Health Analytics Challenge Lab: NHANES

Precision Analysis for Chronic Diseases

Authors/Group Members:

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Abstract:

The concept of precision health is a very promising field. Using the information from National Health and Examination Survey (NHANES), certain variables were extracted with respect to physical activity and nutritional data, and then this data was combined with data related to the chronic diseases: systolic blood pressure, diastolic blood pressure, coronary heart disease, breast cancer, and type 2 diabetes. This data was ingested into the risk analyzer and an analysis was made regarding the risk factors present for certain diseases with respect to bmi groups, gender, and ethnicity for the general population.

Introduction/Background:

There are a number of chronic diseases that plague our society but it's often hard to track the factors or maybe the specific cause for an individual who developed it. For our work we focused on data regarding nutritional intake and physical activity from the National Health and Nutrition Examination Survey (NHANES). NHANES takes surveys every year to gather information on a number of subjects to get a good view of how the population looks with respect to health and lifestyles.

First the physical data was cleaned, by removing all columns had more than 80% missingness. Afterwards the physical activity data was broken down into three groups using the missingness heatmap that was given in dataconstruction.Rmd. Categorical variables with low answers were removed, since they had very high log odds ratio and only represented a small portion of the population. A data dictionary as well as table was created in order.

Problems Tackled:

The goal of this project is to develop a tool to perform dynamic precision analysis of chronic diseases and find risk factors for subpopulations and the population as a whole.

Questions addressed:

- What physical activity/nutritional variables are associated with chronic diseases?
- How can we update the app interface for a more user friendly environment?

Data Description:

NHANES data

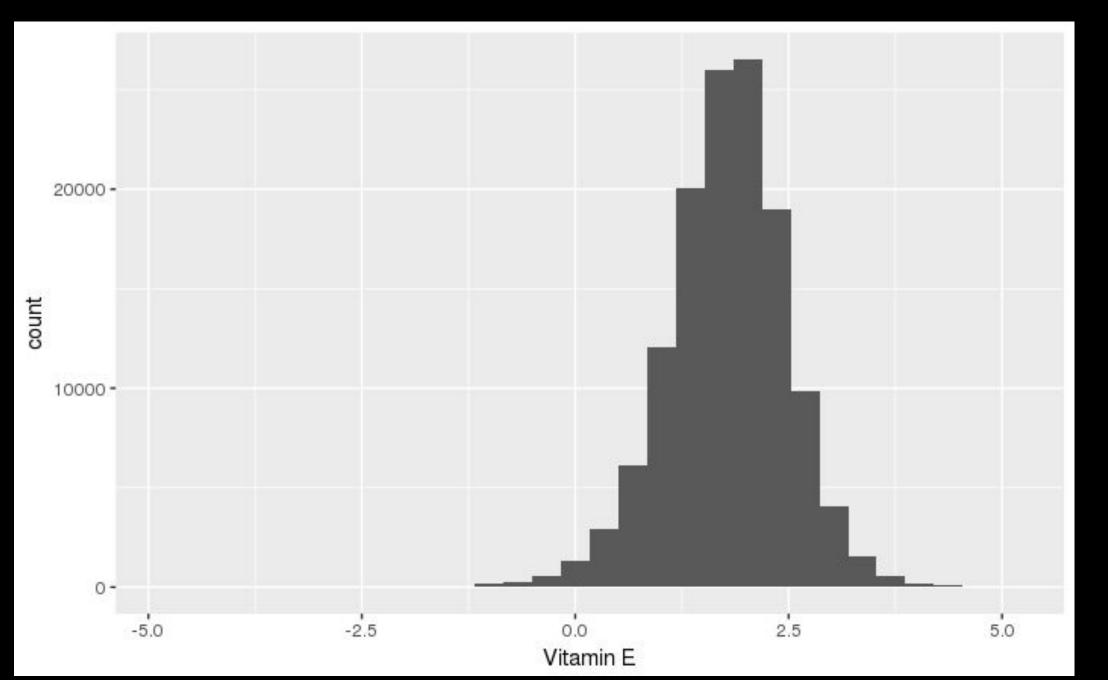
- Categorical and continuous values
- Taken on the entire population
- Stratified data collection
- Information on nutritional intake as well as physical activity
- added 16 new physical activity variables and 17 new nutritional variables



