DZ11-Marcec-Lea.R

Lea

2022-01-13

```
data(iris)
str(iris)
```

```
## 'data.frame': 150 obs. of 5 variables:
## $ Sepal.Length: num 5.1 4.9 4.7 4.6 5 5.4 4.6 5 4.4 4.9 ...
## $ Sepal.Width : num 3.5 3 3.2 3.1 3.6 3.9 3.4 3.4 2.9 3.1 ...
## $ Petal.Length: num 1.4 1.4 1.3 1.5 1.4 1.7 1.4 1.5 1.4 1.5 ...
## $ Petal.Width : num 0.2 0.2 0.2 0.2 0.2 0.4 0.3 0.2 0.2 0.1 ...
## $ Species : Factor w/ 3 levels "setosa", "versicolor", ..: 1 1 1 1 1 1 1 1
1 ...
```

```
# priprema podataka
df <- iris
df</pre>
```

##	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
## 1	5.1	3.5	1.4	0.2	setosa
## 2	4.9	3.0	1.4	0.2	setosa
## 3	4.7	3.2	1.3	0.2	setosa
## 4	4.6	3.1	1.5	0.2	setosa
## 5	5.0	3.6	1.4	0.2	setosa
## 6	5.4	3.9	1.7	0.4	setosa
## 7	4.6	3.4	1.4	0.3	setosa
## 8	5.0	3.4	1.5	0.2	setosa
## 9	4.4	2.9	1.4	0.2	setosa
## 1(4.9	3.1	1.5	0.1	setosa
## 13	5.4	3.7	1.5	0.2	setosa
## 12	4.8	3.4	1.6	0.2	setosa
## 13	4.8	3.0	1.4	0.1	setosa
## 14	4.3	3.0	1.1	0.1	setosa
## 15	5.8	4.0	1.2	0.2	setosa
## 16	5.7	4.4	1.5	0.4	setosa
## 17	5.4	3.9	1.3	0.4	setosa
## 18	5.1	3.5	1.4	0.3	setosa
## 19	5.7	3.8	1.7	0.3	setosa
## 20	5.1	3.8	1.5	0.3	setosa
## 21	5.4	3.4	1.7	0.2	setosa
## 22	5.1	3.7	1.5	0.4	setosa
## 23	4.6	3.6	1.0	0.2	setosa
## 24	5.1	3.3	1.7	0.5	setosa
## 25	4.8	3.4	1.9	0.2	setosa
## 26	5.0	3.0	1.6	0.2	setosa
## 27	5.0	3.4	1.6	0.4	setosa
## 28	5.2	3.5	1.5	0.2	setosa
## 29	5.2	3.4	1.4	0.2	setosa
## 30	4.7	3.2	1.6	0.2	setosa
## 31	4.8	3.1	1.6	0.2	setosa

