

Actionable AI_2

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Project Status: Completed

Project Description:

The project starts by first importing the coefficient csv file as saving it as a dataframe called “df”. some slight process of the dataframe occurred where in the code column, the value “Intercept” was added. besides the coefficient donating the value of intercept.

from that dataframe, another dataframe was created called “df_2”, where the only columns present were those of interest for the project. the columns are as follow:

- The “Response” column
- The “strata” column
- The “coef” column

df_2 was then pivoted, where each strata is now a column instead of a row. this was saved in another variable called “df_3”.

After that we created the “coefficient missing” table/dataframe. This is a table/dataframe where if a certain feature is missing in the history csv, we should take coefficient values from this table not from the original coefficient table, since the original coefficient table assumes that each feature that was used to do its regression is present. However, as directed by vlad, that will not always be the case. The original “coefficient missing” file should be provided by the missing values project teams, but until they do provide it, we were instructed to create a mock up file.

the coefficient missing file was created by iterating over the features of each drug and removing one feature at a time, essentially creating a dataframe, “coeff_missing”, where each drug has all coefficient values for each of its features except the one feature that was removed. the coefficient values were then randomized between the minimum value present and the maximum value.

Next was creating a history csv. a csv file simulating patients. the csv file was in binarized form, indicating what features the patient had and which he did not. it was randomized with one feature in each row being NaN to simulate missing data.

After the history csv was created, code was done that would calculate the probability for each drug for each patient. the coefficient values used would first detect if there is any features missing for a given drug for that patient, and then accordingly would take the coefficient from the correct dataframe(the coeff_missing if there is a feature missing, df_2 if not).

After that, four dictionaries were created:

- pat_drug_dict: basically a dictionary here the key is the patient's ID and the value is dataframe where each row is a drug and it's probability
- pat_features: a dictionary where the key is the patient ID and the values are the features/strata that the patient presented with
- links_dict: a dictionary where the key is the drug and the value is the link of the drug on the NCBI
- pat_rec: a dictionary where the key is the patient ID and the value is a tuple containing the NCBI link and the drug name

A plotting function was created which graphs the relative probabilities of each drug in a horizontal bar chart

and finally the communicating function, which is a code that takes in the patient id, a code that summarizes the features that the patient has been presented with, the drug recommendation and it's NCBI link, the plot of the relative probabilities of each drug and finally some text that we were tasked to display.

```
[1]: ### Importing the Modules required
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
from pathlib import Path
import math
import random
from functools import reduce
import requests
```

```
[2]: ### Importing the coefficient Excel File
excel_path = Path("C:\Sem 3\HAP 786\coefficients.xlsx")
df = pd.read_excel(excel_path)
```

```
C:\Users\Bido_\AppData\Roaming\Python\Python311\site-
packages\openpyxl\worksheet\header_footer.py:48: UserWarning: Cannot parse
header or footer so it will be ignored
warn("""Cannot parse header or footer so it will be ignored""")
```

```
[3]: #### Processing the dataframe to account for the intercept
df["code"] = df["code"].fillna("Intercept")
df["strata"] = df["strata"].fillna("Intercept")
```

```
[4]: df[df["strata"] == "Intercept"]
```

```
[4]:
```

	Response	var_id	cgrp	ctype	\
419	FLUOXETINE	0	NaN	NaN	
420	MIRTAZAPINE	0	NaN	NaN	
421	NORTRIPTYLINE	0	NaN	NaN	
488	PAROXETINE	0	NaN	NaN	
489	ROPINIROLE	0	NaN	NaN	

490		SERTRALINE	0	NaN	NaN
491		TRAZODONE	0	NaN	NaN
587		DOXEPIN	0	NaN	NaN
588		DULOXETINE	0	NaN	NaN
589		ESCITALOPRAM	0	NaN	NaN
621		CITALOPRAM	0	NaN	NaN
622		DESVENLAFAXINE	0	NaN	NaN
630		VENLAFAXINE	0	NaN	NaN
631	Less common antidepressants or combination of ...		0	NaN	NaN
1414		BUPROPION	0	NaN	NaN
1439		AMITRIPTYLINE	0	NaN	NaN

	code	coef	likelir	vif	description	notes	Proper	\
419	Intercept	-0.10	NaN	NaN	Intercept	Intercept	Intercept	
420	Intercept	-1.60	NaN	NaN	Intercept	Intercept	Intercept	
421	Intercept	-2.59	NaN	NaN	Intercept	Intercept	Intercept	
488	Intercept	-0.35	NaN	NaN	Intercept	Intercept	Intercept	
489	Intercept	-0.94	NaN	NaN	Intercept	Intercept	Intercept	
490	Intercept	0.04	NaN	NaN	Intercept	Intercept	Intercept	
491	Intercept	-4.08	NaN	NaN	Intercept	Intercept	Intercept	
587	Intercept	-1.61	NaN	NaN	Intercept	Intercept	Intercept	
588	Intercept	-0.55	NaN	NaN	Intercept	Intercept	Intercept	
589	Intercept	-1.66	NaN	NaN	Intercept	Intercept	Intercept	
621	Intercept	-0.04	NaN	NaN	Intercept	Intercept	Intercept	
622	Intercept	0.28	NaN	NaN	Intercept	Intercept	Intercept	
630	Intercept	0.02	NaN	NaN	Intercept	Intercept	Intercept	
631	Intercept	-0.71	NaN	NaN	Intercept	Intercept	Intercept	
1414	Intercept	-3.12	NaN	NaN	Intercept	Intercept	Intercept	
1439	Intercept	-4.09	NaN	NaN	Intercept	Intercept	Intercept	

	Revised	strata	antidep_link
419	Intercept	Intercept	https://www.ncbi.nlm.nih.gov/books/NBK459223/
420	Intercept	Intercept	https://www.ncbi.nlm.nih.gov/books/NBK519059/
421	Intercept	Intercept	https://www.ncbi.nlm.nih.gov/books/NBK482214/
488	Intercept	Intercept	https://www.ncbi.nlm.nih.gov/books/NBK526022/
489	Intercept	Intercept	https://www.ncbi.nlm.nih.gov/books/NBK554532/
490	Intercept	Intercept	https://www.ncbi.nlm.nih.gov/books/NBK547689/
491	Intercept	Intercept	https://www.ncbi.nlm.nih.gov/books/NBK470560/
587	Intercept	Intercept	https://www.ncbi.nlm.nih.gov/books/NBK542306/
588	Intercept	Intercept	https://www.ncbi.nlm.nih.gov/books/NBK549806/
589	Intercept	Intercept	https://www.ncbi.nlm.nih.gov/books/NBK557734/
621	Intercept	Intercept	https://www.ncbi.nlm.nih.gov/books/NBK482222/
622	Intercept	Intercept	https://www.ncbi.nlm.nih.gov/books/NBK534829/
630	Intercept	Intercept	https://www.ncbi.nlm.nih.gov/books/NBK548799/
631	Intercept	Intercept	https://www.ncbi.nlm.nih.gov/books/NBK538182/
1414	Intercept	Intercept	https://www.ncbi.nlm.nih.gov/books/NBK470212/
1439	Intercept	Intercept	https://www.ncbi.nlm.nih.gov/books/NBK537225/

```
[5]: ##### Taking only the the required columns
df_2 = df[["Response"] + ["strata"] + ["coef"]]
df_2 = df_2.reset_index(drop = True)
df_2[df["strata"] == "Intercept"]
```

```
[5]:
```

