

Supplement

Two-Hour Algorithm for Early Diagnosis of Acute Myocardial Infarction Using A High-sensitivity Cardiac Troponin I Assay

Supplemental Methods

Derivation Cohort (APACE)

Patient Population

Advantageous Predictors of Acute Coronary Syndrome Evaluation (APACE) is an ongoing prospective international multicenter study with 12 centers in 5 European countries (Switzerland, Spain, Italy, Poland, Czech Republic) designed to contribute to improving the management of patients with MI. (ClinicalTrials.gov registry, number NCT00470587).(1)(2)(3)(4)(5) Adult patients presenting to the emergency department (ED) with symptoms suggestive of AMI (such as acute chest discomfort and angina pectoris) with an onset or peak within the last 12 hours were recruited. Enrollment was independent from renal function at presentation, while patients with terminal renal failure on chronic dialysis were excluded. The study was carried out according to the principles of the Declaration of Helsinki and approved by the local ethics committees. Written informed consent was obtained from all patients.

Routine Clinical Assessment

All patients underwent a clinical assessment that included standardized and detailed medical history including assessment of 34 predefined chest pain characteristics, vital signs, physical examination, 12-lead ECG, continuous ECG rhythm monitoring, pulse oximetry, standard blood test, and chest radiography if indicated. Concentrations of

25 cardiac troponins (cTn) including high-sensitivity cTn (hs-cTn) in some centers were
26 measured at presentation and serially thereafter as long as clinically indicated.
27 Treatment of patients was left to discretion of the attending physician. Clinical suspicion
28 for acute coronary syndrome (ACS) as the cause of the presenting symptom was
29 quantified 90 min after presentation by the treating emergency department (ED)
30 physician using a visual analog scale (VAS). At this time point, the ED physician had
31 completed his/her clinical assessment, including patient history, chest pain
32 characteristics, and detailed physical examination including vital signs, and reviewed
33 the ECG and the first local cTn measurement. We considered a concentration of \geq
34 80% as high pretest probability for ACS.(6)

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36 **Reference Standard: Adjudicated Final Diagnosis**

37 AMI was defined and cTn concentrations interpreted as recommended in current
38 guidelines.(7)(8–10) In brief, AMI was diagnosed when there was evidence of
39 myocardial necrosis with a significant rise and/or fall in a clinical setting consistent with
40 myocardial ischemia. Patients with AMI were further subdivided into type 1 AMI
41 (primary coronary events) and type 2 AMI (ischemia due to increased demand or
42 decreased supply, for example tachyarrhythmia or hypertensive crisis).(8,11)

43 The adjudication of final diagnoses was performed centrally in the core lab
44 (University Hospital Basel) for all patients incorporating concentrations of hs-cTnT
45 (primary analysis). More specifically, two independent cardiologists not directly
46 involved in patient care reviewed all available medical records (including patient
47 history, physical examination, results of laboratory testing including hs-cTnT
48 concentrations, radiologic testing, ECG, echocardiography, cardiac exercise test,
49 lesion severity and morphology in coronary angiography, discharge summary)

50 pertaining to the patient from the time of ED presentation to 90-day follow-up. In
51 general, serial sampling was performed until at least 3-6h after presentation to the ED
52 or onset of chest pain. In situations of diagnostic disagreement, cases were reviewed
53 and adjudicated in conjunction with a third cardiologist. While discharge diagnoses
54 often were correct and in agreement with the final adjudicated diagnosis, there were
55 also cases where the discharge diagnosis needed to be revised, most often because
56 more information became available from medical testing during early follow-up, and
57 more rarely, because the discharge diagnosis was not in agreement with the Universal
58 Definition of AMI.

59 The 99th percentile (hs-cTnT: 14ng/L) was used as cut-off for myocardial necrosis.
60 Absolute cTn changes were used to determine significant changes based on the
61 diagnostic superiority of absolute over relative changes.(12–17) Based on studies of
62 the biological variation of cTn.(18,19) as well as on data from previous chest pain
63 cohort studies,(12,20) a significant absolute change was defined as a rise or fall of at
64 least 10ng/L within six hours, or, in an assumption of linearity, as an absolute change
65 of 6ng/L within three hours. Predefined alternative diagnoses included “unstable
66 angina” (UA), “Cardiac symptoms of origin other than coronary artery disease” and
67 “non-cardiac chest pain”.

68

69 **Clinical Care: The (hs)-cTn assays and cut-off concentrations used for local** 70 **clinical care**

71 Routine clinical care comprised five different cTn assays at the different hospitals
72 throughout the whole recruitment period. The cTn assays used clinically in most of the
73 participating institutions changed during the study from a conventional cTn assay to
74 the hs-cTnT assay. In order to take advantage of the higher sensitivity and higher
75 overall diagnostic accuracy offered by the hs-cTnT assay (primary analysis), patients

76 were adjudicated using the hs-cTnT concentrations obtained from study specific blood
77 samples in addition to the clinically used (hs)-cTn concentrations.

78 The following conventional cTn assays were used: For the Roche cTnT 4th generation
79 assay, the 10% CV concentration is 0.035µg/L. The laboratories of the participating
80 sites reported only two decimals; therefore 0.04µg/L was used as a cut-off for
81 myocardial necrosis. In order to fulfil the criteria of a significant change (30% of 99th
82 percentile or 10% CV concentration), a patient would e.g. need to have a concentration
83 of <0.01µg/L at presentation and 0.04µg/L at 6h. A patient would also qualify if the first
84 concentration is 0.02µg/L and the second 0.04µg/L. A patient would not fulfil the criteria
85 if the first concentration is 0.03µg/L and the second is 0.04µg/L. If the first
86 concentration is 0.04µg/L, the second concentration needs to be at least 0.06µg/L.

87 For the Abbott AxSYM cTnI ADV, the 10% CV concentration is 0.16µg/L. A patient
88 having 0.16µg/L at presentation would meet the criteria for significant change if the
89 second was ≥0.21µg/L. A patient having <0.12µg/L at presentation (limit of detection)
90 would qualify if the second is >0.16µg/L.

91 For the Beckmann Coulter Accu cTnI, the 10% CV concentration is 0.06µg/L. A patient
92 having 0.06µg/L at presentation would qualify if the second is ≥0.08µg/L. A patient
93 having 0.05µg/L at presentation would qualify if the second is 0.07µg/L, but not
94 0.06µg/L. A patient having undetectable cTnI (cTnI <0.01µg/L) at presentation would
95 qualify if the second is ≥0.06µg/L.

96 For the Siemens Dimension Vista s-cTnI, the 10% CV concentration is 40ng/L. The
97 limit of detection is 15ng/L and the 99th percentile is 45ng/L. An absolute change of
98 20ng/L or more within 3-6h was considered significant.

99 For Elecsys hs-cTnT measured clinically, the same change criteria were applied as for
100 hs-cTnT measured from the study blood samples.

101

102 **Follow-up**

103 Patients were contacted 3, 12 and 24 months after discharge by telephone calls or in
104 written form. Information regarding death during follow up was furthermore obtained
105 from the patient's hospital notes, the family physician's records and the national
106 registry on mortality.

