1	External model validation of binary clinical risk prediction models in
2	cardiovascular and thoracic surgery
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21	CENTRAL PICTURE
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30	External validation of binary clinical risk-prediction models is vital. We provide
31	strategies for accomplishing this.
32	

### 34 INTRODUCTION

35 Clinical risk-prediction models (CRPMs, also known as prognostic models or risk score models) serve an important role in healthcare,<sup>1</sup> particularly for binary 36 adverse events (in-hospital, 30-day, or operative mortality) after cardiac, thoracic, 37 and vascular surgery. These models may be applied to 3 different objectives: 1) to 38 assess patient risk, which surgeons and patients can then factor in to healthcare 39 40 decisions; 2) to stratify risk, both for clinical decision-making and inclusion criteria in a controlled randomized trial.<sup>2</sup> and 3) to assess and compare healthcare outcomes 41 among providers (benchmarking). The comparison of observed with expected 42 43 outcomes, accounting for statistical uncertainty, can identify underperforming healthcare providers for quality improvement interventions.<sup>3</sup> 44

The wide-ranging importance of CRPMs in the cardiovascular specialty means that stakeholders must have confidence in them. A poorly performing model can lead to suboptimal decision-making, misinformed patients, false reassurance of a healthcare provider's performance, or false stigmatization of the provider.

49 Confidence is established by validating the model.<sup>4</sup>

Model validation can be internal, temporal, or external. Internal model 50 51 validation is one element of CRPM development, usually published alongside the model to confirm the model performs well for the training data. External validation, 52 which evaluates the generalizability (or transportability) of the model to other groups 53 of patients, is fundamental to demonstrating a model is appropriate for adoption in 54 clinical practice.<sup>4</sup> In cardiovascular and thoracic surgery, the majority of CRPMs 55 56 encountered will predict binary outcomes, which were created using multivariable regression techniques, in particular logistic regression. Therefore, we focus our 57 discussion to this area. However, the general principles and need for external 58

- validation apply to other outcome types and models, e.g. time-to-event data,<sup>5,6</sup> as
  well as to non-regression techniques, e.g. machine learning approaches.<sup>7</sup>
- 61

## 62 MODEL PERFORMANCE CONCEPTS

63 Performance of CRPMs is typically based on assessing two important
 64 features: calibration and discrimination.<sup>6</sup>

*Calibration* is the accuracy of the model for predicting events relative to
observed events in groups of patients. For example, if the mean predicted event
occurrence is 5% in a patient group, but the observed event occurrence is 10%, then
we conclude the model is not well calibrated because it underpredicts.

*Discrimination* is the ability of a model to distinguish between patients who 69 70 experienced the event and those who did not. Discrimination is measured using the 71 area under the receiver-operating-characteristic curve (AUROC), also referred to as the concordance (*c*-)statistic or *c*-index.<sup>5</sup> This value has a meaningful interpretation. 72 73 If we randomly select 2 patients, 1 who experienced the event and 1 who did not, 74 then the AUROC is equivalent to the probability that the risk score attributed to the former is greater than that attributed to the latter. An AUROC of 1 indicates perfect 75 classification; a value of 0.5 is equivalent to tossing a fair coin. 76

Other aspects of performance assessment include clinical usefulness,
 impact,<sup>8</sup> and overall performance measures such as the Brier score<sup>9</sup> and

79 concordance index, particularly for time-related events.

80

# 81 DESIGNING AND REPORTING AN EXTERNAL VALIDATION

82 When designing a validation study, thought must be given to several key 83 elements.

84 Selection of patients. The selection of patients used to externally validate a CRPM might differ from those used to develop the model. These differences might 85 be temporal or geographical, or related to clinical setting, inclusion or exclusion 86 87 criteria, definitions, diagnostic techniques, or inherent baseline case-mix differences between the two populations. It is important to highlight any differences that might 88 affect model transportability between the validation and original study sample, 89 90 particularly with validation of general all-surgery models (e.g. the EuroSCORE) within procedural<sup>10</sup> or operative subgroups.<sup>11</sup> 91

92 *Risk factor data*. It goes without saying that calculating a risk score requires access to all variables that comprise the risk score. One potential issue is conflict in 93 94 variable definitions. For example, a registry that only collects binary data on whether 95 pulmonary artery (PA) systolic pressure is >60 mmHg (a risk factor in the logistic 96 EuroSCORE model) would not be able to compute the EuroSCORE II risk score, which includes model coefficients for PA systolic pressures of 31 - 55 mmHg and 97 98 >55 mmHq. This is primarily an issue for retrospective validation studies, as clinical 99 registries can be updated to capture contemporary risk-score data.

