

Mushrooms

Ina Ding

Introduction

This section includes an introduction to the project motivation, data, and research question.

The research question and motivation are clearly stated in the introduction, including citations.

The primary goal of this project is to best predict whether or not a mushroom is poisonous depending on various physical characteristics, rarity, and habitat of the fungus. The data set consists of 8124 hypothetical samples, constructed from the Audobon Society Field Guide. The samples correspond to 23 species of mushrooms from the Agaricus and Lepiota Families. Each mushroom is categorized as either poisonous or edible, with mushrooms ‘not recommended for eating’ or of unknown edibility are counted as poisonous. Though the observations are hypothetical mushrooms, analyzing them can still provide beneficial results that can be applied to help identify the edibility of the near 14,000 existing species of mushrooms.

1. cap-shape:	bell=b,conical=c,convex=x,flat=f, knobbed=k,sunken=s
2. cap-surface:	fibrous=f,grooves=g,scaly=y,smooth=s
3. cap-color:	brown=n,buff=b,cinnamon=c,gray=g,green=r, pink=p,purple=u,red=e,white=w,yellow=y
4. bruises?:	bruises=t,no=f
5. odor:	almond=a,anise=l,creosote=c,fishy=y,foul=f, musty=m,none=n,pungent=p,spicy=s
6. gill-attachment:	attached=a,descending=d,free=f,notched=n
7. gill-spacing:	close=c,crowded=w,distant=d
8. gill-size:	broad=b,narrow=n
9. gill-color:	black=k,brown=n,buff=b,chocolate=h,gray=g, green=r,orange=o,pink=p,purple=u,red=e, white=w,yellow=y
10. stalk-shape:	enlarging=e,tapering=t

11. stalk-root:	bulbous=b,club=c,cup=u,equal=e, rhizomorphs=z,rooted=r,missing
12. stalk-surface-above-ring:	fibrous=f,scaly=y,silky=k,smooth=s
13. stalk-surface-below-ring:	fibrous=f,scaly=y,silky=k,smooth=s
14. stalk-color-above-ring:	brown=n,buff=b,cinnamon=c,gray=g,orange=o, pink=p,red=e,white=w,yellow=y
15. stalk-color-below-ring:	brown=n,buff=b,cinnamon=c,gray=g,orange=o, pink=p,red=e,white=w,yellow=y
16. veil-type:	partial=p,universal=u
17. veil-color:	brown=n,orange=o,white=w,yellow=y
18. ring-number:	none=n,one=o,two=t
19. ring-type:	cobwebby=c,evanescent=e,flaring=f,large=l, none=n,pendant=p,sheathing=s,zone=z
20. spore-print-color:	black=k,brown=n,buff=b,chocolate=h,green=r, orange=o,purple=u,white=w,yellow=y
21. population:	abundant=a,clustered=c,numerous=n, scattered=s,several=v,solita
22. habitat:	grasses=g,leaves=l,meadows=m,paths=p, urban=u,waste=w,woods=d

Citations Mushroom. (1987). UCI Machine Learning Repository. <https://doi.org/10.24432/C5959T>.

```
install.packages("leaps")
```

```
mushrooms <- read.csv("Mushrooms - Sheet1.csv")
library(tidymodels)
library(tidyverse)
library(leaps)
library(glmnet)
```

```
no_veil <- subset(mushrooms, select = -c(veil))
sapply(lapply(no_veil, unique), length)
```

poisonous	cap.shape	cap.surface	cap.color	bruises
2	6	4	10	2
odor	gill.attachment	gill.spacing	gill.size	gill.color
9	2	2	2	12
stalk.shape	stalk.root	ss.above	ss.below	sc.above
2	5	4	4	9
sc.below	veil.color	ring.num	ring.type	spore.color
9	4	3	5	9
population	habitat			
6	7			

```

no_veil$poisonous[no_veil$poisonous == 'p'] <- 1
no_veil$poisonous[no_veil$poisonous == 'e'] <- 0

no_veil$poisonous <- as.numeric(as.character(no_veil$poisonous))

##m1 <- glm(poisonous ~ .,
            ## data = no_veil,
            ## family = "binomial")
##summary(m1)

m_all <- regsubsets(poisonous ~ .,
                    data = no_veil,
                    nbest = 1, nvmax = 5, really.big=T)

```

Reordering variables and trying again:

```
m_all
```

Subset selection object

Call: regsubsets.formula(poisonous ~ ., data = no_veil, nbest = 1,
nvmax = 5, really.big = T)