## 32	5.4	3.4	1.5	0.4 setosa	
## 33	5.2	4.1	1.5	0.1 setosa	
## 34	5.5	4.2	1.4	0.2 setosa	
## 35	4.9	3.1	1.5	0.2 setosa	
## 36	5.0	3.2	1.2	0.2 setosa	
## 37	5.5	3.5	1.3	0.2 setosa	
## 38	4.9	3.6	1.4	0.1 setosa	
## 39	4.4	3.0	1.3	0.2 setosa	
## 40	5.1	3.4	1.5	0.2 setosa	
## 41	5.0	3.5	1.3	0.3 setosa	
## 42	4.5	2.3	1.3	0.3 setosa	
## 43	4.4	3.2	1.3	0.2 setosa	
## 44	5.0	3.5	1.6	0.6 setosa	
## 45	5.1	3.8	1.9	0.4 setosa	
## 46	4.8	3.0	1.4	0.3 setosa	
## 47	5.1	3.8	1.6	0.2 setosa	
## 48	4.6	3.2	1.4	0.2 setosa	
## 49	5.3	3.7	1.5	0.2 setosa	
## 50	5.0	3.3	1.4	0.2 setosa	
## 51	7.0	3.2	4.7	1.4 versicolor	
## 52	6.4	3.2	4.5	1.5 versicolor	
## 53	6.9	3.1	4.9	1.5 versicolor	
## 54	5.5	2.3	4.0	1.3 versicolor	
## 55	6.5	2.8	4.6	1.5 versicolor	
## 56	5.7	2.8	4.5	1.3 versicolor	
## 57	6.3	3.3	4.7	1.6 versicolor	
## 58	4.9	2.4	3.3	1.0 versicolor	
## 59	6.6	2.9	4.6	1.3 versicolor	
## 60	5.2	2.7	3.9	1.4 versicolor	
## 61	5.0	2.0	3.5	1.0 versicolor	
## 62	5.9	3.0	4.2	1.5 versicolor	
## 63	6.0	2.2	4.0	1.0 versicolor	
## 64	6.1	2.9	4.7	1.4 versicolor	
## 65	5.6	2.9	3.6	1.3 versicolor	
## 66	6.7	3.1	4.4	1.4 versicolor	
## 67	5.6	3.0	4.5	1.5 versicolor	
## 68	5.8	2.7	4.1	1.0 versicolor	
## 69	6.2	2.2	4.5	1.5 versicolor	
## 70	5.6	2.5	3.9	1.1 versicolor	
## 71	5.9	3.2	4.8	1.8 versicolor	
## 72	6.1	2.8	4.0	1.3 versicolor	
## 73	6.3	2.5	4.9	1.5 versicolor	
## 74	6.1	2.8	4.7	1.2 versicolor	
## 75	6.4	2.9	4.3	1.3 versicolor	
## 76	6.6	3.0	4.4	1.4 versicolor	
## 77	6.8	2.8	4.8	1.4 versicolor	
## 78	6.7	3.0	5.0	1.7 versicolor	
## 79	6.0	2.9	4.5	1.5 versicolor	
## 80	5.7	2.6	3.5	1.0 versicolor	
## 81	5.5	2.4	3.8	1.1 versicolor	
## 82	5.5	2.4	3.7	1.0 versicolor	
## 83	5.8	2.7	3.9	1.2 versicolor	
## 84	6.0	2.7	5.1	1.6 versicolor	
## 85	5.4	3.0	4.5	1.5 versicolor	
## 86	6.0	3.4	4.5	1.6 versicolor	
## 87	6.7	3.1	4.7	1.5 versicolor	
## 88	6.3	2.3	4.4	1.3 versicolor	
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## 89	5.6	3.0	4.1	1.3 versicolor
## 90	5.5	2.5	4.0	1.3 versicolor
## 91	5.5	2.6	4.4	1.2 versicolor
## 92	6.1	3.0	4.6	1.4 versicolor
## 93	5.8	2.6	4.0	1.2 versicolor
## 94	5.0	2.3	3.3	1.0 versicolor
## 95	5.6	2.7	4.2	1.3 versicolor
## 96	5.7	3.0	4.2	1.2 versicolor
## 97	5.7	2.9	4.2	1.3 versicolor
		2.9	4.3	1.3 versicolor
	6.2			
## 99	5.1	2.5	3.0	1.1 versicolor
## 100	5.7	2.8	4.1	1.3 versicolor
## 101	6.3	3.3	6.0	2.5 virginica
## 102	5.8	2.7	5.1	1.9 virginica
## 103	7.1	3.0	5.9	2.1 virginica
## 104	6.3	2.9	5.6	1.8 virginica
## 105	6.5	3.0	5.8	2.2 virginica
## 106	7.6	3.0	6.6	2.1 virginica
## 107	4.9	2.5	4.5	1.7 virginica
## 108	7.3	2.9	6.3	1.8 virginica
## 109	6.7	2.5	5.8	1.8 virginica
## 110	7.2	3.6	6.1	2.5 virginica
## 111	6.5	3.2	5.1	2.0 virginica
## 112	6.4	2.7	5.3	1.9 virginica
## 113	6.8	3.0	5.5	2.1 virginica
## 114	5.7	2.5	5.0	2.0 virginica
## 115	5.8	2.8	5.1	2.4 virginica
## 116	6.4	3.2	5.3	2.3 virginica
## 117	6.5	3.0	5.5	1.8 virginica
## 118	7.7	3.8	6.7	2.2 virginica
## 119	7.7	2.6	6.9	2.3 virginica
## 120	6.0	2.2	5.0	1.5 virginica
## 121	6.9	3.2	5.7	2.3 virginica
## 122	5.6	2.8	4.9	2.0 virginica
## 123	7.7	2.8	6.7	2.0 virginica
## 124	6.3	2.7	4.9	1.8 virginica
## 125	6.7	3.3	5.7	2.1 virginica
## 126	7.2	3.2	6.0	1.8 virginica
## 127	6.2	2.8	4.8	1.8 virginica
## 128	6.1	3.0	4.9	1.8 virginica
## 129	6.4	2.8	5.6	2.1 virginica
## 130	7.2	3.0	5.8	1.6 virginica
## 131	7.4	2.8	6.1	1.9 virginica
## 132	7.9	3.8	6.4	2.0 virginica
## 133	6.4	2.8	5.6	2.2 virginica
## 134	6.3	2.8	5.1	1.5 virginica
## 135	6.1	2.6	5.6	1.4 virginica
## 136	7.7	3.0	6.1	2.3 virginica
## 137	6.3	3.4	5.6	2.4 virginica
## 138	6.4	3.1	5.5	1.8 virginica
## 139	6.0	3.0	4.8	1.8 virginica
## 140	6.9	3.1	5.4	2.1 virginica
## 141	6.7	3.1	5.6	2.4 virginica
## 142	6.9	3.1	5.1	2.3 virginica
## 143	5.8	2.7	5.1	1.9 virginica
## 144	6.8	3.2	5.9	2.3 virginica
## 145	6.7	3.3	5.7	2.5 virginica
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```
## 146
            6.7
                       3.0
                                   5.2
                                              2.3 virginica
                                   5.0
                                              1.9 virginica
## 147
             6.3
                       2.5
                                   5.2
## 148
            6.5
                                              2.0 virginica
                        3.0
## 149
             6.2
                        3.4
                                   5.4
                                              2.3 virginica
                                   5.1 1.8 virginica
## 150
             5.9
                        3.0
str(df)
## 'data.frame': 150 obs. of 5 variables:
## $ Sepal.Length: num 5.1 4.9 4.7 4.6 5 5.4 4.6 5 4.4 4.9 ...
## $ Sepal.Width : num 3.5 3 3.2 3.1 3.6 3.9 3.4 3.4 2.9 3.1 ...
## $ Petal.Length: num 1.4 1.4 1.3 1.5 1.4 1.7 1.4 1.5 1.4 1.5 ...
## $ Petal.Width : num 0.2 0.2 0.2 0.2 0.4 0.3 0.2 0.2 0.1 ...
## $ Species : Factor w/ 3 levels "setosa", "versicolor", ...: 1 1 1 1 1 1 1 1 1
1 ...
df$Species <- factor(df$Species, levels = c("setosa", "versicolor", "virginica"),</pre>
labels = c("Setosa", "Versicolor", "Virginica"))
# slucajan odabir skupa za ucenje i skupa za testiranje
set.seed(1234)
train <- sample(nrow(df), 0.7*nrow(df))</pre>
# skup za ucenje
df.train <- df[train,]</pre>
# skup za testiranje
df.validate <- df[-train,]</pre>
# tablica frevencija - originalan skup podataka
table(df$Species)
##
     Setosa Versicolor Virginica
##
##
      50 50
                        50
# tablica frekvenvija - skup za ucenje
table(df.train$Species)
##
##
    Setosa Versicolor Virginica
      39 29 37
# tablica frekvencija - skup za testiranje
table(df.validate$Species)
##
##
     Setosa Versicolor Virginica
      11 21 13
##
```

