

Logistic Regression Analysis

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1 Load required and new packages

```
if (!require("pacman")) install.packages("pacman")
```

```
## Loading required package: pacman
```

```
library(pacman)
pacman::p_load("here", "glue", "crayon", "readxl", "writexl", "dplyr", "tidyr", "rstatix")
pacman::p_load("DataExplorer")

`%ni%` = Negate(`%in%`)
```

2 Set data paths and details

```
main.path = here::here()
data.path = file.path(main.path, "02 Data")
output.path = file.path(main.path, "04 Outputs")

file.name = "Final Data.xlsx"
sheet.name = "FINAL"
output.name = paste0("OUTPUT_", format(Sys.Date(), "%m%d%y"), ".xlsx")
```

3 Load dataset

```
df = readxl::read_excel(file.path(data.path, file.name),
                        sheet = sheet.name) %>%
  dplyr::select(-c(id, id_mrn, gender, admission, seizures, enceph_type, comorbidities, risk_fac
tors, neuroimaging_findings, discharge_destination:eeg_triphasic_waves, new_classification, mrs,
mrs_classification))
```

4 Process data

```
summary(df)
```

##	gender_female	gender_male	age	admission_room
##	Min. :0.0000	Min. :0.0000	Min. :20.00	Min. :0.0000
##	1st Qu.:0.0000	1st Qu.:0.0000	1st Qu.:60.50	1st Qu.:0.0000
##	Median :1.0000	Median :0.0000	Median :71.50	Median :1.0000
##	Mean :0.5417	Mean :0.4583	Mean :68.56	Mean :0.5556
##	3rd Qu.:1.0000	3rd Qu.:1.0000	3rd Qu.:82.50	3rd Qu.:1.0000
##	Max. :1.0000	Max. :1.0000	Max. :97.00	Max. :1.0000
##	admission_icu	seizure_focal_motor	seizure_generalized	
##	Min. :0.0000	Min. :0.0000	Min. :0.00000	
##	1st Qu.:0.0000	1st Qu.:0.0000	1st Qu.:0.00000	
##	Median :0.0000	Median :0.0000	Median :0.00000	
##	Mean :0.4444	Mean :0.1111	Mean :0.09722	
##	3rd Qu.:1.0000	3rd Qu.:0.0000	3rd Qu.:0.00000	
##	Max. :1.0000	Max. :1.0000	Max. :1.00000	
##	seizure_myoclonic_jerks	seizure_none	enceph_anorexic_ischemic	
##	Min. :0.00000	Min. :0.00	Min. :0.000	
##	1st Qu.:0.00000	1st Qu.:0.75	1st Qu.:0.000	
##	Median :0.00000	Median :1.00	Median :0.000	
##	Mean :0.04167	Mean :0.75	Mean :0.125	
##	3rd Qu.:0.00000	3rd Qu.:1.00	3rd Qu.:0.000	
##	Max. :1.00000	Max. :1.00	Max. :1.000	
##	enceph_drug_induced	enceph_electrolyte_disturbance	enceph_endocrine	
##	Min. :0.00000	Min. :0.0000	Min. :0.00000	
##	1st Qu.:0.00000	1st Qu.:0.0000	1st Qu.:0.00000	
##	Median :0.00000	Median :0.0000	Median :0.00000	
##	Mean :0.02778	Mean :0.1806	Mean :0.05556	
##	3rd Qu.:0.00000	3rd Qu.:0.0000	3rd Qu.:0.00000	
##	Max. :1.00000	Max. :1.0000	Max. :1.00000	
##	enceph_hepatic	enceph_hypercapnic	enceph_septic	enceph_uremic
##	Min. :0.0000	Min. :0.00000	Min. :0.0000	Min. :0.00000
##	1st Qu.:0.0000	1st Qu.:0.00000	1st Qu.:0.0000	1st Qu.:0.00000
##	Median :0.0000	Median :0.00000	Median :1.0000	Median :0.00000
##	Mean :0.1111	Mean :0.01389	Mean :0.5972	Mean :0.09722
##	3rd Qu.:0.0000	3rd Qu.:0.00000	3rd Qu.:1.0000	3rd Qu.:0.00000
##	Max. :1.0000	Max. :1.00000	Max. :1.0000	Max. :1.00000
##	enceph_withdrawal	com_cancer	com_ckd	com_diabetes
##	Min. :0.00000	Min. :0.0000	Min. :0.0000	Min. :0.0000
##	1st Qu.:0.00000	1st Qu.:0.0000	1st Qu.:0.0000	1st Qu.:0.0000
##	Median :0.00000	Median :0.0000	Median :0.0000	Median :0.0000
##	Mean :0.02778	Mean :0.1389	Mean :0.1806	Mean :0.3056
##	3rd Qu.:0.00000	3rd Qu.:0.0000	3rd Qu.:0.0000	3rd Qu.:1.0000
##	Max. :1.00000	Max. :1.0000	Max. :1.0000	Max. :1.0000
##	com_hypertension	com_liver_disease	com_previous_stroke	risk_alcohol
##	Min. :0.0000	Min. :0.0000	Min. :0.0000	Min. :0.0000
##	1st Qu.:0.0000	1st Qu.:0.0000	1st Qu.:0.0000	1st Qu.:0.0000
##	Median :1.0000	Median :0.0000	Median :0.0000	Median :0.0000
##	Mean :0.5833	Mean :0.1389	Mean :0.2361	Mean :0.1389
##	3rd Qu.:1.0000	3rd Qu.:0.0000	3rd Qu.:0.0000	3rd Qu.:0.0000
##	Max. :1.0000	Max. :1.0000	Max. :1.0000	Max. :1.0000
##	risk_drug	risk_smoking	risk_none	nf_cerebral_edema
##	Min. :0.00000	Min. :0.0000	Min. :0.0000	Min. :0.00000
##	1st Qu.:0.00000	1st Qu.:0.0000	1st Qu.:0.0000	1st Qu.:0.00000

```

## Median :0.00000 Median :0.0000 Median :1.0000 Median :0.00000
## Mean :0.01389 Mean :0.4306 Mean :0.5278 Mean :0.02778
## 3rd Qu.:0.00000 3rd Qu.:1.0000 3rd Qu.:1.0000 3rd Qu.:0.00000
## Max. :1.00000 Max. :1.0000 Max. :1.0000 Max. :1.00000
## nf_cerebrocerebellar_atrophy nf_chronic_infarct nf_white_matter_changes
## Min. :0.0000 Min. :0.0000 Min. :0.0000
## 1st Qu.:0.0000 1st Qu.:0.0000 1st Qu.:0.0000
## Median :1.0000 Median :0.0000 Median :1.0000
## Mean :0.5833 Mean :0.4167 Mean :0.5139
## 3rd Qu.:1.0000 3rd Qu.:1.0000 3rd Qu.:1.0000
## Max. :1.0000 Max. :1.0000 Max. :1.0000
## nf_unremarkable malignance classification_benign
## Min. :0.000 Min. :0.0000 Min. :0.0000
## 1st Qu.:0.000 1st Qu.:0.0000 1st Qu.:0.0000
## Median :0.000 Median :0.0000 Median :1.0000
## Mean :0.125 Mean :0.3472 Mean :0.6528
## 3rd Qu.:0.000 3rd Qu.:1.0000 3rd Qu.:1.0000
## Max. :1.000 Max. :1.0000 Max. :1.0000
## classification_malignant classification_highly_malignant mrs_good
## Min. :0.0000 Min. :0.00000 Min. :0.000
## 1st Qu.:0.0000 1st Qu.:0.00000 1st Qu.:0.000
## Median :0.0000 Median :0.00000 Median :0.000
## Mean :0.2778 Mean :0.06944 Mean :0.375
## 3rd Qu.:1.0000 3rd Qu.:0.00000 3rd Qu.:1.000
## Max. :1.0000 Max. :1.00000 Max. :1.000
## mrs_poor mrs_death
## Min. :0.0000 Min. :0.0000
## 1st Qu.:0.0000 1st Qu.:0.0000
## Median :0.0000 Median :0.0000
## Mean :0.3056 Mean :0.3194
## 3rd Qu.:1.0000 3rd Qu.:1.0000
## Max. :1.0000 Max. :1.0000

