1 A comparative study of machine learning algorithms in predicting severe complications

2 after bariatric surgery

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20 Abstract

21	Accurate models to predict severe postoperative complications could be of value in the
22	preoperative assessment of potential candidates for bariatric surgery. Traditional statistical
23	methods have so far failed to produce high accuracy. To find a useful algorithm to predict the
24	risk for severe complication after bariatric surgery, we trained and compared 29 supervised
25	machine learning (ML) algorithms using information from 37,811 patients operated with a
26	bariatric surgical procedure between 2010 and 2014 in Sweden. The algorithms were then
27	tested on 6,250 patients operated in 2015. Most ML algorithms showed high accuracy (>90%)
28	and specificity (>0.9) in both the training and test data. However, none achieved an acceptable
29	sensitivity in the test data. ML methods may improve accuracy of prediction but we did not
30	yet identify one with a high enough sensitivity that can be used in clinical praxis in bariatric
31	surgery. Further investigation on deeper neural network algorithms is needed.

33 Introduction

34	Morbid obesity is a global public health threat of growing proportions(Ng et al., 2014).
35	Bariatric surgery offers the best chance for long-term weight-loss and resolution of
36	comorbidities(Sjostrom et al., 2004). Although modern bariatric surgery is considered to be
37	safe, severe postoperative complications still occur(Finks et al., 2011; Stenberg et al., 2014).
38	Accurate prediction models for severe postoperative complications could aid preoperative
39	decision making for surgeons, anesthesiologists and patients. These models could also serve
40	as basis for case-mix comparisons between different centers. Some prediction models based
41	on linear regression of patient-specific data allow for relatively simple and interpretable
42	inference; however, they have so far been proven inaccurate and can thus not be used in
43	clinical practice(Geubbels et al., 2015; Stenberg et al., 2018).
44	In contrast, some machine learning (ML) methods have been shown to provide quite accurate
45	predictions, and have increasingly been used in diagnosis and prognosis of different diseases
46	and health conditions(Anderin et al., 2015; Kourou et al., 2015; Pan et al., 2017). ML
47	methods are data-driven analytic approaches that specialize in the integration of multiple risk
48	factors into a predictive algorithm(Passos et al., 2016). Over the past several decades, ML
49	tools have become more and more popular for medical researchers. A variety of ML
50	algorithms, including artificial neural networks, decision trees, Bayesian networks, and
51	support vector machines (SVMs) have been widely applied with the aims to detect key
52	features of the patient conditions and to model the disease progression after treatment from
53	complex health information and medical datasets. The application of different ML methods
54	for feature selection and classification in multidimensional heterogeneous data can provide
55	promising tools for inference in medical practices. These highly nonlinear approaches have
56	been utilized in medical research for the development of predictive models, resulting in
57	effective and accurate decision making(Ali, 2017; Jiang et al., 2017).

58 Although new and improved software packages have significantly eased the implementation 59 burden for many ML methods in recent years, few studies have used ML methods to examine the risk factors or predict the prognosis after bariatric surgery, including diabetes 60 remission(Hayes et al., 2011; Pedersen et al., 2016), complication(Razzaghi et al., 2017), 61 weight status(Piaggi et al., 2010; Thomas et al., 2017), and adverse events and death(Ehlers et 62 63 al., 2017). Even though there is evidence that the use of ML methods can improve our 64 understanding of postoperative progression of bariatric surgery, an appropriate level of validation is needed in order for these methods to be considered in the clinical practice. 65 In this study, we compared different conventional supervised ML algorithms in the modeling 66 67 of severe postoperative complication after bariatric surgery. The study was based on the data from the Scandinavian Obesity Surgery Registry (SOReg). The SOReg is a national quality 68 and research register, covering virtually all bariatric surgical procedures performed in Sweden 69 since 2010. The register has been described in detail elsewhere(Hedenbro et al., 2015; 70 Stenberg et al., 2014), and a prediction model based on logistic regression for the same group 71 72 of patients has been described previously(Stenberg et al., 2018). The aim of the current study 73 was to find an algorithm or algorithms that perform well not only on the training data but also on the test data that were not used to train the algorithms. 74

