



From the [Artemis home page](#):

Artemis is a free genome viewer and annotation tool that allows visualization of sequence features and the results of analyses within the context of the sequence, and its six-frame translation. Artemis is written in Java, and is available for UNIX, Macintosh and Windows systems. It can read [EMBL](#) and [GENBANK](#) database entries or sequence in [FASTA](#) or raw format. Extra sequence features can be in EMBL, GENBANK or [GFF](#) format.

Artemis is not part of the GMOD project, but Artemis does work with [Chado](#) (see the [Artemis-Chado Integration Tutorial](#)), and the Artemis group is working closely with [Apollo](#) developers on [Gbol](#). [Apollo](#) is GMOD's official genome [annotation](#) editor.