

# Predicting the incidence of postoperative nausea and vomiting

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# Introduction

PONV incidence is generally 20-40%

Negative effects on health and well-being of patients

Financially costly to healthcare providers

Prophylaxis has negative side effects and financial costs

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# Challenge

Develop a scoring system for accurate prescription

Prophylaxis for patients at high risk of PONV

Predictive model that is neither too conservative nor too liberal

Balancing act known as “prevent-or-cure dilemma”

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# Scoring systems

Several well-documented models for predicting PONV

Guide prudent administration of anti-emetic prophylaxis

Logistic regression and stepwise backward elimination typically used for variable selection

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# Performance evaluation

Discrimination and calibration

- Most common measures of validity

AUC is most common measure of discriminating power

- Area under receiver operating characteristic curve (ROC)
- Values range from 0.61 to 0.785 across studies

Slope and squared correlation ( $R^2$ ) for line in calibration plot

- Calibration slopes range from 0.3 to 1.71
- Squared correlation ranges from 0.763 to 0.99

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# Purpose

Analyze data set of 461 patients from anesthesiologist Jelena Velickovic, MD, in Belgrade, Serbia

Develop predictive model for PONV

Performance comparable to or better than previous models published in medical literature

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Software

R statistical computing

R Markdown interface in R Studio

Additional packages installed to R

- dplyr - rename variables
- alr4 - marginal model plots
- leaps - regression subset plots
- car - regression subset plots
- rms - logistic regression
- pROC - area under ROC curve
- caret - data splitting, resampling

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Data set

Raw

- No missing values
- 916 rows and 26 columns

Cleaned

- Removed 93 duplicates
- 823 rows and 26 columns

Sample size

- Removed 362 records of patients who received prophylaxis
- 461 rows and 26 columns

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Variables

Original 26 variables encoded for analysis

- ID variable, 9 response variables, 16 predictor variables

Response variable selected

- Y = PONV0to24 (binary) = incidence of PONV within 24 hours of operation

Predictor variables selected

- Excluded 8 variables
  - Anesthetic and postoperative risk factors
- Remaining 8 variables for full model
  - Preoperative patient risk factors

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Predictors

$x_1$  = Age (integer)

$x_2$  = Gender (binary)

$x_3 \dots x_{27}$  = Diagnosis (categorical with 26 levels)

$x_{28} \dots x_{34}$  = Surgery (categorical with 8 levels)

$x_{35}$  = BMI (real)

$x_{36}$  = Nonsmoker (binary)

$x_{37}$  = Kinetosis history (binary)

$x_{38}$  = PONV history (binary)

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Full model

38 total variables

- When factors with more than 2 levels are taken into full account
- 25 dummy variables (for the 26 levels of *Diagnosis*)
- 7 dummy variables (for 8 levels of *Surgery*)

10 variables for full model

- ID variable
- Response variable
- 8 predictor variables

Incidence of PONV for this data set is 37%

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Exploratory data analysis

Full logistic regression model

Fitted using generalized linear method of least squares

3 predictors have estimated coefficients that are statistically significant

- PONV history
- Gender
- Nonsmoker

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Standardized deviance residuals

Skewness present in response and all 8 predictors

• Most are right-skewed

Dependency of log odds on each skewed predictors

• Linear function

• Log transformation

Residual plots are difficult to interpret for binary data

Examined marginal model plots instead

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Marginal model plots

Curves fitted for observed and predicted responses

Reasonable agreement between both fits in each of the marginal model plots for BMI and linear predictor

Lack of fit for Age, parabolic curvature for the observed response

Resolved by adding a quadratic term for Age

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Marginal model plots

• Reasonable agreement between observed and predicted fits

• Age, Age<sup>2</sup>, BMI, and linear predictor

• Indicates that current model is an adequate fit for the data

Marginal Model Plots

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Leverage values and standardized deviance residuals

None of leverage points exceed 2.5 standard deviations

Six points exceed two standard deviations

• Should be investigated

• Only 1% of 461 values in data set

Continued with assumption that current model is an adequate fit for the data

Proceeded next to variable selection

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Modeling techniques for logistic regression