```
# LOGISTICKA REGRESIJA
# varijabla Species je zavisna varijabla, a varijable Sepal.Length, Sepal.Width, P
etal.Length, Petal.Width su prediktori
fit.logit <- glm(Species~., data = df.train, family=binomial())</pre>
## Warning: glm.fit: algorithm did not converge
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
# analiza modela
summary(fit.logit)
##
## Call:
## glm(formula = Species ~ ., family = binomial(), data = df.train)
## Deviance Residuals:
              1Q Median
   Min
                                             3Q
                                                       Max
## -2.124e-05 -2.110e-08 2.110e-08 2.110e-08 2.105e-05
##
## Coefficients:
               Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                -10.934 351258.549 0
                  -3.339 92833.858
                                         0
                                                   1
## Sepal.Length
## Sepal.Width
                 -11.549 64471.626
                                         0
                                                  1
                 22.570 80185.641
## Petal.Length
                                         0
                                                   1
## Petal.Width
                  3.230 163837.188
                                         0
##
\#\# (Dispersion parameter for binomial family taken to be 1)
##
     Null deviance: 1.3854e+02 on 104 degrees of freedom
## Residual deviance: 1.8975e-09 on 100 degrees of freedom
## AIC: 10
##
## Number of Fisher Scoring iterations: 25
# testiranje modela sa novim podatcima
prob <- predict(fit.logit, df.validate, type = "response")</pre>
logit.pred <- factor(prob >.5, levels = c(FALSE, TRUE), labels = c("LAZ", "ISTINA"
# predikcija modela i stvarni rezultat
logit.perf <- table(df.validate$Species, logit.pred, dnn = c("Stvarni rezultat",</pre>
"Predikcija"))
# tablicni prikaz rezultata
logit.perf
                  Predikcija
## Stvarni rezultat LAZ ISTINA
##
       Setosa 11 0
```

