analize_2

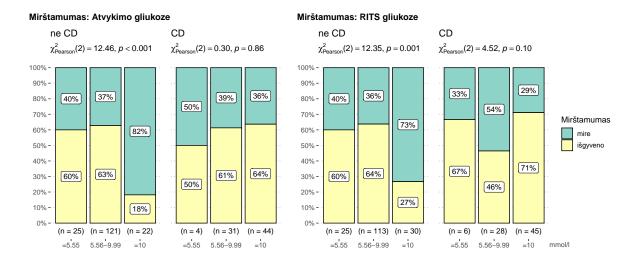
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1 Duomenu analize

Warning: NAs introduced by coercion

1.1 Glikemijos verčių ir mirštamumo vertinimas

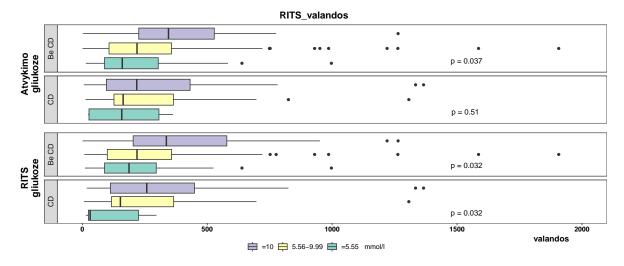


1.2 Glikemijos verčių ir hospitalizacijos trukmės vertinimas

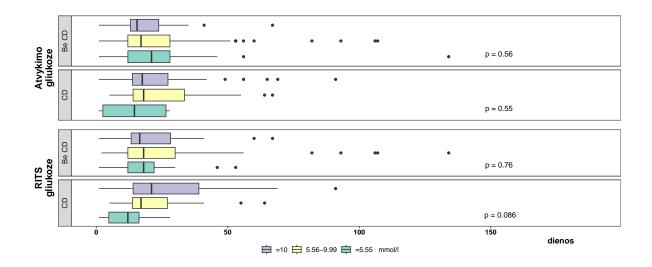
Grupinis

Bendras

 $RITS_valandos$

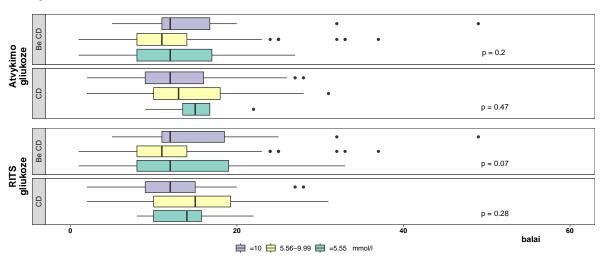


Lovadieniai

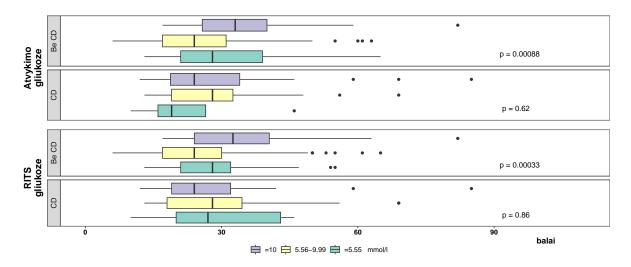


1.3 Glikemija ir prognozes skales

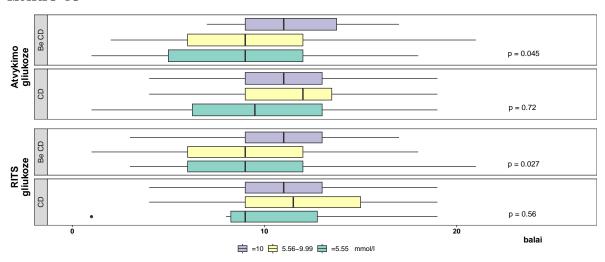
APACHEII



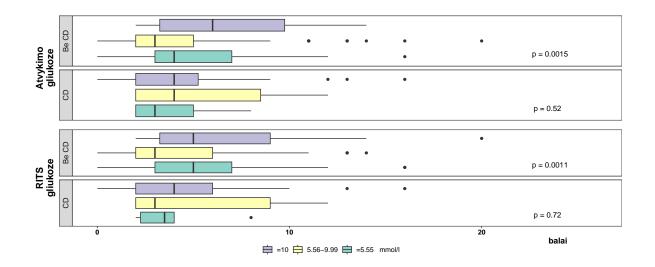
SAPS



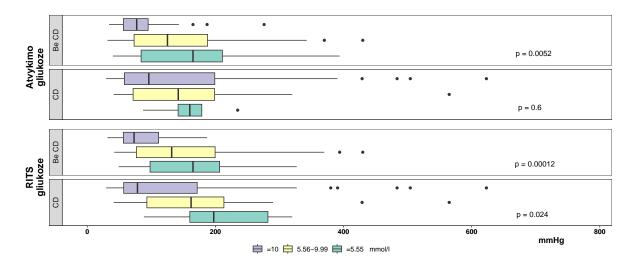
ISARIC 4C



SOFA

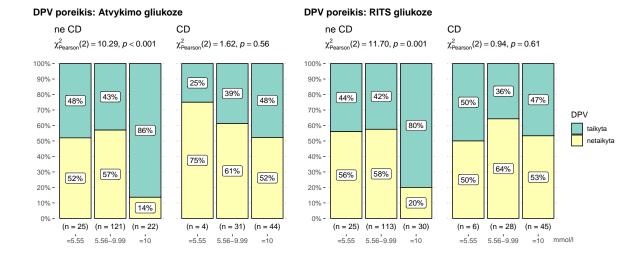


1.4 Hiperglikemija ir kvėpavimo funkcija

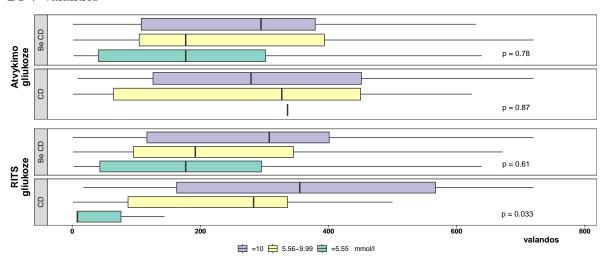


1.5 Hiperglikemija ir kvėpavimo funkcijos nepakankamumo gydymo būdai

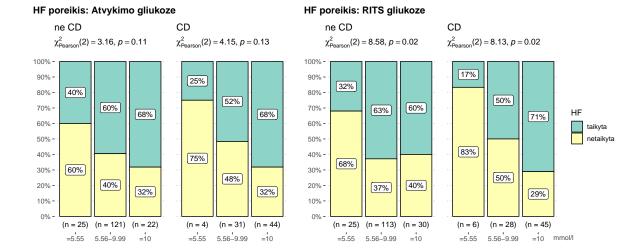
DPV poreikis



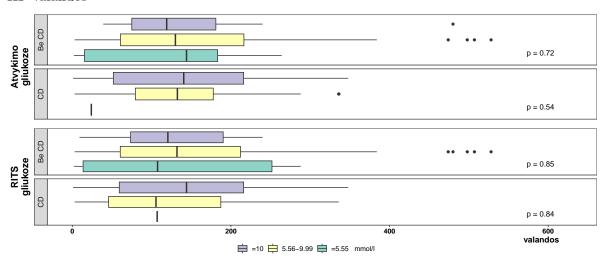
DPV valandos



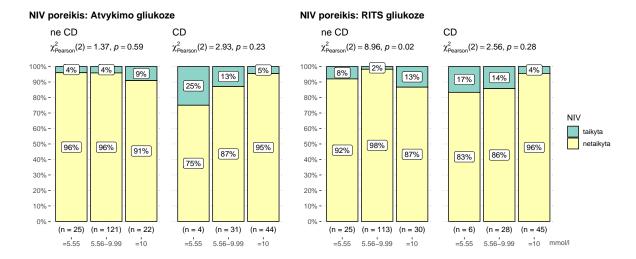
HF poreikis



