

Resource Summary Report

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PretextView

RRID:SCR_022024

Type: Tool

Proper Citation

PretextView (RRID:SCR_022024)

Resource Information

URL: <https://github.com/wtsi-hpag/PretextView>

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Description: Software tool to enable visualization from PretextView output. Desktop application for viewing pretext contact maps. OpenGL Powered PretextView Contact Map Viewer.

Synonyms: Wellcome Sanger Institute's High-Performance Assembly Group's Paired REad TEXTure Viewer, REad TEXTure Viewer

Resource Type: software application, software resource

Keywords: visualization from PretextView output, viewing pretext contact maps, map viewer

Availability: Free, Available for download, Freely available

Resource Name: PretextView

Resource ID: SCR_022024

License URLs: <https://github.com/wtsi-hpag/PretextView/blob/master/LICENCE.txt>

Record Creation Time: 20220421T050138+0000

Record Last Update: 20260810T040809+0000

Ratings and Alerts

No rating or validation information has been found for PretextView.

No alerts have been found for PretextView.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 448 mentions in open access literature.

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

Pennington LK, et al. (2026) A reference genome for Colusa grass, *Neostapfia colusana*, a threatened and endangered California vernal pool plant. *The Journal of heredity*, 117(3), 459.

Nguyen PT, et al. (2026) A genome assembly of the California poppy, *Eschscholzia californica*. *The Journal of heredity*, 117(2), 309.

Martínez-Gómez J, et al. (2026) The reference genome of a Sierra Nevada endemic, the cut-leaved monkeyflower, *Mimulus laciniatus* (syn. *Erythranthe lacinata*). *The Journal of heredity*, 117(2), 318.

Pezzi PH, et al. (2026) Chromosome-Level Reference Genome of the Beach False Foxglove, *Agalinis fasciculata* (Orobanchaceae). *Genome biology and evolution*, 18(5).

Payne CY, et al. (2026) A genome assembly for Lahontan Cutthroat Trout, *Oncorhynchus clarkii henshawi*. *The Journal of heredity*, 117(1), 159.

Vasileiadou K, et al. (2026) ERGA-BGE reference genome of *Diadema setosum*: the Black Longspine Urchin invading the Mediterranean sea. *Open research Europe*, 6, 75.

Kozak KM, et al. (2026) A highly contiguous genome assembly for the California vole, *Microtus californicus*, provides insight into phylogenetic relationships and patterns of synteny among voles. *The Journal of heredity*, 117(2), 347.

Richmond JQ, et al. (2026) A chromosome-level reference genome assembly for Gilbert's skink *Plestiodon gilberti*. *The Journal of heredity*, 117(1), 115.

Johns J, et al. (2026) A near chromosome-level assembly of the serpentine endemic columbine, *Aquilegia eximia*. *The Journal of heredity*, 117(1), 85.

Hernández-Roco D, et al. (2026) Chromosome-level genome assembly of the common tenrec, *Tenrec ecaudatus* (Schreber, 1778), a new model for early placental mammal evolution. *BMC genomics*, 27(1).

Linn IJ, et al. (2026) A high-quality reference genome for the ornate shrew (*Sorex ornatus*). *The Journal of heredity*, 117(3), 450.

Dorsey BL, et al. (2026) A genome assembly of Douglas' meadowfoam, *Limnanthes douglasii* R. BR. *The Journal of heredity*, 117(3), 592.

Ballare KM, et al. (2026) Genome assemblies of two bumble bee species: *Bombus sonorus* and *Bombus vosnesenskii*. *The Journal of heredity*, 117(3), 611.

Franchini P, et al. (2026) Highly contiguous chromosome-level assembly of the rock goby (*Gobius paganellus*) genome. *Scientific data*, 13(1).

Mochales-Riaño G, et al. (2025) Chromosome-level reference genome for the medically important Arabian horned viper (*Cerastes gasperettii*). *GigaScience*, 14.

Martínez P, et al. (2025) Screening a new European hake (*Merluccius merluccius*) chromosome-level genome assembly suggests an XX/XY sex-determining system driven by the SRY-box transcription factor 3 (*sox3*). *G3 (Bethesda, Md.)*, 15(8).

Aurelle D, et al. (2025) ERGA-BGE reference genome of *Eunicella cavolini*, an IUCN Near Threatened Gorgonian of the Mediterranean Sea. *Open research Europe*, 5, 323.

Genetti M, et al. (2025) Secrets of the Goo: The genome assembly of the Pacific banana slug, *Ariolimax columbianus*. *The Journal of heredity*, 116(3), 335.

Barclay MVL, et al. (2025) The genome sequence of a beetle, *Brachypterus glaber* (Newman, 1834). *Wellcome open research*, 10, 10.

Supple MA, et al. (2025) A chromosome-level genome assembly of the mountain lion, *Puma concolor*. *The Journal of heredity*, 116(4), 479.