This is a density plot of carotene intake with diabetics and non diabetics. this displays a significant difference that can be used to determine if someone may be or will become diabetic

Data Cleaning

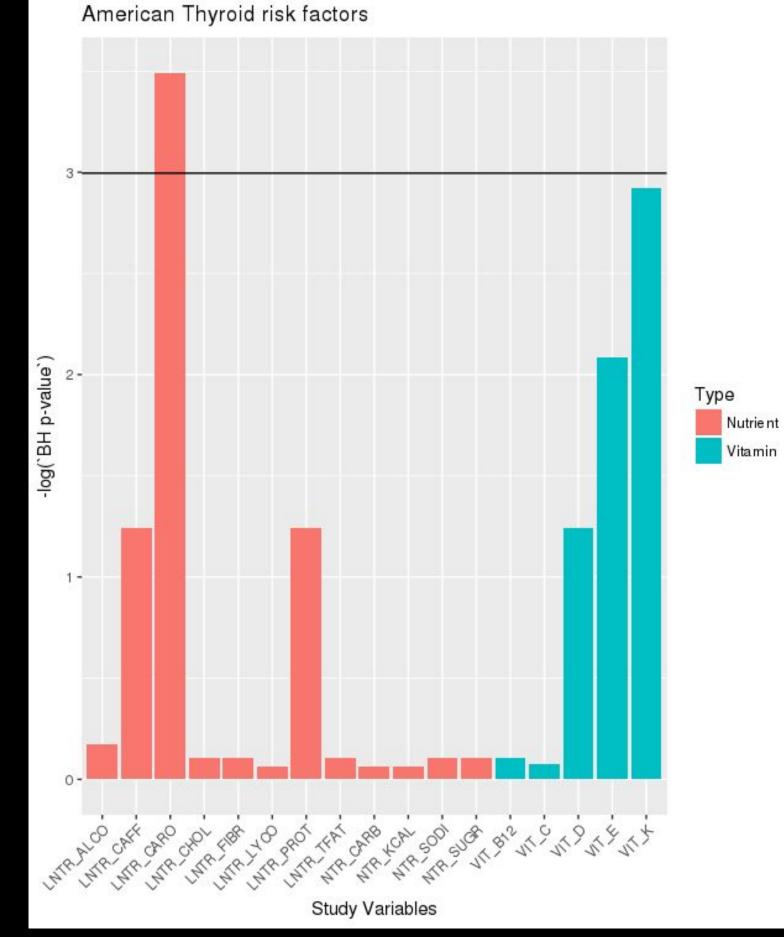
- Account for missing values
- Stitch together similar variables
- Transform skewed data
- Remove unnecessary variables
- Adjusted survey weights to account for multiple years of surveys put together



a histogram showing the distribution of vitamin E intake among the population. it is fairly normal so no transform is needed and it can be packaged up for the analysis by the risk analyzer.

Data Analytics Methods:

First the physical data was cleaned, by removing all columns that had more than 80% missingness. Afterwards the physical activity data was broken down into three groups using the missingness heatmap that was given in dataconstruction.Rmd. Categorical variables with low answers were removed, since they had very high log odds ratio and only represented a small portion of the population. A data dictionary as well as table was created in order to differntiate between categorical and numerivcal variables in the risk analyzer. With the nutritional data, after checking for important nutritional variables from the NHANES documentation, logistic regression was used to determine important factors. Afterwards the nutritional data were grouped by similar missingness using the documentation on the NHANES website.