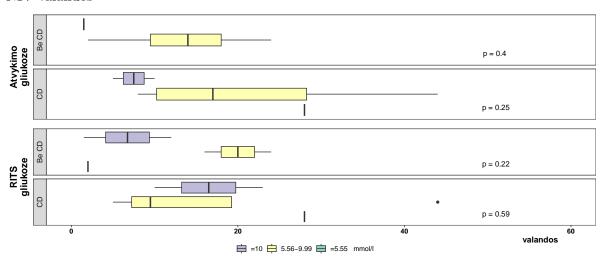
HF valandos



NIV poreikis

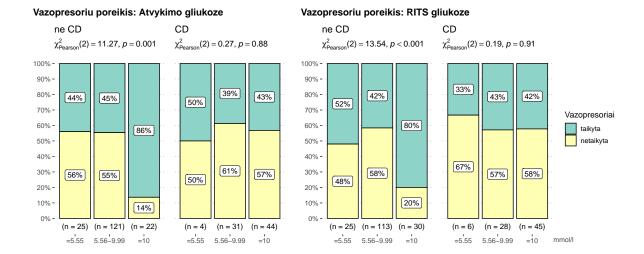


NIV valandos



1.6 Hiperglikemija ir kraujotakos palaikymo būdai

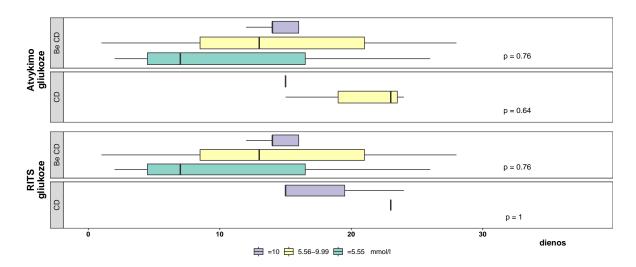
Vazopresoriu poreikis



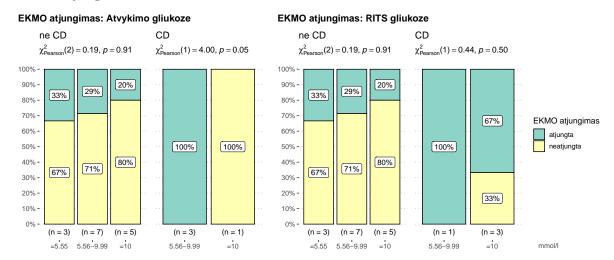
ECMO

EKMO poreikis: Atvykimo gliukoze EKMO poreikis: RITS gliukoze ne CD ne CD CD $\chi^2_{Pearson}(2) = 8.37, \, \rho = 0.03$ $\chi^2_{Pearson}(2) = 4.54, p = 0.17$ $\chi^2_{Pearson}(2) = 2.30, p = 0.32$ $\chi^2_{Pearson}(2) = 0.69, \, p = 0.71$ 100% -100% -4% 7% 6% 6% 10% 12% 12% 17% 90% -90% -23% 80% -80% -70% -70% -EKMO 60% -60% taikyta 50% -100% 50% -100% netaikyta 98% 96% 94% 94% 93% 88% 90% 88% 83% 40% -40% -77% 30% -30% -10% -10% -(n = 25) (n = 121) (n = 22) (n = 4) (n = 31) (n = 44) (n = 25) (n = 113) (n = 30) (n = 6) (n = 28) (n = 45) =5.55 5.56-9.99 =10 =5.55 5.56-9.99 =10 =5.55 5.56-9.99 =10 =5.55 5.56-9.99 =10 mmol/l

ECMO dienos

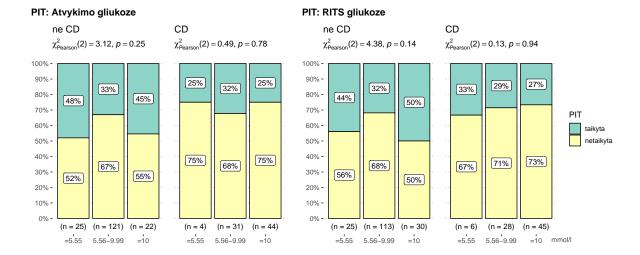


ECMO atjungta

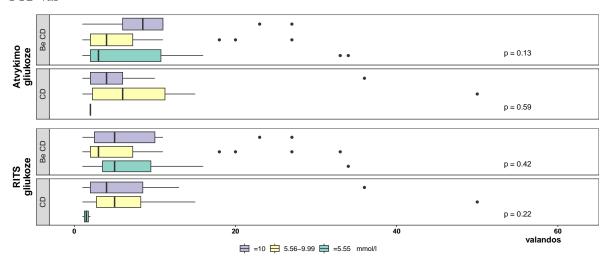


