# **Resource Summary Report**

Generated by dkNET on Apr 16, 2025

## **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:** software resource, data analysis service, software application, service resource, analysis service resource, production service resource

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

**Funding:** 

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:** 20250416T063537+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 dkNET.

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

Garzón-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.