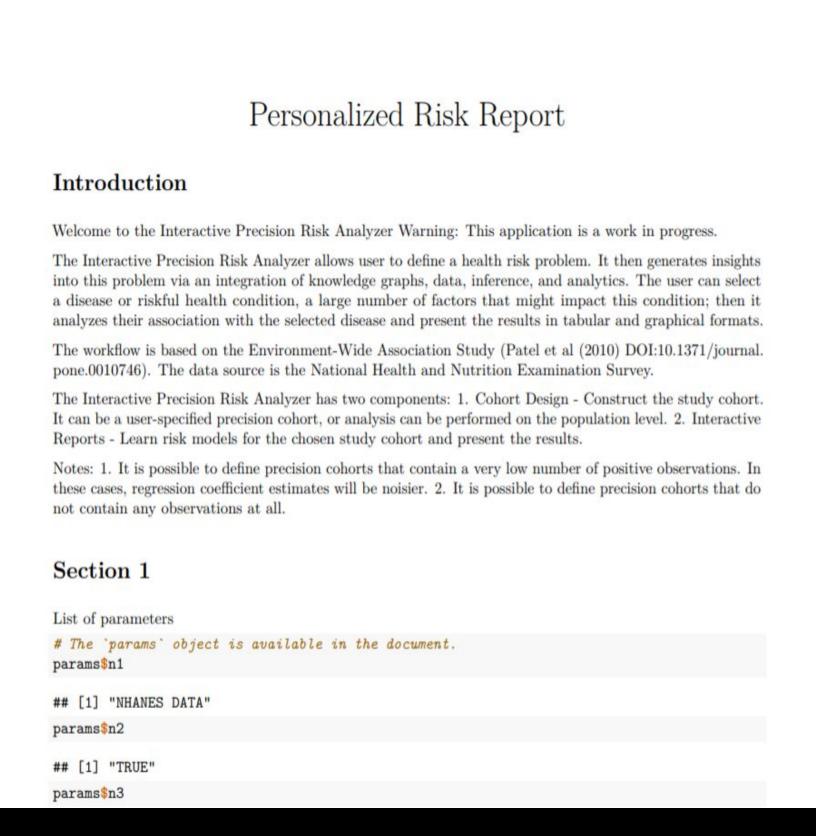


This is a bar graph shows relative BH p-values of diffent nutritional factors and their relation to thyroid condition on the population as whole. plots like these are made in the risk analyzer and you can see what factors have the highest association and a line is given on the graph to help evaluate what is considered highly associated. These plots are then made for other subpopulations that will be analyzed in the same way and comparisons are made that we can draw higher level conclusions about with regards to how different groups of people are affected with chronic illnesses.

Experimental Results:

The risk analyzer is used to find risk factors for diseases thyroid condition, type 2 diabetes, systolic blood pressure, diastolic blood pressure, breast cancer, and coronary heart disease given control variables, such as age, income, ethnicity, BMI, gender, serum cotinine, and menopausal status. One of which was based on certain subpopulations. In order to see what variables are important we made use of the *p*-values, and set the threshold to 0.05. Afterwards the risk factors were shown for the certain diseases with respect to the whole population. The results of what was found is shown in the table below. Later we found risk factors for certain diseases with respect to subpopulations like bmi groups, ethnicity groups, age

groups, and gender.



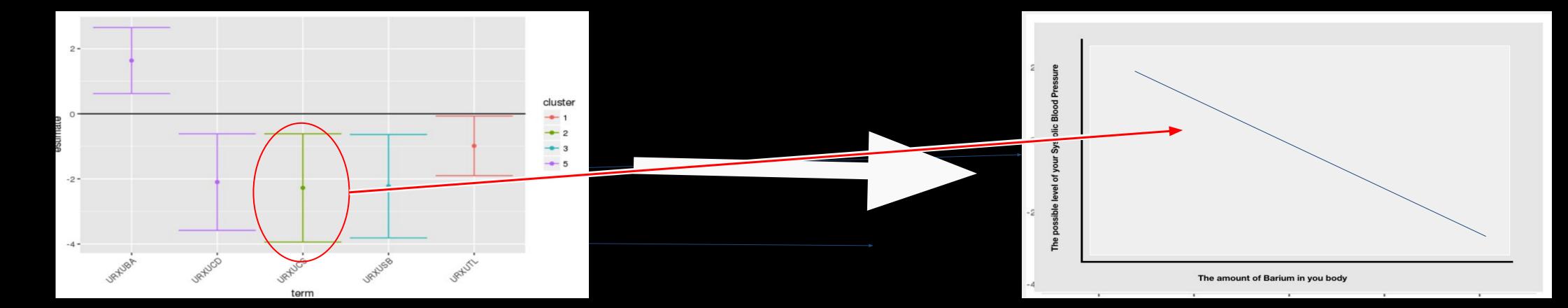
	most people your age	active past 30 days than in past 12 months		
Protein	Unable to do tasks around home/yard for at least 10 minutes in the last 30 days		Alcohol	Protein
Total Sugars	Unable to do muscle strengthening activities for at least 10 minutes over the past 30 days.		Energy (kcal)	Alcohol
Less active than most people your age	unable to do moderately intense activities for at least 10 minutes over past 30 days			Caffeine
Do not engage in vigorous intensity sports for at least 10 minutes on a daily basis	unable to do vigorous activities over the past 30 days for at least 10 minutes.			Total Fat
	do not walk or use bicycle for at least 10 minutes to get to places			Carbohydrat e
	Hours watching tv and computer use in past 30 days			Sugars
				Energy(kcal)

Personal risk report

Discussion:

These are some examples of risk factors for chronic illnesses for the population as a whole

The findings of the risk analyzer and secondary analysis of those outputs can be important for determining how a sub population is affected by certain lifestyle factors causing chronic illness. The tests can be run on a wide array of factors and the personalized output method can make the response important for a common person or even a healthcare professional.



