

Resource Summary Report

Generated by [NIF](#) on Oct 20, 2026

SPLINK

RRID:SCR_009414

Type: Tool

Proper Citation

SPLINK (RRID:SCR_009414)

Resource Information

URL: <https://watson.hgen.pitt.edu/docs/splink108.html>

Proper Citation: SPLINK (RRID:SCR_009414)

Description: Software application for linkage analysis using affected sib pairs (entry from Genetic Analysis Software)

Synonyms: affected Sib Pairs LINKage analysis

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, c++, unix, sunos, ms-dos

Resource Name: SPLINK

Resource ID: SCR_009414

Alternate IDs: nlx_154659

Old URLs: <http://www-gene.cimr.cam.ac.uk/clayton/software/>

Record Creation Time: 20220129T080252+0000

Record Last Update: 20261017T203554+0000

Ratings and Alerts

No rating or validation information has been found for SPLINK.

No alerts have been found for SPLINK.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.