

Clinical Infectious Diseases

Development of a Risk Prediction Model for Carbapenem-Resistant Enterobacteriaceae Infection after Liver Transplantation: A Multinational Cohort Study --Manuscript Draft--

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| Manuscript Number: | CID-105646R1 |
| Full Title: | Development of a Risk Prediction Model for Carbapenem-Resistant Enterobacteriaceae Infection after Liver Transplantation: A Multinational Cohort Study |
| Short Title: | CRE infection after LT in carriers |
| Article Type: | Major Article |
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| Manuscript Region of Origin: | ITALY |
| Abstract: | <p>Background. Patients colonized with carbapenem resistant Enterobacteriaceae (CRE) are at higher risk of developing CRE infection after liver transplantation (LT) with associated high morbidity and mortality. Prediction model for CRE infection after LT among carriers could be useful to target preventive strategies.</p> <p>Methods. Multinational multicenter cohort study of consecutive adult patients underwent LT and colonized with CRE before or after LT, from January 2010 to December 2017. Risk factors for CRE infection were analyzed by univariate analysis and by Fine-Gray sub-distribution hazard model, with death as competing event. A nomogram to predict 30- and 60-day CRE infection risk was created.</p> <p>Results. 840 LT recipients found to be colonized with CRE before (n=203) or after (n=637) LT were enrolled. CRE infection was diagnosed in 250 (29.7%) patients within 19 (IQR 9-42) days after LT. Pre-and post-LT colonization, multisite post-LT colonization, prolonged mechanical ventilation, acute renal injury, and surgical re-intervention were retained in the prediction model. Median 30 and 60-day predicted risk was 15% (IQR 11-24%) and 21% (IQR 15-33%), respectively. Discrimination and prediction accuracy for CRE infection was acceptable on derivation (AUC 74.6, Brier index 16.3) and bootstrapped validation dataset (AUC 73.9, Brier index 16.6). Decision-curve analysis suggested net benefit of model-directed intervention over default strategies (treat all, treat none) when CRE infection probability exceeded 10%. The risk prediction model is freely available as mobile application at https://idbologna.shinyapps.io/CREPostOLTPredictionModel/ .</p> <p>Conclusions. Our clinical prediction tool could enable better targeting interventions for CRE infection after transplant.</p> |
| Response to Reviewers: | <p>Ajit P. Limaye, M.D., Special Section Editor Clinical Infectious Diseases email: cid.editorialoffice@idsociety.org</p> <p>Bologna, December 2020</p> <p>Dear Dr. Limaye:</p> <p>Many thanks for your comments on our manuscript CID-105646 entitled "Development of a Risk Prediction Model for Carbapenem-Resistant Enterobacteriaceae Infection after Liver Transplantation: A Multinational Cohort Study". We have carefully considered every comment and suggestion raised by the reviewers and changed our manuscript accordingly. We attach the answers to all the items raised and two copies of the revised manuscript, one of them with all the changes clearly marked in yellow.</p> <p>REVIEWER COMMENTS</p> <p>Reviewer #1: Giannella and colleagues present a multinational cohort study from which a risk prediction model for post-transplant CRE infection was developed. This is a topic</p> |

of significant importance given the morbidity and mortality associated with CRE infections in liver and other solid organ transplant recipients. Further, while CRE colonization has been associated with CRE infections following liver transplantation, there is a paucity of data which helps stratify the risk for infection among those colonized patients. Overall, this is a high-yield manuscript of interest to the transplant community. The authors do highlight limitations of the study and certainly future studies would be required to confirm these findings.

AUTHORS' RESPONSE: Thanks for the kind comment.

One minor point would be the following: All patients were screened for rectal CRE carriage at pre-defined time points. However, screening for CRE colonization at other sites was not uniformly performed. While it would make sense that patients with multi-site colonization are at higher risk for invasive infection, it would be important to highlight how this may impact the results.

AUTHORS' RESPONSE: We agree with reviewer, we have commented this limitation in the discussion (Page 13, line 318-322).

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Major comments:

1. There were a number of notable between-group differences for those colonized pre-transplant as opposed to post-transplant, including higher rates of moderate-severe renal disease, more frequent prior transplantation, higher MELD scores, higher rates of hepato-renal syndrome, higher rates of grade III/refractory ascites, more frequent prior GI bleeding, higher rates of hepatic encephalopathy, more frequent preceding ICU admission, more frequent acute-on-chronic liver failure, more frequent occurrence of any infection, more frequent choledochojejunostomy, higher rates of prolonged surgery, or more frequent use of maintenance steroids in the pre-transplant CRE colonization groups. Despite that, the observed CRE infection rate was similar overall between pre- and post-liver transplant colonized patients. One would predict an increased CRE infection rate in the pre-transplant colonized group. The lack of observed difference in CRE rates may be explained by the observation that 25.1% of patients colonized pre-transplant with CRE received targeted perioperative prophylaxis. Thus, it would be of interest to know if the rate of CRE infection differed between those who did and did not receive CRE-targeted prophylaxis. Despite this, I believe that most transplant centers would target antimicrobial prophylaxis to cover CRE in known colonized patients undergoing liver transplantation.