	Response	strata	coef
419	FLUOXETINE	Intercept	-0.10
420	MIRTAZAPINE	Intercept	-1.60
421	NORTRIPTYLINE	Intercept	-2.59
488	PAROXETINE	Intercept	-0.35
489	ROPINIROLE	Intercept	-0.94
490	SERTRALINE	Intercept	0.04
491	TRAZODONE	Intercept	-4.08
587	DOXEPIN	Intercept	-1.61
588	DULOXETINE	Intercept	-0.55
589	ESCITALOPRAM	Intercept	-1.66
621	CITALOPRAM	Intercept	-0.04
622	DESVENLAFAXINE	Intercept	0.28
630	VENLAFAXINE	Intercept	0.02
631	Less common antidepressants or combination of ...	Intercept	-0.71
1414	BUPROPION	Intercept	-3.12
1439	AMITRIPTYLINE	Intercept	-4.09

```
[6]: ### Pivoting the table so that each feature becomes a column to ease data
      ↪ processing
df_3 = df_2.pivot_table(index = "Response", columns = "strata", values =
      ↪ "coef", aggfunc = "sum")
df_3
```

```
[6]: strata
```

	Intercept \
Response	
AMITRIPTYLINE	-4.09
BUPROPION	-3.12
CITALOPRAM	-0.04
DESVENLAFAXINE	0.28
DOXEPIN	-1.61
DULOXETINE	-0.55
ESCITALOPRAM	-1.66
FLUOXETINE	-0.10
Less common antidepressants or combination of a...	-0.71
MIRTAZAPINE	-1.60
NORTRIPTYLINE	-2.59
PAROXETINE	-0.35
ROPINIROLE	-0.94
SERTRALINE	0.04
TRAZODONE	-4.08
VENLAFAXINE	0.02

strata	aa:dem:ager_1319 \
Response	
AMITRIPTYLINE	-0.71
BUPROPION	NaN
CITALOPRAM	-0.53
DESVENLAFAXINE	-0.32
DOXEPIN	-0.36
DULOXETINE	-0.40
ESCITALOPRAM	-0.19
FLUOXETINE	-0.54
Less common antidepressants or combination of a...	-0.46
MIRTAZAPINE	-0.38
NORTRIPTYLINE	-0.54
PAROXETINE	-0.47
ROPINIROLE	NaN
SERTRALINE	-0.48
TRAZODONE	NaN
VENLAFAXINE	-0.40

strata	aa:dem:ager_2040 \
Response	
AMITRIPTYLINE	-0.40
BUPROPION	-0.10
CITALOPRAM	-0.17
DESVENLAFAXINE	-0.13
DOXEPIN	-0.25
DULOXETINE	-0.10
ESCITALOPRAM	-0.08
FLUOXETINE	-0.18
Less common antidepressants or combination of a...	-0.21
MIRTAZAPINE	-0.17
NORTRIPTYLINE	-0.20
PAROXETINE	-0.13
ROPINIROLE	-0.30
SERTRALINE	-0.16
TRAZODONE	NaN
VENLAFAXINE	-0.06

strata	aa:dem:ager_6579 \
Response	
AMITRIPTYLINE	NaN
BUPROPION	NaN
CITALOPRAM	NaN
DESVENLAFAXINE	NaN
DOXEPIN	0.22
DULOXETINE	-0.13

ESCITALOPRAM	-0.13
FLUOXETINE	0.10
Less common antidepressants or combination of a...	0.14
MIRTAZAPINE	NaN
NORTRIPTYLINE	NaN
PAROXETINE	0.04
ROPINIROLE	NaN
SERTRALINE	0.03
TRAZODONE	NaN
VENLAFAXINE	NaN

strata	aa:dem:ager_8089 \
Response	
AMITRIPTYLINE	-0.40
BUPROPION	NaN
CITALOPRAM	-0.35
DESVENLAFAXINE	NaN
DOXEPIN	NaN
DULOXETINE	-0.50
ESCITALOPRAM	-0.43
FLUOXETINE	-0.24
Less common antidepressants or combination of a...	0.05
MIRTAZAPINE	-0.33
NORTRIPTYLINE	-0.57
PAROXETINE	-0.18
ROPINIROLE	NaN
SERTRALINE	-0.24
TRAZODONE	NaN
VENLAFAXINE	-0.24

strata	aa:dem:gender_F \
Response	
AMITRIPTYLINE	-0.15
BUPROPION	-0.07
CITALOPRAM	NaN
DESVENLAFAXINE	NaN
DOXEPIN	NaN
DULOXETINE	0.06
ESCITALOPRAM	-0.02
FLUOXETINE	0.04
Less common antidepressants or combination of a...	0.08
MIRTAZAPINE	-0.15
NORTRIPTYLINE	-0.20
PAROXETINE	NaN
ROPINIROLE	-0.15
SERTRALINE	0.03
TRAZODONE	NaN

VENLAFAXINE	0.04
strata aa:meas:adrm_1SR \	
Response	
AMITRIPTYLINE	NaN
BUPROPION	1.91
CITALOPRAM	0.29
DESVENLAFAXINE	NaN
DOXEPIN	0.75
DULOXETINE	0.45
ESCITALOPRAM	1.33
FLUOXETINE	0.24
Less common antidepressants or combination of a...	0.70
MIRTAZAPINE	1.25
NORTRIPTYLINE	1.84
PAROXETINE	0.43
ROPINIROLE	0.56
SERTRALINE	0.21
TRAZODONE	NaN
VENLAFAXINE	0.27
strata aa:meas:adrm_2SN \	
Response	
AMITRIPTYLINE	-1.08
BUPROPION	-0.84
CITALOPRAM	-0.91
DESVENLAFAXINE	-0.70
DOXEPIN	-1.16
DULOXETINE	-0.90
ESCITALOPRAM	-0.99
FLUOXETINE	-0.90
Less common antidepressants or combination of a...	-0.35
MIRTAZAPINE	-0.83
NORTRIPTYLINE	-1.12
PAROXETINE	-0.95
ROPINIROLE	-1.12
SERTRALINE	-0.92
TRAZODONE	-0.90
VENLAFAXINE	-0.85
strata aa:meas:adrm_3DR \	
Response	
AMITRIPTYLINE	0.71
BUPROPION	0.77
CITALOPRAM	0.37
DESVENLAFAXINE	0.39
DOXEPIN	0.34

DULOXETINE	0.39
ESCITALOPRAM	0.61
FLUOXETINE	0.41
Less common antidepressants or combination of a...	0.43
MIRTAZAPINE	0.80
NORTRIPTYLINE	0.55
PAROXETINE	0.39
ROPINIROLE	NaN
SERTRALINE	0.36
TRAZODONE	0.91
VENLAFAXINE	0.37

strata	aa:meas:adrm_4DN	...	\
Response			...
AMITRIPTYLINE	-0.36	...	
BUPROPION	-0.54	...	
CITALOPRAM	-0.18	...	
DESVENLAFAXINE	-0.22	...	
DOXEPIN	-0.11	...	
DULOXETINE	-0.28	...	
ESCITALOPRAM	-0.26	...	
FLUOXETINE	-0.19	...	
Less common antidepressants or combination of a...	-0.27	...	
MIRTAZAPINE	-0.26	...	
NORTRIPTYLINE	-0.34	...	
PAROXETINE	-0.19	...	
ROPINIROLE	-0.30	...	
SERTRALINE	-0.19	...	
TRAZODONE	-0.79	...	
VENLAFAXINE	-0.26	...	

strata	rx:GEN:G12753	\
Response		
AMITRIPTYLINE	NaN	
BUPROPION	NaN	
CITALOPRAM	NaN	
DESVENLAFAXINE	NaN	
DOXEPIN	NaN	
DULOXETINE	NaN	
ESCITALOPRAM	NaN	
FLUOXETINE	NaN	
Less common antidepressants or combination of a...	-0.03	
MIRTAZAPINE	NaN	
NORTRIPTYLINE	NaN	
PAROXETINE	-0.07	
ROPINIROLE	NaN	
SERTRALINE	NaN	