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108 **Validation Cohort (ADAPT and IMPACT)**

109 **Patient Population**

110 Accelerated Diagnostic Protocol to Assess patients with chest Pain symptoms using
111 contemporary Troponins as the only biomarker (ADAPT) was a multicenter,
112 diagnostic study enrolling patients between November 2007 and February 2011 in
113 two study centers in Australia and New Zealand.(21) The current study only uses
114 data from the Australian site. The Improved Assessment of Chest pain trial (IMPACT)
115 was a single center intervention study enrolling patients in the same Australian site
116 between February 2011 and March 2014.(22) Criteria for enrollment were the same
117 for ADAPT and IMPACT and included ≥ 18 years of age, with at least 5 minutes of
118 symptoms where the attending physician planned to perform serial s-cTnI tests. The
119 American Heart Association case definitions for possible cardiac symptoms were
120 used (i.e., acute chest, epigastric, neck, jaw, or arm pain; or discomfort or pressure
121 without an apparent non cardiac source).(23) Patients were excluded for any of the
122 following: a clear cause other than acute coronary syndrome for the symptoms at
123 presentation (e.g., examination findings of pneumonia), inability to provide informed
124 consent, staff considered recruitment to be inappropriate (e.g., receiving palliative
125 treatment), transfer from another hospital, pregnancy, previous enrollment, or inability
126 to be contacted after discharge. Patients with STEMI on presentation were also

127 removed from the current analysis. Perceived high risk was not used as an exclusion
128 criterion. The studies were carried out according to the principles of the Declaration
129 of Helsinki and approved by the local ethics committees.

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131 **Routine Clinical Assessment**

132 All patients underwent a clinical assessment that included standardized and detailed
133 medical history including vital signs, physical examination, 12-lead ECG, continuous
134 ECG rhythm monitoring, pulse oximetry, standard blood test, and chest radiography if
135 indicated. Concentrations of s-cTnI were measured at presentation and serially
136 thereafter as long as clinically indicated.

137

138 **Reference Standard: Adjudicated Final Diagnosis**

139 In the ADAPT and IMPACT cohorts, 30-day outcomes and final diagnoses were
140 adjudicated independently by local cardiologists using predefined standardized
141 reporting guidelines according to the current universal definition of myocardial
142 infarction. Cardiologists had knowledge of the clinical record, ECG, s-cTnI results and
143 objective testing from standard care. A second cardiologist conducted a blind review
144 of all acute coronary syndrome (ACS) cases and 10% of non-ACS cases. In cases of
145 disagreement, endpoints were agreed by consensus. Consensus was achieved for all
146 endpoints. Late samples were available for adjudication of final diagnosis in all
147 patients of the validation cohort, which means that serial sampling was performed
148 until at least 6h after presentation to the ED in all patients. For ADAPT, adjudication
149 of final diagnosis included s-cTnI measurements from blood samples obtained at
150 presentation and 6 to 12h thereafter. For IMPACT, adjudication included s-cTnI

151 measurements from blood samples obtained at presentation and at least 2 h
152 thereafter.

153 The Beckman Coulter 2nd generation AccuTnI assay (Beckman Coulter, Chaska,
154 MN) was used at the Australian site. This assay has a LoD of 10 ng/L, a 99th
155 percentile of 40ng/L and a 10% CV of 60ng/L. The 99th percentile (40ng/L) was used
156 as cut-off for myocardial necrosis. A delta of $\geq 20\%$ was used to detect a rising or
157 falling pattern and at least one value above the 99th percentile was necessary to
158 qualify for AMI diagnosis.

159

160 **Follow-up and clinical endpoints**

161 Patients were contacted at 6 weeks and 12 months by telephone calls or in written
162 form. 6 weeks follow-up was conducted for all patients, while 12 months follow-up
163 information was only available where the patient provided consent to further contact.
164 Information regarding death at twelve months was furthermore obtained from the
165 patient's hospital notes, the family physician's records and the national registry on
166 mortality. The primary prognostic endpoint was 30 days all-cause mortality.

167

168 **Derivation and validation of the hs-cTnI-Access 0/2h-algorithm**

169 The 0/2h-algorithm incorporates hs-cTnI concentration concentrations at presentation
170 and absolute hs-cTnI- changes within 2h as well as time since chest pain onset in order
171 to reflect the concept of the current hs-cTn 0/1h-algorithms suggested by the ESC. The
172 rule-out criteria for hs-cTnI are defined as an undetectable (very low) concentration at
173 presentation or a low baseline hs-cTnI concentration together with no relevant absolute
174 change within the first 2 hours Rule-in is defined by a high baseline hs-cTnI

175 concentration or a relevant absolute change in hs-cTnI within the first 2 hours. The
176 remaining patients were triaged towards the observe zone The hs-cTnI-Access 0/2h-
177 algorithm was developed in the derivation cohort (APACE) in selected patients with
178 available hs-cTnI-Access measurements at baseline and after 2h. The 0/2h-algorithm
179 incorporates hs-cTnI-Access concentrations at presentation and absolute hs-cTnI-
180 Access changes within 2h. (**Supplemental Figure 2**). Selection of these parameters
181 was based on the very high diagnostic accuracy of the combination of blood
182 concentrations at presentation with absolute changes for rule-out and rule-in of
183 AMI.(12,24–31) Derived thresholds for rule-out were selected to allow for a minimal
184 sensitivity and negative predictive value (NPV) of 99.5% and sensitivity of 99.0%.
185 Derived thresholds for rule-in were obtained based on a classification and regression
186 tree (CART) analysis targeting a minimal positive predictive value (PPV) of 70%.(32)
187 Nodes in the CART tree were constrained to have a minimal number of cases of 20 in
188 parent and child nodes. If a predefined target performance was missed in the derivation
189 sample using the CART-derived thresholds, thresholds were changed stepwise until
190 the predefined performance was fulfilled. The algorithm developed in the derivation
191 sample was tested for its diagnostic accuracy in an independent validation sample
192 consisting of the remaining subjects. The decision values derived in the derivation
193 sample were rounded to give whole values in ng/L. The whole algorithm developed in
194 the derivation cohort was then tested for diagnostic accuracy in the validation cohort
195 (ADAPT and IMPACT).

196

197 **Direct comparison of the hs-cTnI-Access 0/2h-algorithm with the 0/2h-** 198 **algorithms using hs-cTnT-Elecsys and hs-cTnI-Architect**

199 The diagnostic performance of the hs-cTnI-Access 0/2h-algorithm in both cohorts was
200 directly compared to the performances of the hs-cTnT-Elecsys 0/2h-algorithm and the

201 hs-cTnI-Architect 0/2h-algorithm in all patients of the validation cohort, who also had a
202 full set of hs-cTnT-Elecsys and hs-cTnI-Architect measurements at 0h and 2h.

203

204 **Follow-up**

205 The co-primary prognostic endpoints were overall survival after 30 days and 2 years
206 in the derivation cohort and 30 days and 1 year in the validation cohort.

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208 **Statistical analysis**

209 Continuous variables are described as mean \pm SD or median with interquartile
210 range (IQR), categorical variables by numbers and percentages. Differences in
211 baseline characteristics between patients with and without AMI as well as between
212 patients in the derivation and validation cohort were assessed using the Mann-Whitney
213 U test for continuous variables and the Pearson Chi-square test for categorical
214 variables.

215 For the assessment of the diagnostic performance of the hs-cTnI-Access 0/2h-
216 algorithm in the derivation and validation cohort, safety was assessed as the NPV and
217 the sensitivity for AMI in the rule-out group, accuracy of the rule-in strategy as the PPV
218 and specificity for AMI in the rule-in group while efficacy was quantified as the
219 percentage of patients triaged towards rule-out or rule-in for AMI within 2h.

220 Survival during 30 days and 365/720 days of follow-up according to the
221 classification provided by the hs-cTnI-Access 0/2h-algorithm was plotted in Kaplan-
222 Meier curves and the log-rank test was used to assess differences in survival between
223 groups. Bootstrapping has been used to calculate confidence intervals.

224 All hypothesis testing was two-tailed and *P*-values <0.05 were considered
225 statistically significant. Statistical analyses were performed using IBM SPSS Statistics
226 for Windows, version 25.0 (SPSS Inc), MedCalc 17.6 (MedCalc Software) and Stata
227 14 (StataCorp).

