

Dear BEST-COST colleagues

To get started with the testing

0. Make sure you have access to the [BEST-COST GitHub repository](#). If you don't have access yet to the BEST-COST GitHub page, please provide us the email address of the GitHub account that we should give access to.
1. Install the package following the steps in the [README file at the landing page of the GitHub repository](#). For those that have already done so, note that now you have to specify `build_vignettes = TRUE` in the `install_github()` call.
2. Consult the *testing* vignette (i.e. documentation on how to do the testing) once you've installed the package by either
 - a. running `browseVignettes("healthiar")` in the console and clicking on HTML on the page that pops up
 - b. going to the "Packages" tab in RStudio, and clicking on *healthiar>User guides, package vignettes and other documentation>HTML*.
3. The testing vignette will guide you step by step through the process of adding a test!

And don't hesitate to reach out to us, we are happy to assist you! :)

Lastly, if you find anything unclear (instructions, function documentation, function argument names, ...) or have a suggestion to improve the package, please let us know, preferably via a GitHub issue (click on the green "New issue" button). The package is still under development, and we are grateful for any feedback!

Additional guidance on using the healthiar package

Prerequisite: *healthiar* installed

For more general guidance on how to use the *healthiar* package you may consult the *intro_to_healthiar* vignette (see step 2. above for how to access it).

For specific guidance on how to use a *healthiar* function either 1) run `"?healthiar::function_name"` (e.g. `?healthiar::attribute_health`) in the console or 2) go to the "Packages" tab in RStudio, click on *healthiar* and scroll to the function you are interested in.