TRAZODONE	NaN
VENLAFAXINE	NaN

strata	rx:GEN:G12775 \
Response	
AMITRIPTYLINE	NaN
BUPROPION	NaN
CITALOPRAM	NaN
DESVENLAFAXINE	NaN
DOXEPIN	NaN
DULOXETINE	0.13
ESCITALOPRAM	NaN
FLUOXETINE	NaN
Less common antidepressants or combination of a...	NaN
MIRTAZAPINE	NaN
NORTRIPTYLINE	NaN
PAROXETINE	NaN
ROPINIROLE	NaN
SERTRALINE	NaN
TRAZODONE	NaN
VENLAFAXINE	NaN

strata	rx:GEN:G12795 \
Response	
AMITRIPTYLINE	NaN
BUPROPION	NaN
CITALOPRAM	0.12
DESVENLAFAXINE	NaN
DOXEPIN	NaN
DULOXETINE	NaN
ESCITALOPRAM	NaN
FLUOXETINE	NaN
Less common antidepressants or combination of a...	NaN
MIRTAZAPINE	NaN
NORTRIPTYLINE	NaN
PAROXETINE	NaN
ROPINIROLE	NaN
SERTRALINE	NaN
TRAZODONE	NaN
VENLAFAXINE	NaN

strata	rx:GEN:G12861 \
Response	
AMITRIPTYLINE	NaN
BUPROPION	NaN
CITALOPRAM	0.20
DESVENLAFAXINE	0.21

DOXEPIN	NaN
DULOXETINE	0.33
ESCITALOPRAM	0.24
FLUOXETINE	0.13
Less common antidepressants or combination of a...	0.30
MIRTAZAPINE	NaN
NORTRIPTYLINE	NaN
PAROXETINE	0.21
ROPINIROLE	-0.21
SERTRALINE	0.20
TRAZODONE	NaN
VENLAFAXINE	0.58

strata	rx:GEN:G12871	\
Response		
AMITRIPTYLINE	NaN	
BUPROPION	NaN	
CITALOPRAM	NaN	
DESVENLAFAXINE	NaN	
DOXEPIN	NaN	
DULOXETINE	NaN	
ESCITALOPRAM	NaN	
FLUOXETINE	0.08	
Less common antidepressants or combination of a...	0.07	
MIRTAZAPINE	NaN	
NORTRIPTYLINE	NaN	
PAROXETINE	0.10	
ROPINIROLE	NaN	
SERTRALINE	NaN	
TRAZODONE	NaN	
VENLAFAXINE	NaN	

strata	rx:GEN:G12883	\
Response		
AMITRIPTYLINE	NaN	
BUPROPION	NaN	
CITALOPRAM	NaN	
DESVENLAFAXINE	NaN	
DOXEPIN	NaN	
DULOXETINE	NaN	
ESCITALOPRAM	NaN	
FLUOXETINE	NaN	
Less common antidepressants or combination of a...	0.43	
MIRTAZAPINE	NaN	
NORTRIPTYLINE	NaN	
PAROXETINE	NaN	
ROPINIROLE	NaN	

SERTRALINE	NaN
TRAZODONE	NaN
VENLAFAXINE	NaN

strata	rx:GEN:G13242 \
Response	
AMITRIPTYLINE	NaN
BUPROPION	NaN
CITALOPRAM	0.05
DESVENLAFAXINE	NaN
DOXEPIN	NaN
DULOXETINE	NaN
ESCITALOPRAM	NaN
FLUOXETINE	NaN
Less common antidepressants or combination of a...	0.04
MIRTAZAPINE	NaN
NORTRIPTYLINE	NaN
PAROXETINE	NaN
ROPINIROLE	NaN
SERTRALINE	NaN
TRAZODONE	NaN
VENLAFAXINE	NaN

strata	rx:GEN:G13348 \
Response	
AMITRIPTYLINE	NaN
BUPROPION	NaN
CITALOPRAM	NaN
DESVENLAFAXINE	NaN
DOXEPIN	NaN
DULOXETINE	NaN
ESCITALOPRAM	NaN
FLUOXETINE	NaN
Less common antidepressants or combination of a...	-0.07
MIRTAZAPINE	NaN
NORTRIPTYLINE	NaN
PAROXETINE	NaN
ROPINIROLE	NaN
SERTRALINE	NaN
TRAZODONE	NaN
VENLAFAXINE	NaN

strata	rx:GEN:G13447 \
Response	
AMITRIPTYLINE	NaN
BUPROPION	NaN
CITALOPRAM	NaN

DESVENLAFAXINE	NaN
DOXEPIN	NaN
DULOXETINE	NaN
ESCITALOPRAM	NaN
FLUOXETINE	NaN
Less common antidepressants or combination of a...	0.09
MIRTAZAPINE	NaN
NORTRIPTYLINE	NaN
PAROXETINE	NaN
ROPINIROLE	NaN
SERTRALINE	NaN
TRAZODONE	NaN
VENLAFAXINE	NaN

strata	rx:GEN:G13461
Response	
AMITRIPTYLINE	NaN
BUPROPION	NaN
CITALOPRAM	NaN
DESVENLAFAXINE	NaN
DOXEPIN	NaN
DULOXETINE	0.04
ESCITALOPRAM	0.05
FLUOXETINE	-0.04
Less common antidepressants or combination of a...	-0.04
MIRTAZAPINE	0.13
NORTRIPTYLINE	NaN
PAROXETINE	NaN
ROPINIROLE	NaN
SERTRALINE	NaN
TRAZODONE	NaN
VENLAFAXINE	NaN

[16 rows x 457 columns]

```
[7]: ##### Creating a coefficient missing file. this file is a mockup of how the real
      ↪ coefficient missing file.
import copy
coeff_missing = pd.DataFrame(columns = ["Response", "strata", "coef"])

def change_value(x):
    return x

for x in set(df_2["Response"]):
    temp_df = df_2[df_2["Response"] == x]
    temp_df = temp_df.reset_index(drop = True)
    #print(temp_df)
```

```

for y in range(len(temp_df)):
    #print(temp_df.iloc[y])
    #print(temp_df["code"].iloc[y])
    key = f"{x} {temp_df['strata'].iloc[y]} missing"
    value = copy.deepcopy(temp_df)
    value = value.drop(y)
    value["Response"] = value["Response"].apply(lambda x: change_value(key))
    #print(value)
    #print(type(value))
    coeff_missing = pd.concat([coeff_missing, value], ignore_index = True)

```

C:\Users\Bido\AppData\Local\Temp\ipykernel_10976\2764937496.py:21:

FutureWarning: The behavior of DataFrame concatenation with empty or all-NA entries is deprecated. In a future version, this will no longer exclude empty or all-NA columns when determining the result dtypes. To retain the old behavior, exclude the relevant entries before the concat operation.

```
coeff_missing = pd.concat([coeff_missing, value], ignore_index = True)
```

[8]: coeff_missing

```

[8]:

```

			Response	strata	coef
0	DULOXETINE	dxo:ICD9:78909	missing	dxo:ICD9:78900	-0.06
1	DULOXETINE	dxo:ICD9:78909	missing	dxo:ICD9:4619	0.04
2	DULOXETINE	dxo:ICD9:78909	missing	dxo:ICD9:30924	-0.12
3	DULOXETINE	dxo:ICD9:78909	missing	aa:dem:ager_1319	-0.40
4	DULOXETINE	dxo:ICD9:78909	missing	dxo:ICD9:30928	-0.13
...		
221377	ESCITALOPRAM	rx:GEN:G13461	missing	dxo:ICD9:2859	-0.05
221378	ESCITALOPRAM	rx:GEN:G13461	missing	dxo:ICD9:56400	-0.07
221379	ESCITALOPRAM	rx:GEN:G13461	missing	dxo:ICD9:4439	-0.10
221380	ESCITALOPRAM	rx:GEN:G13461	missing	dxo:ICD9:61610	-0.09
221381	ESCITALOPRAM	rx:GEN:G13461	missing	dxo:ICD9:5990	-0.06

[221382 rows x 3 columns]

```

[9]: ## Randomizing the coefficient missing values
coeff_missing["coef"] = coeff_missing["coef"].apply(lambda x : random.
    ↪uniform(min(coeff_missing["coef"]), max(coeff_missing["coef"])))

