

Resource Summary Report

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PacmonSTR

RRID:SCR_002796

Type: Tool

Proper Citation

PacmonSTR (RRID:SCR_002796)

Resource Information

URL: <https://github.com/alibashir/pacmonstr>

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Description: Software that implements a reference-based probabilistic approach to identify the Tandem Repeat (TR) region and estimate the number of these TR elements in long DNA reads.

Synonyms: PacmonSTR - Tandem Repeat Detection for Long Read Sequence Data

Resource Type: software resource

Defining Citation: [PMID:25028725](#)

Keywords: standalone software, pacific biosciences

Availability: Free, Freely available, Available for download

Resource Name: PacmonSTR

Resource ID: SCR_002796

Alternate IDs: OMICS_05208

Record Creation Time: 20220129T080215+0000

Record Last Update: 20261017T202518+0000

Ratings and Alerts

No rating or validation information has been found for PacmonSTR.

No alerts have been found for PacmonSTR.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.