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.
- **Resource Type:** Resource, software resource
- **Keywords:** bio.tools
- **Website Status:** Last checked up
- **Abbreviations:** BitSeq
- **Resource Name:** BitSeq
- **Resource ID:** SCR_009904
- **Alternate IDs:** OMICS_01269, biotools:bitseq
- **Alternate URLs:** [https://bio.tools/bitseq](https://bio.tools/bitseq)

---

**Ratings and Alerts**

No rating or validation information has been found for BitSeq.

No alerts have been found for BitSeq.
Usage and Citation Metrics

We found 15 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [FDI Lab - SciCrunch.org](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), 2179-2191.


