Peer Models Network API Guide

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Introduction

Peer Models Network is an initiative to make models for healthcare decision-making accessible, transparent and interpretable to all British Columbians. With support from the Canada Foundation for Innovation, BC SUPPORT Unit and BC Academic Health Science Network, the Peer Models Network (PMN) aims to increase engagement between the modelling community and policymakers, interdisciplinary scholars, patients, journalists, and other members of the public.

The PMN centres around a software infrastructure called the Programmable Interface for Statistical and Simulation Models (PRISM). PRISM hosts models on the cloud and provides standard Application Programming Interfaces (APIs) that allows users of different levels of technical backgrounds to interact with models hosted on a cloud. Users can submit queries to cloud-hosted models through a multitude of software platforms, including Microsoft Excel, R, Python, bash scripts, Javascript and custom web apps. This enables direct access to users who lack either the software or hardware infrastructure or the technical expertise to set up models on their own devices. It also enables more technical users to integrate modes with other pieces of softwar

In its current implementation, PRISM can hosts models that are developed in R, including those that use Cpp and STAN.

Synchronous APIs calls are currently implemented and support for asynchronous calls that are required by extra long processing times is under active development.

ACCEPT

| Field | Value |
|----------------|---|
| Model Name | Acute COPD Exacerbation Prediction |
| | Tool (ACCEPT) |
| Modelling Team | RESP |
| Publication | doi:10.1016/S2213-2600(19)30397-2 |
| Purpose | Predict probability, rate, and severity |
| | of COPD exacerbations |
| Video | The ACCEPT Model in 90 Seconds |
| Interviews | Amin Adibi on the ACCEPT Model |
| | for COPD: Why and How |
| Media | COPD exacerbations: finally, a more |
| | than ACCEPTable risk score |
| Web App | ACCEPT web App |
| R Package | accept |
| Excel Sheet | PC Version |
| API User Guide | Link |

ADO

| Field | Value |
|----------------|---|
| Model Name | The ADO (age, dyspnoea, airflow obstruction) Index |
| Modelling Team | Puhan et al. |
| Publication | 10.1016/S0140-6736(09)61301-5 |
| Outcome | Mortality |
| Excel Sheet | PC Version |
| R Package | ado |
| API User Guide | Link |

BODE

| Field | Value |
|----------------|------------------------------------|
| Model Name | The Body-Mass Index, Airflow |
| | Obstruction, Dyspnea, and Exercise |
| | Capacity Index |
| Modelling Team | Celli et al. |
| Publication | 10.1056/NEJMoa021322 |
| Outcome | Mortality Risk |
| Excel Sheet | PC Version |
| R Package | bode |
| API User Guide | Link |

CFMortality

| Field | Value |
|-------------------------------|--|
| Model Name | Mortality prediction models in cystic fibrosis |
| Modelling Team Publication | Stanojevic et al. 10.1183/13993003.00224-2019 |
| Outcome | Mortality Risk |
| Excel Sheet | PC Version |
| R Package | cfmortality |
| API User Guide | Link |

CHDWilson

| Field | Value |
|----------------|--|
| Model Name | Prediction of Coronary Heart Disease Using Risk Factor Categories |
| Modelling Team | Wilson et al. |
| Publication | 10.1161/01.CIR.97.18.1837 |
| Outcome | Coronary Heart Disease |
| Excel Sheet | PC Version |
| R Package | chdwilson |
| API User Guide | Link |

CODEXCOPD

| Field | Value |
|----------------|------------------------------------|
| Model Name | The CODEX (comorbidity, |
| | obstruction, dyspnea, and previous |
| | severe exacerbations) Index |
| Modelling Team | Almagro et al. |
| Publication | 10.1378/chest.13-1328 |
| Outcome | Survival and readmission at both 3 |
| | months and 1 year after hospital |
| | discharge for a COPD exacerbation |
| R Package | codexcopd |
| API User Guide | Link |

