

# Resource Summary Report

Generated by [RRID](#) on Aug 4, 2026

## [kent](#)

RRID:SCR\_026686

Type: Tool

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### Proper Citation

kent (RRID:SCR\_026686)

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### Resource Information

**URL:** <https://github.com/ucscGenomeBrowser/kent>

**Proper Citation:** kent (RRID:SCR\_026686)

**Description:** UCSC Genome Browser source tree. Stable branch: "beta".

**Resource Type:** software resource, source code

**Keywords:** UCSC Genome Browser source tree

**Availability:** Free, Available for download, Freely available

**Resource Name:** kent

**Resource ID:** SCR\_026686

**License:** MIT License

**License URLs:** <https://github.com/ucscGenomeBrowser/kent?tab=License-1-ov-file#readme>

**Record Creation Time:** 20250401T060330+0000

**Record Last Update:** 20260804T043920+0000

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### Ratings and Alerts

No rating or validation information has been found for kent.

No alerts have been found for kent.

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### Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Wooldridge TB, et al. (2025) Chromosome-scale Genomes Show Rapid Diversification and Ancient Gene Flow Among Bear Species. *Genome biology and evolution*, 17(11).

Teyssier E, et al. (2025) A plant Lysin Motif Receptor-Like Kinase plays an ancestral function in mycorrhiza. *Proceedings of the National Academy of Sciences of the United States of America*, 122(24), e2426063122.

Vanhaeren T, et al. (2025) Enhancing R-loop prediction with high-throughput sequencing data. *NAR genomics and bioinformatics*, 7(2), lqaf077.