

Living HTA

Automating HTA with R

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Living HEOR: Automating HTA with R

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Background The process of updating economic models is time-consuming and expensive, and often involves the transfer of sensitive data between parties. Here, we demonstrate how HEOR can be conducted in a way that allows companies to retain full control of their data, while automating reporting as new information becomes available.

Method We developed an automated analysis and reporting pipeline for health economic modelling and made the source code openly available on a GitHub repository. It consists of three parts:

- An economic model is constructed by the consultant using pseudo data (i.e. random data, which has the same format as the real data).
- On the company side, an application programming interface (API), generated using the R package plumber, is hosted on a server. An automated workflow is created. This workflow sends the economic model to the company API. The model is then run within the company server. The results are sent back to the consultant, and a (PDF) report is automatically generated using RMarkdown.
- This API hosts all sensitive data, so that data does not have to be provided to the consultant.

Results & Discussion The method is relatively complex, and requires a strong understanding of R, APIs, RMarkdown and GitHub Actions. However, the result is a process, which allows the consultant to conduct health economic (or any other) analyses on company data, without having direct access – the company does not need to share their sensitive data. The workflow can be scheduled to run at defined time points (e.g. monthly), or when triggered by an event (e.g. an update to the underlying data or model code). Results are generated automatically and wrapped into a full report. Documents no longer need to be revised manually.

Conclusions This example demonstrates that it is possible, within a HEOR setting, to separate the health economic model from the data, and automate the main steps of the analysis pipeline. We believe this is the first application of this procedure for a HEOR project.

Agenda

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1.

Background

What problems exist?

Background

There are several changes occurring simultaneously in health economics & decision science.

1) New technologies - shift to open-source packages (e.g. Rmarkdown, Shiny, BCEA, Heemod).

Previous presentations on Shiny to make R more accessible, but why R? Why not Python. APIs help mitigate this.

2) Big data - Increased use of large datasets, and concerns around data-security.

We will show a method of allowing companies to retain control of their data.

3) Big simulation – incorporating uncertainty in health economic models.

By running models on remote servers, we can easily scale up computing power.

4) Shift from recommendations from ‘static’ to ‘dynamic’ (Living)

“The ability to link real-world evidence with evidence-based practice will drive a shift from recommendations being produced at a single ‘static’ point in time to more dynamic, living guidance, and from health technology assessment to health technology management.” [NICE strategy 2021 to 2026: Dynamic, Collaborative, Excellent](#)



2.

Concept

How to implement this ...

