Title: **Exploring biodiversity data is tidier than ever with {galah}**

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Over 112 million records of more than 150,000 Australian species are held within the Atlas of Living Australia (ALA), used by many for monitoring and conservation. Despite R’s popularity as a statistical tool in ecology and biology, constructing a coherent data query in R to search, filter and return data from biodiversity databases (including the ALA) has been fairly unintuitive, especially for new users. Translating queries into simple R syntax is difficult, so these R packages have often required a steeper learning curve to use them. {galah}, the ALA’s latest R package, makes query building simpler. It allows users to build queries just like how they wrangle their datasets with the{dplyr} package and the tidyverse. {galah} improves the transparency of shared code, promoting that data downloads are clear and reproducible. Here, we’ll demonstrate how {galah} makes use of pipes to make filtering and downloading biodiversity data easier, and discuss its implications for public and scientific communities.