Stepwise selection using all possible subsets of variables

Repeated sampling using  $k$ -fold cross-validation

Repeated sampling using bootstrap

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Plots of adjusted  $R^2$  against subset size

• Picks best subset of each size

• Squared and adjusted value of correlation coefficient

• Increases with number of predictors

• May lead to overfitting of model to data set

• Predictor with smallest  $p$ -value for its estimated coefficient was added to each subset to obtain the next subset

Adjusted R-squared

Adjusted R-squared

Subset Size

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Stepwise selection using all possible subsets

Minimum BIC value corresponds to predictor subset of size two

- PONVhistory, Surgery

Minimum AIC and AIC<sub>c</sub> values correspond to predictor subset of size four

- PONVhistory, Surgery, Gender, and Nonsmoker
- Includes same three predictors as full logistic regression
- PONVhistory, Gender, and Nonsmoker
- Before and after adding squared term for Age
- Has higher R<sup>2</sup> value than predictor subset of size two

Surgery excluded because none of its coefficients were statistically significant

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Parsimonious logistic regression model from stepwise selection

$$Y = g(\beta_0 + \beta_1 \text{PONVhistory} + \beta_2 \text{Gender} + \beta_3 \text{Nonsmoker} + e)$$

where  $e \sim \text{iid } N(0,1)$

Logit function used to model binary response variable:

$$g^{-1}(Y) = \left( \frac{\theta(Y)}{1 - \theta(Y)} \right)$$
$$= \beta_0 + \beta_1 \text{PONVhist} + \beta_2 \text{Gender} + \beta_3 \text{Nonsmoker} + e$$

where  $\theta(Y) = \frac{(Y)}{1+(Y)} = \frac{1}{1+(-Y)}$

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Estimated coefficients

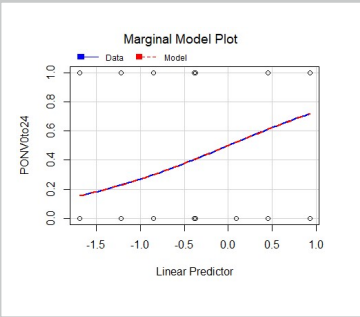
- Intercept and all three predictors are statistically significant

```
## Call:
## glm(formula = PONVhtc24 ~ PONVhistory + Gender + Nonsmoker, family =
## binomial,
## data = ponv)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -1.588  -1.023  -0.721   1.348   1.929
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  -1.6911     0.3827  -5.587 2.32e-08 ***
## PONVhistory1  1.3820     0.3132   4.569 3.19e-05 ***
## Gender1       0.8481     0.2883   2.997  0.00273 **
## Nonsmoker1    0.4766     0.2159   2.288  0.02724 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##    Null deviance: 689.86 on 460 degrees of freedom
## Residual deviance: 573.95 on 457 degrees of freedom
## AIC: 581.95
##
## Number of Fisher Scoring iterations: 4
```

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Linear predictor

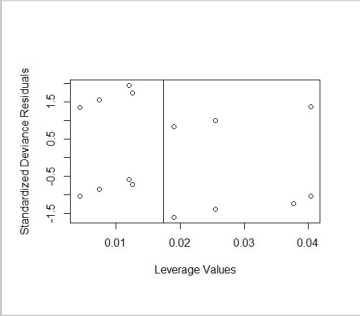
- Reasonable agreement between actual and predicted fits
- Model is an adequate fit for the data set



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Leverage values and standardized deviance residuals

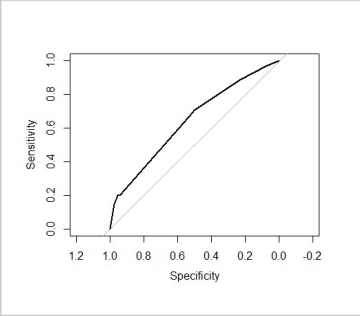
- All points within two standard deviations
- No bad leverage points
- Valid model for prediction



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Area under ROC curve

- Receiver operating characteristic
- The area under ROC curve (AUC) is 0.639
- ROC curve is insensitive to class imbalance
- ROC curve is a function of both sensitivity and specificity