COVIDSEIR

| Field | Value |
|----------------|---|
| Model Name | Bayesian SEIR model to estimate physical-distancing effects |
| Modelling Team | Anderson et al. |
| Publication | 10.1101/2020.04.17.20070086 |
| Outcome | COVID-19 Cases |
| R Package | covidseir |
| API User Guide | Link |

CVDAnderson

| Field | Value |
|----------------|--|
| Model Name | Prediction of Coronary Heart Disease Using Risk Factor Categories |
| Modelling Team | Anderson et al. |
| Publication | 10.1016/0002-8703(91)90861-B |
| Outcome | CHD, MI, CHD Mortality, Stroke, |
| | CVD, CVD Mortalitye |
| R Package | cvdanderson |
| API User Guide | Link |

DOSE

| Field | Value |
|----------------|---|
| Model Name | The DOSE (dyspnoea, obstruction, smoking, exacerbation) Index |
| Modelling Team | Jones et al. |
| Publication | 10.1164/rccm.200902-0271OC |
| Outcome | Mortality |
| Excel Sheet | PC Version |
| R Package | dose |
| API User Guide | Link |

EPIC

| Field | Value |
|----------------|---|
| Model Name | Evaluation Platform in COPD (EPIC) |
| Modelling Team | RESP |
| Publication | doi:10.1177%2F0272989X18824098 |
| Outcome | Patient-level outcomes, as well as mortality, prevalence, QALYs, costs, |
| | etc. |
| Video | The EPIC Model in 2 Minutes |
| Interviews | Mohsen Sadatsafavi on the EPIC |
| | Model |
| R Package | epicR |
| Excel Sheet | PC Version |
| API User Guide | Link |

FEV1

| Field | Value |
|----------------|----------------------------------|
| Model Name | Individualized prediction of |
| | lung-function decline in COPD |
| Modelling Team | RESP |
| Publication | doi:10.1503/cmaj.151483 |
| Outcome | Lung function over next 11 years |
| Web App | FEV1 web App |
| Excel Sheet | PC Version |
| R Package | fev1 |
| API User Guide | Link |

${\bf Framing ham FEV 1 Predictor}$

| Field | Value |
|----------------|--|
| Model Name | An Individualized Prediction Model for |
| | Long-term Lung Function Trajectory |
| | and Risk of COPD in the General |
| | Population |
| Modelling Team | RESP |
| Publication | doi:10.1016/j.chest.2019.09.003 |
| Outcome | Lung function over next 20 years |
| Web App | Lung Function Predictor for the |
| | General Population |
| API User Guide | Link |

ISARIC4C

| Field | Value |
|----------------|--|
| Model Name | prediction of in-hospital mortality or |
| | in-hospital clinical deterioration |
| | probability (defined as any requirement |
| | of ventilatory support or critical care, |
| | or death) for hospitalised COVID-19 |
| | patients |
| Modelling Team | ISARIC4C consortium |
| Publication | 10.1016/S2213-2600(20)30559-2 |
| Outcome | In-hospital mortality or in-hospital |
| | clinical deterioration probability |
| Excel Sheet | PC Version |
| R Package | isaric4c |
| API User Guide | Link |

MMRCEA

| Field | Value |
|----------------|---------------------------------------|
| Model Name | Cost-effectiveness evaluation of |
| | province-wide universal mismatch |
| | repair (MMR) reflex testing for lynch |
| | syndrome (LS) in British Columbia |
| Modelling Team | Centre for Clinical Epidemiology & |
| | Evaluation (C2E2) |
| Outcome | Costs, QALYs and life-years for the |
| | prevention and management of |
| | colorectal and endometrial cancers |
| | patients and their families |
| Excel Sheet | Pending |
| API User Guide | Link |