76 **Results**

Baseline characteristics of the patients in the training data and the test data are presented in 77 Tables 1 and 2. The percentages of severe complication in the two data sets are 3.2% and 78 79 3.0%, respectively. No statistically significant difference was found for percentages of severe 80 complication between the two data sets (Pearson chi-square = 0.8283, p = 0.363). Univariable analyses indicate that differences of mean age, mean body mass index (BMI), 81 median HbA1c, percentages of comorbidities for hypertension, diabetes, dyslipidaemia, and 82 previous venous thromboembolism, and percentage of revisional surgery between the patients 83 84 presenting and without severe complication are statistically significant in the training data (Table 1). In the test data, the statistically significant differences were found for age, waist 85 circumference (WC), HbA1c, dyslipidaemia, and revisional surgery (Table 2). 86 87 Multivariable logistic regression analysis for the same data was published elsewhere(Stenberg et al., 2018). In brief, revisional surgery, age, low BMI, operation year, WC, and dyspepsia 88 were associated with the an increased risk for severe postoperative complication, however, the 89 90 performance of the multivariable logistic regression model for predicting the risk in individual patient case was poor. Validation of the model tested on patients operated in 2015 resulted in 91 an area under the receiver operating characteristic (ROC) curve of only 0.53, a Hosmer-92 Lemshow goodness of fit 17.91 (p=0.056) and Nagelkerke R² 0.013(Stenberg et al., 2018). 93 94 In current study, 19 supervised machine learning algorithms were compared and ten of them 95 were also trained using the synthetic minority oversampling technique (SMOTE), resulting in 29 ML algorithms. Most of the machine learning algorithms shown high accuracy (>90%) and 96 97 specificity (>0.9) for both training data and test data (Table 3), except that bagging linear discriminant analysis (LDA), bagging quadratic discriminant analysis (QDA), adaptive 98 boosting (AdaBoost) support vector machine (SVM), and multilayer perceptron (MLP) shown 99

low accuracy (<60%) for SMOTE training data, and oversampling-based bagging QDA 100 101 shown low accuracy for test data (accuracy = 56.1%) (Table 3). 102 Although most of the algorithms shown low sensitivity for both the training data and the test 103 data, some of them exhibited promising prediction ability in the training data. Sensitivities of 104 oversampling-based bagging QDA, random forest, AdaBoost extremely randomized (AdaExtra) trees, AdaBoost gradient regression (AdaGradient) trees, bagging k-nearest 105 106 neighbor (KNN), and deep learning neural network (NN) are 0.707, 0.965, 0.980, 0.968, 0.996, and 0.757 for SMOTE training data, respectively (Table 3). Even for test data, 107 108 oversampling-based bagging QDA and AdaBoost SVM show significant higher prediction 109 ability than other algorithms. The sensitivities of the two algorithms are 0.417 and 0.364, 110 respectively (Table 3). However, they still do not achieve an acceptable level for practical application. 111 When considering sensitivity and specificity together, most of the algorithms did not show 112 113 better prediction ability than a random predictor, i.e. an area under ROC curve of 0.5. The 114 areas under the ROC curves for all the algorithms, except for oversampling-based random 115 forest, AdaExtra trees, and adaGradient trees, and KNN, are around 0.5 (Figures 1 - 4). Although oversampling-based random forest, AdaExtra trees, AdaGradient trees, and KNN 116 117 show outstanding prediction ability on the SMOTE training data (areas under ROC curves are above 0.9), their performance on the test data are not optimistic (Figures 2 and 3). 118 119 The performance of the three regression-based algorithms (logistic regression, LDA, QDA), SVM, and the two neural network-based algorithms (MLP and deep learning NN) was poor in 120 any situation. However the bagging MLP and deep learning NN outperforms the tree-based 121 algorithms (Figures 2 and 4) for test data, their areas under ROC curves for the test data are 122 0.58 and 0.56, respectively (Figure 4) that are greatest among all the algorithms. 123

124 Discussion

Historically, laparoscopic gastric bypass has for a long time been the most common bariatric 125 procedure in Sweden, although laparoscopic sleeve gastrectomy has increased in popularity 126 127 over more recent years(Stenberg et al., 2014; The international federation for the surgery of obesity and metabolic disorders, 2017). The surgical technique is highly standardized with 128 more than 99% of all gastric bypass procedure being the antecolic, antegastric, laparoscopic 129 130 gastric bypass (so called Lönnroth technique)(Olbers et al., 2003). Virtually all patients receive pharmacologic prophylaxis for deep venous thrombosis and intraoperative antibiotic 131 prophylaxis(Hedenbro et al., 2015; Stenberg et al., 2014). Patients who have bariatric surgery 132 133 are exposed to the risk of having postoperative complications, which may increase the complexity of managing safety and healthcare costs. 134 135 Previous studies on postoperative complications of bariatric surgery have mainly used scoring

136 for identifying patients who are more likely to have complications after surgery. However,

these methods are not sensitive enough for clinical application(Geubbels et al., 2015;

138 Stenberg et al., 2018). The potential of ML tools as clinical decision support in identifying

risk factors and predicting health outcomes is therefore worth investigation on complications

140 associated with bariatric surgery. To our knowledge, there is only one study that compared the

141 performance of different ML algorithms in predicting the postoperative complications in

142 imbalanced bariatric surgery data set(Razzaghi et al., 2017). Although the study indicates that

the combination of a suitable feature selection method with ensemble ML algorithm equipped

144 with SMOTE can achieve higher performance in predictive models for bariatric surgery risks,

the ML algorithms were not validated using external test data. After all, for prediction

