



First there was an idea:

Find interesting dataset
Apply gathered knowledge
Make something usefull

No1

We found **mushrooms.csv**
from **kaggle**

(<https://www.kaggle.com/uciml/mushroom-classification?select=mushrooms.csv>)

No missing values
No duplicates
23 columns and 8124 unique rows
Each row = mushroom
Each column = mushroom's parameter

Columns provide information such as:
shape and color of different parts of mushroom
where mushroom may be found
how big is the population
odor
bruices
and main column - "class" (edible or poisonous)

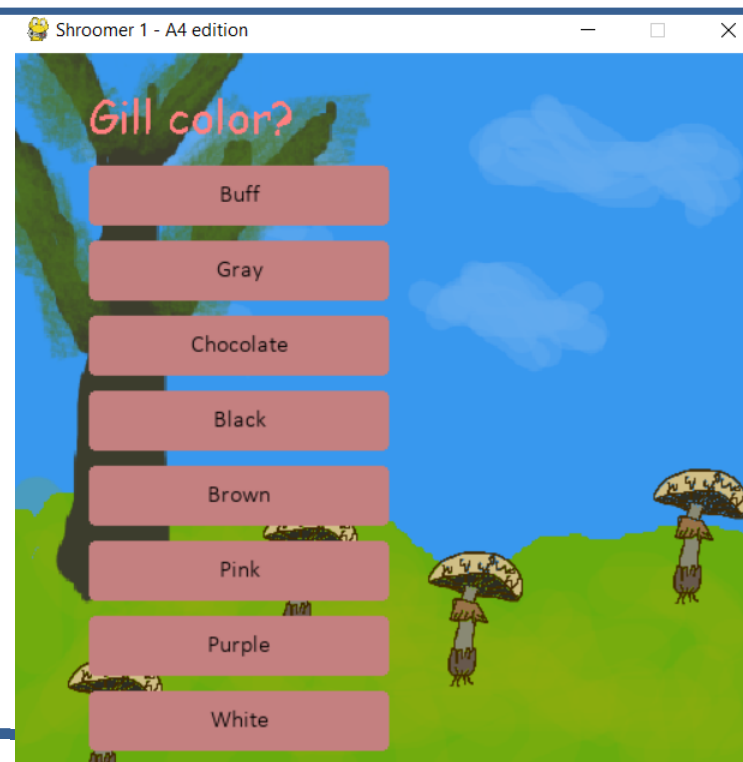
All information is categorical.

"Hmm... it is good data set for project" - we thought

And decided to:

Clear the data (we did not know that this is already clear)
Train some models to predict edability
Find strongest correlations between edability and other parameters
Find interesting correlations among other attributes
Write a programm for mushroom gatherers based on trained model

No3

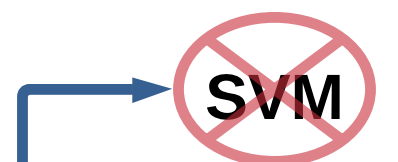


All tests led us to developing simple but usefull
programm written on python with use of pygame
Programm asks questions about mushroom
parameter and then predicts edability of
mushroom using trained model



No2

- 1) Import data and check for duplicates and missing values
there was non of those
- 2) Use hot-encoding to be able to train ML models
map function and dummies turned our 23 columns into 114 columns
- 3) Use apriori algorithm and visualized decision tree to find interesting connections between attributes
we found, that if a mushroom has no odor and no green spore prints - it is probably eadible.
while this rule may be useful for some novice shroomers, we had decided to not stop there, since
those 2 attributes are in our opinion somewhat hard to measure.
- 3) Run Lasso and Ridge regression with best alphas and watch, which attributes had high coefficients
regression does not really help with classification, but model still search for useful corellations and try
to give values as close as possible to 1 (edible) or 0 (poisonous)
Both regression models was agree on 'odor' and 'spore color' parameters, but Lasso prefers 'stalk
color' as 3rd most valuable parameter, Ridge preferred 'ring type'
- 4) Run KNN and RandomForest models, trained only on parameters with high coefficients from Lasso
and Ridge regression. This led us to model with 99.7%(on 3 valuable parameters from Lasso) and
99.5% (on 3 valuable parameters from Ridge) accuracy.



We did not even try SVM
because encoded dataset
is too high dimensional

Based on 'odor', 'spore-print-color' and 'stalk-color-below-ring'

Knn prediction score: 0.9975381585425899

Random Forest prediction score: 0.9975381585425899

Based on 'odor', 'spore-print-color' and 'ring-type'

Knn prediction score: 0.9950763170851797

Random Forest prediction score: 0.9950763170851797

which was pretty good, but not very usefull for mushroom gatherers. Odor is too subjective and
spore color too hard to notice

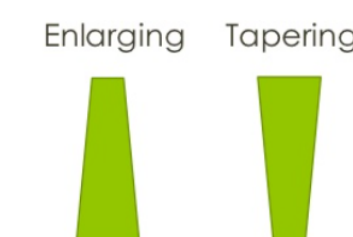
- 5) Return to original data set and drop columns with parameters that are hard to define.
such as 'odor', 'spore print color', 'population', etc. 13 columns left
- 6) Run Lasso and Ridge regression with parameters that are left after cleaning
models were agree on 3 parameters, that are very easy to define, that are:

Gill colors

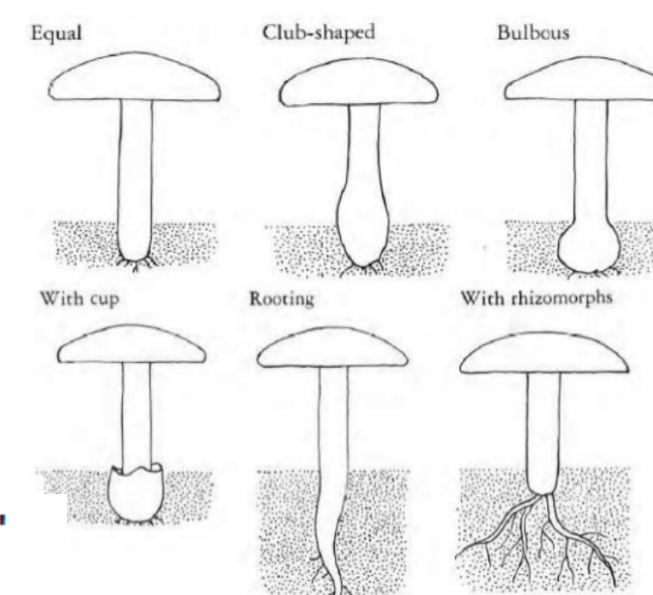
brown, orange,
white, yellow



Stalk shapes



Stalk root shapes



And prediction score
reduces by less than 5%

Based on 'stalk-shape', 'gill-color' and 'stalk-root'

Knn prediction score: 0.9533932951757972

Random Forest prediction score: 0.9533932951757972