```

```
df.final = df %>%
  dplyr::mutate(dplyr::across(-c(age), ~ .x[!is.na(.x)] %>% as.factor()))

# set.seed(102899)

df.benign = df.final %>%
  dplyr::filter(classification_benign == 1) %>%
  dplyr::sample_n(47, replace = T)

# set.seed(061400)

df.malignant = df.final %>%
  dplyr::filter(classification_malignant == 1) %>%
  dplyr::sample_n(20, replace = T)

# set.seed(082624)

df.hmalignant = df.final %>%
  dplyr::filter(classification_highly_malignant == 1) %>%
  dplyr::sample_n(5, replace = T)

df.final = dplyr::bind_rows(df.final, df.benign, df.malignant, df.hmalignant)

summary(df.final)
```

```

## gender_female gender_male age admission_room admission_icu
## 0:67 0:77 Min. :20.00 0:65 0:79
## 1:77 1:67 1st Qu.:58.00 1:79 1:65
## Median :69.50
## Mean :67.59
## 3rd Qu.:84.00
## Max. :97.00
## seizure_focal_motor seizure_generalized seizure_myoclonic_jerks seizure_none
## 0:126 0:130 0:138 0: 38
## 1: 18 1: 14 1: 6 1:106
##
##
##
## enceph_anorexic_ischemic enceph_drug_induced enceph_electrolyte_disturbance
## 0:127 0:140 0:121
## 1: 17 1: 4 1: 23
##
##
##
## enceph_endocrine enceph_hepatic enceph_hypercapnic enceph_septic enceph_uremic
## 0:133 0:126 0:142 0:61 0:124
## 1: 11 1: 18 1: 2 1:83 1: 20
##
##
##
## enceph_withdrawal com_cancer com_ckd com_diabetes com_hypertension
## 0:142 0:122 0:116 0:99 0:67
## 1: 2 1: 22 1: 28 1:45 1:77
##
##
##
## com_liver_disease com_previous_stroke risk_alcohol risk_drug risk_smoking
## 0:123 0:113 0:124 0:142 0:79
## 1: 21 1: 31 1: 20 1: 2 1:65
##
##
##
## risk_none nf_cerebral_edema nf_cerebrocerebellar_atrophy nf_chronic_infarct
## 0:71 0:140 0:64 0:90
## 1:73 1: 4 1:80 1:54
##
##
##
## nf_white_matter_changes nf_unremarkable malignance classification_benign
## 0:68 0:127 0:94 0:50
## 1:76 1: 17 1:50 1:94

```

```
##
##
##
##
## classification_malignant classification_highly_malignant mrs_good mrs_poor
## 0:104                0:134                0:87    0:97
## 1: 40                1: 10                1:57    1:47
##
##
##
##
## mrs_death
## 0:104
## 1: 40
##
##
##
##
```