100 *Missing data*. One cannot calculate a risk score without access to data for variables that comprise the CRPM. If a model contains a risk factor such as 101 102 preoperative serum creatinine, but these data are sparsely available in the dataset, 103 then in many cases the risk score cannot be calculated. Case-complete analyses-104 those that delete subjects with missing data for required variables-might lead to bias if those subjects are not representative of the whole population.<sup>12</sup> In certain 105 106 cases, reasonable estimates and assumptions can be made based on clinical expertise or additional information in the dataset. For example, a number of variables 107 108 in Society of Thoracic Surgeons (STS) risk models have coefficients set to 0 for

109 some variables in some models; if one is validating such a model, missing data for such a variable is of no consequence. Alternatively, statistical imputation or subset 110 analysis techniques might be applied to compensate.<sup>13,14</sup> If a validation study 111 specifically excludes certain groups of patients (for example, emergency surgery, 112 reoperations, or endocarditis), imputation of 0 is an accurate and appropriate 113 substitution, but the validation is only partial. In any case, it is always necessary to 114 115 summarize the frequency of missing data and present methods for managing it and its assumptions. 116

117 Sample size. Considerations regarding sample size should not be limited to randomized control trials. Single-center validation studies will often have a limited 118 pool of subjects, especially for subgroup analyses, and increasing the sample size 119 120 will require widening the study period, which could come at a price (see comment on calibration drift below). When designing a study, sample size (number of subjects) 121 alone is not enough; one must also consider effective sample size (number of 122 events). Relatively little attention has been given to this matter, but some studies 123 have recommended a minimum of 100 events and 100 non-events for validation 124 studies, and in certain applications, larger effective sizes will be required to obtain 125 adequate power.<sup>15,16</sup> 126

Outcome definitions. Many well-known CRPMs in cardiac surgery predict
 early or operative mortality, including the logistic EuroSCORE<sup>17</sup> and STS Cardiac
 Surgery Risk Models.<sup>18–20</sup> Operative mortality is generally accepted to mean death
 within 30 days (or later if the patient has not been discharged within 30 days).<sup>21</sup>
 However, other definitions of mortality exist, such as in-hospital mortality.<sup>22</sup> Two
 large databases reported operative mortality to be 4.63% and 3.57%, compared with
 in-hospital mortality of 4.02% and 2.94%, respectively.<sup>23,24</sup> In both cases, in-hospital

134 mortality was approximately 0.6% lower. In-hospital mortality is generally easier to robustly measure, whereas 30-day mortality requires post-discharge follow-up for 135 most patients.<sup>25</sup> Therefore, it is common to see models validated against in-hospital 136 137 mortality. In this example, we would expect the model to over-predict mortality relative to the observed data. It is reasonable to assess the model performance for 138 this similar endpoint; however, this subtlety should be borne in mind when designing 139 140 a study, particularly if the objective of the study is to compare models that have different outcome definitions. Similar considerations apply to cases where the 141 142 definition of a major postoperative complication used for model development differs from that in the validation dataset. 143

Large study windows. One simple way to increase sample size in a 144 145 validation study is to widen the study window. However, validation of a CRPM over a 146 substantially wide period can introduce a number of complexities. One potential issue is calibration drift.<sup>26,27</sup> Multiple studies demonstrated that the ratio of observed 147 148 mortality to mean logistic EuroSCORE was decreasing with time. Changing risk profiles, other variables influencing mortality, and changes in the association of risk 149 factors with outcome can all contribute to this phenomenon. This prompted the 150 introduction of the EuroSCORE II model<sup>23</sup> and the series of contemporary STS 151 models.<sup>18–20</sup> Researchers should be aware of this, particularly when validating 152 153 cardiac surgery CRPMs.

TRIPOD statement. In recent years, reporting of biomedical research has
 been improved with guidelines such as the CONSORT statement<sup>28</sup> for randomized
 trials and the PRISMA statement<sup>29</sup> for systematic reviews and meta-analyses.
 Prompted by evidence of poor quality reporting in the CRPM literature, the recent
 TRIPOD statement describes reporting guidelines for studies developing, validating,

or updating a prediction model.<sup>30</sup> We strongly encourage researchers to follow these
 guidelines and make use of the checklist for validating models. Examples of good
 practice and additional details have been previously published.<sup>31</sup>

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## 163 METHODS FOR ASSESSING CALIBRATION

Hosmer-Lemeshow test. The Hosmer-Lemeshow test is a frequently 164 reported statistical test for assessing calibration in CRPMs. However, it has a 165 number of drawbacks.<sup>31–35</sup> First, it is not easily interpreted; that is, it does not provide 166 167 a measure of the magnitude of any miscalibration. Second, for slight deviations in calibration, the test is sensitive to sample size. Third, the classical version of the test 168 is dependent on arbitrary groupings of patients. In some cases, the Hosmer-169 170 Lemeshow test remains a useful adjunct statistic, but should only be included as part of a more comprehensive assessment. Typically, the Hosmer-Lemeshow test refers 171 to a test based on 10 groups composed by deciles of risk. However, authors should 172 173 be aware that there are variations on the test with regard to groupings (quantiles vs. fixed cut-points), number of groups (g), degrees of freedom of the chi-squared 174 statistic (g-2 for internal vs. g for external validation), and software 175 implementations.<sup>35,36</sup> While q is typically selected to be 10, one must ensure the cell 176 counts are sufficient to justify the distributional approximation. Including a table of 177 178 observed and expected events by binning group provides a useful summary, and allows for inspection of each term for fit, as recommended by Hosmer and 179 Lemeshow (p. 188).<sup>36</sup> 180

*Calibration plot*. If a standard Hosmer-Lemeshow test is performed, then a
 simple graph—the calibration plot—is a straightforward next step (Figure).<sup>4</sup> Within
 each of the *g* groups, observed events are plotted against expected events. If the

model is well calibrated, then these points should be close to the 45° line. The
calibration plot can be augmented by overlaying a non-parametric smoothing curve
(e.g. loess) through the observed and predicted data<sup>37</sup> or a calibration curve.<sup>38</sup>
Contrary to the Hosmer-Lemeshow test and basic calibration plot, these additional
fits are not dependent on arbitrary groupings.