PLCOM2012

| Field | Value |
|----------------|---|
| Model Name | Prediction of 6-year probability of lung cancer in an individual person |
| Modelling Team | Tammemag et al. |
| Publication | 10.1056/NEJMoa1211776 |
| Outcome | Lung cancer risk score |
| Excel Sheet | PC Version |
| R Package | plcom 2012 |
| API User Guide | Link |

Chapter 17

QRISK3

| Field | Value |
|----------------|-------------------------------------|
| Model Name | 10-Year Cardiovascular Disease Risk |
| | Calculator |
| Modelling Team | Hippisley-Cox et al. |
| Publication | 10.1136/bmj.j2099 |
| Outcome | 10-yr risk of heart attack/stroke |
| R Package | QRISK3 |
| API User Guide | Link |

Chapter 18

API Users' Guide

18.1 ACCEPT API

Cloud Access through R

Users can access models on the Peer Models Network using the peermodels R package, available on GitHub. The following code snippet illustrates how you can run the model for example patients provided in the accept package:

```
remotes::install_github (resplab/peermodels)
library(peermodels)
input <- get_default_input("accept", api_key = YOUR_API_KEY)
results <- model_run(input)</pre>
```

Cloud Access through Python

```
import json
import requests
url = 'https://prism.peermodelsnetwork.com/route/accept/run'
headers = {'x-prism-auth-user': YOUR_API_KEY}
model_run = requests.post(url, headers=headers,
json = {"func":["prism_model_run"],"model_input":[{"ID": "10001","male": 1,"age": 57,"smoker": 0,
print(model_run)
results = json.loads(model_run.text)
print(results)
```

Cloud Access through Linux Bash

```
curl \
-X POST \
-H "x-prism-auth-user: REPLACE_WITH_API_KEY" \
-H "Content-Type: application/json" \
-d '{"func":["prism_model_run"],"model_input":[{"ID": "10001","male": 1,"age": 57,"smo.
https://prism.peermodelsnetwork.com/route/accept/run
```

18.2 BODE API

Cloud Access through R

Users can access models on the Peer Models Network using the peermodels R package, available on GitHub. The following code snippet illustrates how you can run the model for example patient provided in the bode package:

```
remotes::install_github (resplab/peermodels)
library(peermodels)
input <- get_default_input("bode", api_key = YOUR_API_KEY)
results <- model_run(input)</pre>
```

Cloud Access through Linux Bash

In Ubuntu, you can call the API with curl:

```
curl \
-X POST \
-H "x-prism-auth-user: REPLACE_WITH_API_KEY" \
-H "Content-Type: application/json" \
-d '{"func":["prism_model_run"],"model_input":[{"FEV1": 40,"mMRC": 3,"BMI": 22,"walk":
https://prism.peermodelsnetwork.com/route/bode/run
```

18.3 EPIC API

Cloud Access through R

Users can access models on the Peer Models Network using the peermodels R package, available on GitHub. The following code snippet illustrates how you can run the model for example patient provided in the epicR package:

```
remotes::install_github (resplab/peermodels)
library(peermodels)
input <- get_default_input("epic", api_key = YOUR_API_KEY)
results <- model_run(input)</pre>
```

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18.4 FEV1 API

Cloud Access through R

Users can access models on the Peer Models Network using the peermodels R package, available on GitHub. The following code snippet illustrates how you can run the model for example patient provided in the fev1 package:

```
remotes::install_github (resplab/peermodels)
library(peermodels)
input <- get_default_input("fev1", api_key = YOUR_API_KEY)
results <- model_run(input)</pre>
```

Cloud Access through Linux Bash

In Ubuntu, you can call the API with curl:

```
curl \
-X POST \
-H "x-prism-auth-user: REPLACE_WITH_API_KEY" \
-H "Content-Type: application/json" \
-d '{"func":["prism_model_run"],"model_input":[{"male":1,"age":70,"smoker":1,"FEV1":2.5,"height":
https://prism.peermodelsnetwork.com/route/fev1/run
```