146 purpose, we are not very interested in whether or not an algorithm accurately predicts severe

147 complication for patients used to train the algorithm, since we already know which of those

patients have severe complications, but are interested in whether the algorithms may 148 149 accurately predict the future patients based on their clinical measurements. 150 Our study compared in total 29 ML algorithms using real world data. Although the 151 sensitivities of the algorithms were generally low, the study indicates that some ML algorithms were able to achieve higher accuracy than tradition logistic regression 152 models(Geubbels et al., 2015; Stenberg et al., 2018). Four of 29 algorithms were able to 153 154 achieve high sensitivity (>0.95) and two achieved moderate sensitivity (>0.70) in the training data, including three tree-based algorithms, bagging KNN, bagging QDA, and deep learning 155 NN. We should notice that all the high or moderate sensitivities were obtained from SMOTE 156 157 training data and/or using ensemble algorithms. Our findings support the previous study that ensemble ML algorithms equipped with SMOTE can achieve higher performance metrics for 158 imbalanced data(Razzaghi et al., 2017). 159

Despite showing promising capability of prediction in training data, none of the 29 ML 160 161 algorithms satisfactorily predicted severe postoperative complication after bariatric surgery in 162 the test data. Why did the algorithms do a poor job of predicting the patients who had severe complication in test data? One potential explanation for this may be related to the limited 163 number of severe postoperative complication in the current dataset, which cannot reveal the 164 165 underlying relationship between risk factors and adverse health outcomes. Although there are several known risk factors, each of them only imposes a small increase in the risk for 166 postoperative complication(Finks et al., 2011; Longitudinal Assessment of Bariatric Surgery 167 et al., 2009; Maciejewski et al., 2012; Stenberg et al., 2014). Another likely explanation may 168 be that preoperatively known variables are insufficient to predict postoperative complications. 169 170 In previous studies, the highest accuracy for prediction of postoperative complication has been models including operation data, mainly intraoperative complication and conversion to 171 open surgery(Geubbels et al., 2015; Stenberg et al., 2014). Although including intraoperative 172

adverse events and conversion to open surgery may improve the accuracy of prediction 173 174 models, such models would not be useful in the preoperative assessment for patients or for case mix comparisons. Furthermore, because the algorithms try to minimize the total error 175 rate out of all classes, irrespective of which class the errors come from, they are not 176 appropriate for imbalanced data such as what we used in our study (Maalouf et al., 2018). 177 Compared with traditional generalized linear predictive models, non-linear ML algorithms are 178 179 more flexible and may achieve higher accuracy but at the expense of less interpretability. Although there are interpretable models such as regression, Naïve Bayes, decision tree and 180 random forests, several models are not designed to be interpretable(James et al., 2013). The 181 182 aim of the methods is to extract information from the trained model to justify their prediction outcome, without knowing how the model works in details. The trade-off between prediction 183 accuracy and model interpretability is always an issue when we have to consider in building a 184 ML algorithm. A common quote on model interpretability is that with an increase in model 185 complexity, model interpretability goes down at least as fast. Fully nonlinear methods such as 186 187 bagging, boosting and support vector machines with nonlinear kernels are highly flexible 188 approaches that are harder to interpret. Deep learning algorithms are notorious for their uninterpretability due to the sheer number of parameters and the complex approach to extracting 189 190 and combining features. Feature importance is a basic (and often free) approach to interpreting the model. Although some nonlinear algorithms such as tree-based algorithms 191 192 (e.g. random forest) may allow to obtain information on the feature importance, we cannot obtain such information from many ML algorithms. 193

Therefore, recent attempts have been made to improve interpretability for the black-box
algorithms even such as deep learning. Local interpretable model-agnostic explanations
(LIME) is one of them to make these complex models at least partly understandable. LIME is
a more general framework that aims to make the predictions of 'any' ML model more

interpretable. In order to remain model-independent, LIME works by modifying the input to
the model locally(Mishra et al., 2017; Ribeiro et al., 2016). So instead of trying to understand
the entire model at the same time, a specific input instance is modified and the impact on the
predictions are monitored.

202 Regarding specific algorithm, though their motivations differ, the logistic regression and LDA or QDA methods are closely connected, therefore we were not surprised that LDA or QDA 203 204 did not show significant improvement in prediction than logistic regression(Stenberg et al., 205 2018). KNN takes a complete different approach from classification which is completely non-206 parametric(James et al., 2013). Therefore, we can expect it to outperform parametric models 207 such as logistic and LDA. However, KNN cannot tell us which predictor are of importance. 208 QDA serves as a compromise between the non-parametric KNN and the LDA and logistic regression. Though not as flexible as KNN, QDA can perform better in the limited training 209 data situation. MLP is a class of feedforward artificial neural network, which consists of at 210 least three layers of nodes. Its multiple layers and non-linear activation can distinguish data 211 212 that is not linearly separable. Deep learning NNs are high-level NNs including convolutional 213 NN and recurrent NN et al. In our study, the deep learning NN with five hidden layers outperforms the conventional MLP with two hidden layers, especially on SMOTE training 214 215 data (areas under ROC curves are 0.67 vs. 0.37), which deserves further investigation in the 216 future.

