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To the editorial staff at Bioinformatics,

We respectfully submit our manuscript, **"JBrowseR: An R Interface to the JBrowse 2 Genome Browser"** by Hershberg, Stevens, Diesh, Xie, De Jesus Martinez, Buels, Stein, and Holmes.

We are requesting your consideration for publication as an Application Note. We believe that JBrowseR constitutes a meaningful advance in genome analysis by providing a native R interface to a highly performant and extensible genome browser. This makes it possible, for the first time, to render a genome browser directly into an R graphics context, such as a Shiny app, an R Markdown document, or an interactive console.

We anticipate considerable option, given the widespread use of the R language in the bioinformatics community, and the fact that JBrowseR has already been downloaded over 200 times after less than a month on CRAN. Based on the scope and broad impact of Bioinformatics, we view the journal as a natural home for our manuscript describing JBrowseR.

Sincerely,

Ian Holmes