

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

ALLELIX

RRID:SCR_009115

Type: Tool

Proper Citation

ALLELIX (RRID:SCR_009115)

Resource Information

URL: <http://www.allelix.net>

Proper Citation: ALLELIX (RRID:SCR_009115)

Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented on September 23, 2013. Software application / data analysis service where one can enter the alleles of commonly used STR by clicking the mouse. The algorithm calculates the paternity index and the Essen-Moeller probability of kinship for the deficiency- and the trio case. Everybody can use the network-software online after registering. The usage on the internet is free. Academic users can ask me to unlock an option to display the details (formulas/frequencies etc.) and to have an export-funktion to MS Word. The program is in German and (non-professional) English. An expansion to other languages is easy, if somebody helps us with the translation. For those who are interested to have the software running on their own intranet (for database security reasons) an individual agreement can be found. (entry from Genetic Analysis Software) (German version is: <http://www.allelix.de>)

Abbreviations: ALLELIX

Synonyms: Allelix - Paternity Linkage Analysis Online

Resource Type: service resource, data analysis service, analysis service resource, software resource, software application, production service resource

Keywords: gene, genetic, genomic, c++, java script, ms-windows, (2000 &iis/me & personal webserver)

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: ALLELIX

Resource ID: SCR_009115

Alternate IDs: nlx_154216

Record Creation Time: 20220129T080251+0000

Record Last Update: 20260728T043316+0000

Ratings and Alerts

No rating or validation information has been found for ALLELIX.

No alerts have been found for ALLELIX.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

Listed below are recent publications. The full list is available at [nidm-terms](#) .

Camargo-Ayala PA, et al. (2018) On the Evolution and Function of Plasmodium vivax Reticulocyte Binding Surface Antigen (pvrbsa). Frontiers in genetics, 9, 372.

Garz??n-Ospina D, et al. (2014) Heterogeneous genetic diversity pattern in Plasmodium vivax genes encoding merozoite surface proteins (MSP) -7E, -7F and -7L. Malaria journal, 13, 495.