FastQ Screen

RRID:SCR_000141
Type: Tool

Proper Citation

FastQ Screen (RRID:SCR_000141)

Resource Information

URL: http://www.bioinformatics.babraham.ac.uk/projects/fastq_screen/

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Description: Software that allows you to screen a library of sequences in FastQ format against a set of sequence databases so you can see if the composition of the library matches with what you expect.

Abbreviations: FastQ Screen

Resource Type: software resource

Defining Citation: PMID:30254741

Keywords: perl, FASEB list

Availability: Free, Freely available

Resource Name: FastQ Screen

Resource ID: SCR_000141

Alternate IDs: OMICS_01042

Ratings and Alerts

No rating or validation information has been found for FastQ Screen.

No alerts have been found for FastQ Screen.
Usage and Citation Metrics

We found 36 mentions in open access literature.

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


Kok YL, et al. (2018) Spontaneous reactivation of latent HIV-1 promoters is linked to the cell cycle as revealed by a genetic-insulators-containing dual-fluorescence HIV-1-based vector. Scientific reports, 8(1), 10204.