Our study demonstrates that ensemble learning may improve predictions by combining several base algorithms. However, usually there are several ensemble methods available, such as bagging, boosting, and stacking(Zhou, 2012). A number of studies have shown that, when decomposing a classifier's error into bias and variance terms, AdaBoost is more effective at reducing bias, bagging is more effective at reducing variance, and stacking may improve predictions in general(Kotsiantis et al., 2007). There is no golden rule on which method works

best. The choice of specific ensemble methods is case by case and depends enormously on thedata.

225 There are some limitations in our study. First, the study was limited to data registered within 226 the SOReg. Cardiovascular and pulmonary comorbidities other than sleep apnea are not 227 mandatory variables within the registry and could thus not be included in the model. Although these comorbidities are known risk factors for postoperative complications(Finks et al., 2011; 228 229 Gupta et al., 2011; Maciejewski et al., 2012), they are not highly prevalent in European 230 studies(Geubbels et al., 2015). Second, although we compared 29 ML algorithms investigated 231 in our study, they are convenient and feasible methods for general medical researchers. 232 Because of computational complexity and less interpretability, many complicated and advanced ML algorithms were not yet investigated in our study. However, our study at least 233 points out a promising way for future investigations, i.e. deep learning NN equipped with 234 SMOTE. Last but not the least, the exhaustive grid search was used in our hyperparameter 235 optimization, which is extremely resource consuming and not optimal for complex ML 236 237 algorithms, therefore other advanced methods such as gradient-based or evolutionary optimization would be considered in the future. 238

239 Conclusion

240 ML algorithms have the potential to improve the accuracy in predicting the severe

postoperative complication among the 44,061 Swedish bariatric surgery patients during 2010

- 2015. Because the imbalance nature of the data where the number of the interested outcome

243 is relative small, oversampling technique needs to be adopted to balance the two outcomes

- 244 (presenting or without severe complication). Ensemble algorithms outperform base
- algorithms. In general, deep learning NN results in better predictions for unseen patients.
- 246

247 Materials and Methods

248 **Patients and features**

Patients registered in the SOReg between 2010 and 2015 were included in the present study. 249 All patients who underwent a bariatric procedure between 2010 and 2014 were used as 250 251 training data in the ML. Data from patients who underwent a bariatric surgical procedure in 2015 were used as test data to validate the algorithm's performance in predicting sever 252 253 postoperative complication within 30 days after surgery. In total, 37,811 and 6,250 bariatric patients from SOReg were included in the training data and test data, respectively. In total 16 254 features were included in ML, including five continuous features (age, HbA1c, body mass 255 index [BMI], waist circumstance [WC]), and operation year) and 11 binary features (sleep 256 apnoea, hypertension, diabetes, dyslipdaemia, dyspepsia, depression, musculoskeletal pain, 257 258 previous venous thromboembolism, revisional surgery, and severe postoperative complication). The last binary feature, i.e. severe postoperative complication, was used as 259 260 output variable for the supervised ML classifiers. All the continuous features were 261 standardized to have mean 0 and standard deviation 1 before they enter the classifier. HbA1c was log transformed before standardization because of its asymmetric distribution. 262

263 Descriptive and inferential statistical methods

264 Demographic and baseline characteristics of the patients were presented using descriptive 265 statistical methods. Continuous variables were portrayed as mean and standard deviation 266 (SD), or median and interquartile range where suitable, while categorical variables were 267 outlined as counts and percentages. The difference between the patient presenting and without 268 severe postoperative complication was tested using the Student's t-test or the Mann-Whitney 269 U test for normally or asymmetrically distributed continuous variables, respectively; and χ^2 270 test was used for binary variables.

271 ML algorithms

- 272 In current study, eight base ML algorithms, i.e. logistic regression, linear discriminant
- analysis (LDA), quadratic discriminant analysis (QDA), decision tree, k-nearest neighbor
- 274 (KNN), support vector machine (SVM), multilayer perceptron (MLP) and deep learning
- 275 neural network (NN), and 11 ensemble algorithms, i.e. adaptive boosting (AdaBoost) logistic
- 276 regression, bagging LDA, bagging QDA, random forest, extremely randomized (Extra) trees,
- 277 AdaBoost Extra trees, gradient regression tree, AdaBoost Gradient trees, bagging KNN,
- 278 AdaBoost SVM, and bagging MLP, were implemented.