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250 **References**

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Online Tables and Figures

Supplemental Table 1: Study specific characteristics in both cohorts

	Derivation cohort (APACE)	Validation cohort (ADAPT and IMPACT)
Study design and study population		
Study design	prospective observational study	prospective observational study (ADAPT) Intervention study (IMPACT)
Study centers	12 centers in 5 European countries (Switzerland, Spain, Italy, Poland, Czech Republic)	1 center in Australia
Inclusion criteria	patients presenting with symptoms suggestive of AMI to an ED, age ≥ 18 years	patients presenting with symptoms suggestive of AMI to an ED, age ≥ 18 years
Informed Consent	Obtained in the emergency department	Obtained in the emergency department or during hospitalization
Chest pain onset	onset or peak within last 12h before presentation	onset or peak within last 12h before presentation
Exclusion criteria	Patients with terminal renal failure on chronic dialysis, no informed consent obtained	Patients with a clear cause other than acute coronary syndrome at presentation, palliative treatment at presentation, pregnant women, transfer from another hospital no informed consent obtained
<p>All 3 studies carried out according to the principles of the Declaration of Helsinki and approved by the local ethics committees.</p> <p>Written informed consent was obtained from all patients.</p>		
Routine clinical assessment	medical history, 34 predefined chest pain characteristics, vital signs, physical examination, 12-lead ECG, continuous ECG rhythm monitoring, pulse oximetry, standard blood tests, chest radiography	medical history, vital signs, physical examination, 12-lead ECG, continuous ECG rhythm monitoring, pulse oximetry, standard blood tests, chest radiography
Routine treatment of patients	at discretion of the attending physician	at discretion of the attending physician
Investigational troponin assays	hs-cTnT Elecsys, hs-cTnI Architect and Hs-cTnI Access	hs-cTnT Elecsys, hs-cTnI Architect and Hs-cTnI Access

Blood sampling	obtained in plasma or EDTA tubes, centrifugated and frozen at -80°C until assayed in a blinded fashion in a dedicated core laboratory.	obtained in plasma or EDTA tubes, centrifugated and frozen at -80°C until assayed in a blinded fashion in a dedicated core laboratory.
Time points of measurement	at presentation and after 1h, 2h, 3h and 6h	at presentation and after 2h and 6h to 12h for ADAPT at presentation and after 2h for low to intermediate risk patients and 6 to 12h for high risk patients in IMPACT
Number of patients with measured hs-cTnT/I (for APACE) and s-cTnI (for ADAPT and Impact) concentration for adjudication	0h: 1131/1131 patients (100%) 1h: 1091/1131 patients (96.5%) 2h: 1129/1131 patients (99.8%) 3-6h: 442/1131 patients (39.1%) Clinically used assays: 0-2h 1115/1132 (98.6%) 3-6h 567/1131 (50.1%)	0h: 1277/1280 (99.8%) 2h: 1275/1280 (99.6%) 6h: 687/1280 (53.7%)
central adjudication of final diagnoses		
Troponin assays used for central adjudication of final diagnoses	hs-cTnT Elecsys Roche Assay, and hs-cTnI Architect Abbott Assay measured at presentation and after 1h to 6h PLUS the clinically used (hs)-cTnT/I assay (hs-cTnT, or s-cTnI-Vista, or cTnI-Axsym, or cTnT) measured at presentation and serially thereafter	cTnI Beckman Coulter 2nd generation AccuTnI assay measured at presentation and after 6h to 12h
Troponin criteria used for central adjudication to qualify for AMI	at least one value above the 99th percentile together with a significant rise and/or fall defined as a rise or fall of at least 10ng/L within six hours, or of 6ng/L within three hours	at least one value above the 99th percentile together with a significant rise and/or fall defined as a relative change of at least 20%
Number of patients who are lost for follow-up	2/1131 patients (0.2%)	No patients were lost to follow-up. However, one-year follow-up only occurred for the 1056/1280 of patients who consented to be contacted at 1 year.

Table Legend: AMI: acute myocardial infarction, ED: emergency department, ECG: electrocardiography, (hs)-cTn: high sensitivity-cardiac troponin

LBBB: left bundle branch block, * = detailed description of investigational assays in the main manuscript

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Section & Topic	No	Item	Reported on page #
TITLE OR ABSTRACT			1,2
	1	Identification as a study of diagnostic accuracy using at least one measure of accuracy (such as sensitivity, specificity, predictive values, or AUC)	3,5,6
ABSTRACT			
	2	Structured summary of study design, methods, results, and conclusions (for specific guidance, see STARD for Abstracts)	2
INTRODUCTION			
	3	Scientific and clinical background, including the intended use and clinical role of the index test	4
	4	Study objectives and hypotheses	4
METHODS			
<i>Study design</i>	5	Whether data collection was planned before the index test and reference standard were performed (prospective study) or after (retrospective study)	5-8, supplement
<i>Participants</i>	6	Eligibility criteria	5-6, supplement
	7	On what basis potentially eligible participants were identified (such as symptoms, results from previous tests, inclusion in registry)	5-6, supplement
	8	Where and when potentially eligible participants were identified (setting, location and dates)	5-6, supplement
	9	Whether participants formed a consecutive, random or convenience series	5-6, supplement
<i>Test methods</i>	10a	Index test, in sufficient detail to allow replication	5-8
	10b	Reference standard, in sufficient detail to allow replication	5-8
	11	Rationale for choosing the reference standard (if alternatives exist)	5-8
	12a	Definition of and rationale for test positivity cut-offs or result categories of the index test, distinguishing pre-specified from exploratory	5-8
	12b	Definition of and rationale for test positivity cut-offs or result categories of the reference standard, distinguishing pre-specified from exploratory	5-8
	13a	Whether clinical information and reference standard results were available to the performers/readers of the index test	5-8
	13b	Whether clinical information and index test results were available to the assessors of the reference standard	5-8
<i>Analysis</i>	14	Methods for estimating or comparing measures of diagnostic accuracy	supplement
	15	How indeterminate index test or reference standard results were handled	supplement
	16	How missing data on the index test and reference standard were handled	supplement
	17	Any analyses of variability in diagnostic accuracy, distinguishing pre-specified from exploratory	supplement
	18	Intended sample size and how it was determined	n/a
RESULTS			
<i>Participants</i>	19	Flow of participants, using a diagram	supplemental Figure 1
	20	Baseline demographic and clinical characteristics of participants	9, Table 1A, 1B
	21a	Distribution of severity of disease in those with the target condition	9-11
	21b	Distribution of alternative diagnoses in those without the target condition	9-11
	22	Time interval and any clinical interventions between index test and reference standard	9-11
<i>Test results</i>	23	Cross tabulation of the index test results (or their distribution) by the results of the reference standard	9-11
	24	Estimates of diagnostic accuracy and their precision (such as 95% confidence intervals)	9-11
	25	Any adverse events from performing the index test or the reference standard	Table 2
DISCUSSION			

	26	Study limitations, including sources of potential bias, statistical uncertainty, and generalisability	14-15
	27	Implications for practice, including the intended use and clinical role of the index test	12-15
OTHER INFORMATION			
	28	Registration number and name of registry	3
	29	Where the full study protocol can be accessed	3
	30	Sources of funding and other support; role of funders	16-18