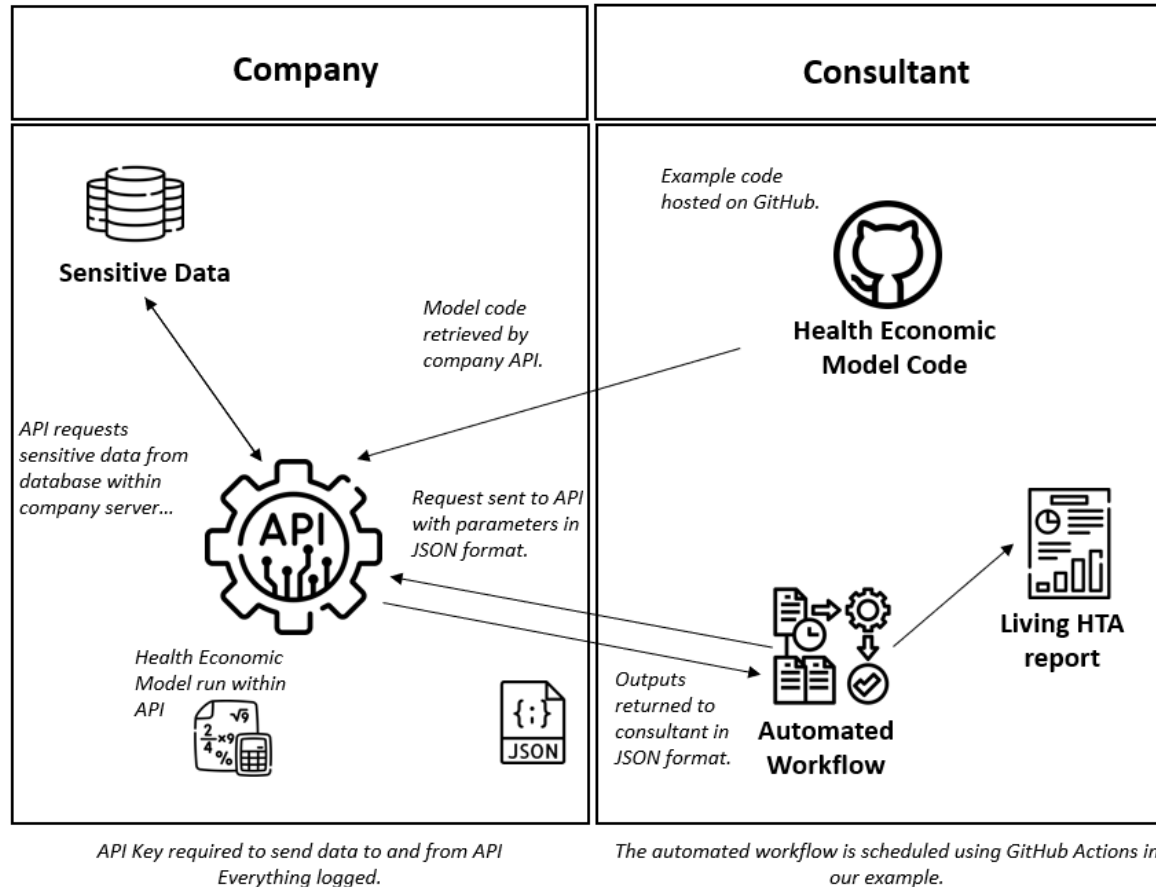
What we will show today ...

Using open-source software to generate an automated HTA workflow...

- Construct a script-based health economic models in R. Although could be Python, Ruby, Julia
- Host health economic model on GitHub repository, provide access to company.
- Host API on company server. The API will:
 - ... have access to sensitive data.
 - ... enable users to run R code on it.
 - ... allow users to pass R code to it.
 - ... return model results from it.
- Write script to pass model inputs & code to company API & generate a PDF report using markdown.
- Create a workflow which runs the script on schedule, or as new evidence emerges (e.g. when a data-base is updated).
- Contain all this functionality within a Shiny app – to enable non-programmers to interact with the API.



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3.

Hosting a HE model in an API

Creating an *Application Programming Interface* to host a health economic model using the plumber R package.