279 Ensemble learning

In order to improve generalizability and robustness over a single ML algorithm, we also used ensemble methods to combine multiple base or ensemble algorithms. Five ensemble methods were applied in our study:

- AdaBoost for logistic regression, Extra trees, gradient regression trees, and
 SVM(Schapire, 2003);
- bagging for LDA, QDA, KNN, and MLP(Kotsiantis et al., 2007);
- random forests for decision tree(Liaw & Wiener, 2002);
- Extra trees for decision tree(Geurts et al., 2006);
- gradient boosted regression trees for decision tree(Friedman, 2002).

289 Initialization and optimization of hyperparameters

- 290 ML algorithms involve a number of hyperparameters that have to be fixed before running the
- algorithms. In contrast to the parameters that are learned by training, hyperparameters
- determine the structure of a ML algorithm and how the algorithm is trained. The initial values
- of the hyperparmeters for each ML algorithm used in our study are the default values

294	specified in the employed software packages based on recommendations or experience(Probst
295	et al., 2018). In KNN algorithm, ten nearest neighbors were used. In MLP algorithm, two
296	hidden layer were used with five and two neurons, respectively. In deep learning NN
297	algorithm, the sequential linear stack of layers was used, with five hidden layers (three dense
298	layers and two dropout regularization layers). For the detailed hyperparameterization of the
299	algorithms, please refer the scikit-learn user manual at http://scikit-
300	learn.org/stable/supervised_learning.html(Pedregosa et al., 2011) and the Keras
301	Documentation at <u>https://keras.io/</u> .
302	The hyperparameter optimization is defined as a tuple of hyperparameters that yields an
303	optimal algorithm which minimizes a predefined loss function (i.e. cross entropy loss function

in our study, see Annex 1) on a held-out validation set of the training data. The most wildly

305 used however exhaustive grid search was used to perform hyperparameter optimization in our

study, which specified subset of the hyperparameter space of a ML algorithm and was

307 evaluated by cross-validation using the training data(Bergstra & Bengio, 2012).

308 Cross validation

309 For training data, k-fold (k = 5 in our analyses) cross-validated predictions were used as predicted values. This approach involves randomly dividing the training data into k groups, or 310 folds, of approximately equal size. Then an algorithm is trained on the k-1 folds and the rest 311 312 one fold is retained as the validation fold for testing the algorithm. The process is repeated 313 until the algorithm is validated on all the k folds. For each patient in the training data, the predicted value that he/she obtained is the prediction when he/she was in the validation fold. 314 Therefore, only cross-validation strategies that assign all patients to a validation fold exactly 315 316 once can be used for the cross-validated prediction(James et al., 2013).

317 **SMOTE**

The bariatric surgery data is extreme imbalanced, i.e. only 1,408 of 44,061 (3.2%) patients 318 319 experienced severe postoperative complication after bariatric surgery. The imbalance often results in serious bias in the performance metrics(Batista et al., 2004). Therefore, we 320 321 performed synthetic minority oversampling technique (SMOTE) to tackle the imbalance(Chawla et al., 2002). SMOTE generates a synthetic instance by interpolating m 322 323 instances (for a given integer value m) of the minority class that lies close enough to each 324 other to achieve the desired ratio between the majority and minority classes. In our study, a 325 1:1 ratio between the patients presenting severe postoperative complication and without severe postoperative complication was achieved in the training data, i.e. SMOTE training 326 327 data. The aforementioned nine of the 11 ensemble ML algorithms and the deep learning NN were also implemented for the SMOTE training data. 328

329 **Performance metrics**

330 The performance of the in total 29 ML algorithms were evaluated using accuracy, sensitivity,

331 specificity, and area under the receiver operating characteristic (ROC) curve. ROC curve

shows the trade-off that the algorithms set the different threshold values for the posteriorprobability for the prediction.

334 Terminology and derivations of accuracy, sensitivity, specificity, and area under the ROC335 curve are given in Annex 1.

336 Software and hardware

337 The descriptive and inferential statistical analyses were performed using Stata 15.1

338 (StataCorp, College Station). ML algorithms were achieved using packages scikit-learn 0.19.1

- 339 (scikit-learn, <u>http://scikit-learn.org/</u>)(Pedregosa et al., 2011) and Keras 2.1.6 (Keras,
- 340 <u>https://keras.io/</u>) in Python 3.6 (Python Software Foundation, <u>https://www.python.org/</u>).

341	All the computation was conducted in a computer with 64-bit Windows 7 Enterprise operation
342	system (Service Pack 1), Intel ® Core TM i5-4210U CPU @ 2.40 GHz, and 16.0 GB installed
343	random access memory.
344	
345	Author contributions
346	Yang Cao, Xin Fang, Conceptualization, Data curation, Software, Formal analysis, Writing-

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- 348 Stenberg Data curation, Conceptualization, Validation, Investigation, Writing-original draft;
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- 358 Ethics
- 359 The study was approved by the Regional Ethics Committee in Stockholm and was conducted
- 360 in accordance with the ethical standards of the Helsinki Declaration (6th revision).

