

Resource Summary Report

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ALOHOMORA

RRID:SCR_009117

Type: Tool

Proper Citation

ALOHOMORA (RRID:SCR_009117)

Resource Information

URL: <http://gmc.mdc-berlin.de/alohomora/>

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Description: Software application designed to facilitate genome-wide linkage studies performed with high-density single nucleotide polymorphism (SNP) marker panels such as the Affymetrix GeneChip(R) Human Mapping 10K Array. (entry from Genetic Analysis Software)

Abbreviations: ALOHOMORA

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, perl, ms-window, linux, unix, solaris, bio.tools

Resource Name: ALOHOMORA

Resource ID: SCR_009117

Alternate IDs: nlx_154219, biotools:alohomora

Alternate URLs: <https://bio.tools/alohomora>

Record Creation Time: 20220129T080251+0000

Record Last Update: 20260727T040451+0000

Ratings and Alerts

No rating or validation information has been found for ALOHOMORA.

No alerts have been found for ALOHOMORA.

Data and Source Information

Usage and Citation Metrics

We found 8 mentions in open access literature.

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

Vona B, et al. (2021) A biallelic variant in CLRN2 causes non-syndromic hearing loss in humans. Human genetics, 140(6), 915.

Doll J, et al. (2020) Novel Loss-of-Function Variants in CDC14A are Associated with Recessive Sensorineural Hearing Loss in Iranian and Pakistani Patients. International journal of molecular sciences, 21(1).

Doll J, et al. (2020) Genetic Spectrum of Syndromic and Non-Syndromic Hearing Loss in Pakistani Families. Genes, 11(11).

Pr??ss H, et al. (2019) Linkage Evidence for a Two-Locus Inheritance of LQT-Associated Seizures in a Multigenerational LQT Family With a Novel KCNQ1 Loss-of-Function Mutation. Frontiers in neurology, 10, 648.

Bouhouche A, et al. (2018) Genetic Analysis of Undiagnosed Juvenile GM1-Gangliosidosis by Microarray and Exome Sequencing. Case reports in genetics, 2018, 8635698.

Davarniya B, et al. (2015) The Role of a Novel TRMT1 Gene Mutation and Rare GRM1 Gene Defect in Intellectual Disability in Two Azeri Families. PloS one, 10(8), e0129631.

Said MB, et al. (2013) Posterior microphthalmia and nanophthalmia in Tunisia caused by a founder c.1059_1066insC mutation of the PRSS56 gene. Gene, 528(2), 288.

Decker E, et al. (2008) PTHR1 loss-of-function mutations in familial, nonsyndromic primary failure of tooth eruption. American journal of human genetics, 83(6), 781.