

Using the DHS NFHS dataset, the objective is to recode the dataset to a usable form(for modeling) and get started on developing a computational statistical model of childhood vaccination outcomes.

GITHUB: https://github.com/masternaveen123/vaccination_coverage

With the email, I have attached the following files:

- Script for recoding the dataset(Recoding.ipynb)
- Metadata of recoded variables(DPT vs Penta - Variables.xlsx)
- Modeling approach(modelling zero_dose.ipynb)

Definitions:

- Zero doses are those data points without the first dose of DPT or pentavalent vaccines.
- Fully vac are those with doses of BCG + OPV 3rd dose + (DPT 3rd dose| PENTA 3rd dose) + MR1 [a total of 8 doses]
- Under vacc is the inverse of full vac.
- NULL refers to empty or missing values

Script for recoding the dataset

The script is written in Python, and it recodes the NFHS 5 dataset to be used for modeling. The script takes NFHS 5 data along with a binary flag as input. If the flag is true, it converts the dataset to binary values only, otherwise, some values are converted to binary whereas some columns contain categorical values. The dataset is filtered to only have records of kids between the age range of 12-23 months(including).

Variables used as inputs:

- Categorical_eliminate_flag (binary)
- Children(location to DTA file containing children's records)
- Household(location to DTA file containing household records)
- Individual(location to DTA file containing individual records)

Variables when flag is set to 1:

'impr_ws', 'unimpr_ws', 'basic_drinking_w', 'limited_drinking_w',
'jmp_w8', 'jmp_w2', 'jmp_w5', 'jmp_s1', 'jmp_s6', 'jmp_s8', 'jmp_s7',
'highest_grade_comp', 'jmp_h1', 'jmp_h2', 'jmp_h3', 'wi_combined_poor',
'wi_ur_poor', 'wi_statewise_poor', 'wi_statewise_ur_rs_poor',
'electricity', 'kaccha_floor', 'kaccha_roof', 'kaccha_walls',
'all_kaccha_house', 'own_house', 'own_agri_land', 'bpl_card',
'insurance', 'clean_fuel_usage', 'caste_General', 'caste_OBC',
'caste_SC', 'caste_ST', 'bank_acc', 'highest_edu_lvl_Higher',
'highest_edu_lvl_No education', 'highest_edu_lvl_Primary',
'highest_edu_lvl_Secondary', 'w_religion_Buddhist / Neo_Buddhist',
'w_religion_Christian', 'w_religion_Hindu', 'w_religion_Jain',
'w_religion_Muslim', 'w_religion_No religion',
'w_religion_Parsi / Zoroastrian', 'w_religion_Sikh', 'child_death',
'w_marital_status_Married', 'w_marital_status_Never in union/marriage',
'w_marital_status_widowed divorced separated deserted',
'residing_husband', 'other_wives', 'mcp_card', 'antenatal_care',
'antenatal_4plus', 'tetanus', 'birth_personnel', 'delivery_place_Home',
'delivery_place_Private', 'delivery_place_Public',
'delivery_financial_assistance', 'delivery_jsy', 'baby_checkup_2mnts',
'modern_contraceptive', 'icds_rec', 'icds_rec_bf', 'any_anaemia',
'preg_wm_any_anem', 'union_before_15', 'union_before_18', 'owns_phone',
'internet', 'stunting', 'stunting_severe', 'wasting', 'wasting_severe',
'underweight', 'underweight_severe', 'zero_dose', 'fully_vac',
'under_vacc', 'bcg', 'bcg_card', 'polio_0', 'polio_0_card', 'dpt_1',
'dpt_1_card', 'pentavalent_1', 'pentavalent_1_card', 'hepatitis_b',
'hepatitis_b_card', 'rotavirus_1', 'rotavirus_1_card', 'dpt_1_booster'

Variables when flag is set to 0:

'impr_ws', 'unimpr_ws', 'basic_drinking_w', 'limited_drinking_w',

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'jmp_w8', 'jmp_w2', 'jmp_w5', 'jmp_s1', 'jmp_s6', 'jmp_s8', 'jmp_s7',
'jmp_h1', 'jmp_h2', 'jmp_h3', 'wi_combined', 'wi_combined_score',
'wi_ur', 'wi_ur_score', 'wi_statewise', 'wi_statewise_ur_rs',
'wealth_q_HV270', 'wealth_q_HV271', 'electricity', 'kaccha_floor',
'kaccha_roof', 'kaccha_walls', 'all_kaccha_house', 'own_house',
'own_agri_land', 'bpl_card', 'insurance', 'clean_fuel_usage', 'caste',
'highest_grade_comp', 'bank_acc', 'highest_edu_lvl', 'highest_edu_year',
'w_religion', 'child_death', 'child_sex_ratio_statewise',
'child_sex_ratio_districtwise', 'child_sex_ratio_clusterwise',
'w_marital_status', 'residing_husband', 'other_wives', 'mcp_card',
'antenatal_care', 'antenatal_4plus', 'tetanus', 'birth_personnel',
'delivery_place', 'delivery_financial_assistance', 'delivery_jsy',
'baby_checkup_2mnts', 'modern_contraceptive', 'icds_rec', 'icds_rec_bf',
'w_age_marr', 'any_anaemia', 'preg_wm_any_anem', 'union_before_15',
'union_before_18', 'owns_phone', 'internet', 'stunting',
'stunting_severe', 'wasting', 'wasting_severe', 'underweight',
'underweight_severe', 'zero_dose', 'fully_vac', 'under_vacc', 'bcg',
'bcg_card', 'polio_0', 'polio_0_card', 'polio_doses', 'dpt_1',
'dpt_1_card', 'dpt_doses', 'pentavalent_1', 'pentavalent_1_card',
'pentavalent_doses', 'hepatitis_b', 'hepatitis_b_card',
'hepatitis_b_doses', 'rotavirus_1', 'rotavirus_1_card',
'rotavirus_doses', 'je_doses', 'measles_doses', 'dpt_1_booster'
```

We have merged the household and childrens record to get the household details for every child datapoint.

Verifying if there is any mismatch during the merging of household records and children's record

```
In [213]: 1 s1[['HHID', 'caseid', 'SHDIST', 'sdist']]
```

Out[213]:

	HHID	caseid	SHDIST	sdist
0	0100101357	0100101357 02	1.0	1
1	0100101358	0100101358 04	1.0	1
2	0100100690	0100100690 02	1.0	1
3	0100102986	0100102986 02	1.0	1
4	0100102945	0100102945 02	1.0	1
...
43176	3700402011	3700402011 02	4.0	4
43177	3700401534	3700401534 10	4.0	4
43178	3700401534	3700401534 14	4.0	4
43179	3700401542	3700401542 04	4.0	4
43180	3700401546	3700401546 02	4.0	4

43181 rows x 4 columns

We can see that the districts on both sides(children and household) are matching.

Metadata of recoded variables

Along with this document, an Excel file is also

attached(<https://docs.google.com/spreadsheets/d/1ypyREMqNoYry7VJ2rSggHw-NGCYxtGLv-OUZNYyAtl/edit#gid=0>). Use the link above for a better viewing experience. This contains all the variables that have been recoded. Information such as the variables used to get the recoded variable, definition, stats, etc is included.

References for where the variables have been taken from have been mentioned in the definition column, in case the reference is not mentioned, you can assume the variables have been created following the guidelines from Mira.

Modeling approach

I have attached a notebook file on a logistic regression modeling approach with this document. Using the flag and setting it to 1, we get a dataset consisting of only binary values. This is taken as input for the script and modeled on this dataset. Certain variables are removed as they have excessive empty values.

Variables used in modeling:

```
'impr_ws', 'unimpr_ws', 'basic_drinking_w', 'limited_drinking_w',  
  'jmp_w8', 'jmp_w2', 'jmp_w5', 'jmp_s1', 'jmp_s6', 'jmp_s8', 'jmp_s7',  
  'highest_grade_comp', 'jmp_h1', 'jmp_h2', 'jmp_h3', 'wi_combined_poor',  
  'wi_ur_poor', 'wi_statewise_poor', 'wi_statewise_ur_rs_poor',  
  'electricity', 'kaccha_floor', 'kaccha_roof', 'kaccha_walls',  
  'all_kaccha_house', 'own_house', 'own_agri_land', 'bpl_card',  
  'insurance', 'clean_fuel_usage', 'caste_General', 'caste_OBC',  
  'caste_SC', 'caste_ST', 'bank_acc', 'highest_edu_lvl_Higher',  
  'highest_edu_lvl_No education', 'highest_edu_lvl_Primary',  
  'highest_edu_lvl_Secondary', 'w_religion_Buddhist / Neo_Buddhist',  
  'w_religion_Christian', 'w_religion_Hindu', 'w_religion_Jain',  
  'w_religion_Muslim', 'w_religion_No religion',  
  'w_religion_Parsi / Zoroastrian', 'w_religion_Sikh', 'child_death',  
  'w_marital_status_Married', 'w_marital_status_Never in union/marriage',  
  'w_marital_status_widowed divorced separated deserted', 'mcp_card', 'antenatal_care',  
  'antenatal_4plus', 'tetanus', 'birth_personnel', 'delivery_place_Home',  
  'delivery_place_Private', 'delivery_place_Public',  
  'delivery_financial_assistance', 'delivery_jsy', 'baby_checkup_2mnts',  
  'modern_contraceptive', 'icds_rec', 'icds_rec_bf', 'any_anaemia',  
  'preg_wm_any_anem', 'union_before_15', 'union_before_18', 'owns_phone',  
  'internet', 'stunting', 'stunting_severe', 'wasting', 'wasting_severe',  
  'underweight', 'underweight_severe', 'zero_dose', 'fully_vac',  
  'under_vacc', 'bcg', 'bcg_card', 'polio_0', 'polio_0_card', 'dpt_1',  
  'dpt_1_card', 'pentavalent_1', 'pentavalent_1_card', 'hepatitis_b',  
  'hepatitis_b_card', 'rotavirus_1', 'rotavirus_1_card', 'dpt_1_booster'
```

The modelling approach used here is logistic regression, and in order to run such a model, the best input is binary input and not continuous input. Hence, I converted a few variables into binary variables. Let's look at the dataset at first. We have a total of 43,181 datapoints. There are many empty values in this, so we remove all rows containing any empty values and end up with 31,457 datapoints. The distribution of the values for zero_dose is 27,498 values as False and 2,770 values as True. We see an imbalance in the ratio of True and False values. To counter this, we will use a few undersampling and oversampling approaches.

Below is a table with the counts after using a few sampling approaches. The samplings have been achieved using imblearn library.

	Dataset	Unsampled Train	Unsampled Test	Undersampling Train	Undersampling Test	Oversampling Train	Oversampling Test	Oversampling Train	Oversampling Test
								SMOTE	
False Datapoint	27498	19237	8261	1950	8261	19237	8261	19237	8261
True Datapoint	2770	1950	820	1950	820	19237	820	19237	820

Using this, a model was trained on logistic regression. Details of it's accuracy is given below.

Test/Train	Accuracy	Precision	Recall	F1	Datapoints
No sampling on True datapoint	0.84	0.32	0.67	0.44	820
No sampling on True datapoint	0.86	0.36	0.70	0.47	1950
Undersampling on True datapoint	0.84	0.32	0.67	0.44	820
Undersampling on True datapoint	0.78	0.84	0.71	0.77	1950
Oversampling on True datapoint	0.84	0.32	0.67	0.43	820
Oversampling on True datapoint	0.79	0.84	0.71	0.77	19237
Oversampling SMOTE on True datapoint	0.84	0.30	0.58	0.39	820
Oversampling SMOTE on True datapoint	0.85	0.86	0.82	0.84	19237

After going through a few parameters in logistic regression, i came across a parameter class_weight in the logistic regression model. This would assign the minority class a higher weight, hence I tried using this and compared the results with my previous approach.

Test/Train	Accuracy	Precision	Recall	F1	Datapoints
No sampling on True datapoint	0.91	0.48	0.07	0.13	820
No sampling on True datapoint	0.91	0.55	0.07	0.13	1950
Undersampling on True datapoint	0.84	0.32	0.67	0.44	820
Undersampling on True datapoint	0.78	0.84	0.71	0.77	1950
Oversampling on True datapoint	0.84	0.32	0.67	0.43	820
Oversampling on True datapoint	0.79	0.84	0.71	0.77	19237
Oversampling SMOTE on True datapoint	0.84	0.30	0.58	0.39	820
Oversampling SMOTE on True datapoint	0.85	0.86	0.82	0.84	19237

I noticed the accuracy to be slightly higher in the latter case.

In order to find out with variables must be included in the analysis, we did run a pearson and cramer V orrelation. It did not assist us much as the correlations weren't strong.