	All	No serious	Having serious	p-value
	N=37,811	complication	complication	
		N=36,591	N=1,220 (3.2%)	
		(96.8%)		
Age in years, mean \pm SD	41.2 ± 11.2	41.1 ± 11.2	$42.9\ \pm 10.7$	< 0.001*
Sex, n (%)				
Female	28,682 (75.9%)	27,766 (75.9%)	916 (75.1%)	0.521^{\dagger}
Male	9,129 (24.1%)	8,825 (24.1%)	304 (24.9%)	
BMI in kg/m ² , mean \pm	42.12 ± 5.66	42.13 ± 5.66	41.79 ± 5.58	0.0355^{*}
SD				
WC in cm, mean ± SD	126.0 ± 14.0	126.0 ± 14.0	126.2 ± 13.8	0.6018^{*}
HbA1c, median (P25,	38 (35, 42)	38 (38, 32)	38 (35, 43)	0.0090‡
P75)				
Comorbidity, n (%)				
Sleep apnoea	3,792 (10.0%)	3,656 (10.0%)	136 (11.2%)	0.186^{\dagger}
Hypertension	9,760 (25.8%)	9,404 (25.7%)	356 (29.2%)	0.006^{\dagger}
Diabetes	5,407 (14.3%)	5,204 (14.2%)	203 (16.6%)	0.018^{\dagger}
Dyslipidaemia	3,802 (10.1%)	3,667 (10.0%)	135(11.1%)	0.233^{\dagger}
Dyspepsia	3,970 (10.5%)	3,803 (10.4%)	167 (13.7%)	< 0.001†
Depression	5,609 (14.8%)	5,409 (14.8%)	200 (16.4%)	0.119 [†]
Musculoskeletal pain	4,905 (13.0%)	4,754 (13.0%)	151 (12.4%)	0.529^{\dagger}
Previous venous	918 (2.4%)	875 (2.39%)	43 (3.52%)	0.011^{\dagger}
thromboembolism				
Revisional surgery	1,367 (3.6%)	1,261 (3.5%)	106 (8.7%)	$< 0.001^{+}$

Table 1. Base line characteristics of the training patients

363 SD: standard deviation; BMI, body mass index; WC, waist circumference; P25, the 25th percentile;

364 P75, the 75th percentile.

365 *t-test was used; $^{\dagger}\chi^2$ test was used; $^{\ddagger}Mann$ -Whitney U test was used.

	All	No serious	Having serious	p-value
	N=6,250	complication	complication	
		N=6,062 (97.0%)	N=188 (3.0%)	
Age in years, mean \pm SD	41.2 ± 11.5	41.2 ± 11.5	42.9 ± 11.8	0.0423*
Sex, n (%)				
Female	4,832 (77.3%)	4,682 (77.2%)	150 (79.8%)	0.411^{\dagger}
Male	1,418 (22.7%)	1,380 (22.8%)	38 (20.2%)	
BMI in kg/m ² , mean \pm SD	41.22 ± 5.87	41.20 ± 5.89	41.95 ± 5.40	0.0848^{*}
WC in cm, mean \pm SD	123.3 ± 14.1	123.2 ± 14.0	126.2 ± 14.7	0.0086^*
HbA1c, median (P25,	37 (34, 41)	37 (34, 41)	38 (35, 44)	0.0017‡
P75)				
Comorbidity, n (%)				
Sleep apnoea	622 (10.0%)	607 (10.0%)	15 (8.0%)	0.359^{\dagger}
Hypertension	1,563 (25.0%)	1,506 (24.8%)	57 (30.3%)	0.088^{\dagger}
Diabetes	761 (12.2%)	734 (12.1%)	27 (14.4%)	0.352^{\dagger}
Dyslipidaemia	518 (8.3%)	493 (8.13%)	25 (13.3%)	0.011^{\dagger}
Dyspepsia	645 (10.3%)	620 (10.2%)	25 (13.3%)	0.173^{\dagger}
Depression	1,096 (17.5%)	1,053 (17.4%)	43 (22.9%)	0.051^{\dagger}
Musculoskeletal pain	1,315 (21.0%)	1,268 (20.9%)	47 (25.0%)	0.176^{\dagger}
Previous venous	182 (2.9%)	177 (2.99%)	5 (2.7%)	0.834^{\dagger}
thromboembolism				
Revisional surgery	61 (1.0%)	54 (0.9%)	7 (3.7%)	${<}0.001^{\dagger}$

Table 2. Base line characteristics of the test patients

367 SD: standard deviation; BMI, body mass index; WC, waist circumference; P25, the 25th percentile;

368 P75, the 75th percentile.

369 *t-test was used; $^{\dagger}\chi^2$ test was used; $^{\ddagger}Mann$ -Whitney U test was used.