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Supplemental Table 2	STARD Checklist for Studies of Diagnostic Accuracy
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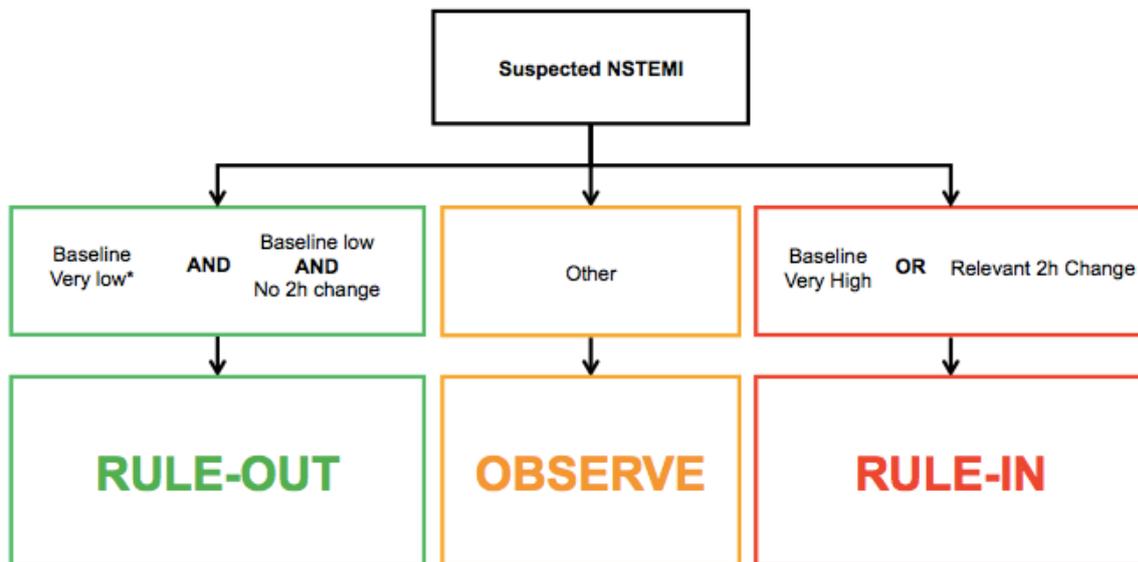
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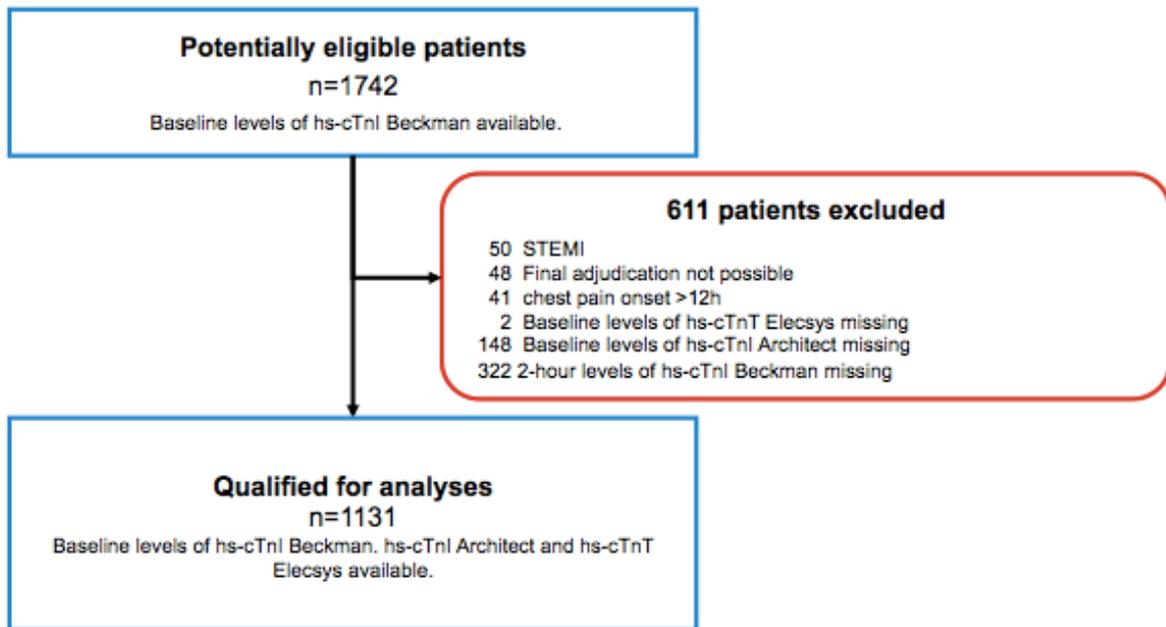
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**Supplemental
Figure 1**

Concept of the hs-cTnI-Access 0/2h-algorithm

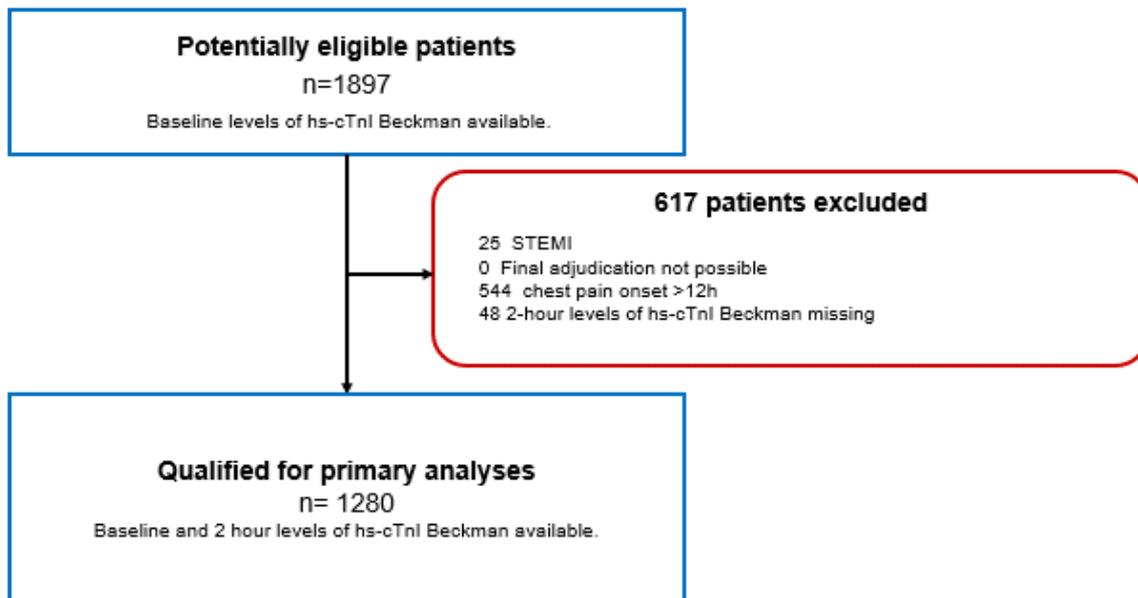
The official concept of the hs-cTn 0/2h-algorithm. Patients are triaged towards rule-out, observe, and rule-in. *if chest pain onset >3h