```
##
       Versicolor 0 21
        Virginica 0
##
                          13
# Stepwise logisticka regresija - uklanjanje prediktora koji nisu statisticki znac
fit.reducedlogit <- step(fit.logit)</pre>
## Start: AIC=10
## Species ~ Sepal.Length + Sepal.Width + Petal.Length + Petal.Width
## Warning: glm.fit: algorithm did not converge
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: algorithm did not converge
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: algorithm did not converge
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: algorithm did not converge
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
##
                 Df Deviance AIC
## - Petal.Width 1 1.9170e-09 8
## - Sepal.Length 1 1.9432e-09 8
## - Sepal.Width 1 2.5184e-09 8
## - Petal.Length 1 2.9548e-09 8
## <none>
                   1.8975e-09 10
## Warning: glm.fit: algorithm did not converge
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
##
## Step: AIC=8
## Species ~ Sepal.Length + Sepal.Width + Petal.Length
## Warning: glm.fit: algorithm did not converge
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: algorithm did not converge
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: algorithm did not converge
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
                Df Deviance AIC
##
## - Sepal.Length 1 1.9435e-09 6
## - Sepal.Width 1 2.6657e-09 6
## - Petal.Length 1 1.0303e-08 6
## <none>
                   1.9170e-09 8
## Warning: glm.fit: algorithm did not converge
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
##
## Step: AIC=6
## Species ~ Sepal.Width + Petal.Length
## Warning: glm.fit: algorithm did not converge
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
##
                 Df Deviance AIC
## - Sepal.Width 1 0.000 4.000
                      0.000 6.000
## <none>
## - Petal.Length 1 95.637 99.637
## Warning: glm.fit: algorithm did not converge
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
##
## Step: AIC=4
## Species ~ Petal.Length
##
               Df Deviance AIC
##
## <none>
                0.00 4.00
## - Petal.Length 1 138.54 140.54
summary(fit.reducedlogit)
##
## Call:
## glm(formula = Species ~ Petal.Length, family = binomial(), data = df.train)
##
## Deviance Residuals:
```

```
10
                           Median
## -3.680e-05 -2.100e-08 2.100e-08 2.100e-08 4.269e-05
##
## Coefficients:
             Estimate Std. Error z value Pr(>|z|)
##
## (Intercept) -78.02 60970.43 -0.001 0.999
## Petal.Length
                29.95 21853.16 0.001
                                          0.999
## (Dispersion parameter for binomial family taken to be 1)
##
     Null deviance: 1.3854e+02 on 104 degrees of freedom
## Residual deviance: 3.1968e-09 on 103 degrees of freedom
## AIC: 4
##
## Number of Fisher Scoring iterations: 25
```

```
prob.reducetlogit <- predict(fit.reducedlogit, df.validate, type = "response")
pred.reducetlogit <- factor(prob.reducetlogit >.5, levels = c(FALSE, TRUE), labels
= c("LAZ", "ISTINA"))
perf.reducedlogit <- table(df.validate$Species, pred.reducetlogit, dnn = c("Stvarn
i rezultat", "Predikcija"))
perf.reducedlogit</pre>
```

```
## Predikcija
## Stvarni rezultat LAZ ISTINA
## Setosa 11 0
## Versicolor 0 21
## Virginica 0 13
```