

Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

Fellowship for Interpretation of Genomes

RRID:SCR_004821

Type: Tool

Proper Citation

Fellowship for Interpretation of Genomes (RRID:SCR_004821)

Resource Information

URL: <http://thefig.info/>

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Description: Fonstein, Kogan, Osterman, Overbeek and Vonstein found the non-profit Fellowship for Interpretation of Genomes (FIG) to follow their vision of seeking advances in understanding through comparative analysis of the growing body of genomic data. The team at FIG began a new open source comparative genomics environment, the SEED. The Project to Annotate 1000 Genomes is a central goal of FIG. FIG is a nonprofit organization devoted to providing support for those analyzing genomes. It focuses on organizing the data needed to support interpretation of genomes, providing the infrastructure needed by the world community in its efforts to achieve understanding. FIG offers the key services required to architect and implement a comparative framework for interpreting genomes. In addition, we pick specific, critical problems and attempt to actively participate in the unraveling of the secrets within these amazing entities. It is only by merging the work of building infrastructure with the applications that use it that we will more deeply understand what is needed at each step.

Abbreviations: FIG

Resource Type: data or information resource, portal, topical portal

Resource Name: Fellowship for Interpretation of Genomes

Resource ID: SCR_004821

Alternate IDs: nlx_80790

Record Creation Time: 20220129T080226+0000

Record Last Update: 20260905T132525+0000

Ratings and Alerts

No rating or validation information has been found for Fellowship for Interpretation of Genomes.

No alerts have been found for Fellowship for Interpretation of Genomes.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.