Calibration curves. Cox's calibration regression fits a logistic regression 189 between the observed event and the log-odds transformed predicted values.<sup>39</sup> A 190 191 perfectly calibrated CRPM (deriving from a logistic regression model) yields an intercept = 0 and a slope = 1. These fitted regression models can be superimposed 192 onto a calibration plot, giving an alternative graphical description of the 193 194 miscalibration. As well as quantifying the degree of miscalibration, one can also simultaneously test whether the estimated parameters reject the null hypothesis of 195 calibration. There are other related null hypotheses that can be tested for assessing 196 calibration also (p. 274).<sup>6</sup> 197

*Other tests.* The Hosmer-Lemeshow is ubiquitous in biomedical CRPM 198 199 literature. However, researchers can take advantage of a wide variety of statistical 200 tests to assess model validation, such as the aforementioned calibration curve test(s), the Spiegelhalter Z-test,<sup>40</sup> and methods proposed by Stallard.<sup>41</sup> Most can be 201 calculated using routine software packages.<sup>6,38</sup> There is no omnibus test of 202 calibration; each approach has different merits and limitations. Therefore, it is 203 204 important that researchers employ a broad repertoire of methods to address the study questions. 205

206

### 207 MODEL UPDATING

A natural extension to the validation of a CRPM is the concept of updating an existing model. This might involve exploring whether a new biomarker improves a model (e.g. using net reclassification improvement measures<sup>42</sup>), recalibrating a model,<sup>43</sup> and, more recently, assessing whether multiple models can be combined to provide a more accurate prediction (e.g. meta-models and model averaging).<sup>44</sup> This expanding research area is especially important in an era of personalized medicine.<sup>45</sup>

215

## 216 CONCLUSIONS

External validation of CRPMs is necessary to demonstrate their predictive 217 accuracy. Available models have likely been validated internally; however, using 218 219 them in different settings, locations, and populations can result in relatively poor performance. CRPMs that have been overfitted during development will also often 220 fail to generalise to the external validation sample. Calibration and discrimination 221 222 must be measured in order to establish validity. There are multiple statistical approaches available to interrogate the calibration, with it being widely accepted that 223 the ubiquitous Hosmer-Lemeshow test has limited utility. Execution of a rigorous 224 CRPM validation study rests in proper study design, application of suitable statistical 225 226 methods, and transparent reporting.

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### 363 FIGURE LEGEND

**Figure.** A calibration plot for simulated data (n = 500). The green triangles denote 364 the mean predicted and observed event probabilities for patients grouped into tenths 365 366 using deciles. The grey dashed line denotes perfect calibration. A smoothing curve (blue dashed line) and the calibration curve (red solid line) are also overlaid. The 367 distribution of calculated predicted probabilities is overlaid along the horizontal axis. 368 A subset of various statistics useful for validating the model are also shown. This 369 figure was generated using standard statistical software: the rms package for R (R 370 371 Core Team, R Foundation for Statistical Computing, Vienna, Austria; version 3.1.2). Further details are given in Harrell (2001)<sup>38</sup> and Harrell (2015).<sup>46</sup> Code to reproduce 372 this plot is given in the Appendix. 373 374

375	APPENDIX
376	R code to produce figure
377	# If `rms' package not install, run command
378	<pre># install.packages("rms")</pre>
379	library(rms)
380	## Simulate fake data:
381	## y = binary outcome
382	## x1, x2, x3 = covariates in the risk model
383	## n = sample size
384	set.seed(1)
385	n <- 1000 # 500 development + 500 validation
386	x1 <- runif(n) # covariate 1
387	x2 <- runif(n) # covariate 2
388	x3 <- runif(n) # covariate 3
389	logit <5 + 0.5*x1 + 2*x2 + 3.5*x3
390	P <- 1 / (1 + exp(-logit))
391	y <- ifelse(runif(n) <= P, 1, 0) # outcomes
392	d <- data.frame(x1, x2, x3, y) # combined dataset
393	
394	## Fit a risk prediction model to first half of the data
395	f <- lrm(y ~ x1 + x2 + x3, subset = 1:500)
396	
397	## Use model to get predictions for second half of data
398	<pre>pred.logit &lt;- predict(f, d[501:1000, ])</pre>
399	<pre>phat &lt;- 1 / (1 + exp(-pred.logit))</pre>
400	
401	## Validate prediction
402	<pre>val.prob(phat, y[501:1000], g = 10, riskdist = "predicted")</pre>