Introduction

Bioinstaller is a comprehensive R package to manage bioinformatics software/script and database based on the R Shiny web application and the GitHub forum. Hundreds of bioinformatics tools/script and database has been included in Bioinstaller.

Feature:
- Easy to use
- User-friendly Shiny application
- Integrative platform of Databases and bioinformatics resources
- Open source and completely free
- One click to download and install Bioinformatics resources (via R, Shiny or OpenComp REST APIs)
- More attention for those software and database resource that have not been by other tools
- Logging
- System monitor
- Task submitting system
- Parallel tasks

Field
- Quality Control
- Alignment And Assembly
- Alternative Splicing
- ChIP-seq analysis
- Gene Expression Data Analysis
- Variant Detection
- Variant Annotation

Files in the PATH

<table>
<thead>
<tr>
<th>files</th>
<th>size</th>
<th>sidir</th>
<th>mtime</th>
<th>ctime</th>
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<tbody>
<tr>
<td>/usr/bin/zto-3</td>
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<td>2017-12-01T22:44:20Z</td>
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<tr>
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<td>2018-07-04T11:02:36Z</td>
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<td>false</td>
<td>2017-12-01T19:39:08Z</td>
<td>2017-12-01T22:44:16Z</td>
</tr>
</tbody>
</table>

Upload

Choose Need Uploaded File

File Type: txt
Genome Version: hg19
Description

Upload and update database successful!