18.5 CVDAnderson API

Cloud Access through R

Users can access models on the Peer Models Network using the peermodels R package, available on GitHub. The following code snippet illustrates how you can run the model for example patient provided in the cvdanderson package:

```
remotes::install_github (resplab/peermodels)
library(peermodels)
input <- get_default_input("cvdanderson", api_key = YOUR_API_KEY)
results <- model_run(input)</pre>
```

Cloud Access through Linux Bash

```
curl \
-X POST \
-H "x-prism-auth-user: REPLACE_WITH_API_KEY" \
-H "Content-Type: application/json" \
-d '{"func":["prism_model_run"],"model_input":[{"age": 33,"gender": 1,"Tchol": 230,"ECG_LVH": 0,'
https://prism.peermodelsnetwork.com/route/cvdanderson/run
```

18.6 CHDWilson API

Cloud Access through R

Users can access models on the Peer Models Network using the peermodels R package, available on GitHub. The following code snippet illustrates how you can run the model for example patient provided in the chdwilson package:

```
remotes::install_github (resplab/peermodels)
library(peermodels)
input <- get_default_input("chdwilson", api_key = YOUR_API_KEY)
results <- model_run(input)</pre>
```

Cloud Access through Linux Bash

In Ubuntu, you can call the API with curl:

```
curl \
-X POST \
-H "x-prism-auth-user: REPLACE_WITH_API_KEY" \
-H "Content-Type: application/json" \
-d '{"func":["prism_model_run"],"model_input":[{"age": 55,"gender": 1,"TChol": 250,"LD:
https://prism.peermodelsnetwork.com/route/chdwilson/run
```

18.7 CFMortality API

Cloud Access through R

Users can access models on the Peer Models Network using the peermodels R package, available on GitHub. The following code snippet illustrates how you can run the model for example patient provided in the chdwilson package:

```
remotes::install_github (resplab/peermodels)
library(peermodels)
input <- get_default_input("cfmortality", api_key = YOUR_API_KEY)
results <- model_run(input)</pre>
```

Cloud Access through Linux Bash

```
curl \
-X POST \
-H "x-prism-auth-user: REPLACE_WITH_API_KEY" \
-H "Content-Type: application/json" \
-d '{"func":["prism_model_run"],"model_input":[{"male": 0,"age": 57,"fvc": 66.7,"fev1"
https://prism.peermodelsnetwork.com/route/cfmortality/run
```

18.8 QRISK3 API

Cloud Access through R

Users can access models on the Peer Models Network using the peermodels R package, available on GitHub. The following code snippet illustrates how you can run the model for example patient provided in the QRISK3 package:

```
remotes::install_github (resplab/peermodels)
library(peermodels)
input <- get_default_input("qrisk3", api_key = YOUR_API_KEY)
results <- model_run(input)</pre>
```

18.9 CODEXCOPD API

Cloud Access through R

Users can access models on the Peer Models Network using the peermodels R package, available on GitHub. The following code snippet illustrates how you can run the model for example patient provided in the codexcopd package:

```
remotes::install_github (resplab/peermodels)
library(peermodels)
input <- get_default_input("codexcopd", api_key = YOUR_API_KEY)
results <- model_run(input)</pre>
```

Cloud Access through Linux Bash

In Ubuntu, you can call the API with curl:

```
curl \
-X POST \
-H "x-prism-auth-user: REPLACE_WITH_API_KEY" \
-H "Content-Type: application/json" \
-d '{"func":["prism_model_run"],"model_input":[{"age":40,"charlson":8,"FEV1":40,"mMRC":3,"exacer&
https://prism.peermodelsnetwork.com/route/codexcopd/run
```