1.7 Hiperglikemija ir pakaitinė inkstų terapija

PIT



PIT val



1.8 Logistinis modelis

1.8.1 SAPS

Confusion Matrix and Statistics

Reference

Prediction X0 X1

XO 84 44

X1 7 29

Accuracy: 0.689

95% CI : (0.6122, 0.7589)

No Information Rate : 0.5549 P-Value [Acc > NIR] : 0.0003035

Kappa : 0.3372

Mcnemar's Test P-Value: 4.631e-07

Precision : 0.8056 Recall : 0.3973

F1: 0.5321

Prevalence: 0.4451
Detection Rate: 0.1768

Detection Prevalence : 0.2195 Balanced Accuracy : 0.6602

'Positive' Class : X1

1.9 SAPS + glu_fac

Confusion Matrix and Statistics

Reference

Prediction X0 X1

XO 83 38

X1 8 35

Accuracy: 0.7195

95% CI: (0.6441, 0.7868)

No Information Rate : 0.5549 P-Value [Acc > NIR] : 1.063e-05

Kappa : 0.4081

Mcnemar's Test P-Value : 1.904e-05

Precision: 0.8140

Recall : 0.4795

F1: 0.6034

Prevalence: 0.4451

Detection Rate : 0.2134
Detection Prevalence : 0.2622
Balanced Accuracy : 0.6958

'Positive' Class : X1

1.10 SOFA

Confusion Matrix and Statistics

Reference

Prediction X0 X1

X0 85 53

X1 6 20

Accuracy : 0.6402

95% CI : (0.5617, 0.7136)