95 Variables (and intercept)

	Forced in	Forced out
cap.shapec	FALSE	FALSE
cap.shapef	FALSE	FALSE
cap.shapek	FALSE	FALSE
cap.shapes	FALSE	FALSE
cap.shapex	FALSE	FALSE
cap.surfaceg	FALSE	FALSE
cap.surfaces	FALSE	FALSE
cap.surfacey	FALSE	FALSE
cap.colorc	FALSE	FALSE
cap.colore	FALSE	FALSE
cap.colorg	FALSE	FALSE
cap.colorn	FALSE	FALSE
cap.colorp	FALSE	FALSE
cap.colorr	FALSE	FALSE
cap.coloru	FALSE	FALSE
cap.colorw	FALSE	FALSE
cap.colory	FALSE	FALSE

sc.belown	FALSE	FALSE
sc.belowp	FALSE	FALSE
sc.beloww	FALSE	FALSE
sc.belowy	FALSE	FALSE
veil.coloro	FALSE	FALSE
ring.numo	FALSE	FALSE
ring.typef	FALSE	FALSE
ring.typep	FALSE	FALSE
spore.colork	FALSE	FALSE
spore.colorn	FALSE	FALSE
spore.coloro	FALSE	FALSE
spore.colorr	FALSE	FALSE
spore.coloru	FALSE	FALSE
spore.colorw	FALSE	FALSE
spore.colory	FALSE	FALSE
populationc	FALSE	FALSE
populationn	FALSE	FALSE
populations	FALSE	FALSE
populationv	FALSE	FALSE
populationy	FALSE	FALSE
habitatg	FALSE	FALSE
habitatl	FALSE	FALSE
habitatm	FALSE	FALSE
habitatp	FALSE	FALSE
habitatu	FALSE	FALSE
sc.abovec	FALSE	FALSE
sc.belowc	FALSE	FALSE
sc.belowo	FALSE	FALSE
veil.colorw	FALSE	FALSE
veil.colory	FALSE	FALSE
ring.numt	FALSE	FALSE
ring.type1	FALSE	FALSE
ring.typen	FALSE	FALSE
spore.colorh	FALSE	FALSE
habitatw	FALSE	FALSE

1 subsets of each size up to 6
Selection Algorithm: exhaustive

```
summary(m_all)$which
```

```
(Intercept) cap.shapec cap.shapef cap.shapek cap.shapes cap.shapex
```

1	TRUE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
2	TRUE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
3	TRUE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
4	TRUE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
5	TRUE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
6	TRUE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE

	cap.surfaceg	cap.surfaces	cap.surfacey	cap.colorc	cap.colore	cap.colorg	
1	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
2	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
3	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
4	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
5	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
6	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE

	cap.colorn	cap.colorp	cap.colorr	cap.coloru	cap.colorw	cap.colory	bruiseest
1	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
2	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
3	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
4	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
5	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
6	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE

	odorc	odorf	odorl	odorm	odorn	odorp	odors	odory	gill.attachmentf
1	FALSE	FALSE	FALSE	FALSE	TRUE	FALSE	FALSE	FALSE	FALSE
2	FALSE	FALSE	FALSE	FALSE	TRUE	FALSE	FALSE	FALSE	FALSE
3	FALSE	FALSE	FALSE	FALSE	TRUE	FALSE	FALSE	FALSE	FALSE
4	FALSE	FALSE	FALSE	FALSE	TRUE	FALSE	FALSE	FALSE	FALSE
5	FALSE	FALSE	FALSE	FALSE	TRUE	FALSE	FALSE	FALSE	FALSE
6	FALSE	FALSE	FALSE	FALSE	TRUE	FALSE	FALSE	FALSE	FALSE

	gill.spacingw	gill.sizen	gill.colore	gill.colorg	gill.colorh	gill.colork	
1	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
2	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
3	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
4	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
5	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
6	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE

	gill.colorn	gill.coloro	gill.colorp	gill.colorr	gill.coloru	gill.colorw	
1	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
2	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
3	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
4	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
5	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
6	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE

	gill.colory	stalk.shapet	stalk.rootb	stalk.rootc	stalk.roote	stalk.rootr	
1	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE

2	FALSE	FALSE	FALSE	TRUE	FALSE	FALSE
3	FALSE	FALSE	FALSE	TRUE	FALSE	TRUE
4	FALSE	FALSE	FALSE	TRUE	FALSE	TRUE
5	FALSE	FALSE	FALSE	TRUE	FALSE	TRUE
6	FALSE	FALSE	FALSE	TRUE	FALSE	TRUE

	ss.abovek	ss.aboves	ss.abovey	ss.belowk	ss.belows	ss.belowy	sc.abovec
1	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
2	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
3	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
4	FALSE	FALSE	FALSE	FALSE	FALSE	TRUE	FALSE
5	FALSE	FALSE	FALSE	FALSE	FALSE	TRUE	FALSE
6	FALSE	FALSE	FALSE	FALSE	FALSE	TRUE	FALSE

	sc.abovee	sc.aboveg	sc.aboven	sc.aboveo	sc.abovep	sc.abovev	sc.abovey
1	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
2	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
3	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
4	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
5	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
6	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE

	sc.belowc	sc.belowe	sc.belowg	sc.belown	sc.belowo	sc.belowp	sc.beloww
1	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
2	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
3	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
4	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
5	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
6	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE

	sc.belowy	veil.coloro	veil.colorw	veil.colory	ring.numo	ring.numt	ring.typef
1	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
2	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
3	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
4	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
5	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
6	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE

	ring.typel	ring.typen	ring.typep	spore.colorh	spore.colork	spore.colorn
1	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
2	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
3	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
4	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
5	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
6	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE

	spore.coloro	spore.colorr	spore.coloru	spore.colorw	spore.colory	populationc
1	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
2	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE

3	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
4	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
5	FALSE	TRUE	FALSE	FALSE	FALSE	FALSE	FALSE
6	FALSE	TRUE	TRUE	FALSE	FALSE	FALSE	FALSE
	populationm	populations	populationv	populationy	habitatg	habitatl	habitatm
1	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
2	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
3	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
4	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
5	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
6	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
	habitatp	habitatv	habitatw				
1	FALSE	FALSE	FALSE				
2	FALSE	FALSE	FALSE				
3	FALSE	FALSE	FALSE				
4	FALSE	FALSE	FALSE				
5	FALSE	FALSE	FALSE				
6	FALSE	FALSE	FALSE				

```
summary(m_all)$rsq
```

```
[1] 0.6170992 0.7850914 0.8701623 0.9041082 0.9387674 0.9614494
```

```
install.packages("glmnet")
```

```
library(glmnet)
```

```
y <- no_veil$poisonous
x <- model.matrix(poisonous ~ .,
                  data = no_veil)
```

```
m_lasso_cv <- cv.glmnet(x, y, alpha = 1)
best_lambda <- m_lasso_cv$lambda.min
best_lambda
```

```
[1] 0.0002523176
```

```
m_best <- glmnet(x, y, alpha = 1, lambda = best_lambda)
m_best$beta
```


96 x 1 sparse Matrix of class "dgCMatrix"

```

              s0
(Intercept)      .
cap.shapec       2.469270e-01
cap.shapef       .
cap.shapek       .
cap.shapes      -4.780328e-03
cap.shapex       .
cap.surfacecg    7.481027e-01
cap.surfaces     .
cap.surfacey     7.228459e-04
cap.colorc      -6.613558e-02
cap.colore       .
cap.colorg       .
cap.colorn      -9.789276e-04
cap.colorp       .
cap.colorr       .
cap.coloru       .
cap.colorw       2.720506e-03
cap.colory       .
bruiseest       6.459105e-04
odorc           9.665179e-01
odorf           7.966668e-01
odorl          -4.046078e-03
odorm           5.659035e-02
odorn          -4.517302e-02
odorp           9.423022e-01
odors           7.963789e-01
odory           7.964045e-01
gill.attachmentf .
gill.spacingw   -1.719982e-02
gill.sizen      .
gill.colore     .
gill.colorg     .
gill.colorh     .
gill.colork     .
gill.colorn     .
gill.coloro     .
gill.colorp     .
gill.colorr     .
gill.coloru     .
gill.colorw     -1.959178e-04
gill.colory     .

```

stalk.shapet	6.117153e-03
stalk.rootb	-7.871970e-03
stalk.rootc	-4.573025e-02
stalk.roote	7.148989e-03
stalk.rootr	-8.618155e-01
ss.abovek	2.321158e-03
ss.aboves	.
ss.abovey	-8.035657e-01
ss.belowk	6.740357e-04
ss.belows	.
ss.belowy	8.175232e-01
sc.abovec	3.588411e-02
sc.abovee	.
sc.aboveg	.
sc.aboven	.
sc.aboveo	-7.569320e-03
sc.abovep	1.198385e-04
sc.abovew	.
sc.abovey	4.736608e-01
sc.belowc	2.978803e-06
sc.belowe	.
sc.belowg	.
sc.belown	-9.342302e-03
sc.belowo	-1.576968e-02
sc.belowp	.
sc.beloww	1.350767e-03
sc.belowy	2.835038e-02
veil.coloro	.
veil.colorw	.
veil.colory	2.771655e-01
ring.numo	9.290282e-02
ring.numt	-4.116992e-02
ring.typef	-1.562118e-01
ring.typel	8.587265e-03
ring.typhen	.
ring.typep	.
spore.colorh	1.555109e-01
spore.colork	.
spore.colorn	-6.014691e-04
spore.coloro	.
spore.colorr	1.135084e+00
spore.coloru	-2.520651e-02
spore.colorw	1.481708e-01

spore.colory	.
populationc	5.019038e-02
populationn	-7.732554e-04
populations	.
populationv	.
populationy	.
habitatg	.
habitatl	.
habitatm	.
habitatp	8.385491e-05
habitatu	.
habitatw	-6.500746e-02