With the current risk analyzer the output is more expert focused and shows a series of regression coefficients for all important factors but for a normal patient a new plot should be made that can easily show the effect of a single factor.

Conclusion & Recommendations:

The whole point of the shiny app that analyzes the data is to make recommendations to provide context and suggestions on what changes can be made to improve the life of an individual or maybe even a larger community as a whole. In the end expanding the functionality of this app will help it be a tool to help people in all sorts of groups.

References/Citations:

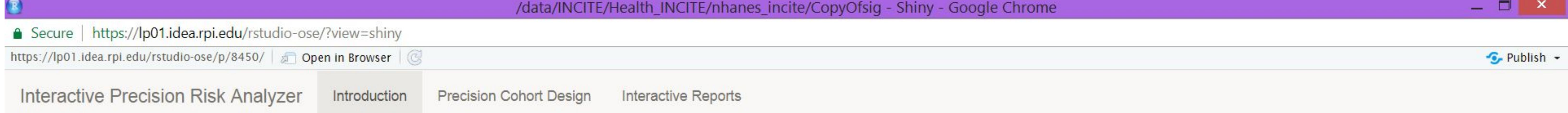
Alexander, Breneman, Curt, Bennett, and Kristin P. "Cadre Modeling: Simultaneously Discovering Subpopulations and Predictive Models." [1402.1128] Long Short-Term Memory Based Recurrent Neural Network Architectures for Large Vocabulary Speech Recognition. February 07, 2018. Accessed August 08, 2018. https://arxiv.org/abs/1802.02500.

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No. J. Sun, J. Hu, and F. Wong, "Demonalized Project Biobank and Project Biobank and Project Biobank and Project Biobank." Advances in Pediatrics and Project Biobank and Project Biobank and Project Biobank and Project Biobank and Project Biobank. Accessed August 08, 2018. https://www.cdc.gov/nchs/nhanes/index.htm.

No. J. Sun, J. Hu, and F. Wong, "Demonalized Project Biobank and Project Biob

Ng, K., J. Sun, J. Hu, and F. Wang. "Personalized Predictive Modeling and Risk Factor Identification Using Patient Similarity." Advances in Pediatrics. March 25, 2015. Accessed August 08, 2018. https://www.ncbi.nlm.nih.gov/pubmed/26306255. Patel, C. J., J. Bhattacharva, and A. J. Butte. "An Environment-Wide Association Study (EWAS) on Type 2 Diabetes Mellitus." Advances in Pediatrics. May 20, 2010. Accessed August 08, 2018. https://www.ncbi.nlm.nih.gov/pubmed/20505766.



Welcome to the Interactive Precision Risk Analyzer

Warning: This application is a work in progress.

The Interactive Precision Risk Analyzer allows user to define a health risk problem. It then generates insights into this problem via an integration of knowledge graphs, data, inference, and analytics. The user can select a disease or riskful health condition, a large number of factors that might impact this condition; then it analyzes their association with the selected disease and present the results in tabular and graphical formats.

The workflow is based on the Environment-Wide Association Study (Patel et al (2010) DOI:10.1371/journal.pone.0010746). The data source is the National Health and Nutrition Examination Survey.

The Interactive Precision Risk Analyzer has two components:

- 1. Cohort Design Construct the study cohort. It can be a user-specified precision cohort, or analysis can be performed on the population level.
- 2. Interactive Reports Learn risk models for the chosen study cohort and present the results.

Notes:

- 1. It is possible to define precision cohorts that contain a very low number of positive observations. In these cases, regression coefficient estimates will be noisier.
- 2. It is possible to define precision cohorts that do not contain any observations at all.