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Confusion matrix

- Sensitivity of 0.94
  - Good at classifying high risk patients
- Specificity of 0.20
  - Poor at classifying low risk patients
- Reflected in the calibration plot

```
## Confusion Matrix and Statistics
##
##      Reference
## Prediction  0   1
##      0 272 138
##      1   17   34
##
##      Accuracy : 0.6638
##      95% CI   : ( 0.6186, 0.7068)
##      No Information Rate : 0.6269
##      P-Value [Acc > NIR] : 0.05527
##
##      Kappa : 0.1619
##
##      Mcnemar's Test P-Value : < 2e-16
##
##      Sensitivity : 0.9412
##      Specificity : 0.1977
##      Pos Pred Value : 0.6634
##      Neg Pred Value : 0.6667
##      Prevalence : 0.6269
##      Detection Rate : 0.5980
##      Detection Prevalence : 0.8894
##      Balanced Accuracy : 0.5694
##
##      'Positive' Class : 0
##
## Setting levels: control = 0, case = 1
## Setting direction: controls < cases
```

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Calibration plot

- Slope
  - Underestimates patients with low PONV risk
  - Overestimates patients with high PONV risk

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Repeated sampling using *k*-fold cross-validation

- 10-fold cross-validation
  - Each training set contains 90% of data set
  - Each test set contains the other 10% of data (left out)
  - Each sample size between 414 and 416
    - Large training sets avoid potential bias and variance issues
- 5 repetitions of process
  - Generate 50 different holdout sets for estimating model accuracy
  - Increase precision of estimates while maintaining small bias

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Repeated sampling using *k*-fold cross-validation

- Performed using the full logistic regression model
- 3 predictor coefficients were statistically significant ( $p < .05$ )
  - PONV history, Gender, and Nonsmoker
- Same as model fitted using all possible subsets on full data set

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Area under ROC curve

- 10-fold cross-validation
- AUC = 0.7358
- Improvement over AUC of 0.64 for baseline model obtained from all possible subsets

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Calibration plot

- 10-fold cross-validation
- Sensitivity of 0.31
  - Poor at classifying high risk patients
- Specificity of 0.83
  - Good at classifying low risk patients
- Not reflected in calibration plot

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Repeated sampling using the bootstrap

Random sample equal to size of data set

Taken with replacement

Repeated 25 times

Performed using the full logistic regression model

3 predictor coefficients were statistically significant ( $p < .05$ )

- PONV history, Gender, and Nonsmoker
- Same as model fitted using all possible subsets on full data set

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Area under ROC curve

- Bootstrap
- AUC = 0.7408 for bootstrap
- Improvement over AUC of 0.64 for baseline model obtained from all possible subsets

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Calibration plot

- Bootstrap
- Sensitivity of 0.34
  - Poor at classifying high risk patients
- Specificity of 0.78
  - Good at classifying low risk patients
- Not reflected in calibration plot

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Predictions of PONV

Logistic regression model obtained from all possible subsets consists of three predictors which are all binary variables

Lacks numerous dummy variables of two models trained with resampling techniques

Chosen as parsimonious model to make predictions

Applied to examples of hypothetical patients

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Predictions of PONV

PONV history	Gender	Nonsmoker	PONV probability
Yes	Female	Yes	0.7168
Yes	Female	No	0.6111
Yes	Male	Yes	0.5221
Yes	Male	No	0.4042
No	Female	Yes	0.4075
No	Female	No	0.2992
No	Male	Yes	0.2289
No	Male	No	0.1556

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Predictions of PONV

The more significant predictors that a patient has, the more likely the patient will have PONV

A patient with all three risk factors in the parsimonious model has a 72% probability of experiencing PONV

A patient with none of the three risk factors has a 16% probability of experiencing PONV

Previously determined to be a valid predictive model

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Conclusion

Simple model may have practical application for patient populations with characteristics like this data set

To choose a threshold for prescription of prophylaxis, healthcare professionals could select one of the two hypothetical predictions in our example having a probability greater than 50%

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