18.10 MMRCEA API

Cloud Access through R

Users can access models on the Peer Models Network using the peermodels R package, available on GitHub. The following code snippet illustrates how you can run the model for example inputs provided by the modelling team:

library(peermodels)

```
input <- get_default_input("mmrcea", api_key = YOUR_API_KEY)</pre>
```

| input\$timeHorizon | <- 5 |
|---|----------|
| input\$discountRate | <- 0.015 |
| input\$averageFamilyReferred | <- 2 |
| input\$familyReferredHCP | <- 1 |
| input\$adherenceIntensiveScreening | <- 0.9 |
| input\$crcmmrDefReferredHCPIntArm | <- 0.3 |
| input\$crcmmrDefReferredHCPmmrbrafCompArmVCHA | <- 0.3 |
| input\$crcmmrDefReferredHCPmmrCompArmVCHA | <- 0.08 |
| input\$crcmmrDefReferredHCPmmrbrafCompArmVIHA | <- 0.08 |
| input\$crcmmrDefReferredHCPmmrCompArmOtherHA | <- 0.08 |
| input\$ecmmrDefReferredHCPIntArm | <- 0.8 |

model_run(input)

18.11 COVIDSEIR API (experimental)

We have recently deployed the Bayesian SEIR model to estimate physical-distancing effects developed by Anderson et al to our cloud infrastructure. The current version of the R package deployed to the PRISM server is covidseir v0.0.0.9006.

The current implementation of the covidseir package is experimental and has some limitations, given the computational intensity of the model. Depending on the number of chains and iterations, the html connection might be dropped for calls that more than a dozen minutes. Currently, fitting the model with default parameters takes about 5 minutes. We are actively working to implement asynchronous support which would email the results back to the user when hours long model runs are requested.

Cloud Access through R

Users can access models on the Peer Models Network using the peermodels R package, available on GitHub. The following code snippet illustrates how you can run the model for example patient provided in the covidseir package:

```
library(peermodels)
input <- get_default_input("covidseir", api_key = YOUR_API_KEY, bypass_router = T)
names(input)
input$daily_cases
results <- model_run(input)
names(results)
draw_plots()</pre>
```

The same workflow can be used to fit and run the model based on custom data. For example, the code snippet below, automatically downloads latest COVID-19 case numbers for British Columbia, and reruns the simulation using different assumptions.

```
library(stringr)
library(dplyr)
library(tidyr)
library(readr)
library(lubridate)
library(peermodels)
url <- "https://docs.google.com/spreadsheets/d/1ad7-09_Jn6AxsdkVPE33T-iLfGpPRmd3piXQqFiVeas/expon
CanadaCases <- read_csv(url)</pre>
covidCases <- CanadaCases %>% rename (name = "prname") %>% rename (Cases = "numconf") %>% mutat
  filter (name!="Canada") %>% filter (date!=today())
bcCases <- covidCases %>% filter (name == "British Columbia")
connect_to_model("covidseir", api_key = "YDbxcnNHmf4XoteSmCFHKx", bypass_router = T)
input <- get_default_input()</pre>
first <- length(bcCases$numtoday)-42+1</pre>
last <- length(bcCases$numtoday)</pre>
input$daily_cases <- bcCases$numtoday[first:last]</pre>
# Example assumed sampling fractions of positive cases:
s1 <- c(rep(0.1, 13), rep(0.2, length(input$daily_cases) - 13))</pre>
samp_frac_seg <- c(rep(1, 13), rep(2, length(input$daily_cases) - 13))</pre>
s2 <- rep(0.07, length(input$daily_cases)) # Assuming 7\% of positive individuals are hospitalized
input$samp_frac_fixed <- rep(0.1, length(input$daily_cases))</pre>
input$fit_iter <- 100
input$chains <- 1
inputf_fixed \leftarrow rep(0.1, 90)
results <- model_run(input)</pre>
draw_plots()
```