**Supplemental
Figure 2A**

Patient flow of enrolled patients in the derivation cohort

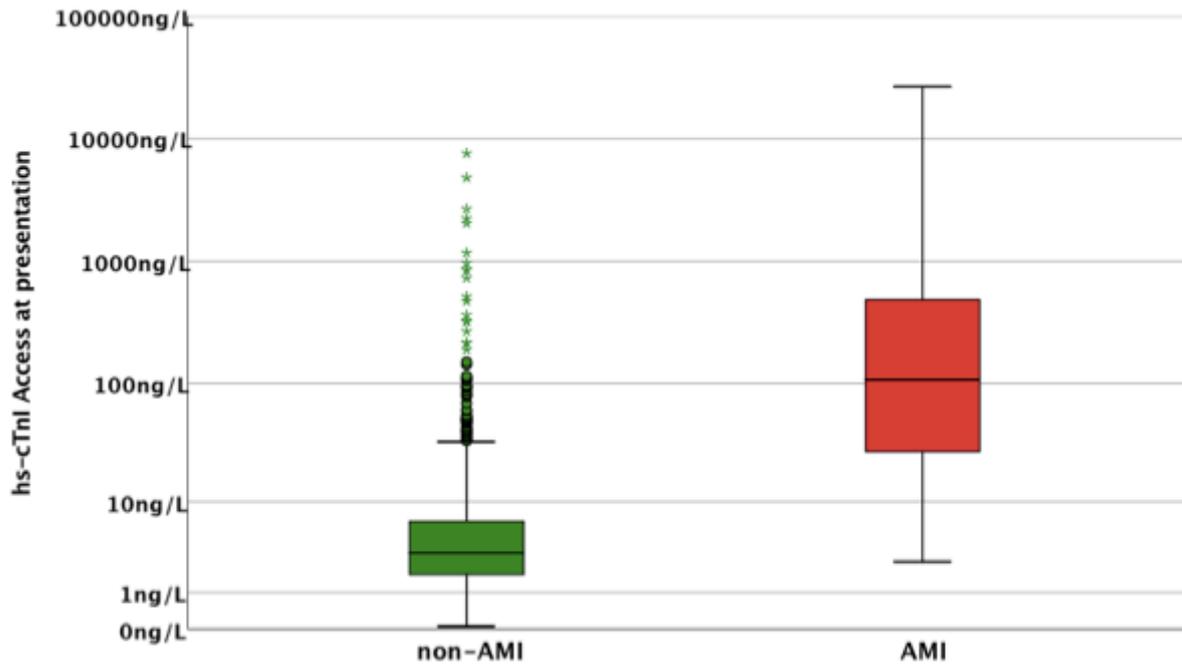
Hs-cTnI denotes high-sensitivity cardiac troponin I; hs-cTnT denotes high-sensitivity cardiac troponin T; STEMI denotes ST-elevation myocardial infarction.



**Supplemental
Figure 2B**

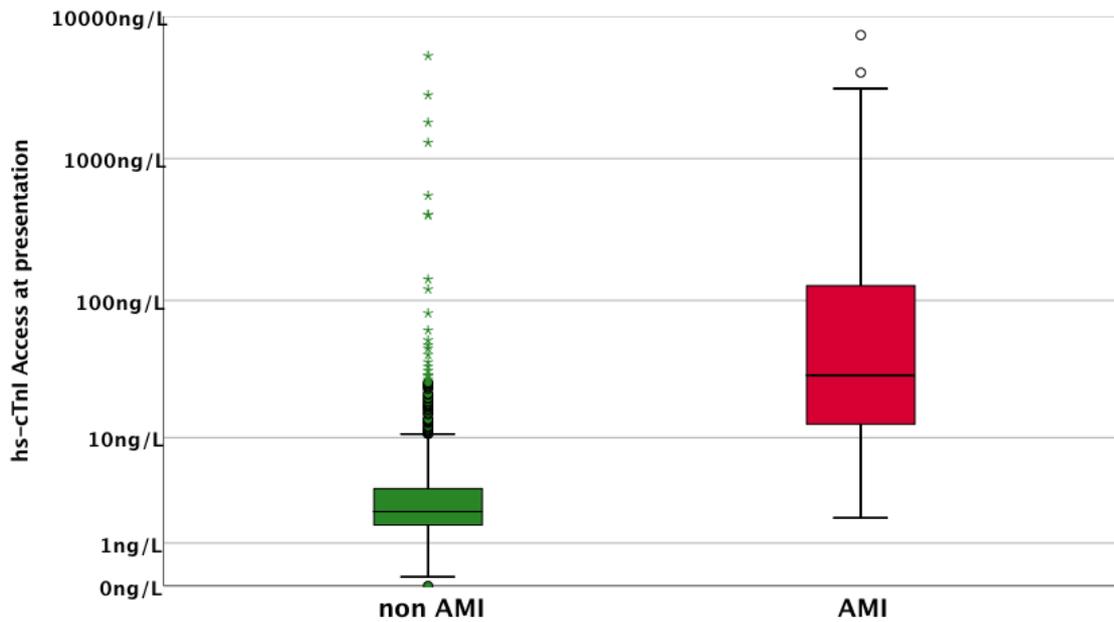
Patient flow of enrolled patients in the validation cohort

Hs-cTnI denotes high-sensitivity cardiac troponin I; hs-cTnT denotes high-sensitivity cardiac troponin T; STEMI denotes ST-elevation myocardial infarction.



Supplemental Figure 3A Boxplots showing Concentrations of hs-cTnI-Access at Presentation according to the final diagnosis the derivation cohort

Boxes represent medians and interquartile ranges (IQRs), while whiskers display the smallest and the largest non-outliers. Rings display outliers further than 1.5 IQRs and stars display outliers further than 3 IQRs from the respective end of the box. AMI denotes acute myocardial infarction; hs-cTnI denotes high-sensitivity cardiac troponin I.

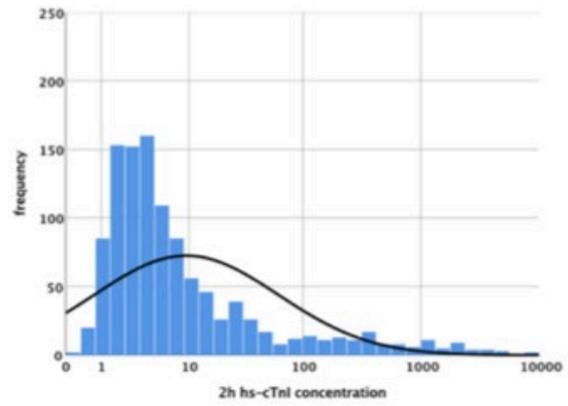
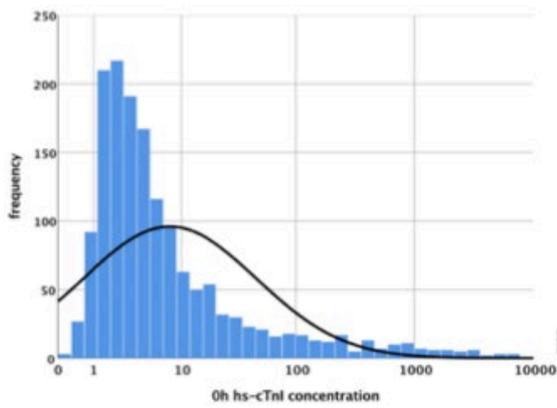


Supplemental Figure 3B Boxplots showing Concentrations of hs-cTnI-Access at Presentation according to the final diagnosis the validation cohort