No Information Rate : 0.5549 P-Value [Acc > NIR] : 0.01637

Kappa : 0.2222

Mcnemar's Test P-Value : 2.115e-09

Precision: 0.7692

Recall : 0.2740

F1 : 0.4040

Prevalence: 0.4451

Detection Rate: 0.1220

 ${\tt Detection\ Prevalence}\ :\ {\tt 0.1585}$

Balanced Accuracy: 0.6040

'Positive' Class : X1

$1.10.1 \text{ SOFA} + \text{glu_fac}$

Confusion Matrix and Statistics

 ${\tt Reference}$

Prediction X0 X1 X0 84 47 X1 7 26

Accuracy : 0.6707

95% CI: (0.5932, 0.742)

No Information Rate : 0.5549 P-Value [Acc > NIR] : 0.001638

Kappa : 0.2952

Mcnemar's Test P-Value : 1.113e-07

Precision: 0.7879 Recall: 0.3562

F1: 0.4906

Prevalence : 0.4451
Detection Rate : 0.1585
Detection Prevalence : 0.2012
Balanced Accuracy : 0.6396

'Positive' Class : X1

1.10.2 ISARIC4C

Confusion Matrix and Statistics

Reference

Prediction X0 X1

X0 78 33

X1 13 40

Accuracy: 0.7195

95% CI : (0.6441, 0.7868)

No Information Rate : 0.5549 P-Value [Acc > NIR] : 1.063e-05

Kappa : 0.4164

Mcnemar's Test P-Value : 0.005088

Precision: 0.7547

Recall : 0.5479

F1 : 0.6349

Prevalence: 0.4451

Detection Rate: 0.2439

Detection Prevalence: 0.3232 Balanced Accuracy: 0.7025

'Positive' Class : X1

1.11 ISARIC4C + gliukoze 2 faktoriais

Confusion Matrix and Statistics

Reference

Prediction X0 X1

X0 80 31

X1 11 42

Accuracy : 0.7439

95% CI: (0.67, 0.8088)

No Information Rate : 0.5549 P-Value [Acc > NIR] : 4.333e-07

Kappa : 0.4671

Mcnemar's Test P-Value : 0.00337

Precision : 0.7925 Recall : 0.5753

F1 : 0.6667

Prevalence: 0.4451

Detection Rate: 0.2561

Detection Prevalence: 0.3232

Balanced Accuracy: 0.7272

'Positive' Class : X1

1.12 APACHEII

Confusion Matrix and Statistics

Reference

Prediction X0 X1

X0 83 37

X1 8 36

Accuracy : 0.7256

95% CI: (0.6506, 0.7923)

No Information Rate: 0.5549 P-Value [Acc > NIR] : 4.994e-06

Kappa : 0.4218

Mcnemar's Test P-Value : 2.993e-05

Precision: 0.8182

Recall : 0.4932

F1: 0.6154

Prevalence: 0.4451

Detection Rate: 0.2195

Detection Prevalence: 0.2683 Balanced Accuracy: 0.7026

'Positive' Class : X1

1.13 APACHEII + gliukozė 2 faktoriais

Confusion Matrix and Statistics

Reference

Prediction X0 X1

X0 82 30

X1 9 43

Accuracy : 0.7622

95% CI : (0.6896, 0.8251)

No Information Rate: 0.5549 P-Value [Acc > NIR] : 2.846e-08 Kappa : 0.5045

Mcnemar's Test P-Value : 0.001362

Precision: 0.8269 Recall: 0.5890

F1: 0.6880

Prevalence: 0.4451
Detection Rate: 0.2622
Detection Prevalence: 0.3171
Balanced Accuracy: 0.7451

'Positive' Class : X1

Call:

NULL

Coefficients:

Estimate Std. Error z value Pr(>|z|)
(Intercept) -3.3805 0.5718 -5.912 3.38e-09 ***
APACHEII 0.2266 0.0420 5.395 6.85e-08 ***
`glu_fac 10` 2.2945 0.6858 3.346 0.000821 ***

Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 225.37 on 163 degrees of freedom Residual deviance: 159.74 on 161 degrees of freedom

AIC: 165.74

Number of Fisher Scoring iterations: 5

1.14 Modelis tik su Atvykimo gliukoze

Call:

glm(formula = Exitus_RITS ~ Atvykimo_glu, family = "binomial",
 data = analize_nocd)

Coefficients:

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 230.01 on 167 degrees of freedom Residual deviance: 219.86 on 166 degrees of freedom

AIC: 223.86

Number of Fisher Scoring iterations: 4