```

Unexpected exception formatting exception. Falling back to standard exception

Traceback (most recent call last):

File "C:\Users\Bido\AppData\Roaming\Python\Python311\site-packages\IPython\core\interactiveshell.py", line 3526, in run_code
 exec(code_obj, self.user_global_ns, self.user_ns)

File "C:\Users\Bido\AppData\Local\Temp\ipykernel_10976\111479983.py", line 2,
 in <module>

```
coeff_missing["coef"] = coeff_missing["coef"].apply(lambda x :
```

```

random.uniform(min(coeff_missing["coef"]), max(coeff_missing["coef"])))
~~~~~
File "C:\Users\Bido\AppData\Roaming\Python\Python311\site-
packages\pandas\core\series.py", line 4760, in apply
    ).apply()
~~~~~
File "C:\Users\Bido\AppData\Roaming\Python\Python311\site-
packages\pandas\core\apply.py", line 1207, in apply
    return self.apply_standard()
~~~~~
File "C:\Users\Bido\AppData\Roaming\Python\Python311\site-
packages\pandas\core\apply.py", line 1287, in apply_standard
    mapped = obj._map_values(
~~~~~
File "C:\Users\Bido\AppData\Roaming\Python\Python311\site-
packages\pandas\core\base.py", line 921, in _map_values
    return algorithms.map_array(arr, mapper, na_action=na_action,
convert=convert)
~~~~~
File "C:\Users\Bido\AppData\Roaming\Python\Python311\site-
packages\pandas\core\algorithms.py", line 1814, in map_array
    return lib.map_infer(values, mapper, convert=convert)
~~~~~
File "lib.pyx", line 2917, in pandas._libs.lib.map_infer
File "C:\Users\Bido\AppData\Local\Temp\ipykernel_10976\111479983.py", line 2,
in <lambda>
    coeff_missing["coef"] = coeff_missing["coef"].apply(lambda x :
random.uniform(min(coeff_missing["coef"]), max(coeff_missing["coef"])))
~~~~~
File "C:\Users\Bido\AppData\Roaming\Python\Python311\site-
packages\pandas\core\frame.py", line 3856, in __getitem__
    def __getitem__(self, key):

```

KeyboardInterrupt

During handling of the above exception, another exception occurred:

Traceback (most recent call last):

```

File "C:\Users\Bido\AppData\Roaming\Python\Python311\site-
packages\IPython\core\interactiveshell.py", line 2120, in showtraceback
    stb = self.InteractiveTB.structured_traceback(
~~~~~
File "C:\Users\Bido\AppData\Roaming\Python\Python311\site-
packages\IPython\core\ultratb.py", line 1435, in structured_traceback
    return FormattedTB.structured_traceback(
~~~~~
File "C:\Users\Bido\AppData\Roaming\Python\Python311\site-

```

```

packages\IPython\core\ultratb.py", line 1326, in structured_traceback
    return VerboseTB.structured_traceback(
        ~~~~~
    File "C:\Users\Bido\AppData\Roaming\Python\Python311\site-
packages\IPython\core\ultratb.py", line 1173, in structured_traceback
    formatted_exception = self.format_exception_as_a_whole(etype, evalue, etb,
number_of_lines_of_context,
~~~~~
    File "C:\Users\Bido\AppData\Roaming\Python\Python311\site-
packages\IPython\core\ultratb.py", line 1063, in format_exception_as_a_whole
    self.get_records(etb, number_of_lines_of_context, tb_offset) if etb else []
    ~~~~~
    File "C:\Users\Bido\AppData\Roaming\Python\Python311\site-
packages\IPython\core\ultratb.py", line 1155, in get_records
    FrameInfo(
    File "C:\Users\Bido\AppData\Roaming\Python\Python311\site-
packages\IPython\core\ultratb.py", line 780, in __init__
    ix = inspect.getsourcelines(frame)
    ~~~~~
    File "C:\Users\Bido\anaconda3\Lib\inspect.py", line 1244, in getsourcelines
    lines, lnum = findsource(object)
    ~~~~~
    File "C:\Users\Bido\anaconda3\Lib\inspect.py", line 1073, in findsource
    raise OSError('source code not available')
OSError: source code not available

```

```
[ ]: coeff_missing
```

```
[ ]: ### saving the missing coefficient file
coeff_missing.to_csv("coeff_missing.csv", index = False)
```

```
[ ]:
```

```
[10]: ### Importing the Modules required
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
from pathlib import Path
import math
import random
from functools import reduce
import requests
```

```
[11]: ### Importing the coefficient Excel File
excel_path = Path("C:\Sem 3\HAP 786\coefficients.xlsx")
df = pd.read_excel(excel_path)
coeff_missing = pd.read_csv("coeff_missing.csv")
```

```

history_csv = pd.read_csv(r"C:\Users\Bido_\Downloads\path_to_save_replaced_file.
↪csv")
df["code"] = df["code"].fillna("Intercept")
df["strata"] = df["strata"].fillna("Intercept")
df_2 = df[["Response"] + ["strata"] + ["coef"]]
df_2 = df_2.reset_index(drop = True)
df_2[df_2["strata"] == "Intercept"]

```

```

C:\Users\Bido_\AppData\Roaming\Python\Python311\site-
packages\openpyxl\worksheet\header_footer.py:48: UserWarning: Cannot parse
header or footer so it will be ignored
    warn("""Cannot parse header or footer so it will be ignored""")

```

```

[11]:
419          FLUOXETINE  Intercept -0.10
420          MIRTAZAPINE  Intercept -1.60
421          NORTRIPTYLINE  Intercept -2.59
488          PAROXETINE  Intercept -0.35
489          ROPINIROLE  Intercept -0.94
490          SERTRALINE  Intercept  0.04
491          TRAZODONE  Intercept -4.08
587          DOXEPIN  Intercept -1.61
588          DULOXETINE  Intercept -0.55
589          ESCITALOPRAM  Intercept -1.66
621          CITALOPRAM  Intercept -0.04
622          DESVENLAFAXINE  Intercept  0.28
630          VENLAFAXINE  Intercept  0.02
631  Less common antidepressants or combination of ...  Intercept -0.71
1414          BUPROPION  Intercept -3.12
1439          AMITRIPTYLINE  Intercept -4.09

```

```

[12]: df_2[df_2["strata"] == "Intercept"]

```

```

[12]:
419          FLUOXETINE  Intercept -0.10
420          MIRTAZAPINE  Intercept -1.60
421          NORTRIPTYLINE  Intercept -2.59
488          PAROXETINE  Intercept -0.35
489          ROPINIROLE  Intercept -0.94
490          SERTRALINE  Intercept  0.04
491          TRAZODONE  Intercept -4.08
587          DOXEPIN  Intercept -1.61
588          DULOXETINE  Intercept -0.55
589          ESCITALOPRAM  Intercept -1.66
621          CITALOPRAM  Intercept -0.04
622          DESVENLAFAXINE  Intercept  0.28
630          VENLAFAXINE  Intercept  0.02
631  Less common antidepressants or combination of ...  Intercept -0.71

```


1414	BUPROPION	Intercept	-3.12
1439	AMITRIPTYLINE	Intercept	-4.09

```
[13]: coeff_missing
```

```
[13]:
```

		Response		strata	coef
0	DESVENLAFAXINE	dxo:ICD9:30924 missing	aa:dem:ager_1319	-3.550513	
1	DESVENLAFAXINE	dxo:ICD9:30924 missing	dxo:ICD9:30928	-2.579408	
2	DESVENLAFAXINE	dxo:ICD9:30924 missing	aa:dem:ager_2040	-3.013912	
3	DESVENLAFAXINE	dxo:ICD9:30924 missing	rx:GEN:G00623	-3.297315	
4	DESVENLAFAXINE	dxo:ICD9:30924 missing	rx:GEN:G00825	2.495287	
...		
221377	NORTRIPTYLINE	rx:GEN:G12405 missing	dxo:ICD9:29635	0.752783	
221378	NORTRIPTYLINE	rx:GEN:G12405 missing	rx:GEN:G08636	-0.518146	
221379	NORTRIPTYLINE	rx:GEN:G12405 missing	aa:meas:nrem_2+	1.408530	
221380	NORTRIPTYLINE	rx:GEN:G12405 missing	px:CPT4:90862	-1.855721	
221381	NORTRIPTYLINE	rx:GEN:G12405 missing	px:CPT4:90834	-3.381366	

