hot scan

RRID:SCR_002840
Type: Tool

Proper Citation

hot scan (RRID:SCR_002840)

Resource Information

URL: https://github.com/itojal/hot_scan

Proper Citation: hot scan (RRID:SCR_002840)

Description: A free software to detect genomic regions unusually rich in translocation breakpoints. More generally, it may be used to detect a region that is unusually rich in a given character of a binary sequence.

Synonyms: hot_scan

Resource Type: software resource

Defining Citation: PMID:24860160

Keywords: software package, perl, r, bio.tools

Availability: GNU General Public License, v2

Resource Name: hot scan

Resource ID: SCR_002840

Alternate IDs: biotools:hot_scan

Alternate URLs: https://bio.tools/hot_scan

Ratings and Alerts

No rating or validation information has been found for hot scan.
No alerts have been found for hot scan.

**Data and Source Information**

**Source:** SciCrunch Registry

**Usage and Citation Metrics**

We found 235 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [RRID](https://doi.org/10.5551/sdr/220).


the underlying mechanisms associated with parallel processing. Psychological research, 82(4), 720-733.


Kuipers JR, et al. (2018) Abstract images and words can convey the same meaning. Scientific reports, 8(1), 7190.