5 Implement methodology

5.1 Overall Logistic Regression

```
overall.model = stats::glm(malignance ~
                           gender_female +
                           age +
                           admission_icu +
                           seizure_focal_motor + seizure_generalized + seizure_myoclonic_jerks
+
                           com_cancer + com_ckd + com_diabetes + com_hypertension + com_liver_
disease + com_previous_stroke +
                           risk_alcohol + risk_drug + risk_smoking +
                           nf_cerebral_edema + nf_cerebrocerebellar_atrophy + nf_chronic_infar
ct + nf_white_matter_changes + nf_unremarkable +
                           mrs_poor + mrs_death,
                           data = df.final,
                           family = "binomial")
overall.model.summary = summary(overall.model)
overall.model.summary
```

```
##
## Call:
## stats::glm(formula = malignance ~ gender_female + age + admission_icu +
## seizure_focal_motor + seizure_generalized + seizure_myoclonic_jerks +
## com_cancer + com_ckd + com_diabetes + com_hypertension +
## com_liver_disease + com_previous_stroke + risk_alcohol +
## risk_drug + risk_smoking + nf_cerebral_edema + nf_cerebrocerebellar_atrophy +
## nf_chronic_infarct + nf_white_matter_changes + nf_unremarkable +
## mrs_poor + mrs_death, family = "binomial", data = df.final)
##
## Coefficients:
##
## Estimate Std. Error z value Pr(>|z|)
## (Intercept) -3.12561 1.57385 -1.986 0.04704 *
## gender_female1 -1.60225 0.62205 -2.576 0.01000 *
## age 0.02482 0.02122 1.170 0.24209
## admission_icu1 1.04771 0.65236 1.606 0.10827
## seizure_focal_motor1 2.48488 0.90988 2.731 0.00631 **
## seizure_generalized1 -2.76252 1.27867 -2.160 0.03074 *
## seizure_myoclonic_jerks1 -0.62197 1.29345 -0.481 0.63061
## com_cancer1 0.19634 0.74639 0.263 0.79251
## com_ckd1 -0.95364 0.73995 -1.289 0.19747
## com_diabetes1 0.64890 0.58835 1.103 0.27007
## com_hypertension1 -0.10607 0.61680 -0.172 0.86346
## com_liver_disease1 1.21302 0.68520 1.770 0.07667 .
## com_previous_stroke1 0.51487 0.80326 0.641 0.52154
## risk_alcohol1 2.37603 1.06163 2.238 0.02521 *
## risk_drug1 -19.66568 2797.44218 -0.007 0.99439
## risk_smoking1 -0.74252 0.63997 -1.160 0.24595
## nf_cerebral_edema1 18.36198 1718.75527 0.011 0.99148
## nf_cerebrocerebellar_atrophy1 -0.30713 0.87200 -0.352 0.72468
## nf_chronic_infarct1 -0.80303 0.78336 -1.025 0.30531
## nf_white_matter_changes1 1.04427 0.85813 1.217 0.22364
## nf_unremarkable1 1.55821 1.03614 1.504 0.13262
## mrs_poor1 -0.27667 0.76357 -0.362 0.71710
## mrs_death1 1.65122 0.73604 2.243 0.02487 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
## Null deviance: 185.96 on 143 degrees of freedom
## Residual deviance: 121.70 on 121 degrees of freedom
## AIC: 167.7
##
## Number of Fisher Scoring iterations: 16
```

```

overall.names = c("Intercept", names(overall.model.summary$contrasts)[1], "age", names(overall.m
odel.summary$contrasts)[2:21])
overall.crude = as.data.frame(overall.model.summary$coefficients) %>%
  dplyr::mutate(VARS = overall.names,
               OR = exp(`Estimate`),
               LL = exp(`Estimate` - (`Std. Error` * 1.96)),
               UL = exp(`Estimate` + (`Std. Error` * 1.96)),
               sig = ifelse(`Pr(>|z|)` <= 0.05, "S", "NS")) %>%
  dplyr::select(VARS, dplyr::everything())

```

5.2 Benign Logistic Regression

```

benign.model = stats::glm(classification_benign ~
                          enceph_anorexic_ischemic + enceph_drug_induced + enceph_electrolyte_d
isturbance + enceph_endocrine + enceph_hepatic + enceph_hypercapnic + enceph_septic + enceph_ure
mic + enceph_withdrawal +
                          mrs_poor + mrs_death,
                          data = df.final,
                          family = "binomial")
benign.model.summary = summary(benign.model)
benign.model.summary

