**GASiC**

**RRID:** SCR_006765  
**Type:** Tool

### Proper Citation

GASiC (RRID:SCR_006765)

### Resource Information

**URL:** [http://sourceforge.net/projects/gasic/](http://sourceforge.net/projects/gasic/)

**Proper Citation:** GASiC (RRID:SCR_006765)

**Description:** A method to correct read alignment results for the ambiguities imposed by similarities of genomes.

**Resource Type:** Resource, software resource

**References:** PMID:22941661

**Keywords:** metagenome, genome, sequence, python

**Parent Organization:** SourceForge

**Availability:** BSD License

**Website Status:** Last checked up

**Abbreviations:** GASiC

**Resource Name:** GASiC

**Resource ID:** SCR_006765

**Alternate IDs:** OMICS_01437

### Ratings and Alerts
No rating or validation information has been found for GASiC.

No alerts have been found for GASiC.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 2 mentions in open access literature.

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