

Resource Summary Report

Generated by SPARC SAWG on Aug 14, 2026

Pyrocleaner

RRID:SCR_000584

Type: Tool

Proper Citation

Pyrocleaner (RRID:SCR_000584)

Resource Information

URL: <https://mulcyber.toulouse.inra.fr/plugins/mediawiki/wiki/ng6/index.php/Pyrocleaner>

Proper Citation: Pyrocleaner (RRID:SCR_000584)

Description: A software module for cleaning 454 pyrosequencing reads using filters such as read duplication, length, complexity, base-pair quality and number of undetermined bases. The software also uses flowgram files of paired-end sequences to validate paired-ends files of another file.

Abbreviations: Pyrocleaner

Resource Type: software resource

Defining Citation: [PMID:21615897](#)

Keywords: pyrosequence, assembly, rna, dna, duplication, length, complexity, bases, flowgram, paired end

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: Pyrocleaner

Resource ID: SCR_000584

Alternate IDs: OMICS_01069

License: GNU General Public License

Record Creation Time: 20220129T080202+0000

Record Last Update: 20260808T185718+0000

Ratings and Alerts

No rating or validation information has been found for Pyrocleaner.

No alerts have been found for Pyrocleaner.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Jérôme M, et al. (2011) Assessment of replicate bias in 454 pyrosequencing and a multi-purpose read-filtering tool. BMC research notes, 4, 149.