

COLLEGE OF VETERINARY MEDICINE AND BIOMEDICAL SCIENCES

Department of Environmental and Radiological Health Sciences

February 18, 2021

To the editors of *PLoS Computational Biology*:

We are pleased to submit an original "Ten Simple Rules" article entitled "Ten simple rules for finding and selecting R packages" for consideration for publication in *PLoS Computational Biology*.

Open-source software provides powerful and free tools to facilitate scientific research, including computational biology. Open-source programming languages like R are created and maintained by international teams, and a massive community of developers contribute extensions through open source packages that build on the language's basic framework. This process has facilitated an expansive set of tools available to researchers who learn R. However, many researchers new to R may find it difficult to navigate the available packages, to choose one that is robust, well-documented, and suited to their task. In this article, we present ten simple rules to help researchers new to R as they navigate this vast ecosystem of available R packages.

This manuscript has not been published and is not under consideration for publication elsewhere. We have not submitted closely related papers elsewhere.

Thank you for your consideration.

Sincerely,

**Brooke Anderson** 

GB Anderson

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