

Resource Summary Report

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HiGlass

RRID:SCR_026687

Type: Tool

Proper Citation

HiGlass (RRID:SCR_026687)

Resource Information

URL: <https://github.com/higlass/higlass>

Proper Citation: HiGlass (RRID:SCR_026687)

Description: Web-based visual exploration and analysis of genome interaction maps.

Resource Type: web application, source code, software resource

Defining Citation: [PMID:30143029](#)

Keywords: visual exploration and analysis, genome interaction maps,

Funding: NCI U01CA200059;
NHGRI R00 HG007583;
NHGRI U54 HG007963

Availability: Free, Available for download, Freely available

Resource Name: HiGlass

Resource ID: SCR_026687

License: MIT license

Record Creation Time: 20250401T060330+0000

Record Last Update: 20260817T040727+0000

Ratings and Alerts

No rating or validation information has been found for HiGlass.

No alerts have been found for HiGlass.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 45 mentions in open access literature.

Listed below are recent publications. The full list is available at [dkNET](#) .

Fletcher C, et al. (2025) The genome sequence of the European smelt, *Osmerus eperlanus* (Linnaeus, 1758). Wellcome open research, 10, 118.

Mitchell R, et al. (2025) The genome sequence of a soldier beetle, *Malthinus seriepunctatus* Kiesenwetter, 1851. Wellcome open research, 10, 80.

Crowley LM, et al. (2025) The genome sequence of the lesser earwig, *Labia minor* (Linnaeus, 1758). Wellcome open research, 10, 90.

Hutchinson F, et al. (2025) The genome sequence of the Maple Pug moth, *Eupithecia inturbata* (Hübner, 1817). Wellcome open research, 10, 141.

Willink B, et al. (2025) The genome sequence of the Tropical Bluetail Damselfly, *Ischnura senegalensis* (Rambur, 1842). Wellcome open research, 10, 104.

Falk S, et al. (2025) The genome sequence of the Coppice Mining Bee, *Andrena helvola* (Linnaeus, 1758). Wellcome open research, 10, 102.

Crowley LM, et al. (2025) The genome sequence of the Twenty-plume Moth, *Alucita hexadactyla* Linnaeus, 1758. Wellcome open research, 10, 70.

Sivell O, et al. (2025) The genome sequence of the Marsh Click-beetle, *Actenicerus siaelandicus* (Müller, O.F., 1764). Wellcome open research, 10, 81.

Barclay MVL, et al. (2025) The genome sequence of a seed weevil, *Aspidapion aeneum* (Fabricius, 1775). Wellcome open research, 10, 41.

Bista I, et al. (2025) The genome sequence of the Antarctic lanternfish, *Electrona antarctica* (Günther, 1878). Wellcome open research, 10, 89.

Fletcher C, et al. (2025) The genome sequence of the Dotted Footman moth, *Pelosia muscerda* (Hufnagel, 1767). Wellcome open research, 10, 106.

O'Brien MF, et al. (2025) The genome sequence of the Common Pochard, *Aythya ferina* (Linnaeus, 1758). Wellcome open research, 10, 152.

Crowley LM, et al. (2025) The genome sequence of a chironomid fly, *Chironomus tentans* Fabricius, 1805. Wellcome open research, 10, 133.

Rammou DL, et al. (2025) The genome sequence of the European ground squirrel, *Spermophilus citellus* (Linnaeus, 1766). Wellcome open research, 10, 184.

Barclay MVL, et al. (2025) The genome sequence of a bark-dwelling beetle, *Silvanus unidentatus* (Olivier, 1790). Wellcome open research, 10, 136.

Sivell O, et al. (2025) The genome sequence of the Common Sheetweb Spider *Linyphia triangularis* (Clerck, 1757). Wellcome open research, 10, 92.

Hesson JC, et al. (2025) A chromosomal reference genome sequence for the northern house mosquito, *Culex pipiens form pipiens*, Linnaeus, 1758. Wellcome open research, 10, 107.

Falk S, et al. (2025) The genome sequence of a long-legged fly, *Scellus notatus* (Fabricius, 1781). Wellcome open research, 10, 117.

Pearce N, et al. (2025) The genome sequence of the European mole, *Talpa europaea* Linnaeus, 1758. Wellcome open research, 10, 98.

Barclay MVL, et al. (2025) The genome sequence of the false flower beetle, *Anaspis frontalis* (Linnaeus, 1758). Wellcome open research, 10, 82.