371

Table 3. Performance of the algorithms

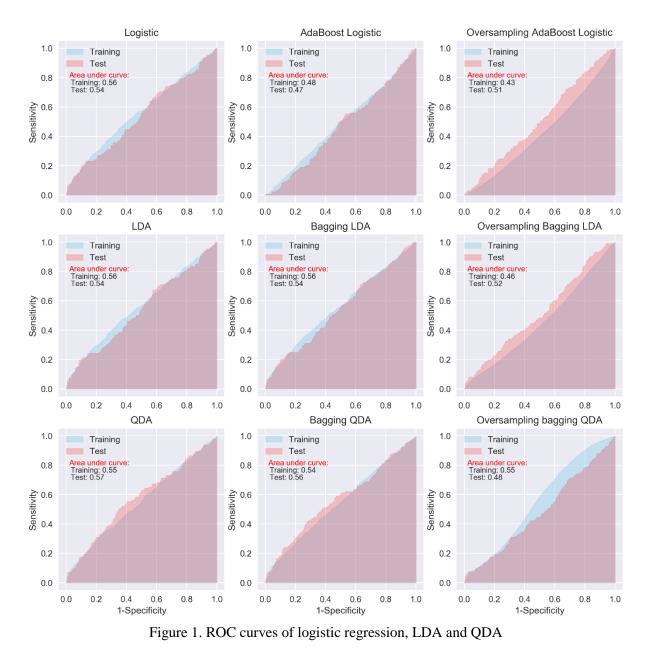
	Training data			Test data		
Algorithm	Accuracy	Specificity	Sensitivity	Accuracy	Specificit	Sensitivit
Logistic	<u>(%)</u> 96.9	1.000	0.000	<u>(%)</u> 97.1	<u>y</u> 1.000	<u>y</u> 0.000
AdaBoost Logistic	96.9	1.000	0.000	97.1	1.000	0.000
Oversampling AdaBoost Logistic	46.5	0.382	0.547	76.9	0.786	0.227
LDA	96.9	1.000	0.000	97.1	1.000	0.000
Bagging LDA	96.9	1.000	0.000	97.1	1.000	0.000
Oversampling Bagging LDA	46.3	0.370	0.556	79.0	0.807	0.212
QDA QDA	92.8	0.954	0.107	94.7	0.973	0.076
Bagging QDA	93.2	0.958	0.103	95.5	0.982	0.068
Oversampling bagging QDA	55.4	0.401	0.707	56.1	0.566	0.417
Decision tree	93.5	0.963	0.038	93.1	0.958	0.045
Random Forest	96.9	1.000	0.000	97.0	1.000	0.000
Oversampling Random Forest	94.5	0.925	0.965	96.6	0.995	0.008
ExtRa Trees	96.6	0.997	0.006	96.7	0.996	0.015
AdaBoost ExtRa Trees	96.6	0.996	0.004	96.6	0.995	0.008
Oversampling AdaExtra Trees	93.0	0.881	0.980	95.3	0.982	0.015
Gradient regression trees	96.9	1.000	0.000	97.1	1.000	0.008
AdaBoost Gradient trees	96.8	0.998	0.000	97.0	0.999	0.000
Oversampling AdaGradient trees	97.0	0.972	0.968	97.0	0.999	0.000
KNN	96.9	1.000	0.000	97.1	1.000	0.000
Bagging KNN	96.9	1.000	0.000	97.1	1.000	0.000
Oversampling Bagging KNN	79.4	0.592	0.996	82.3	0.841	0.235
SVM	96.9	1.000	0.000	97.1	1.000	0.000
AdaBoost SVM	96.9	1.000	0.000	97.1	1.000	0.000
Oversampling AdaBoost SVM	53.6	0.397	0.675	60.6	0.614	0.364
MLP	96.9	1.000	0.000	97.1	1.000	0.000
Bagging MLP	96.9	1.000	0.000	97.1	1.000	0.000
Oversampling bagging MLP	45.7	0.226	0.687	96.6	0.994	0.015
Deep learning NN	96.9	1.000	0.000	97.1	1.000	0.000
Oversampling deep learning NN	62.1	0.484	0.757	93.3	0.959	0.068

