



Antimicrobial Resistance (AMR) Data Analysis in R

Works on Windows, macOS and Linux
with all versions of R since R-3.0 (April 2013)

Completely dependency-free, highly suitable
for environments with limited resources

Provides the full microbiological taxonomy, and
reference data of all antimicrobial drugs

Integrates with EARS-Net, WHONET, ATC,
PubChem, LOINC and SNOMED CT