

S7 Text. Details on software packages used

How geographic access to care shapes disease burden: the current impact of post-exposure prophylaxis and potential for expanded access to prevent human rabies deaths in Madagascar

Rajeev et al. 2021✉

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Our analyses relied on a number of open-source software packages for the R programming language [1] including the following: coda [2], cowplot [3], data.table [4], doParallel [5], doRNG [6], dplyr [7], fasterize [8], forcats [9], foreach [10], gdistance [11], ggbeeswarm [12], ggforce [13], ggh4x [14], ggplot2 [15], ggridges [16], ggsn [17], glue [18], here [19], igraph [20], iterators [21], lubridate [22], magrittr [23], Matrix [24], patchwork [25], raster [26], readr [27], rgdal [28], rjags [29], rmapshaper [30], Rmpi [31], rngtools [32], scales [33], sf [34], sp [35], tidyr [36], triangle [37], RLang2019 [1]. For full details on implementation please see the archived repositories on [Zenodo](#) or on [Github](#).

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