[221382 rows x 3 columns]

```
[14]: ### Importing patient csv
history_csv = pd.read_csv(r"C:\Users\Bido_\Downloads\path_to_save_replaced_file.
↳csv")
history_csv
```

```
[14]:
```

	Patient_ID	aa:dem:ager_1319	aa:dem:ager_2040	aa:dem:ager_6579	\
0	38206	0	1	0	
1	55455	0	0	0	
2	77350	0	0	1	
3	65055	0	0	0	
4	77161	1	0	0	
5	52723	0	1	0	
6	68954	1	0	0	
7	24275	0	0	0	
8	91938	0	0	1	
9	25199	1	0	0	
10	47681	0	1	0	
11	35268	0	0	1	
12	31835	1	0	0	
13	92205	1	0	0	
14	65268	0	0	1	
15	60529	1	0	0	
16	1111	0	0	0	

	aa:dem:ager_8089	aa:dem:gender_F	aa:meas:adrm_1SR	aa:meas:adrm_2SN	\
0	0	1	1	0	
1	1	0	0	1	

2	0	0	0	1
3	1	0	0	0
4	0	0	1	1
5	0	1	1	0
6	0	0	1	1
7	1	0	0	1
8	0	0	1	1
9	0	0	0	0
10	0	0	0	0
11	0	1	0	0
12	0	0	1	1
13	0	0	0	1
14	0	1	1	0
15	0	0	1	0
16	0	0	0	0

	aa:meas:adrm_3DR	aa:meas:adrm_4DN	...	rx:GEN:G12753	rx:GEN:G12775	\
0	0	0	...	0	0	
1	1	0	...	1	1	
2	1	0	...	0	1	
3	0	1	...	1	0	
4	0	0	...	0	0	
5	1	0	...	0	1	
6	1	1	...	0	0	
7	1	0	...	1	1	
8	1	0	...	1	1	
9	1	0	...	1	1	
10	0	1	...	0	0	
11	0	1	...	1	0	
12	1	1	...	0	0	
13	1	0	...	0	1	
14	0	1	...	1	0	
15	1	0	...	1	0	
16	0	0	...	0	0	

	rx:GEN:G12795	rx:GEN:G12861	rx:GEN:G12871	rx:GEN:G12883	rx:GEN:G13242	\
0	0	1	1	1	1	
1	0	1	1	0	0	
2	1	0	1	1	0	
3	1	1	0	0	0	
4	1	1	1	0	0	
5	1	0	1	1	0	
6	1	1	1	1	1	
7	0	0	1	0	0	
8	0	1	1	0	0	
9	0	0	0	1	0	
10	1	1	1	1	1	

11	1	1	1	0	1
12	0	0	0	0	1
13	1	1	1	0	1
14	1	0	1	0	0
15	1	1	0	0	0
16	0	0	0	0	0

	rx:GEN:G13348	rx:GEN:G13447	rx:GEN:G13461
0	1	1	1
1	0	0	1
2	1	0	0
3	1	0	0
4	1	0	0
5	0	1	1
6	1	1	0
7	1	0	1
8	0	1	1
9	1	0	0
10	1	0	0
11	1	0	0
12	0	1	0
13	1	1	1
14	0	1	0
15	0	1	1
16	0	0	0

[17 rows x 457 columns]

```
[15]: ### Calculating the probabilities of each drug for each patient
def calculating_prob(pat_result, df, df_missing):
    pat_df = pd.DataFrame(pat_result).T
    columns = list(pat_df.columns)[1:]
    zero_col = []
    one_col = []
    nan_col = []
    for x in columns:
        value = pat_df[x].item()
        if pd.isna(value):
            nan_col.append(x)
        elif value != 0.0 and isinstance(value, (int, float)):
            one_col.append(x)
        else:
            zero_col.append(x)
    #df_3 = df_2.pivot_table(index = "Response", columns = "strata", values = 
    ↪ "coef", aggfunc = "sum")
    drug_probs_dict = {}
    for drug in df["Response"].unique().tolist():
```

```

temp_df = df[df["Response"] == drug]
val = temp_df[temp_df["strata"] == "Intercept"]["coef"].item()
columns_temp = list(temp_df.pivot_table(index = "Response", columns =
↳ "strata", values = "coef").columns)
if not any(x in nan_col for x in columns_temp):
    eq = sum([temp_df[temp_df["strata"] == x]["coef"].item() *
↳ pat_df[x].item() for x in one_col if x in columns_temp])
else:
    temp_df = df[df["Response"] == drug]
    val = temp_df[temp_df["strata"] == "Intercept"]["coef"].item()
    temp_df = df_missing[
        df_missing["Response"].str.contains(fr"\b{drug}", na =
↳ False) &
        df_missing["Response"].str.contains(fr"{nan_col[0]}", na =
↳ False)
    ]
    eq = sum([temp_df[temp_df["strata"] == x]["coef"].item() *
↳ pat_df[x].item() for x in one_col if x in columns_temp])

eq += val
odds = math.exp(eq)
probs = round(odds / (1 + odds), 2)
drug_probs_dict[drug] = probs

return (pd.DataFrame(drug_probs_dict, index = ["Probs"]).T.reset_index(drop
↳ False).rename(columns = {'index': 'Drug'}),
        (pat_result["Patient_ID"], one_col))

```

```

[16]: ## Creating a patient-drug probabilities dictionary
pat_drug_dict = {}
pat_features = {}
for pat in history_csv.iterrows():
    _, features = calculating_prob(pat[1], df_2, coeff_missing)
    pat_drug_dict[pat[1]["Patient_ID"]] = _
    pat_features[features[0]] = features[1]

```

```

[17]: pat_drug_dict

```

```

[17]: {38206.0:

```

	Drug	Probs
0	CITALOPRAM	0.00
1	Less common antidepressants or combination of ...	0.00
2	DULOXETINE	0.00
3	SERTRALINE	0.00
4	ESCITALOPRAM	0.00
5	FLUOXETINE	0.00
6	PAROXETINE	0.00
7	VENLAFAXINE	0.00

8		AMITRIPTYLINE	0.83	
9		DESVENLAFAXINE	0.00	
10		DOXEPIN	0.00	
11		MIRTAZAPINE	0.00	
12		NORTRIPTYLINE	0.53	
13		BUPROPION	0.98	
14		ROPINIROLE	0.00	
15		TRAZODONE	0.00,	
55455.0:			Drug	Probs
0		CITALOPRAM	0.01	
1	Less common antidepressants or combination of ...		0.35	
2		DULOXETINE	0.02	
3		SERTRALINE	0.00	
4		ESCITALOPRAM	0.23	
5		FLUOXETINE	0.03	
6		PAROXETINE	0.02	
7		VENLAFAXINE	0.01	
8		AMITRIPTYLINE	0.81	
9		DESVENLAFAXINE	0.04	
10		DOXEPIN	0.01	
11		MIRTAZAPINE	0.19	
12		NORTRIPTYLINE	0.10	
13		BUPROPION	0.84	
14		ROPINIROLE	0.04	
15		TRAZODONE	0.67,	
77350.0:			Drug	Probs
0		CITALOPRAM	0.00	
1	Less common antidepressants or combination of ...		0.02	
2		DULOXETINE	0.00	
3		SERTRALINE	0.01	
4		ESCITALOPRAM	0.01	
5		FLUOXETINE	0.00	
6		PAROXETINE	0.00	
7		VENLAFAXINE	0.00	
8		AMITRIPTYLINE	0.16	
9		DESVENLAFAXINE	0.00	
10		DOXEPIN	0.00	
11		MIRTAZAPINE	0.02	
12		NORTRIPTYLINE	0.04	
13		BUPROPION	0.32	
14		ROPINIROLE	0.01	
15		TRAZODONE	0.05,	
65055.0:			Drug	Probs
0		CITALOPRAM	0.01	
1	Less common antidepressants or combination of ...		0.00	
2		DULOXETINE	0.01	
3		SERTRALINE	0.00	