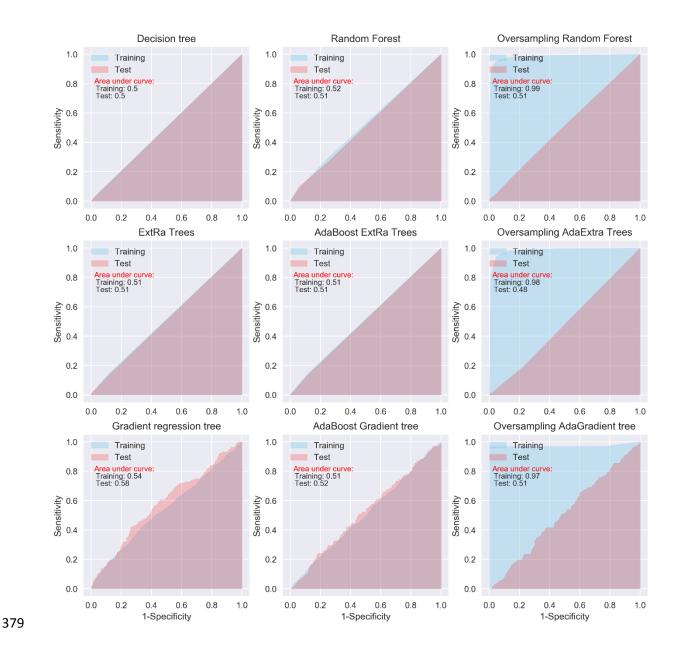
372 AdaBoost, adaptive boosting; LDA, linear discriminant analysis; QDA, quadratic discriminant

analysis; ExtRa, extremely randomized; AdaExtra, adaptive boosting extremely randomized;

AdaGradient, adaptive boosting gradient; KNN, k-nearest neighbor; SVM, support vector machine;

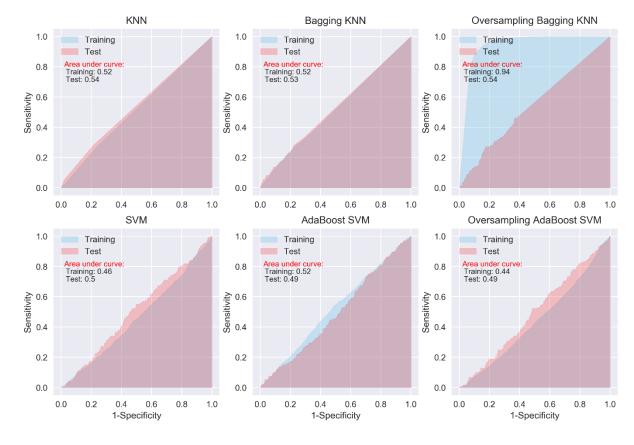
375 MLP, multilayer perceptron; NN, neural network





380

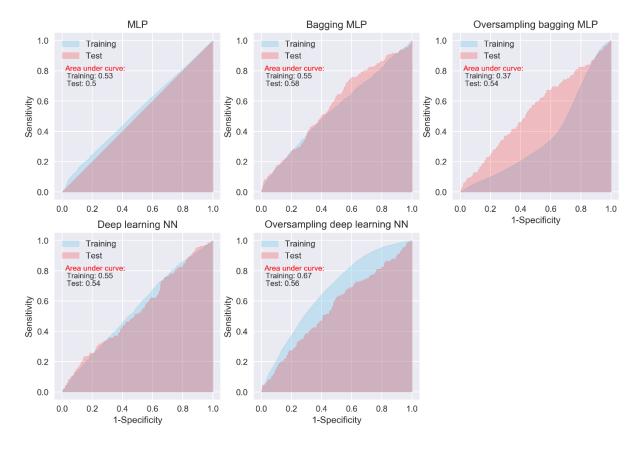
Figure 2. ROC curves of tree-based algorithms





381

Figure 3. ROC curves of KNN and SVM



383 384

Figure 4. ROC curves of neural network algorithms

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488

490	Annex 1: Terminology and derivations
491	Cross Entropy loss (Log loss):
492	$V(f(\vec{x}), y) = -y \ln(f(\vec{x})) - (1 - y) \ln(1 - f(\vec{x}))$
493	where y is true classifier $\in \{0, 1\}$ and $f(\vec{x})$ is predicted value.
494	
495	True or false refers to the predicted outcome being correct or wrong, while positive or negative refers
496	to presenting severe complication or no severe complication.
497	ACC: accuracy
498	SEN: sensitivity
499	SPE: specificity
500	TP: number of true positives, i.e. patient presenting severe complication correctly predicted as positive
501	TN: number of true negatives, i.e. patient without severe complication correctly predicted as negative
502	FP: number of false positives, i.e. patient without severe complication wrongly predicted as positive
503	FN: number of false negatives, i.e. patient presenting severe complication wrongly predicted as
504	negative
505	Total: total number of the patients, i.e. TP+TN+FP+FN
506	P: number of patients presenting severe complication, i.e. TP+FN
507	N: number of patients without severs complication, i.e. TN+FP
508	AUC: area under the receiver operating characteristic (ROC) curve for binary outcome
509	T: threshold for a patient is classified as presenting severe complication if X>T, where X is predicted
510	probability of a patients presenting severe complication by an algorithm.

511 ACC =
$$\frac{TP + TN}{Total} \times 100\%$$

512 SEN =
$$\frac{TP}{P}$$

513 SPE =
$$\frac{TN}{N}$$

514 AUC =
$$\int_0^1 SEN_T (1 - SPE)_T dT$$