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Regarding the last sentence. We agree with the need/opportunity to intervene at transplantation with targeted prophylaxis in patients known to be colonized with CRE. However, the effectiveness of targeted prophylaxis and which agents should be used are not well established. Even in pre-OLT carriers, the colonization data is often only acquired in the immediate post-operative period. Furthermore, most patients are found to be colonized with CRE during the first weeks after OLT. Finally, as shown in previous studies including unselected OLT recipients, both colonization at OLT and

colonization after OLT were the strongest predictors of CRE infection (Freire et al Transplantation 2017, Giannella et al Clin Microbiol Infect 2019). For these reasons, we believe that an individualized risk stratification for colonized patients, either pre or post OLT, could help better targeting preventive strategies not only in the perioperative but also in the post-operative period.

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- The inclusion of KPC-producing isolates in the multivariate model is also a bit unusual given that there was no statistically significant difference between the group not developing CRE infection and those developing CRE infection (70.8% vs. 72.8%, $p=0.61$) in the presence of KPC being the identified genotypic strain of CRE present. Regardless, including colonization in general, multisite colonization, and then colonization with a KPC-producing isolate seems run the risk of counting a similar variable repeatedly and over-inflating an individual's risk of CRE infection.

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- With all the previous considerations, I would like to suggest a new predictive model with additional justification of the reasons to exclude some variables:

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- Reintervention should be included and will reflect a more complicated course of liver transplantation. Re-laparotomy may facilitate CRE infection in colonized patients. Also, could predict the length of ICU stay.

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AUTHOR RESPONSE: We agree with this comment, unfortunately we collected such variable only as binary variable asking for the need of prolonged (>48 h) MV before the development of CRE infection. We have addressed this limitation in the discussion (Page 14, line 325-327).

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and should not be limited in time.

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•Regarding the transplant timeline of CRE colonization, without considering multisite colonization as this variable can be biased by more intensive screening in ICU patients, it looks like pre-transplant CRE colonization has more impact on CRE infections than colonization post-transplant. Therefore, I would suggest including only pre-transplant colonization by CRE.

AUTHORS' RESPONSE: We have appreciated this comment; eventually we have leaved in the model pre- and post-transplant colonization and multisite colonization after LT for the reasons explained below.

•Would it be possible to evaluate the robustness and predictive power of a model including these variables: Reoperation, mechanical ventilation for more than 48 h in the preceding 2 weeks, acute kidney injury and pre-transplant CRE colonization? If this proposed model predicts well the occurrence of CRE infection, then would potentially be applicable to other transplant programs with high prevalence of CRE. In addition, it should be clearly stated that the model will only work in patients with proven CRE colonization. I suspect the selection of some variables may have decreased the predictive power of the model.

AUTHOR'S RESPONSE: We have re-performed the analysis following the suggestions of the reviewer. The table below shows indices of the model discrimination (Harrell's C statistic), accuracy (Brier Index) as well as improvements in the prediction accuracy versus a null model that contains only CRE colonization prior to transplant. As seen below, the more parsimonious model proposed by the reviewer displayed poorer discrimination and accuracy for identifying patients who develop CRE infection within 60 days post-transplant.

ModelC-Statistic (AUC)

(higher number is better)Brier decomposition score

(lower scores are better)Improvements in prediction accuracy vs. null model -CRE col pre 60d

(higher percentages are better)

Full model:

- 1.ATG
- 2.Reintervention
- 3.Mechanical ventilation
- 4.Acute renal failure
- 5.KPC carbapenem.
- 6.CRE col pre 60d
- 7.Multisite colon. Pre trans
- 8.CRE col post 60d
- 9.Multisite colon post trans

75.1
(71.3-78.9)16.1

(6.6-25.6)14.8%

Reviewer suggested model:

- 1.Reintervention
- 2.Mechanical ventilation
- 3.Acute renal failure
- 4.CRE col pre 60d

69.2
[65.2-73.2)17.4

(7.6-27.2)8.21%

However, we agree with the reviewer that a simpler model may be possible. Therefore, using a base model of CRE colonization within 60 days prior to transplant, we examined the degree to which including each variable from our original model improved model performance. Our analysis, in part, supported the reviewer's suggestion that ATG, KPC carbapenemase, and multisite colonization prior to transplantation could be removed from model without marked changes in prediction accuracy.