4	ESCITALOPRAM	0.04	
5	FLUOXETINE	0.01	
6	PAROXETINE	0.00	
7	VENLAFAXINE	0.01	
8	AMITRIPTYLINE	0.14	
9	DESVENLAFAXINE	0.02	
10	DOXEPIN	0.01	
11	MIRTAZAPINE	0.03	
12	NORTRIPTYLINE	0.12	
13	BUPROPION	0.35	
14	ROPINIROLE	0.05	
15	TRAZODONE	0.11,	
77161.0:		Drug	Probs
0	CITALOPRAM	0.00	
1	Less common antidepressants or combination of ...	0.00	
2	DULOXETINE	0.01	
3	SERTRALINE	0.00	
4	ESCITALOPRAM	0.08	
5	FLUOXETINE	0.00	
6	PAROXETINE	0.00	
7	VENLAFAXINE	0.00	
8	AMITRIPTYLINE	0.11	
9	DESVENLAFAXINE	0.00	
10	DOXEPIN	0.02	
11	MIRTAZAPINE	0.11	
12	NORTRIPTYLINE	0.44	
13	BUPROPION	0.84	
14	ROPINIROLE	0.05	
15	TRAZODONE	0.59,	
52723.0:		Drug	Probs
0	CITALOPRAM	0.01	
1	Less common antidepressants or combination of ...	0.00	
2	DULOXETINE	0.02	
3	SERTRALINE	0.00	
4	ESCITALOPRAM	0.09	
5	FLUOXETINE	0.00	
6	PAROXETINE	0.00	
7	VENLAFAXINE	0.00	
8	AMITRIPTYLINE	0.74	
9	DESVENLAFAXINE	0.00	
10	DOXEPIN	0.30	
11	MIRTAZAPINE	0.30	
12	NORTRIPTYLINE	0.90	
13	BUPROPION	0.97	
14	ROPINIROLE	0.51	
15	TRAZODONE	0.94,	
68954.0:		Drug	Probs

0		CITALOPRAM	0.03	
1	Less common antidepressants or combination of ...		0.07	
2		DULOXETINE	0.00	
3		SERTRALINE	0.05	
4		ESCITALOPRAM	0.32	
5		FLUOXETINE	0.01	
6		PAROXETINE	0.01	
7		VENLAFAXINE	0.03	
8		AMITRIPTYLINE	0.13	
9		DESVENLAFAXINE	0.01	
10		DOXEPIN	0.03	
11		MIRTAZAPINE	0.27	
12		NORTRIPTYLINE	0.49	
13		BUPROPION	0.74	
14		ROPINIROLE	0.03	
15		TRAZODONE	0.52,	
24275.0:				Drug Probs
0		CITALOPRAM	0.00	
1	Less common antidepressants or combination of ...		0.00	
2		DULOXETINE	0.00	
3		SERTRALINE	0.00	
4		ESCITALOPRAM	0.04	
5		FLUOXETINE	0.00	
6		PAROXETINE	0.00	
7		VENLAFAXINE	0.00	
8		AMITRIPTYLINE	0.77	
9		DESVENLAFAXINE	0.01	
10		DOXEPIN	0.21	
11		MIRTAZAPINE	0.28	
12		NORTRIPTYLINE	0.72	
13		BUPROPION	0.81	
14		ROPINIROLE	0.04	
15		TRAZODONE	0.87,	
91938.0:				Drug Probs
0		CITALOPRAM	0.00	
1	Less common antidepressants or combination of ...		0.00	
2		DULOXETINE	0.00	
3		SERTRALINE	0.00	
4		ESCITALOPRAM	0.00	
5		FLUOXETINE	0.00	
6		PAROXETINE	0.00	
7		VENLAFAXINE	0.00	
8		AMITRIPTYLINE	0.94	
9		DESVENLAFAXINE	0.03	
10		DOXEPIN	0.14	
11		MIRTAZAPINE	0.18	
12		NORTRIPTYLINE	0.78	

13	BUPROPION	0.97	
14	ROPINIROLE	0.07	
15	TRAZODONE	0.84,	
25199.0:		Drug	Probs
0	CITALOPRAM	0.00	
1	Less common antidepressants or combination of ...	0.12	
2	DULOXETINE	0.02	
3	SERTRALINE	0.02	
4	ESCITALOPRAM	0.11	
5	FLUOXETINE	0.00	
6	PAROXETINE	0.01	
7	VENLAFAXINE	0.00	
8	AMITRIPTYLINE	0.63	
9	DESVENLAFAXINE	0.02	
10	DOXEPIN	0.02	
11	MIRTAZAPINE	0.09	
12	NORTRIPTYLINE	0.44	
13	BUPROPION	0.97	
14	ROPINIROLE	0.14	
15	TRAZODONE	0.73,	
47681.0:		Drug	Probs
0	CITALOPRAM	0.01	
1	Less common antidepressants or combination of ...	0.00	
2	DULOXETINE	0.03	
3	SERTRALINE	0.04	
4	ESCITALOPRAM	0.00	
5	FLUOXETINE	0.00	
6	PAROXETINE	0.02	
7	VENLAFAXINE	0.01	
8	AMITRIPTYLINE	0.75	
9	DESVENLAFAXINE	0.01	
10	DOXEPIN	0.01	
11	MIRTAZAPINE	0.08	
12	NORTRIPTYLINE	0.49	
13	BUPROPION	0.23	
14	ROPINIROLE	0.32	
15	TRAZODONE	0.35,	
35268.0:		Drug	Probs
0	CITALOPRAM	0.08	
1	Less common antidepressants or combination of ...	0.00	
2	DULOXETINE	0.11	
3	SERTRALINE	0.00	
4	ESCITALOPRAM	0.23	
5	FLUOXETINE	0.00	
6	PAROXETINE	0.00	
7	VENLAFAXINE	0.00	
8	AMITRIPTYLINE	0.86	

9	DESVENLAFAXINE	0.20	
10	DOXEPIN	0.18	
11	MIRTAZAPINE	0.26	
12	NORTRIPTYLINE	0.35	
13	BUPROPION	0.55	
14	ROPINIROLE	0.17	
15	TRAZODONE	0.26,	
31835.0:		Drug	Probs
0	CITALOPRAM	0.00	
1	Less common antidepressants or combination of ...	0.03	
2	DULOXETINE	0.00	
3	SERTRALINE	0.00	
4	ESCITALOPRAM	0.11	
5	FLUOXETINE	0.00	
6	PAROXETINE	0.00	
7	VENLAFAXINE	0.00	
8	AMITRIPTYLINE	0.56	
9	DESVENLAFAXINE	0.00	
10	DOXEPIN	0.01	
11	MIRTAZAPINE	0.11	
12	NORTRIPTYLINE	0.35	
13	BUPROPION	0.00	
14	ROPINIROLE	0.01	
15	TRAZODONE	0.28,	
92205.0:		Drug	Probs
0	CITALOPRAM	0.00	
1	Less common antidepressants or combination of ...	0.00	
2	DULOXETINE	0.00	
3	SERTRALINE	0.00	
4	ESCITALOPRAM	0.24	
5	FLUOXETINE	0.00	
6	PAROXETINE	0.00	
7	VENLAFAXINE	0.00	
8	AMITRIPTYLINE	0.65	
9	DESVENLAFAXINE	0.15	
10	DOXEPIN	0.17	
11	MIRTAZAPINE	0.41	
12	NORTRIPTYLINE	0.67	
13	BUPROPION	0.67	
14	ROPINIROLE	0.16	
15	TRAZODONE	0.93,	
65268.0:		Drug	Probs
0	CITALOPRAM	0.00	
1	Less common antidepressants or combination of ...	0.00	
2	DULOXETINE	0.31	
3	SERTRALINE	0.00	
4	ESCITALOPRAM	0.78	

