

Resource Summary Report

Generated by SPARC SAWG on Sep 17, 2026

ASPEX

RRID:SCR_008414

Type: Tool

Proper Citation

ASPEX (RRID:SCR_008414)

Resource Information

URL: <http://aspex.sourceforge.net/>

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Description: A set of programs for performing multipoint exclusion mapping of affected sibling pair data for discrete traits. (entry from Genetic Analysis Software)

Abbreviations: ASPEX

Synonyms: Affected Sib Pairs EXclusion map

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, c, tcl, perl, (the graphic script requires xmgr, available from ftp://ftp.teleport.com/pub/users/pturner/acegr), unix, (solaris/sunos/irix/osf-1), linux, bio.tools

Resource Name: ASPEX

Resource ID: SCR_008414

Alternate IDs: nlx_154234, biotools:aspex

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

Record Creation Time: 20220129T080247+0000

Record Last Update: 20260912T200237+0000

Ratings and Alerts

No rating or validation information has been found for ASPEX.

No alerts have been found for ASPEX.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 9 mentions in open access literature.

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

Adabavazeh Z, et al. (2017) Effect of Adding Cerium on Microstructure and Morphology of Ce-Based Inclusions Formed in Low-Carbon Steel. Scientific reports, 7, 46503.

Chang TJ, et al. (2015) Genetic polymorphisms of PCSK2 are associated with glucose homeostasis and progression to type 2 diabetes in a Chinese population. Scientific reports, 5, 14380.

Ramagopalan SV, et al. (2008) Parental non-inherited HLA resistance alleles do not confer protection against multiple sclerosis. Journal of neuroimmunology, 196(1-2), 170.

Ramagopalan SV, et al. (2007) The inheritance of resistance alleles in multiple sclerosis. PLoS genetics, 3(9), 1607.

Hinrichs AL, et al. (2006) Functional variant in a bitter-taste receptor (hTAS2R16) influences risk of alcohol dependence. American journal of human genetics, 78(1), 103.

Poznik GD, et al. (2006) A novel framework for sib pair linkage analysis. American journal of human genetics, 78(2), 222.

Jorgenson E, et al. (2005) Ethnicity and human genetic linkage maps. American journal of human genetics, 76(2), 276.

McCauley JL, et al. (2005) Genome-wide and Ordered-Subset linkage analyses provide support for autism loci on 17q and 19p with evidence of phenotypic and interlocus genetic correlates. BMC medical genetics, 6, 1.

Rosche B, et al. (2004) The role of the polio virus receptor and the herpesvirus entry mediator B genes for the development of MS. Journal of neuroimmunology, 156(1-2), 171.