Hosting a HE model in an API

```
1 #####
2
3 library(dapack)
4 library(readr)
5 library(assertthat)
6
7 #* @apiTitle Client API hosting sensitive data
8 #*
9 #* @apiDescription This API contains sensitive data, the client does not
10 #* want to share this data but does want a consultant to build a health
11 #* economic model using it, and wants that consultant to be able to run
12 #* the model for various inputs
13 #* (while holding certain inputs fixed and leaving them unknown).
14
15 # Run the DARTH model
16 # @serializer csv
17 # @param path_to_psa_inputs is the path of the csv
18 # @param model_functions gives the github repo to source the model code
19 # @param param_updates gives the parameter updates to be run
20 # @post /runDARTHmodel
21 function(path_to_psa_inputs = "parameter_distributions.csv",
22          model_functions = paste0("https://raw.githubusercontent.com/",
23                                   "breesRed/plumberHE/main/R/darth_funcs.R"),
24          param_updates = data.frame(
25            parameter = c("p_RS1", "p_S18"),
26            distribution = c("beta", "beta"),
27            v1 = c(25, 50),
28            v2 = c(150, 70)
29          )) {
30
31
32   # source the model functions from the shared GitHub repo...
33   source(model_functions)
34
35   # read in the csv containing parameter inputs
36   psa_inputs <- as.data.frame(readr::read_csv(path_to_psa_inputs))
37
38   # for each row of the data-frame containing the variables to be changed...
39   for(n in 1:nrow(param_updates)){
40
41     # update parameters from API input
42     psa_inputs <- overwrite_parameter_value(
43       existing_df = psa_inputs,
44       parameter = param_updates[n,"parameter"],
45       distribution = param_updates[n,"distribution"],
46       v1 = param_updates[n,"v1"],
47       v2 = param_updates[n,"v2"])
48   }
49
50   # run the model using the single run-model function.
51   results <- run_model(psa_inputs)
52
53   # check that the model results being returned are the correct dimensions
54   # here we expect a single dataframe with 6 columns and 1000 rows
55   assertthat::assert_that(
56     all(dim(x = results) == c(1000, 6)),
57     class(results) == "data.frame",
58     msg = "Dimensions or type of data are incorrect,
59     please check the model code is correct or contact an administrator.
60     This has been logged"
61   )
62
63   # check that no data matching the sensitive csv data is included in the output
64   # searches through the results data-frame for any of the parameter names,
65   # if any exist they will flag a TRUE, therefore we assert that all = F
66   assertthat::assert_that(all(psa_inputs[, 1] %in%
67     as.character(unlist(x = results,
68                       recursive = T)) == F))
69
70   return(results)
71
72 }
```

Load necessary packages

Describe the API, used in Swagger

Roxygen style documentation for function, what are the inputs...

Source the model functions from GitHub

Overwrite default data with non-sensitive inputs

Run the model

Check the results object doesn't contain sensitive data

Return the results object



4.

Calling the API from R

Passing parameters and code to the health economic model & returning results...

Running the model – calling the API

```
1 # remove all existing data from the environment.
2 rm(list = ls())
3
4 library(ggplot2)
5 library(jsonlite)
6 library(httr)
7
8 # run the model using the connect server API
9 results <- httr::content(
10   httr::POST(
11     # the Server URL can also be kept confidential, but will leave here for now
12     url = "https://connect.bresmed.com",
13     # path for the API within the server URL
14     path = "rhta2022/runDARTHmodel",
15     # code is passed to the client API from GitHub.
16     query = list(model_functions =
17       paste0("https://raw.githubusercontent.com/",
18         "BresMed/plumberHE/main/R/darth_funcs.R")),
19     # set of parameters to be changed ...
20     # we are allowed to change these but not some others
21     body = list(
22       param_updates = jsonlite::toJSON(
23         data.frame(parameter = c("p_HS1", "p_S1H"),
24           distribution = c("beta", "beta"),
25           v1 = c(25, 50),
26           v2 = c(150, 100))
27       )
28     ),
29     # we include a key here to access the API ... like a password protection
30     config = httr::add_headers(Authorization = paste0("Key ",
31       Sys.getenv("CONNECT_KEY")))
32   )
33 )
34
35 # write the results as a csv to the outputs folder...
36 write.csv(x = results,
37   file = "outputs/darth_model_results.csv")
38
39 source("report/makeCEAC.R")
40 source("report/makeCEPlane.R")
41
42 # render the markdown document from the report folder,
43 # passing the results dataframe to the report.
44 rmarkdown::render(input = "report/darthreport.Rmd",
45   params = list("df_results" = results),
46   output_dir = "outputs")
```

Load necessary packages

Call the API:

- URL and path combine to identify where the API is.
- Query and body both allow for inputs to be provided to the API... We convert the data-frame of inputs to JSON first.
- Config allows us to add the KEY – which is hidden as an environment variable.
- **Result of the API is stored as an object (results).**

Write the results to a csv... not strictly necessary.

Render an Rmarkdown document based on the results of the API call, store the document in 'outputs' directory.



5.

Automating health economic model updates

Automating health economic model updates using GitHub Actions and R-markdown.