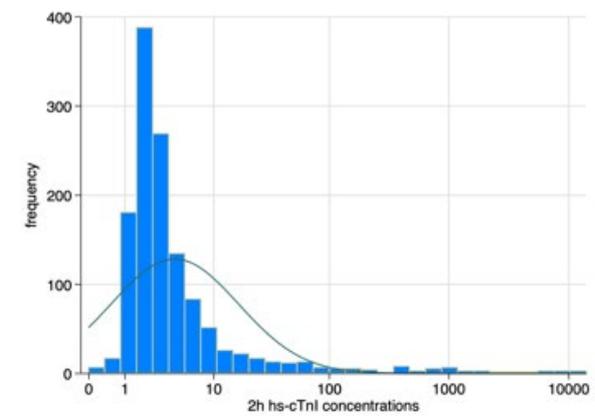
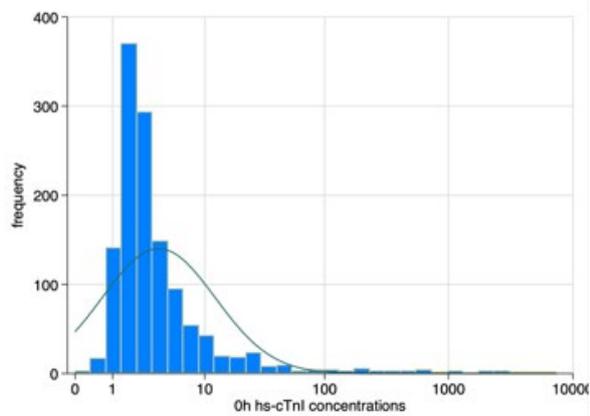
Boxes represent medians and interquartile ranges (IQRs), while whiskers display the smallest and the largest non-outliers. Rings display outliers further than 1.5 IQRs and stars display outliers further than 3 IQRs from the respective end of the box. AMI denotes acute myocardial infarction; hs-cTnI denotes high-sensitivity cardiac troponin

|

Derivation cohort

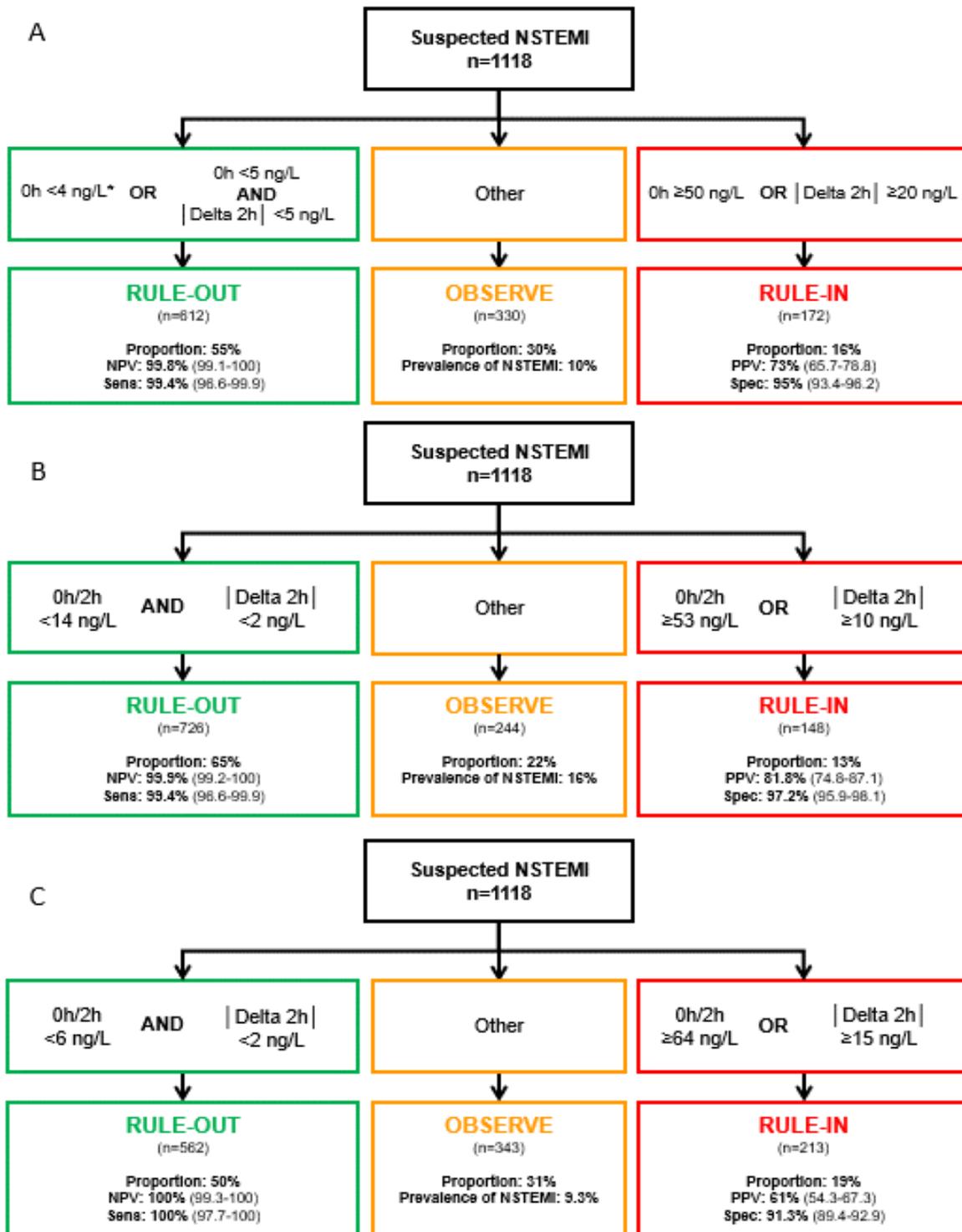


Validation cohorts



Supplemental Figure 4 Histograms

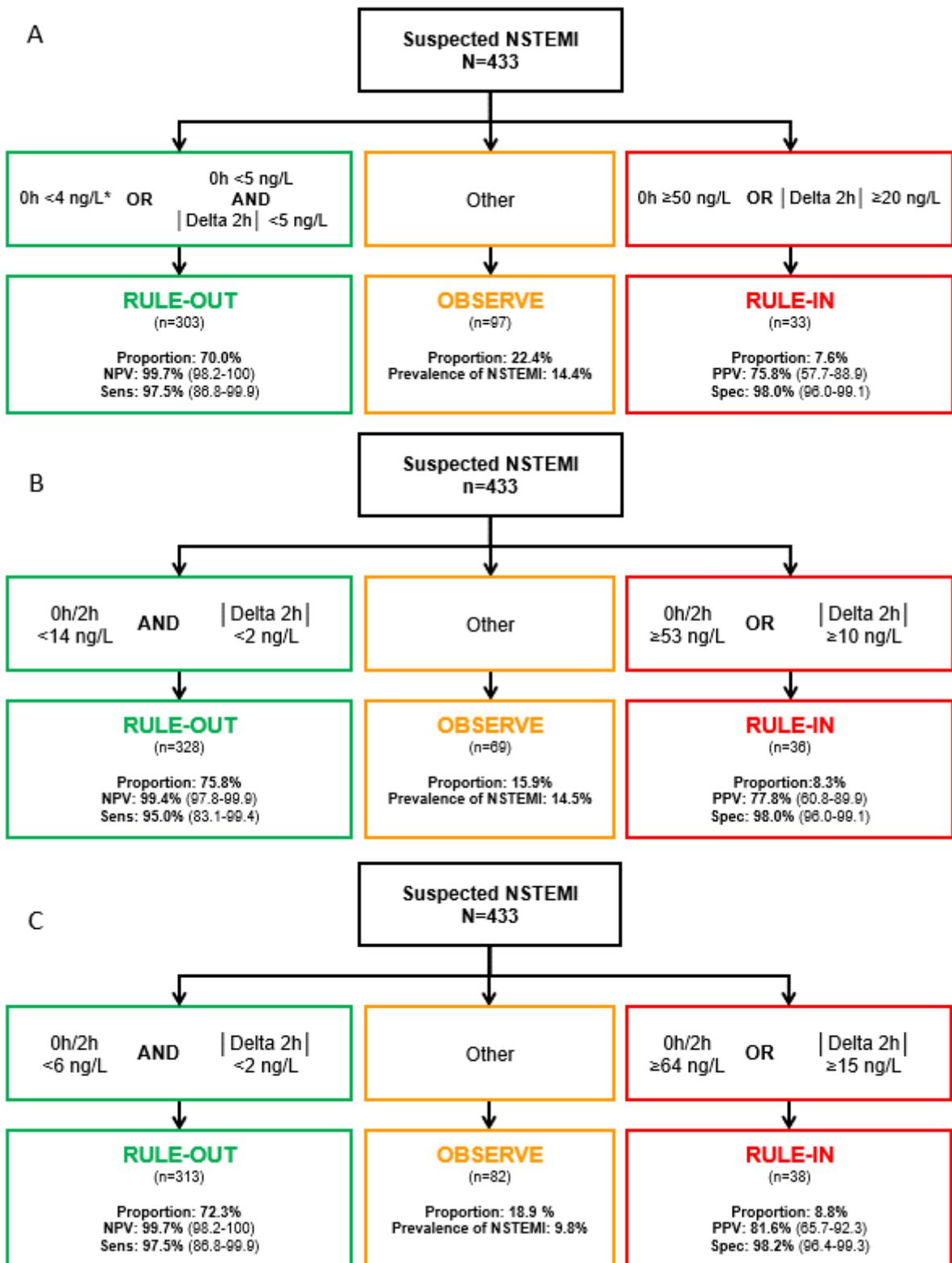
Histograms highlighting the distribution of hs-cTnI concentrations at presentation and after 2h in the derivation and the validations cohorts.