```



```
##
## Call:
## stats::glm(formula = classification_benign ~ enceph_anorexic_ischemic +
##   enceph_drug_induced + enceph_electrolyte_disturbance + enceph_endocrine +
##   enceph_hepatic + enceph_hypercapnic + enceph_septic + enceph_uremic +
##   enceph_withdrawal + mrs_poor + mrs_death, family = "binomial",
##   data = df.final)
##
## Coefficients:
##               Estimate Std. Error z value Pr(>|z|)
## (Intercept)      2.4656     0.7279   3.387 0.000706 ***
## enceph_anorexic_ischemic1 -0.8132     0.7768  -1.047 0.295166
## enceph_drug_induced1     -3.3138     1.3408  -2.471 0.013456 *
## enceph_electrolyte_disturbance1 -1.5644     0.6295  -2.485 0.012948 *
## enceph_endocrine1      -2.1552     0.8450  -2.550 0.010760 *
## enceph_hepatic1       -1.8259     0.7728  -2.363 0.018136 *
## enceph_hypercapnic1     15.4428    1696.7344   0.009 0.992738
## enceph_septic1        -1.1156     0.5826  -1.915 0.055530 .
## enceph_uremic1         1.1119     0.7501   1.482 0.138255
## enceph_withdrawal1     14.1005    1696.7345   0.008 0.993369
## mrs_poor1             -0.2268     0.5115  -0.443 0.657508
## mrs_death1           -1.2833     0.5467  -2.347 0.018906 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##   Null deviance: 185.96  on 143  degrees of freedom
## Residual deviance: 157.90  on 132  degrees of freedom
## AIC: 181.9
##
## Number of Fisher Scoring iterations: 15
```

```
benign.names = c("Intercept", names(benign.model.summary$contrasts))
benign.crude = as.data.frame(benign.model.summary$coefficients) %>%
  dplyr::mutate(VARS = benign.names,
    `Pr(>|z|)` = `Pr(>|z|)`,
    OR = exp(`Estimate`),
    LL = exp(`Estimate` - (`Std. Error` * 1.96)),
    UL = exp(`Estimate` + (`Std. Error` * 1.96)),
    sig = ifelse(`Pr(>|z|)` <= 0.05, "S", "NS")) %>%
  dplyr::select(VARS, dplyr::everything())

benign.model.step = MASS::stepAIC(benign.model, direction = "both")
```

```

## Start: AIC=181.9
## classification_benign ~ enceph_anorexic_ischemic + enceph_drug_induced +
##     enceph_electrolyte_disturbance + enceph_endocrine + enceph_hepatic +
##     enceph_hypercapnic + enceph_septic + enceph_uremic + enceph_withdrawal +
##     mrs_poor + mrs_death
##
##
##           Df Deviance    AIC
## - mrs_poor           1   158.10 180.10
## - enceph_withdrawal   1   158.21 180.21
## - enceph_hypercapnic  1   158.99 180.99
## - enceph_anorexic_ischemic 1   159.06 181.06
## <none>                157.90 181.90
## - enceph_uremic       1   160.36 182.36
## - enceph_septic       1   162.06 184.06
## - mrs_death           1   163.70 185.70
## - enceph_hepatic      1   163.93 185.93
## - enceph_electrolyte_disturbance 1   164.42 186.42
## - enceph_endocrine    1   164.97 186.97
## - enceph_drug_induced 1   165.38 187.38
##
## Step: AIC=180.1
## classification_benign ~ enceph_anorexic_ischemic + enceph_drug_induced +
##     enceph_electrolyte_disturbance + enceph_endocrine + enceph_hepatic +
##     enceph_hypercapnic + enceph_septic + enceph_uremic + enceph_withdrawal +
##     mrs_death
##
##
##           Df Deviance    AIC
## - enceph_withdrawal   1   158.46 178.46
## - enceph_hypercapnic  1   159.10 179.10
## - enceph_anorexic_ischemic 1   159.15 179.15
## <none>                158.10 180.10
## - enceph_uremic       1   161.26 181.26
## + mrs_poor            1   157.90 181.90
## - enceph_septic       1   162.10 182.10
## - mrs_death           1   164.12 184.12
## - enceph_hepatic      1   164.13 184.13
## - enceph_electrolyte_disturbance 1   164.76 184.76
## - enceph_endocrine    1   165.00 185.00
## - enceph_drug_induced 1   165.38 185.38
##
## Step: AIC=178.46
## classification_benign ~ enceph_anorexic_ischemic + enceph_drug_induced +
##     enceph_electrolyte_disturbance + enceph_endocrine + enceph_hepatic +
##     enceph_hypercapnic + enceph_septic + enceph_uremic + mrs_death
##
##
##           Df Deviance    AIC
## - enceph_hypercapnic  1   159.45 177.45
## - enceph_anorexic_ischemic 1   159.64 177.64
## <none>                158.46 178.46
## - enceph_uremic       1   161.54 179.54
## + enceph_withdrawal   1   158.10 180.10
## + mrs_poor            1   158.21 180.21

```

```

## - enceph_septic                1    163.04 181.04
## - mrs_death                    1    164.57 182.57
## - enceph_hepatic               1    164.95 182.95
## - enceph_electrolyte_disturbance 1    165.64 183.64
## - enceph_endocrine             1    165.70 183.70
## - enceph_drug_induced          1    166.05 184.05
##
## Step: AIC=177.45
## classification_benign ~ enceph_anorexic_ischemic + enceph_drug_induced +
##   enceph_electrolyte_disturbance + enceph_endocrine + enceph_hepatic +
##   enceph_septic + enceph_uremic + mrs_death
##
##                                Df Deviance    AIC
## - enceph_anorexic_ischemic    1    160.66 176.66
## <none>                        159.45 177.45
## + enceph_hypercapnic          1    158.46 178.46
## - enceph_uremic               1    162.48 178.48
## + enceph_withdrawal           1    159.10 179.10
## + mrs_poor                    1    159.30 179.30
## - enceph_septic               1    163.95 179.95
## - mrs_death                   1    165.88 181.88
## - enceph_hepatic              1    166.12 182.12
## - enceph_electrolyte_disturbance 1    166.94 182.94
## - enceph_endocrine            1    166.95 182.95
## - enceph_drug_induced         1    167.23 183.23
##
## Step: AIC=176.66
## classification_benign ~ enceph_drug_induced + enceph_electrolyte_disturbance +
##   enceph_endocrine + enceph_hepatic + enceph_septic + enceph_uremic +
##   mrs_death
##
##                                Df Deviance    AIC
## <none>                        160.66 176.66
## + enceph_anorexic_ischemic    1    159.45 177.45
## + enceph_hypercapnic          1    159.64 177.64
## - enceph_septic               1    163.98 177.98
## + enceph_withdrawal           1    160.19 178.19
## + mrs_poor                    1    160.59 178.59
## - enceph_uremic               1    165.04 179.04
## - enceph_hepatic              1    166.15 180.15
## - enceph_electrolyte_disturbance 1    167.00 181.00
## - enceph_drug_induced         1    167.62 181.62
## - enceph_endocrine            1    168.28 182.28
## - mrs_death                   1    170.81 184.81

```

```

benign.model.step.summary = summary(benign.model.step)
benign.model.step.summary