Automating health economic model updates

```
1 on:
2   push:
3     branches:
4       - main
5   schedule:
6     - cron: '1 1 1 * *'
7
8 name: Run DARTH model on client API
9 jobs:
10  createPullRequest:
11    runs-on: windows-2019
12    env:
13      GITHUB_PAT: ${ secrets.GITHUB_TOKEN }
14    # Load repo and install R
15    steps:
16      - uses: actions/checkout@master
17      - uses: r-lib/actions/setup-r@master
18
19      - name: Setup pandoc
20        uses: r-lib/actions/setup-pandoc@v2
21        with:
22          pandoc-version: '2.17.1.1'
23
24      - name: Install TinyTeX
25        uses: r-lib/actions/setup-tinytex@v2
26        env:
27          # install full prebuilt version
28          TINYTEX_INSTALLER: TinyTeX
29
30      - name: Install dependencies
31        run: |
32          install.packages(
33            c("reshape2", "jsonlite", "httr", "readr", "rmarkdown", "markdown")
34          )
35          install.packages(
36            "scales", dependencies = TRUE, repos = 'http://cran.rstudio.com/'
37          )
38          install.packages(
39            "ggplot2", dependencies = TRUE, repos = 'http://cran.rstudio.com/'
40          )
41        shell: Rscript {0}
42
43      - name: Run the model from API and create report
44        env:
45          CONNECT_KEY: ${ secrets.PLUMBER_SECRET }
46        run: |
47          source("scripts/run_darthAPI.R")
48        shell: Rscript {0}
49
50      - name: Create Pull Request
51        uses: peter-evans/create-pull-request@v3
52        with:
53          token: ${ secrets.GITHUB_TOKEN }
54          commit-message: Automated Model Run from API
55          title: 'Living HTA Automated Model Run'
56          body: >
57            Automated model run
58          labels: report, automated pr
```

Schedule jobs based upon a **push to the main branch, or at a scheduled time** (00:01 on 1st of the month).

Set-up code:

- Start running a Windows 2019 server.
- Checkout the repository.
- Set up R.
- Install all dependencies

Run the script querying the **API** and creating a report with Rmarkdown.

Create a **pull request** to the repository with the new results csv and markdown report included.



6.

Discussion

What are the implications of these methods...

What are the pros and cons of this framework.

Advantages of this framework

- **Security** - Data owners retain control of their data. No data need leave the data-owner's servers.
- **Transparency** - Separating the model code from the data can significantly improve the transparency of the health economic model. Many models could be passed to the data, not just one!
- **Computational Power** - The computational burden of the model is handled on a remote server.
- **Storage** – Larger datasets can be analyzed than would be possible on a laptop.
- **Living analysis** - API calls can be made at any time. A decision maker can see a report that will always reflect the data held by the company.

Disadvantages of this framework

- **Security** - Likely to remain concerns about data security, even with the authentication procedures built into the API functionality.
- **Transparency** - Risk that running the model remotely will result in the perception that the model is a 'black box' (I'd disagree!).
- **Coding practice** - The model code needs to be versatile enough to manage unknown data updates. *Proper testing will help mitigate these risks.*
- **Technical skillset** - This is not commonly implemented, or a common skill-set among health economists. Most models are not built in R.

- 1) What are the needs of different stakeholders? The company, the consultant, the HTA body, the ERG?
- 2) When is this likely to be most useful?
- 3) Is this too big a leap – can we pull HEOR this far?

Thank you

More information about this presentation can be found at:

Open-source code: [RobertASmithBresMed/plumberHE: Health Economics using Plumber APIs \(github.com\)](https://github.com/RobertASmithBresMed/plumberHE)

Draft paper: [plumberHE/academicpaper.pdf at main · RobertASmithBresMed/plumberHE \(github.com\)](https://github.com/RobertASmithBresMed/plumberHE/blob/main/plumberHE/academicpaper.pdf)

More information about the author's organizations can be found at:

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