Cloud Access through Python

Users can use Python to access covidseir model on the cloud. The simple code snippet below runs the model with default inputs.

```
import json
import requests
url = 'http://model-covidseir.cp.prism-ubc.linaralabs.com/ocpu/library/covidseirPrism/?
headers = {'x-prism-auth-user': 'YOUR_API_KEY'}
default_input = requests.post(url, headers=headers,json = {"func":["get_default_input"]
# json_default_input = json.loads(default_input.json()[0]) # OR:
json_default_input = json.loads(json.loads(default_input.text)[0])
response = requests.post(url, headers=headers,json = {"func":["prism_model_run"], "moders=leaders]
```

Cloud Access through Linux Bash

In Ubuntu, you can call the API with curl:

```
curl \
-X POST \
-H "x-prism-auth-user: YOUR_API_KEY" \
-H "Content-Type: application/json" \
-d '{"func":["prism_model_run"],"model_input":{"daily_cases":[0,0,1,3,1,8,0,6,5,0,7,7,}
} ' \
http://model-covidseir.cp.prism-ubc.linaralabs.com/ocpu/library/covidseirPrism/R/gateway
```

18.12 PLCOM2012 API

Cloud Access through R

Users can access models on the Peer Models Network using the peermodels R package, available on GitHub. The following code snippet illustrates how you can run the model for example patient provided in the PLCOM2012 package:

```
remotes::install_github (resplab/peermodels)
library(peermodels)
input <- get_default_input("PLCOm2012", api_key = YOUR_API_KEY)
results <- model_run(input)</pre>
```

Cloud Access through Linux Bash

```
curl \
-X POST \
```

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18.13 ADO API

Cloud Access through R

Users can access models on the Peer Models Network using the peermodels R package, available on GitHub. The following code snippet illustrates how you can run the model for example patient provided in the ADO package:

```
remotes::install_github (resplab/peermodels)
library(peermodels)
input <- get_default_input("ado", api_key = YOUR_API_KEY)
results <- model_run(input)</pre>
```

Cloud Access through Linux Bash

In Ubuntu, you can call the API with curl:

```
curl \
-X POST \
-H "x-prism-auth-user: REPLACE_WITH_API_KEY" \
-H "Content-Type: application/json" \
-d '{"func":["prism_model_run"],"model_input":[{"fev":65, "mmrc":1, "age":41 }]}' \
http://model-ado.cp.prism-ubc.linaralabs.com/ocpu/library/adoPrism/R/gateway/json
```

18.14 DOSE API

Cloud Access through R

Users can access models on the Peer Models Network using the peermodels R package, available on GitHub. The following code snippet illustrates how you can run the model for example patient provided in the DOSE package:

```
remotes::install_github (resplab/peermodels)
library(peermodels)
input <- get_default_input("dose", api_key = YOUR_API_KEY)
results <- model_run(input)</pre>
```

Cloud Access through Linux Bash

In Ubuntu, you can call the API with curl:

```
curl \
-X POST \
-H "x-prism-auth-user: REPLACE_WITH_API_KEY" \
-H "Content-Type: application/json" \
-d '{"func":["prism_model_run"],"model_input":[{"mmrc":1, "fev":55, "smoking_status":1}
http://model-dose.cp.prism-ubc.linaralabs.com/ocpu/library/dosePrism/R/gateway/json
```

18.15 ISARIC4C API

Cloud Access through R

Users can access models on the Peer Models Network using the peermodels R package, available on GitHub. The following code snippet illustrates how you can run the model for example patient provided in the ISARIC4C package:

```
remotes::install_github (resplab/peermodels)
library(peermodels)
input <- get_default_input("isaric4c", api_key = YOUR_API_KEY)
results <- model_run(input)</pre>
```

Cloud Access through Linux Bash

```
curl \
-X POST \
-H "x-prism-auth-user: REPLACE_WITH_API_KEY" \
-H "Content-Type: application/json" \
-d '{"func":["prism_model_run"],"model_input":[{"age":69, "sex":1, "num_comorbidities"
http://model-dose.cp.prism-ubc.linaralabs.com/ocpu/library/isaric4cPrism/R/gateway/json
```