**Supplemental
Figure 5A**

Direct Comparison of the hs-cTnI-Access 0/2h-algorithm with the hs-cTnT-Elecsys and hs-cTnI-Architect 0/2h-algorithms in the derivation cohort

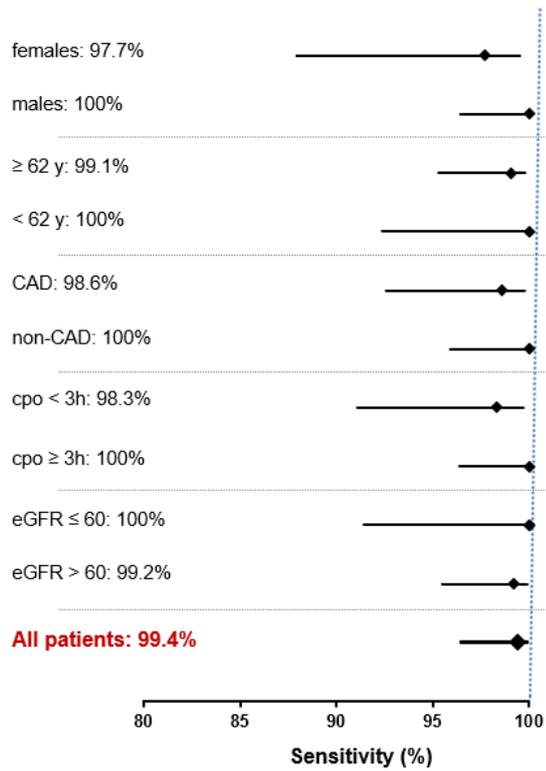
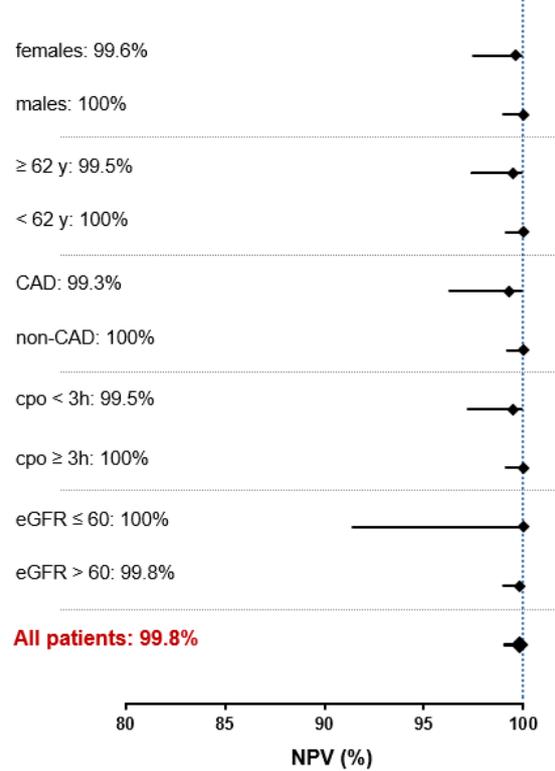
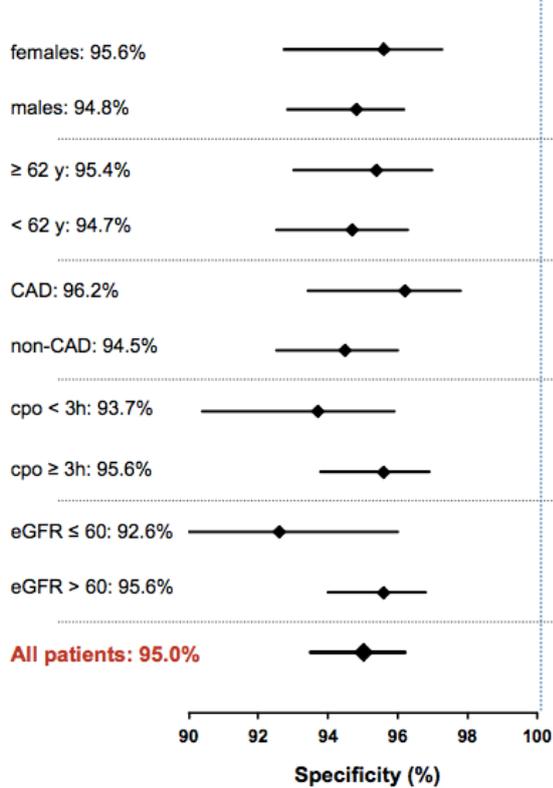
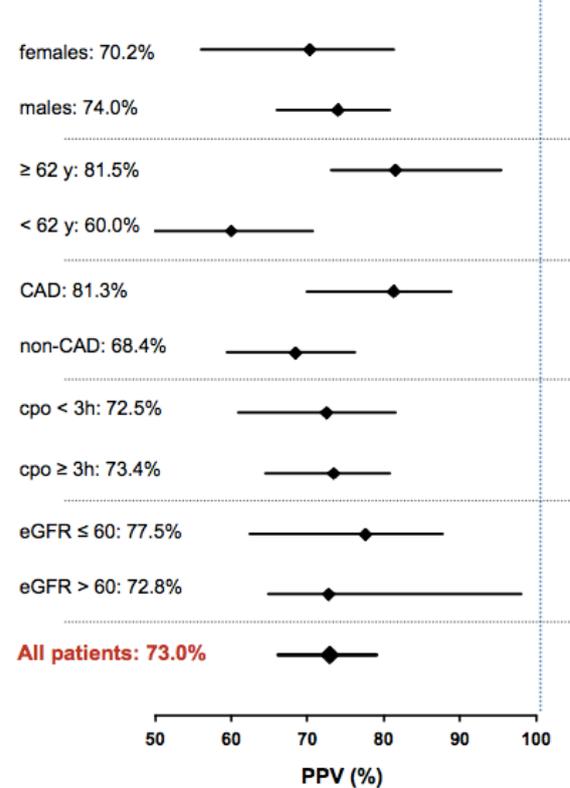
Direct comparison of the diagnostic performance of (A) the hs-cTnI-Access 0/2h-algorithm with (B) the hs-cTnT-Elecsys 0/2h-algorithm and (C) the hs-cTnI-Architect 0/2h-algorithm in the derivation cohort. NSTEMI denotes non-ST-elevation myocardial infarction; 0h/2h denotes based on zero-hour or two-hour blood sample obtained at presentation to the ED and after two-hours; $\Delta 2h$ | denotes absolute (unsigned) change of high-sensitivity cardiac troponin I within two hours; NPV denotes negative predictive value; Sens. denotes sensitivity; PPV denotes positive predictive value; Spec. denotes specificity; * denotes chest pain onset <2 hours