```

```
##
## Call:
## stats::glm(formula = classification_benign ~ enceph_drug_induced +
##           enceph_electrolyte_disturbance + enceph_endocrine + enceph_hepatic +
##           enceph_septic + enceph_uremic + mrs_death, family = "binomial",
##           data = df.final)
##
## Coefficients:
##
##               Estimate Std. Error z value Pr(>|z|)
## (Intercept)         2.1116     0.5322   3.968 7.25e-05 ***
## enceph_drug_induced1 -3.0317     1.2568  -2.412  0.01586 *
## enceph_electrolyte_disturbance1 -1.4313     0.5749  -2.490  0.01278 *
## enceph_endocrine1    -2.0724     0.7586  -2.732  0.00630 **
## enceph_hepatic1     -1.6045     0.7052  -2.275  0.02290 *
## enceph_septic1      -0.8523     0.4829  -1.765  0.07754 .
## enceph_uremic1       1.3593     0.7123   1.908  0.05637 .
## mrs_death1         -1.4166     0.4581  -3.092  0.00199 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 185.96  on 143  degrees of freedom
## Residual deviance: 160.66  on 136  degrees of freedom
## AIC: 176.66
##
## Number of Fisher Scoring iterations: 4
```

```
benign.step.names = c("Intercept", names(benign.model.step.summary$contrasts))
benign.adj = as.data.frame(benign.model.step.summary$coefficients) %>%
  dplyr::mutate(VARS = benign.step.names,
               OR = exp(`Estimate`),
               LL = exp(`Estimate` - (`Std. Error` * 1.96)),
               UL = exp(`Estimate` + (`Std. Error` * 1.96)),
               sig = ifelse(`Pr(>|z|)` <= 0.05, "S", "NS")) %>%
  dplyr::select(VARS, dplyr::everything())
```

5.3 Malignant Logistic Regression

```
malignant.model = stats::glm(classification_malignant ~
                             enceph_anorexic_ischemic + enceph_drug_induced + enceph_electrolyte_
disturbance + enceph_endocrine + enceph_hepatic + enceph_hypercapnic + enceph_septic + enceph_ur
emic + enceph_withdrawal +
                             mrs_poor + mrs_death,
                             data = df.final,
                             family = "binomial")
malignant.model.summary = summary(malignant.model)
malignant.model.summary
```

```
##
## Call:
## stats::glm(formula = classification_malignant ~ enceph_anorexic_ischemic +
##   enceph_drug_induced + enceph_electrolyte_disturbance + enceph_endocrine +
##   enceph_hepatic + enceph_hypercapnic + enceph_septic + enceph_uremic +
##   enceph_withdrawal + mrs_poor + mrs_death, family = "binomial",
##   data = df.final)
##
## Coefficients:
##               Estimate Std. Error z value Pr(>|z|)
## (Intercept)      -2.2375     0.7237  -3.092  0.00199 **
## enceph_anorexic_ischemic1      0.5935     0.8070   0.735  0.46206
## enceph_drug_induced1          3.1954     1.3055   2.448  0.01438 *
## enceph_electrolyte_disturbance1  1.3329     0.6560   2.032  0.04217 *
## enceph_endocrine1            2.1074     0.8462   2.490  0.01276 *
## enceph_hepatic1              2.1521     0.8285   2.598  0.00939 **
## enceph_hypercapnic1        -16.9700    4612.2020  -0.004  0.99706
## enceph_septic1              1.0793     0.5998   1.799  0.07195 .
## enceph_uremic1             -17.7720    1374.1345  -0.013  0.98968
## enceph_withdrawal1         -16.3286    4612.2020  -0.004  0.99718
## mrs_poor1                 -0.4379     0.5358  -0.817  0.41379
## mrs_death1                 0.5278     0.6053   0.872  0.38325
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##   Null deviance: 170.16  on 143  degrees of freedom
## Residual deviance: 134.78  on 132  degrees of freedom
## AIC: 158.78
##
## Number of Fisher Scoring iterations: 17
```

```
malignant.names = c("Intercept", names(malignant.model.summary$contrasts))
malignant.crude = as.data.frame(malignant.model.summary$coefficients) %>%
  dplyr::mutate(VARS = malignant.names,
               OR = exp(`Estimate`),
               LL = exp(`Estimate` - (`Std. Error` * 1.96)),
               UL = exp(`Estimate` + (`Std. Error` * 1.96)),
               sig = ifelse(`Pr(>|z|)` <= 0.05, "S", "NS")) %>%
  dplyr::select(VARS, dplyr::everything())

malignant.model.step = MASS::stepAIC(malignant.model, direction = "both")
```

```

## Start: AIC=158.78
## classification_malignant ~ enceph_anorexic_ischemic + enceph_drug_induced +
##     enceph_electrolyte_disturbance + enceph_endocrine + enceph_hepatic +
##     enceph_hypercapnic + enceph_septic + enceph_uremic + enceph_withdrawal +
##     mrs_poor + mrs_death
##
##
##      Df Deviance    AIC
## - enceph_withdrawal      1   135.16 157.16
## - enceph_anorexic_ischemic 1   135.34 157.34
## - mrs_poor                1   135.44 157.44
## - enceph_hypercapnic      1   135.49 157.49
## - mrs_death              1   135.54 157.54
## <none>                   134.78 158.78
## - enceph_septic          1   138.38 160.38
## - enceph_electrolyte_disturbance 1 138.91 160.91
## - enceph_endocrine       1 141.41 163.41
## - enceph_drug_induced     1 142.00 164.00
## - enceph_hepatic         1 142.07 164.07
## - enceph_uremic          1 146.72 168.72
##
## Step: AIC=157.16
## classification_malignant ~ enceph_anorexic_ischemic + enceph_drug_induced +
##     enceph_electrolyte_disturbance + enceph_endocrine + enceph_hepatic +
##     enceph_hypercapnic + enceph_septic + enceph_uremic + mrs_poor +
##     mrs_death
##
##
##      Df Deviance    AIC
## - mrs_poor                1   135.75 155.75
## - enceph_anorexic_ischemic 1   135.82 155.82
## - enceph_hypercapnic      1   135.88 155.88
## - mrs_death              1   136.01 156.01
## <none>                   135.16 157.16
## + enceph_withdrawal      1   134.78 158.78
## - enceph_septic          1   139.39 159.39
## - enceph_electrolyte_disturbance 1 139.64 159.64
## - enceph_endocrine       1 142.30 162.30
## - enceph_drug_induced     1 142.81 162.81
## - enceph_hepatic         1 142.92 162.92
## - enceph_uremic          1 146.91 166.91
##
## Step: AIC=155.75
## classification_malignant ~ enceph_anorexic_ischemic + enceph_drug_induced +
##     enceph_electrolyte_disturbance + enceph_endocrine + enceph_hepatic +
##     enceph_hypercapnic + enceph_septic + enceph_uremic + mrs_death
##
##
##      Df Deviance    AIC
## - enceph_anorexic_ischemic 1   136.49 154.49
## - enceph_hypercapnic      1   136.62 154.62
## - mrs_death              1   137.75 155.75
## <none>                   135.75 155.75
## + mrs_poor                1   135.16 157.16
## + enceph_withdrawal      1   135.44 157.44