5	FLUOXETINE	0.00	
6	PAROXETINE	0.00	
7	VENLAFAXINE	0.12	
8	AMITRIPTYLINE	0.54	
9	DESVENLAFAXINE	0.30	
10	DOXEPIN	0.23	
11	MIRTAZAPINE	0.68	
12	NORTRIPTYLINE	0.91	
13	BUPROPION	1.00	
14	ROPINIROLE	0.44	
15	TRAZODONE	0.86,	
60529.0:		Drug	Probs
0	CITALOPRAM	0.07	
1	Less common antidepressants or combination of ...	0.19	
2	DULOXETINE	0.10	
3	SERTRALINE	0.00	
4	ESCITALOPRAM	0.31	
5	FLUOXETINE	0.02	
6	PAROXETINE	0.03	
7	VENLAFAXINE	0.04	
8	AMITRIPTYLINE	0.71	
9	DESVENLAFAXINE	0.01	
10	DOXEPIN	0.12	
11	MIRTAZAPINE	0.35	
12	NORTRIPTYLINE	0.71	
13	BUPROPION	0.65	
14	ROPINIROLE	0.31	
15	TRAZODONE	0.24,	
1111.0:		Drug	Probs
0	CITALOPRAM	0.49	
1	Less common antidepressants or combination of ...	0.33	
2	DULOXETINE	0.37	
3	SERTRALINE	0.51	
4	ESCITALOPRAM	0.16	
5	FLUOXETINE	0.48	
6	PAROXETINE	0.41	
7	VENLAFAXINE	0.50	
8	AMITRIPTYLINE	0.02	
9	DESVENLAFAXINE	0.57	
10	DOXEPIN	0.17	
11	MIRTAZAPINE	0.17	
12	NORTRIPTYLINE	0.07	
13	BUPROPION	0.04	
14	ROPINIROLE	0.28	
15	TRAZODONE	0.02}	

```
[18]: ### Creating a drug link dictionary
links = df[["Response"] + ["antidep_link"]]
links
links_dict = {y["Response"]: y["antidep_link"] for x, y in links.iterrows()}

[48]: ### Creating patient-recommendation dictionary
pat_rec = {}
for pat_id in pat_drug_dict:
    temp_df = pat_drug_dict[pat_id]
    max_value = temp_df["Probs"].max()
    print(max_value)
    sec_value = max_value - 0.05
    max_row = temp_df[temp_df["Probs"] == max_value]
    sec_drug = temp_df[(temp_df["Probs"] >= sec_value) & (temp_df["Probs"] <
↪max_value)]
    print(sec_drug)
    max_row = pd.concat([max_row, sec_drug], ignore_index = False)
    rec_links = tuple([(links_dict[x], x) for x in max_row["Drug"].tolist()])
    pat_rec[pat_id] = rec_links

pat_rec
rec_links
```

0.98

Empty DataFrame

Columns: [Drug, Probs]

Index: []

0.84

Drug Probs

8 AMITRIPTYLINE 0.81

0.32

Empty DataFrame

Columns: [Drug, Probs]

Index: []

0.35

Empty DataFrame

Columns: [Drug, Probs]

Index: []

0.84

Empty DataFrame

Columns: [Drug, Probs]

Index: []

0.97

Drug Probs

15 TRAZODONE 0.94

0.74

Empty DataFrame

Columns: [Drug, Probs]

```

Index: []
0.87
Empty DataFrame
Columns: [Drug, Probs]
Index: []
0.97
      Drug  Probs
8  AMITRIPTYLINE  0.94
0.97
Empty DataFrame
Columns: [Drug, Probs]
Index: []
0.75
Empty DataFrame
Columns: [Drug, Probs]
Index: []
0.86
Empty DataFrame
Columns: [Drug, Probs]
Index: []
0.56
Empty DataFrame
Columns: [Drug, Probs]
Index: []
0.93
Empty DataFrame
Columns: [Drug, Probs]
Index: []
1.0
Empty DataFrame
Columns: [Drug, Probs]
Index: []
0.71
Empty DataFrame
Columns: [Drug, Probs]
Index: []
0.57
Empty DataFrame
Columns: [Drug, Probs]
Index: []

```

```
[48]: ((('https://www.ncbi.nlm.nih.gov/books/NBK534829/', 'DESVENLAFAXINE'),)
```

```
[49]: ### Creating a list of the unique patient IDs
      pat_ids = list(pat_drug_dict.keys())

      ### Creating a plotting function
```

```

def plot_values(pat_dict, pat_id):
    temp_df = pat_dict[pat_id]
    temp_df = temp_df.sort_values(by="Probs", ascending = True)
    print(temp_df)
    plt.figure(figsize=(10, 6))
    ax = temp_df.set_index("Drug").plot(kind = "barh")
    for container in ax.containers:
        ax.bar_label(container, label_type='edge', padding=3)
    max_length = 15
    labels = [label.get_text() for label in ax.get_yticklabels()]
    shortened_labels = [label if len(label) <= max_length else label[:
↪max_length] + "..." for label in labels]
    ax.set_yticklabels(shortened_labels)
    xlim = ax.get_xlim()
    ax.set_xlim(xlim[0], xlim[1] * 1.05)

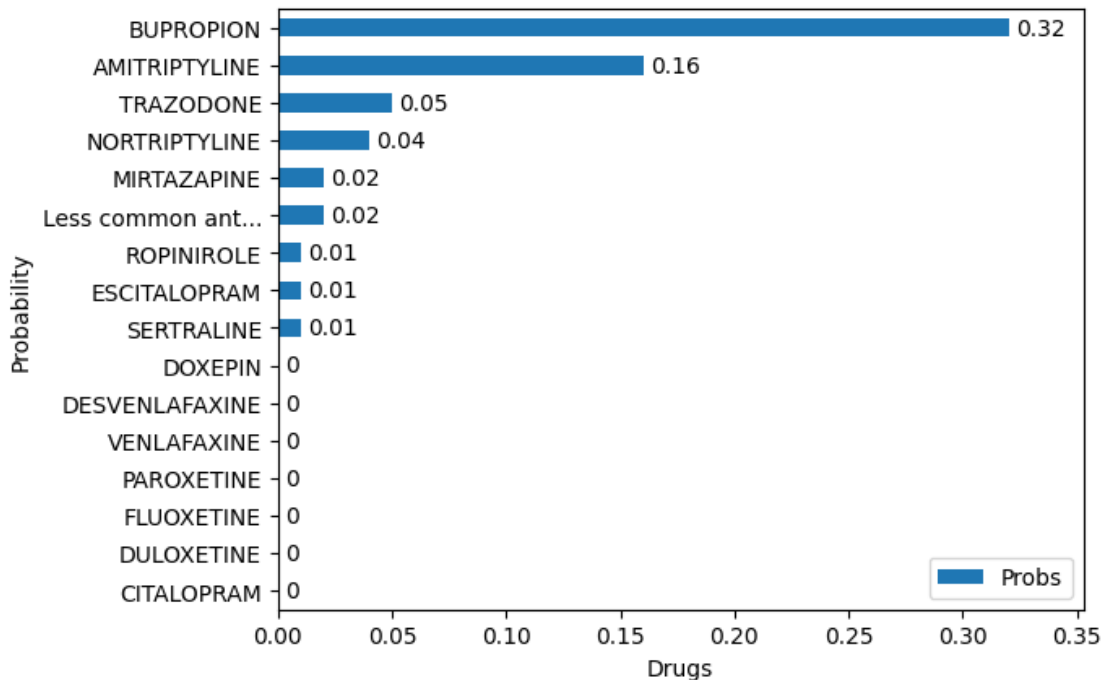
    ax.legend(loc='lower right')
    plt.xlabel("Drugs")
    plt.ylabel("Probability")
    plt.show()

