

# Resource Summary Report

Generated by [nidm-terms](#) on Jul 31, 2026

## MOON

RRID:SCR\_018005

Type: Tool

### Proper Citation

MOON (RRID:SCR\_018005)

### Resource Information

**URL:** <http://www.diploid.com/moon>

**Proper Citation:** MOON (RRID:SCR\_018005)

**Description:** Software package that autonomously diagnoses rare diseases from next generation sequencing NGS data using artificial intelligence by Diploid.

**Resource Type:** data processing software, software toolkit, data analysis software, software resource, software application

**Keywords:** Diagnosis, rare disease, next generation sequencing, NGS, data, artificial intelligence, analysis, Diploid

**Availability:** Restricted

**Resource Name:** MOON

**Resource ID:** SCR\_018005

**Record Creation Time:** 20220129T080338+0000

**Record Last Update:** 20260731T043018+0000

### Ratings and Alerts

No rating or validation information has been found for MOON.

No alerts have been found for MOON.

### Data and Source Information

**Source:** [SciCrunch Registry](#)

## Usage and Citation Metrics

We found 6 mentions in open access literature.

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

Pantea CL, et al. (2025) Genetic landscape of Romanian children with inborn errors of immunity via gene panels, exome, and genome sequencing. Scientific reports, 15(1), 18830.

Johansson J, et al. (2024) Gustavson syndrome is caused by an in-frame deletion in RBMX associated with potentially disturbed SH3 domain interactions. European journal of human genetics : EJHG, 32(3), 333.

Bahena P, et al. (2022) Unraveling the genetic complexities of combined retinal dystrophy and hearing impairment. Human genetics, 141(3-4), 785.

Johansson J, et al. (2022) Loss of Nexilin function leads to a recessive lethal fetal cardiomyopathy characterized by cardiomegaly and endocardial fibroelastosis. American journal of medical genetics. Part A, 188(6), 1676.

Balicza P, et al. (2020) Novel dominant MPAN family with a complex genetic architecture as a basis for phenotypic variability. Neurology. Genetics, 6(5), e515.

Peeters S, et al. (2020) DNA Methylation Profiling and Genomic Analysis in 20 Children with Short Stature Who Were Born Small for Gestational Age. The Journal of clinical endocrinology and metabolism, 105(12).