```

```

## - enceph_electrolyte_disturbance 1 140.04 158.04
## - enceph_septic 1 140.26 158.26
## - enceph_hepatic 1 143.20 161.20
## - enceph_drug_induced 1 144.06 162.06
## - enceph_endocrine 1 144.21 162.21
## - enceph_uremic 1 146.92 164.92
##
## Step: AIC=154.49
## classification_malignant ~ enceph_drug_induced + enceph_electrolyte_disturbance +
## enceph_endocrine + enceph_hepatic + enceph_hypercapnic +
## enceph_septic + enceph_uremic + mrs_death
##
## Df Deviance AIC
## - enceph_hypercapnic 1 137.38 153.38
## <none> 136.49 154.49
## + enceph_anorexic_ischemic 1 135.75 155.75
## + mrs_poor 1 135.82 155.82
## - mrs_death 1 140.11 156.11
## + enceph_withdrawal 1 136.12 156.12
## - enceph_electrolyte_disturbance 1 140.16 156.16
## - enceph_septic 1 140.26 156.26
## - enceph_hepatic 1 143.24 159.24
## - enceph_drug_induced 1 144.24 160.24
## - enceph_endocrine 1 145.24 161.24
## - enceph_uremic 1 149.05 165.05
##
## Step: AIC=153.38
## classification_malignant ~ enceph_drug_induced + enceph_electrolyte_disturbance +
## enceph_endocrine + enceph_hepatic + enceph_septic + enceph_uremic +
## mrs_death
##
## Df Deviance AIC
## <none> 137.38 153.38
## + enceph_hypercapnic 1 136.49 154.49
## + mrs_poor 1 136.54 154.54
## + enceph_anorexic_ischemic 1 136.62 154.62
## + enceph_withdrawal 1 137.01 155.01
## - enceph_septic 1 141.03 155.03
## - enceph_electrolyte_disturbance 1 141.27 155.27
## - mrs_death 1 141.28 155.28
## - enceph_hepatic 1 144.26 158.26
## - enceph_drug_induced 1 145.30 159.30
## - enceph_endocrine 1 146.41 160.41
## - enceph_uremic 1 149.87 163.87

```

```

malignant.model.step.summary = summary(malignant.model.step)
malignant.model.step.summary

```

```
##
## Call:
## stats::glm(formula = classification_malignant ~ enceph_drug_induced +
##           enceph_electrolyte_disturbance + enceph_endocrine + enceph_hepatic +
##           enceph_septic + enceph_uremic + mrs_death, family = "binomial",
##           data = df.final)
##
## Coefficients:
##
##               Estimate Std. Error z value Pr(>|z|)
## (Intercept)      -2.3759      0.5722  -4.153 3.29e-05 ***
## enceph_drug_induced1      3.2789      1.2740   2.574  0.0101 *
## enceph_electrolyte_disturbance1      1.2321      0.6209   1.984  0.0472 *
## enceph_endocrine1      2.3116      0.7816   2.958  0.0031 **
## enceph_hepatic1      1.9020      0.7468   2.547  0.0109 *
## enceph_septic1      0.9572      0.5232   1.829  0.0673 .
## enceph_uremic1     -17.6910    1376.1824  -0.013  0.9897
## mrs_death1      0.9855      0.4996   1.972  0.0486 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 170.16  on 143  degrees of freedom
## Residual deviance: 137.38  on 136  degrees of freedom
## AIC: 153.38
##
## Number of Fisher Scoring iterations: 17
```

```
malignant.step.names = c("Intercept", names(malignant.model.step.summary$contrasts))
malignant.adj = as.data.frame(malignant.model.step.summary$coefficients) %>%
  dplyr::mutate(VARS = malignant.step.names,
               OR = exp(`Estimate`),
               LL = exp(`Estimate` - (`Std. Error` * 1.96)),
               UL = exp(`Estimate` + (`Std. Error` * 1.96)),
               sig = ifelse(`Pr(>|z|)` <= 0.05, "S", "NS")) %>%
  dplyr::select(VARS, dplyr::everything())
```

5.4 Highly Malignant Logistic Regression

```
hmalignant.model = stats::glm(classification_highly_malignant ~
                              enceph_anorexic_ischemic + enceph_drug_induced + enceph_electroly
te_disturbance + enceph_endocrine + enceph_hepatic + enceph_hypercapnic + enceph_septic + enceph
_uremic + enceph_withdrawal +
                              mrs_poor + mrs_death,
                              data = df.final,
                              family = "binomial")
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```



```
hmalignant.model.summary = summary(hmalignant.model)
hmalignant.model.summary
```