**Supplemental
Figure 5B**

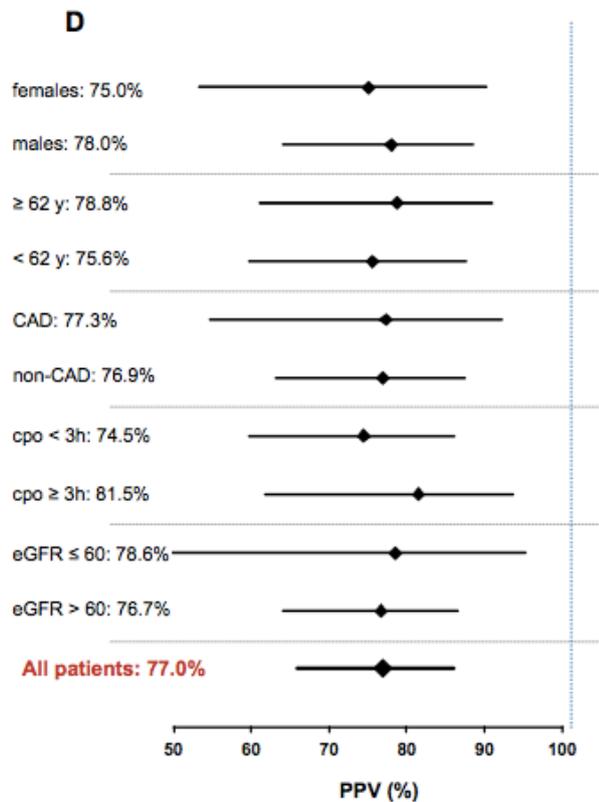
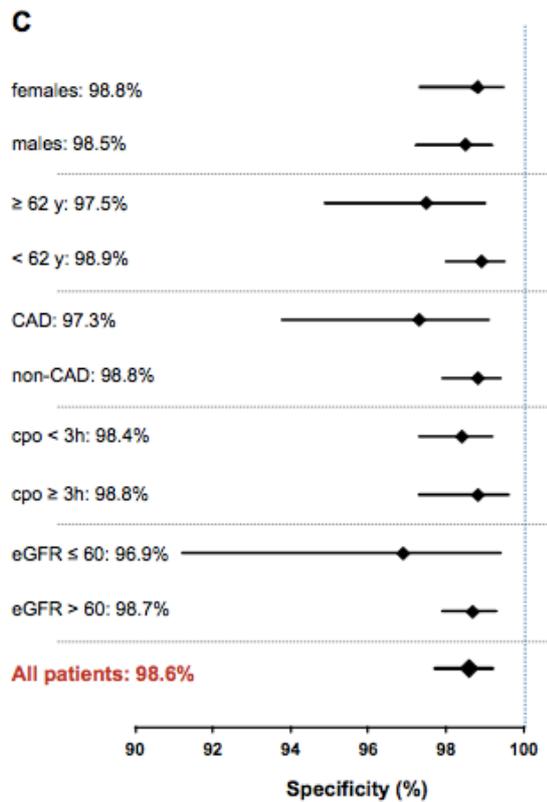
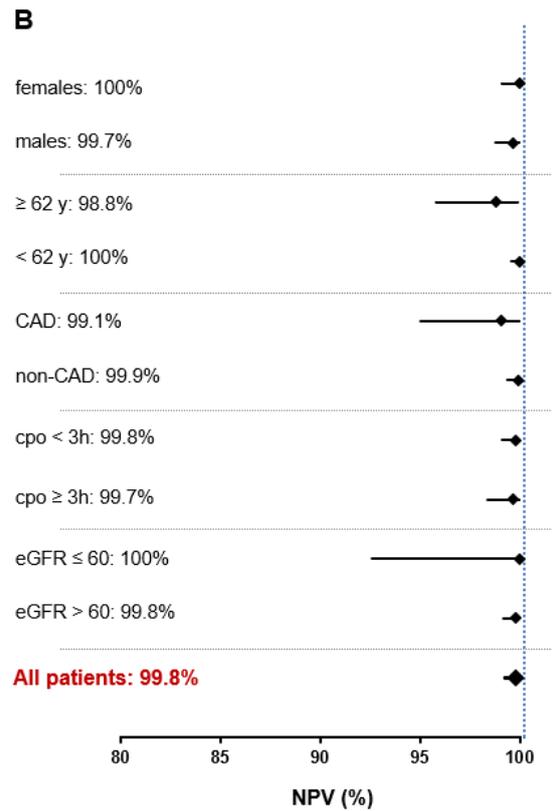
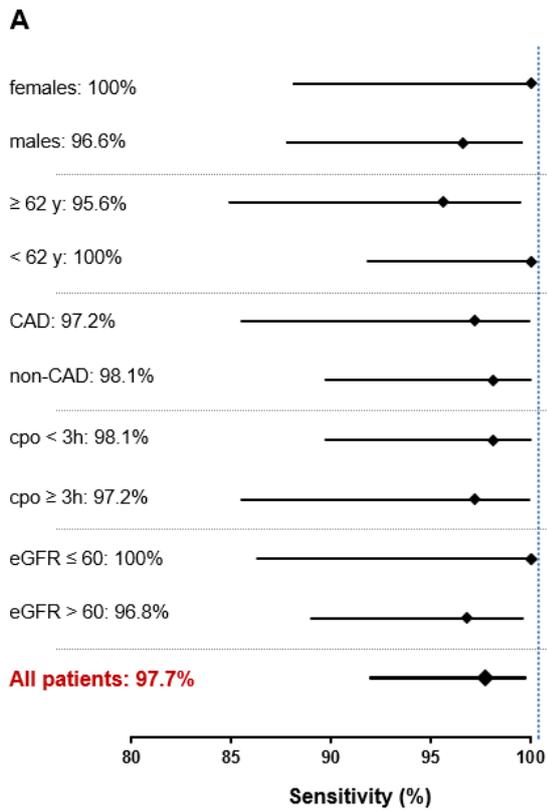
Direct Comparison of the hs-cTnI-Access 0/2h-algorithm with the hs-cTnT-Elecsys and hs-cTnI-Architect 0/2h-algorithms in the validation cohort

Direct comparison of the diagnostic performance of (A) the hs-cTnI-Access 0/2h-algorithm with (B) the hs-cTnT-Elecsys 0/2h-algorithm and (C) the hs-cTnI-Architect 0/2h-algorithm in the derivation cohort. NSTEMI denotes non-ST-elevation myocardial infarction; 0h/2h denotes based on zero-hour or two-hour blood sample obtained at presentation to the ED and after two-hours; $\Delta 2h$ | denotes absolute (unsigned) change of high-sensitivity cardiac troponin I within two hours; NPV denotes negative predictive value; Sens. denotes sensitivity; PPV denotes positive predictive value; Spec. denotes specificity; * denotes chest pain onset <2 hours

A**B****C****D**

Supplemental Figure 6A Forest plots for algorithm performance in predefined subgroup analyses in the derivation cohort.

Forest plots indicating sensitivity (A), negative predictive value (NPV, B), specificity (C) and positive predictive value (PPV, D) for predefined subgroup analyses. CAD indicates coronary artery disease; cpo, chest pain onset; and eGFR, estimated glomerular filtration rate.



Supplemental Figure 6B Forest plots for algorithm performance in predefined subgroup analyses in the validation cohort.

Forest plots indicating sensitivity (A), negative predictive value (NPV, B), specificity (C) and positive predictive value (PPV, D) for predefined subgroup analyses. CAD indicates coronary artery disease; cpo, chest pain onset; and eGFR, estimated glomerular filtration rate.