

# Package: GenomeNexus (via r-universe)

September 12, 2024

**Title** Expose the GenomeNexus API as an R package

**Version** 0.0.0.9000

**Description** This package provides an R interface to the GenomeNexus API. GenomeNexus is a web application that annotates cancer variants with information from multiple databases. The GenomeNexus API provides programmatic access to the data and annotations available in the GenomeNexus web application.

**License** Artistic-2.0

**Depends** AnVIL

**Imports** methods

**Encoding** UTF-8

**Roxygen** list(markdown = TRUE)

**RoxygenNote** 7.3.1

**Repository** <https://link-ny.r-universe.dev>

**RemoteUrl** <https://github.com/LiNk-NY/GenomeNexus>

**RemoteRef** HEAD

**RemoteSha** 63383df693c1b2d26a2f1a3892789902c33ef991