```

```
[50]: plot_values(pat_drug_dict, pat_ids[2])
```

	Drug	Probs
0	CITALOPRAM	0.00
2	DULOXETINE	0.00
5	FLUOXETINE	0.00
6	PAROXETINE	0.00
7	VENLAFAXINE	0.00
9	DESVENLAFAXINE	0.00
10	DOXEPIN	0.00
3	SERTRALINE	0.01
4	ESCITALOPRAM	0.01
14	ROPINIROLE	0.01
1	Less common antidepressants or combination of ...	0.02
11	MIRTAZAPINE	0.02
12	NORTRIPTYLINE	0.04
15	TRAZODONE	0.05
8	AMITRIPTYLINE	0.16
13	BUPROPION	0.32

<Figure size 1000x600 with 0 Axes>



```
[51]: ### Creating a communicating function to deliver the patient recommendations
def communicating(pat_features, pat_id, pat_dict, links_dict):
    if pat_id not in pat_features:
        print("Patient ID is not found. Please type in a valid ID")
    else:
        print("We are assuming that you have major depressive disorder. If you
        ↪do not, our advice is not appropriate for you.\n"
              "If you have bipolar disorder, our advice is not appropriate for you.
        ↪Among the variables you mentioned during the medical history intake,\n"
              "the following are relevant in selection of appropriate antidepressants:
        ↪")
        for x in pat_features[pat_id]:
            print(f"-{x}")
        plot_values(pat_dict, pat_id)
        max_value = pat_dict[pat_id]["Probs"].max()
        if max_value < 0.10:
            print("In our data, we could not find an antidepressant that works
            ↪well for patients with your characteristics.\n"
                  "All 15 antidepressants we examined have less than 10 percent
            ↪chance of success for a patient with your characteristics.\n"
                  "Your primary care provider might be able to make a more
            ↪informed advice. In addition, it may be appropriate to look for other
            ↪options besides oral antidepressants for treatment of
            ↪depression.\n")
```

```

else:
    temp_df = pat_drug_dict[pat_id]
    max_row = temp_df[temp_df["Probs"] == max_value]
    sec_value = max_value - 0.05
    sec_drug = temp_df[(temp_df["Probs"] >= sec_value) &
↳(temp_df["Probs"] < max_value)]
    max_row = pd.concat([max_row, sec_drug], ignore_index = False)
    rec_links = tuple([(links_dict[x], x) for x in max_row["Drug"]
↳tolist()])
    for x in rec_links:
        print(f"Recommended drug: {x[1]}, NCBI site for drug: {x[0]}")

    print("take this information to your primary care provider and
↳discuss if there is a need for change in your medications.\n")
    "Do not change your medications suddenly as abrupt changes in
↳antidepressants could be dangerous\n")

```

```

[52]: ## Testing the function
communicating(pat_features, pat_ids[-1], pat_drug_dict, links_dict)

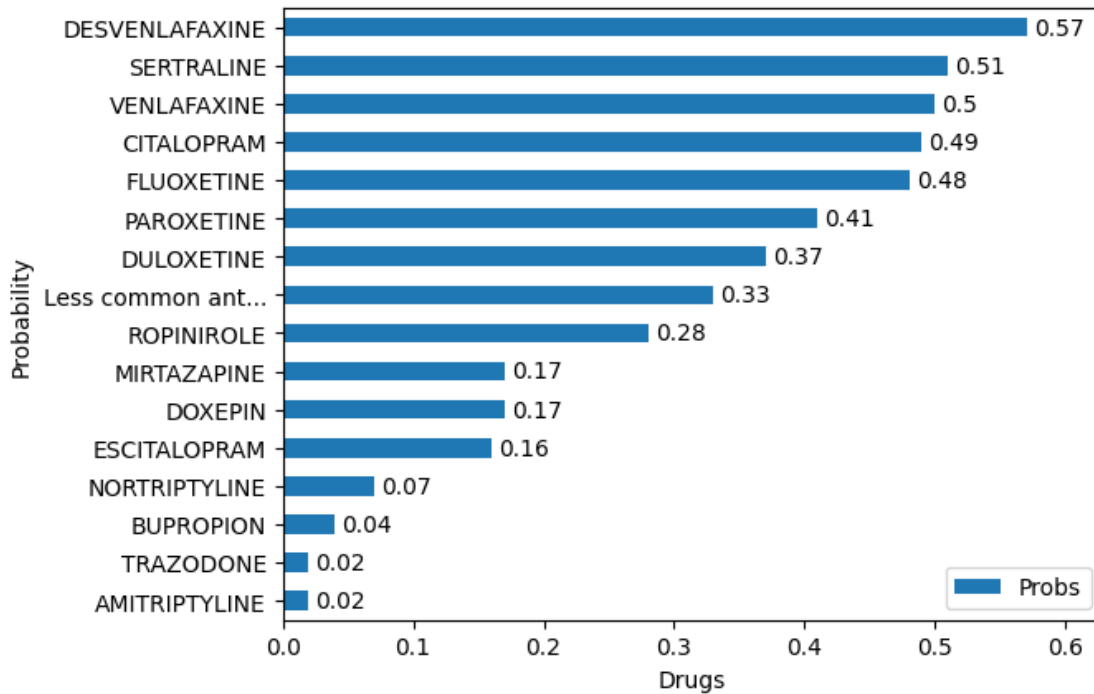
```

We are assuming that you have major depressive disorder. If you do not, our advice is not appropriate for you.

If you have bipolar disorder, our advice is not appropriate for you. Among the variables you mentioned during the medical history intake, the following are relevant in selection of appropriate antidepressants:

	Drug	Probs
8	AMITRIPTYLINE	0.02
15	TRAZODONE	0.02
13	BUPROPION	0.04
12	NORTRIPTYLINE	0.07
4	ESCITALOPRAM	0.16
10	DOXEPIN	0.17
11	MIRTAZAPINE	0.17
14	ROPINIROLE	0.28
1	Less common antidepressants or combination of ...	0.33
2	DULOXETINE	0.37
6	PAROXETINE	0.41
5	FLUOXETINE	0.48
0	CITALOPRAM	0.49
7	VENLAFAXINE	0.50
3	SERTRALINE	0.51
9	DESVENLAFAXINE	0.57

<Figure size 1000x600 with 0 Axes>



Recommended drug: DESVENLAFAXINE, NCBI site for drug:

<https://www.ncbi.nlm.nih.gov/books/NBK534829/>

take this information to your primary care provider and discuss if there is a need for change in your medications.

Do not change your medications suddenly as abrupt changes in antidepressants could be dangerous

[53]: coeff_missing

```
[53]:
```

		Response	strata	coef
0	DESVENLAFAXINE	dxo:ICD9:30924 missing	aa:dem:ager_1319	-3.550513
1	DESVENLAFAXINE	dxo:ICD9:30924 missing	dxo:ICD9:30928	-2.579408
2	DESVENLAFAXINE	dxo:ICD9:30924 missing	aa:dem:ager_2040	-3.013912
3	DESVENLAFAXINE	dxo:ICD9:30924 missing	rx:GEN:G00623	-3.297315
4	DESVENLAFAXINE	dxo:ICD9:30924 missing	rx:GEN:G00825	2.495287
...	
221377	NORTRIPTYLINE	rx:GEN:G12405 missing	dxo:ICD9:29635	0.752783
221378	NORTRIPTYLINE	rx:GEN:G12405 missing	rx:GEN:G08636	-0.518146
221379	NORTRIPTYLINE	rx:GEN:G12405 missing	aa:meas:nrem_2+	1.408530
221380	NORTRIPTYLINE	rx:GEN:G12405 missing	px:CPT4:90862	-1.855721
221381	NORTRIPTYLINE	rx:GEN:G12405 missing	px:CPT4:90834	-3.381366

[221382 rows x 3 columns]

[]: