

Resource Summary Report

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MSG

RRID:SCR_004161

Type: Tool

Proper Citation

MSG (RRID:SCR_004161)

Resource Information

URL: <http://genomics.princeton.edu/AndolfattoLab/MSG.html>

Proper Citation: MSG (RRID:SCR_004161)

Description: A pipeline of scripts to assign ancestry to genomic segments using next-gen sequence data. This method can identify recombination breakpoints in a large number of individuals simultaneously at a resolution sufficient for most mapping purposes, such as quantitative trait locus (QTL) mapping and mapping of induced mutations.

Abbreviations: MSG

Synonyms: Multiplexed shotgun genotyping, Multiplexed shotgun genotyping (MSG), MSG: Multiplexed Shotgun Genotyping

Resource Type: software resource

Defining Citation: [PMID:21233398](#)

Keywords: next generation sequencing, genotyping, genetic mapping, ancestry, genome

Resource Name: MSG

Resource ID: SCR_004161

Alternate IDs: OMICS_01551

Record Creation Time: 20220129T080223+0000

Record Last Update: 20261017T202609+0000

Ratings and Alerts

No rating or validation information has been found for MSG.

No alerts have been found for MSG.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.