MIRA

RRID:SCR_010731
Type: Tool

Proper Citation

MIRA (RRID:SCR_010731)

Resource Information

<table>
<thead>
<tr>
<th><strong>URL</strong></th>
<th><a href="http://sourceforge.net/p/mira-assembler/wiki/Home/">http://sourceforge.net/p/mira-assembler/wiki/Home/</a></th>
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<tbody>
<tr>
<td><strong>Description</strong></td>
<td>Sequence assembler and mapper for whole genome shotgun and EST/RNASeq sequencing data.</td>
</tr>
<tr>
<td><strong>Resource Name</strong></td>
<td>MIRA</td>
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<td><strong>Proper Citation</strong></td>
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<td><strong>Resource Type</strong></td>
<td>Resource, software resource</td>
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<td><strong>Resource ID</strong></td>
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<td><strong>Parent Organization</strong></td>
<td>SourceForge</td>
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<td><strong>References</strong></td>
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<td><strong>Website Status</strong></td>
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<td><strong>Alternate IDs</strong></td>
<td>OMICS_00023</td>
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<tr>
<td><strong>Abbreviations</strong></td>
<td>MIRA</td>
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<td><strong>Mentions Count</strong></td>
<td>624</td>
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</tbody>
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Ratings and Alerts

No rating or validation information has been found for MIRA.

No alerts have been found for MIRA.
Usage and Citation Metrics

We found 624 mentions in open access literature.

**Listed below are recent publications.** The full list is available at FDI Lab - SciCrunch Infrastructure.


Warmerdam DO, et al. (2020) PHF6 promotes non-homologous end joining and G2 checkpoint recovery. EMBO reports, 21(1), e48460.


Ball JW, et al. (2020) Fine-scale species distribution modelling and genotyping by sequencing to examine hybridisation between two narrow endemic plant species. Scientific reports, 10(1), 1562.


