Resource Summary Report

Generated by dkNET on Apr 16, 2025

CYRILLIC

RRID:SCR_001823
Type: Tool

Proper Citation

CYRILLIC (RRID:SCR_001823)

Resource Information

URL: http://www.cyrillicsoftware.com

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Description: Software application for pedigree drawing with fully integrated risk analysis and support for industry standard databases (MS Access and Corel Paradox). It is designed for genetic counselors and others who work with patients. Cyrillic 2 draws pedigrees, works with genetic marker data, lets you do haplotyping and allows exports to a range of linkage analysis packages.

Abbreviations: Cyrillic

Synonyms: CyrillicSoftware

Resource Type: software resource, commercial organization, software application

Defining Citation: PMID:1973333

Keywords: gene, genetic, genomic, visual c++, ms-windows, pedigree, linkage analysis, risk analysis, FASEB list

Funding:

Availability: Apache License, Commercial license

Resource Name: CYRILLIC

Resource ID: SCR_001823

Alternate IDs: nlx_154279, OMICS_00208

Record Creation Time: 20220129T080209+0000

Record Last Update: 20250416T063246+0000

Ratings and Alerts

No rating or validation information has been found for CYRILLIC.

No alerts have been found for CYRILLIC.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 52 mentions in open access literature.

Listed below are recent publications. The full list is available at dkNET.

Bekarystankyzy A, et al. (2024) Multilingual end-to-end ASR for low-resource Turkic languages with common alphabets. Scientific reports, 14(1), 13835.

Pavlovski D, et al. (2024) How older adults experience the age-friendliness of Skopje: Results of the validation of the AFCCQ for use in North Macedonia and a representative survey. Heliyon, 10(9), e30372.

Khan H, et al. (2024) Biallelic variants identified in 36 Pakistani families and trios with autism spectrum disorder. Scientific reports, 14(1), 9230.

Khan J, et al. (2024) Mutational spectrum associated with oculocutaneous albinism and Hermansky-Pudlak syndrome in nine Pakistani families. BMC ophthalmology, 24(1), 345.

, et al. (2024) Commodity risk assessment of plants of 12 selected Prunus species from Moldova. EFSA journal. European Food Safety Authority, 22(3), e8647.

Al-Mutairi DA, et al. (2024) Novel pathogenic variants of DNAH5 associated with clinical and genetic spectra of primary ciliary dyskinesia in an Arab population. Frontiers in genetics, 15, 1396797.

Chen M, et al. (2023) Phenotype, genotype, and management of congenital fibrosis of extraocular muscles type 1 in 16 Chinese families. Graefe's archive for clinical and experimental ophthalmology = Albrecht von Graefes Archiv fur klinische und experimentelle Ophthalmologie, 261(3), 879.

Martín-Luengo B, et al. (2023) Do pictures influence memory and metamemory in Chinese vocabulary learning? Evidence from Russian and Colombian learners. PloS one, 18(11), e0286824.

Bermúdez-Margaretto B, et al. (2022) Ultra-rapid and automatic interplay between L1 and L2 semantics in late bilinguals: EEG evidence. Cortex; a journal devoted to the study of the nervous system and behavior, 151, 147.

Guo YH, et al. (2022) KLF13 Loss-of-Function Mutations Underlying Familial Dilated Cardiomyopathy. Journal of the American Heart Association, 11(22), e027578.

Ljaji? A, et al. (2022) Uncovering the Reasons Behind COVID-19 Vaccine Hesitancy in Serbia: Sentiment-Based Topic Modeling. Journal of medical Internet research, 24(11), e42261.

Heidari M, et al. (2021) Identification of a novel homozygous mutation in the DDR2 gene from a patient with spondylo-meta-epiphyseal dysplasia by whole exome sequencing. Iranian journal of basic medical sciences, 24(2), 191.

Heidari M, et al. (2021) Identification of Two Novel Mutations in PKHD1 Gene from Two Families with Polycystic Kidney Disease by Whole Exome Sequencing. Current genomics, 22(3), 232.

Daneshpour MS, et al. (2021) Chromosomal regions strongly associated with waist circumference and body mass index in metabolic syndrome in a family-based study. Scientific reports, 11(1), 6082.

Saba N, et al. (2021) Congenital cataract: An ocular manifestation of classical homocystinuria. Molecular genetics & genomic medicine, 9(9), e1742.

Vajen B, et al. (2021) Psychological Distress and Coping Ability of Women at High Risk of Hereditary Breast and Ovarian Cancer before Undergoing Genetic Counseling-An Exploratory Study from Germany. International journal of environmental research and public health, 18(8).

Guo XJ, et al. (2021) PRRX1 Loss-of-Function Mutations Underlying Familial Atrial Fibrillation. Journal of the American Heart Association, 10(23), e023517.

Heidari M, et al. (2020) Association of a novel homozygous mutation in the HMGCS2 gene with an HMGCSD in an Iranian patient. Molecular genetics & genomic medicine, 8(11), e1507.

Zhexenova Z, et al. (2020) A Comparison of Social Robot to Tablet and Teacher in a New

Script Learning Context. Frontiers in robotics and AI, 7, 99.

Heidari M, et al. (2019) Identification of Two Novel Mutations in the ATM Gene from Patients with Ataxia-Telangiectasia by Whole Exome Sequencing. Current genomics, 20(7), 531.