SCR_009904)

**Resource Information**

**URL:** [http://code.google.com/p/bitseq/](http://code.google.com/p/bitseq/)

**Proper Citation:** BitSeq (RRID:SCR_009904)

**Description:** A software application for inferring expression levels of individual transcripts from sequencing (RNA-Seq) data and estimating differential expression (DE) between conditions.

**Abbreviations:** BitSeq

**Resource Type:** software resource

**Defining Citation:** [DOI:10.1093/bioinformatics/btv483](https://doi.org/10.1093/bioinformatics/btv483), [DOI:10.1093/bioinformatics/bts260](https://doi.org/10.1093/bioinformatics/bts260)

**Keywords:** bio.tools

**Resource Name:** BitSeq

**Resource ID:** SCR_009904

**Alternate IDs:** OMICS_01269, biotools:bitseq

**Alternate URLs:** https://bio.tools/bitseq, https://sources.debian.org/src/bitseq/

**Record Creation Time:** 20220129T080255+0000

**Record Last Update:** 20240424T182855+0000

**Ratings and Alerts**
No rating or validation information has been found for BitSeq.

No alerts have been found for BitSeq.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 17 mentions in open access literature.

Listed below are recent publications. The full list is available at FDI Lab - SciCrunch.org.


Ustianenko D, et al. (2016) TUT-DIS3L2 is a mammalian surveillance pathway for aberrant
structured non-coding RNAs. The EMBO journal, 35(20), 2179.