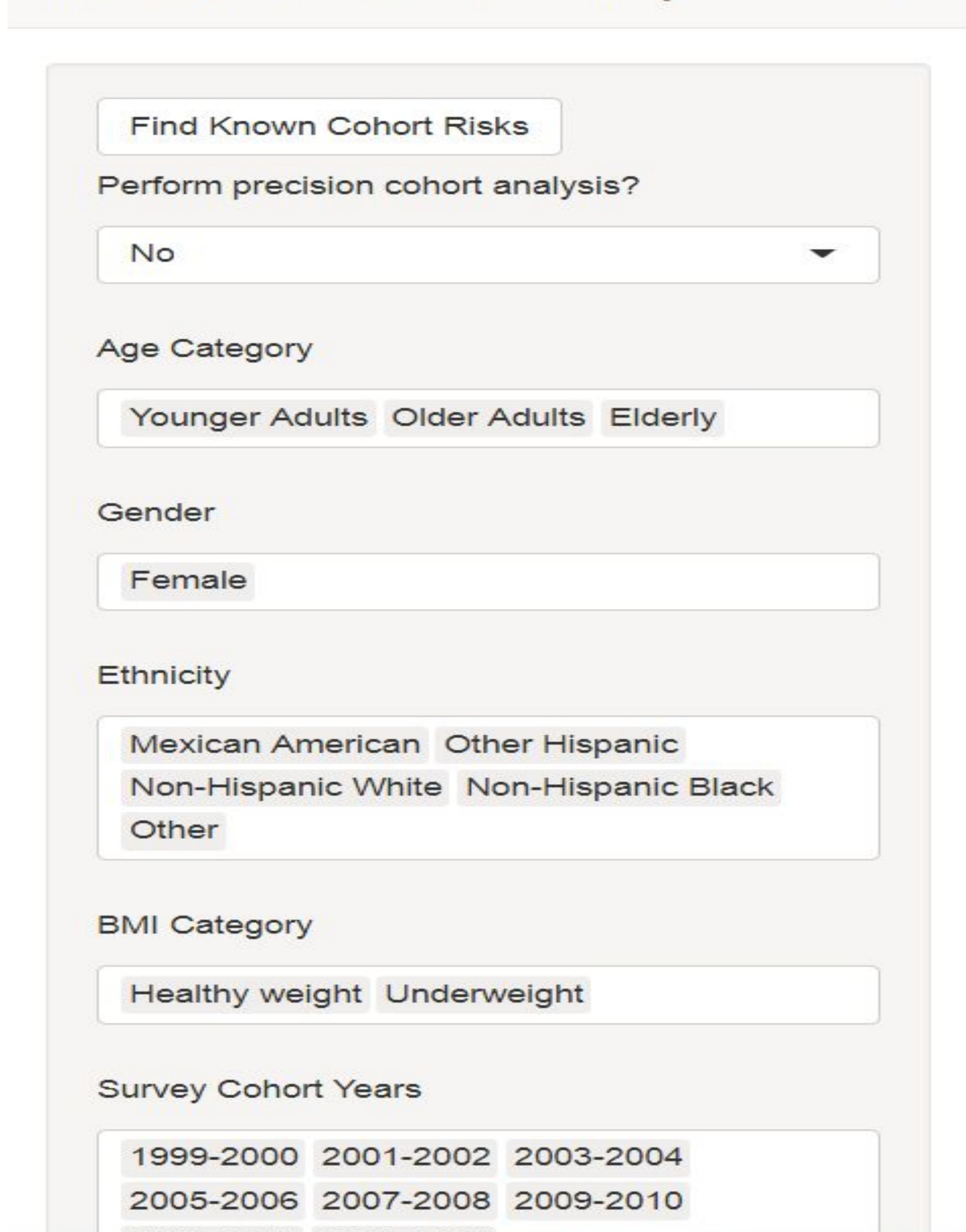
Secure https://lp01.idea.rpi.edu/rstudio-ose/?view=shiny

https://lp01.idea.rpi.edu/rstudio-ose/p/8450/ @ Open in Browser

Interactive Precision Risk Analyzer Introduction

Precision Cohort Design

Interactive Reports



Personalized Risk Report

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Section 1

```
List of parameters

# The `params` object is available in the document.

params$n1

## [1] "NHANES DATA"

params$n2
```

[1] "TRUE"

params\$n3

Significant Study Variable Regression Coefficients With 95% Confidence In