```
##
## Call:
## stats::glm(formula = classification_highly_malignant ~ enceph_anorexic_ischemic +
##   enceph_drug_induced + enceph_electrolyte_disturbance + enceph_endocrine +
##   enceph_hepatic + enceph_hypercapnic + enceph_septic + enceph_uremic +
##   enceph_withdrawal + mrs_poor + mrs_death, family = "binomial",
##   data = df.final)
##
## Coefficients:
##
##               Estimate Std. Error z value Pr(>|z|)
## (Intercept)      -22.8579   2097.8130  -0.011   0.9913
## enceph_anorexic_ischemic1      2.6041     1.8938   1.375   0.1691
## enceph_drug_induced1     -16.2412   6867.7931  -0.002   0.9981
## enceph_electrolyte_disturbance1    1.0995     1.0684   1.029   0.3034
## enceph_endocrine1     -15.0000   4186.5351  -0.004   0.9971
## enceph_hepatic1        1.1482     1.5313   0.750   0.4534
## enceph_hypercapnic1    -18.1381  12537.2648  -0.001   0.9988
## enceph_septic1         0.9445     1.0648   0.887   0.3751
## enceph_uremic1         3.0074     1.5453   1.946   0.0516
## enceph_withdrawal1      2.2919  12711.5628   0.000   0.9999
## mrs_poor1             19.4855   2097.8125   0.009   0.9926
## mrs_death1            18.5424   2097.8121   0.009   0.9929
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##    Null deviance: 72.633  on 143  degrees of freedom
## Residual deviance: 54.604  on 132  degrees of freedom
## AIC: 78.604
##
## Number of Fisher Scoring iterations: 19
```

```
hmalignant.names = c("Intercept", names(hmalignant.model.summary$contrasts))
hmalignant.crude = as.data.frame(hmalignant.model.summary$coefficients) %>%
  dplyr::mutate(VARS = hmalignant.names,
               OR = exp(`Estimate`),
               LL = exp(`Estimate` - (`Std. Error` * 1.96)),
               UL = exp(`Estimate` + (`Std. Error` * 1.96)),
               sig = ifelse(`Pr(>|z|)` <= 0.05, "S", "NS")) %>%
  dplyr::select(VARS, dplyr::everything())

hmalignant.model.step = MASS::stepAIC(hmalignant.model, direction = "both")
```

```
## Start:  AIC=78.6
## classification_highly_malignant ~ enceph_anorexic_ischemic +
##     enceph_drug_induced + enceph_electrolyte_disturbance + enceph_endocrine +
##     enceph_hepatic + enceph_hypercapnic + enceph_septic + enceph_uremic +
##     enceph_withdrawal + mrs_poor + mrs_death
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

##		Df	Deviance	AIC
##	- enceph_withdrawal	1	54.604	76.604
##	- enceph_endocrine	1	54.733	76.733
##	- enceph_drug_induced	1	54.771	76.771
##	- enceph_hypercapnic	1	54.931	76.931
##	- enceph_hepatic	1	55.130	77.130
##	- enceph_septic	1	55.371	77.371
##	- enceph_electrolyte_disturbance	1	55.617	77.617
##	<none>		54.604	78.604
##	- enceph_anorexic_ischemic	1	56.675	78.675
##	- enceph_uremic	1	59.177	81.177
##	- mrs_death	1	62.612	84.612
##	- mrs_poor	1	63.298	85.298

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
##
## Step:  AIC=76.6
## classification_highly_malignant ~ enceph_anorexic_ischemic +
##     enceph_drug_induced + enceph_electrolyte_disturbance + enceph_endocrine +
##     enceph_hepatic + enceph_hypercapnic + enceph_septic + enceph_uremic +
##     mrs_poor + mrs_death
```


	Df	Deviance	AIC
## - enceph_drug_induced	1	54.896	72.896
## - enceph_hypercapnic	1	55.053	73.053
## - enceph_hepatic	1	55.345	73.345
## - enceph_septic	1	55.683	73.683
## - enceph_electrolyte_disturbance	1	55.911	73.911
## <none>		54.733	74.733
## - enceph_anorexic_ischemic	1	57.044	75.044
## + enceph_endocrine	1	54.604	76.604
## + enceph_withdrawal	1	54.733	76.733
## - enceph_uremic	1	59.506	77.506
## - mrs_poor	1	64.166	82.166
## - mrs_death	1	64.308	82.308

##

Step: AIC=72.9

classification_highly_malignant ~ enceph_anorexic_ischemic +

enceph_electrolyte_disturbance + enceph_hepatic + enceph_hypercapnic +

enceph_septic + enceph_uremic + mrs_poor + mrs_death

Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred

Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred

	Df	Deviance	AIC
## - enceph_hypercapnic	1	55.206	71.206
## - enceph_hepatic	1	55.527	71.527
## - enceph_septic	1	55.831	71.831
## - enceph_electrolyte_disturbance	1	56.112	72.112
## <none>		54.896	72.896
## - enceph_anorexic_ischemic	1	57.210	73.210
## + enceph_drug_induced	1	54.733	74.733
## + enceph_endocrine	1	54.771	74.771
## + enceph_withdrawal	1	54.896	74.896
## - enceph_uremic	1	59.652	75.652
## - mrs_poor	1	64.337	80.337
## - mrs_death	1	64.505	80.505

##

Step: AIC=71.21

classification_highly_malignant ~ enceph_anorexic_ischemic +

enceph_electrolyte_disturbance + enceph_hepatic + enceph_septic +

enceph_uremic + mrs_poor + mrs_death

Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred

Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred

	Df	Deviance	AIC
## - enceph_hepatic	1	55.874	69.874
## - enceph_septic	1	56.116	70.116
## - enceph_electrolyte_disturbance	1	56.496	70.496
## <none>		55.206	71.206
## - enceph_anorexic_ischemic	1	57.526	71.526
## + enceph_hypercapnic	1	54.896	72.896
## + enceph_drug_induced	1	55.053	73.053
## + enceph_endocrine	1	55.087	73.087
## + enceph_withdrawal	1	55.206	73.206
## - enceph_uremic	1	59.931	73.931
## - mrs_poor	1	64.444	78.444
## - mrs_death	1	64.790	78.790

##

Step: AIC=69.87

classification_highly_malignant ~ enceph_anorexic_ischemic +
 ## enceph_electrolyte_disturbance + enceph_septic + enceph_uremic +
 ## mrs_poor + mrs_death

Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred

Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred

	Df	Deviance	AIC
## - enceph_septic	1	56.258	68.258
## - enceph_electrolyte_disturbance	1	56.638	68.638
## - enceph_anorexic_ischemic	1	57.526	69.526
## <none>		55.874	69.874
## + enceph_hepatic	1	55.206	71.206
## + enceph_hypercapnic	1	55.527	71.527
## + enceph_endocrine	1	55.671	71.671
## + enceph_drug_induced	1	55.703	71.703
## + enceph_withdrawal	1	55.874	71.874
## - enceph_uremic	1	59.947	71.947
## - mrs_poor	1	64.469	76.469
## - mrs_death	1	64.835	76.835

##

Step: AIC=68.26

classification_highly_malignant ~ enceph_anorexic_ischemic +
 ## enceph_electrolyte_disturbance + enceph_uremic + mrs_poor +
 ## mrs_death

Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred

Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred

```
##                                Df Deviance    AIC
## - enceph_electrolyte_disturbance 1   56.699 66.699
## - enceph_anorexic_ischemic       1   57.531 67.531
## <none>                             56.258 68.258
## + enceph_septic                   1   55.874 69.874
## - enceph_uremic                   1   59.947 69.947
## + enceph_hypercapnic              1   55.949 69.949
## + enceph_endocrine               1   55.949 69.949
## + enceph_drug_induced             1   56.105 70.105
## + enceph_hepatic                  1   56.116 70.116
## + enceph_withdrawal               1   56.258 70.258
## - mrs_poor                        1   64.633 74.633
## - mrs_death                       1   65.258 75.258
##
## Step:  AIC=66.7
## classification_highly_malignant ~ enceph_anorexic_ischemic +
##      enceph_uremic + mrs_poor + mrs_death
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
##                                Df Deviance    AIC
## - enceph_anorexic_ischemic       1   57.858 65.858
## <none>                             56.699 66.699
## + enceph_electrolyte_disturbance 1   56.258 68.258
## + enceph_hypercapnic              1   56.335 68.335
## + enceph_endocrine               1   56.335 68.335
## + enceph_drug_induced             1   56.519 68.519
## + enceph_septic                   1   56.638 68.638
## + enceph_hepatic                  1   56.643 68.643
## + enceph_withdrawal               1   56.699 68.699
## - enceph_uremic                   1   61.140 69.140
## - mrs_poor                        1   66.322 74.322
## - mrs_death                       1   66.756 74.756
##
## Step:  AIC=65.86
## classification_highly_malignant ~ enceph_uremic + mrs_poor +
##      mrs_death
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

	Df	Deviance	AIC
## <none>		57.858	65.858
## + enceph_anorexic_ischemic	1	56.699	66.699
## - enceph_uremic	1	61.177	67.177
## + enceph_hypercapnic	1	57.494	67.494
## + enceph_endocrine	1	57.494	67.494
## + enceph_electrolyte_disturbance	1	57.531	67.531
## + enceph_drug_induced	1	57.678	67.678
## + enceph_septic	1	57.790	67.790
## + enceph_hepatic	1	57.851	67.851
## + enceph_withdrawal	1	57.858	67.858
## - mrs_poor	1	66.444	72.444
## - mrs_death	1	69.621	75.621

```
halignant.model.step.summary = summary(halignant.model.step)
halignant.model.step.summary
```

```
##
## Call:
## stats::glm(formula = classification_highly_malignant ~ enceph_uremic +
##   mrs_poor + mrs_death, family = "binomial", data = df.final)
##
## Coefficients:
##             Estimate Std. Error z value Pr(>|z|)
## (Intercept)   -21.1787   2249.8831  -0.009   0.9925
## enceph_uremic1    1.7579     0.9493    1.852   0.0641 .
## mrs_poor1       18.8038   2249.8832    0.008   0.9933
## mrs_death1      18.9100   2249.8831    0.008   0.9933
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
## Null deviance: 72.633 on 143 degrees of freedom
## Residual deviance: 57.858 on 140 degrees of freedom
## AIC: 65.858
##
## Number of Fisher Scoring iterations: 19
```

```
halignant.step.names = c("Intercept", names(halignant.model.step.summary$contrasts))
halignant.adj = as.data.frame(halignant.model.step.summary$coefficients) %>%
  dplyr::mutate(VARS = halignant.step.names,
               OR = exp(`Estimate`),
               LL = exp(`Estimate` - (`Std. Error` * 1.96)),
               UL = exp(`Estimate` + (`Std. Error` * 1.96)),
               sig = ifelse(`Pr(>|z|)` <= 0.05, "S", "NS")) %>%
  dplyr::select(VARS, dplyr::everything())
```

6 Export necessary data

```
export.list = list(Overall = overall.crude,
                  Benign_Crude = benign.crude,
                  Benign_Adj = benign.adj,
                  Malignant_Crude = malignant.crude,
                  Malignant_Adj = malignant.adj,
                  Highly_Malignant_Crude = h malignant.crude,
                  Highly_Malignant_Adj = h malignant.adj)

if(length(export.list) != 0){
  if (!file.exists(file.path(output.path, output.name))) {
    writexl::write_xlsx(export.list, file.path(output.path, output.name))
    cat(crayon::green("File successfully written. "))
  } else {
    cat(crayon::red(glue::glue("Filename already used: {output.name}")))
    overwrite = readline(prompt = "Overwrite (1 for Yes, 0 for No): ")
    if (overwrite == "1") {
      writexl::write_xlsx(export.list, file.path(output.path, output.name))
      cat(crayon::green("File successfully overwritten"))
    } else {
      cat(crayon::red("File not overwritten"))
    }
  }
}
```

```
## Filename already used: OUTPUT_071025.xlsxOverwrite (1 for Yes, 0 for No):
## File not overwritten
```