Addition of predictor to base model (CRE col pre 60d)C-Statistic, AUC

(95%)Improvements in prediction accuracy vs. null model

(CRE col pre 60d) (higher percentages are better)

ATG53.8 (50.5-57.2)0.71%

Reintervention54.5 (51.0-58.0)2.63%

Mechanical ventilation59.7 (55.5-63.8)7.45%

Acute renal failure 61.7 (57.5-65.8) 3.63%
 KPC carbapenemase 53.9 (49.7-58.1) 0.72%
 Multisite col. pre trans 53.9 (50.5-57.2) 0.62%
 CRE colonisation post-transplant 57.6 (54.1-61.0) 1.71%
 Multisite colon post-transplant 62.6 (58.7-66.5) 6.85%

However, colonization documented post-transplant and colonization at additional sites (an indicator of higher inoculum even though not systematically assessed in all patients) were both associated with improved model performance. Therefore, we have propose a revised model that addresses the reviewer's suggestion as best as possible: Table 3. Final multivariable analysis for predicting risk of CRE infection after liver transplantation

| Variable | Subhazard ratio (95% CI) | β -coefficient | P value |
|--|--------------------------|----------------------|---------|
| Reintervention | 1.37 (1.04-1.80) | 0.30 (0.02-0.57) | 0.02 |
| Prolonged mechanical ventilation | 2.01 (1.49-2.71) | 0.70 (0.40-0.99) | <0.0001 |
| Acute renal injury | 1.69 (1.26-2.27) | 0.52 (0.23-0.82) | 0.001 |
| CRE colonisation 60 days prior to transplant | 2.17 (1.33-3.53) | 0.78 (0.29-1.26) | 0.002 |
| CRE colonization within 60 days after transplant | 1.41 (0.91-2.19) | 0.35 (-0.09-0.78) | 0.12 |
| Multisite colonization within 60 days after transplant | 2.87 (1.98-4.20) | 1.06 (0.68-1.43) | <0.0001 |

Abbreviations: CI confidence interval, CRE carbapenem-resistant Enterobacteriaceae, KPC Klebsiella pneumoniae carbapenemase-producing.

The discrimination and calibration of this revised simpler model (6 variables) is nearly identical the originally proposed model (9 variables) for predicting CRE infection within 60 days post-transplant as shown in the figure below of the model fit for the (a) derivation dataset; and (b) Bootstrapped-resampled dataset.

Therefore, we have revised all of the corresponding tables, figures and nomogram (in the manuscript and online app) to reflect the new simpler model.

Minor comments:

- Anti-thymoglobulin is used throughout and perhaps should be changed to anti-thymocyte globulin.

AUTHORS' RESPONSE: We have done as suggested.

- BLBLI acronym on page 5 should be spelled out on first use.

AUTHORS' RESPONSE: We have added the spelling.

- Line 16 on page 7 should perhaps replace "revise" with "review".

AUTHORS' RESPONSE: We have done as suggested.

- Page 10, last paragraph: "Considering the competing risks of death or CRE infection..." CRE infection should be removed as it is not acting as a competing event when evaluating the risk of CRE infection.

AUTHORS' RESPONSE: We have done as suggested.

- Figure 1 on page 24 has "participating" misspelled.

AUTHORS' RESPONSE: We corrected the error.

Should you or any of the reviewers wish any further clarification, please, do not hesitate to contact us. We hope that our manuscript may be now found suitable for publication in your prestigious journal.

Looking forward to hearing from you

Best regards,

Maddalena Giannella

Editor-in-Chief of Clinical Infectious Diseases

December, 2020

Dear Editor,

We have electronically re-submitted the revised manuscript CID-105646 entitled: “Development of a Risk Prediction Model for Carbapenem-Resistant Enterobacteriaceae Infection after Liver Transplantation: A Multinational Cohort Study”. We have carefully considered every comment and suggestion raised by the reviewers and changed our manuscript accordingly.

We attach a response letter with the answers to all the items raised by the reviewers and two copies of the revised manuscript, one of them with all the changes clearly marked in yellow.

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Looking forward to hearing from you

Best regards,

The corresponding authors

Maddalena Giannella, Matteo Rinaldi

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Ajit P. Limaye, M.D., Special Section Editor
Clinical Infectious Diseases
email: cid.editorialoffice@idsociety.org

Bologna, December 2020

Dear Dr. Limaye:

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| Model | C-Statistic (AUC) (higher number is better) | Brier decomposition score (lower scores are better) | Improvements in prediction accuracy vs. null model -CRE col pre 60d) (higher percentages are better) |
|---|---|---|--|
| Full model: 1. ATG 2. Reintervention 3. Mechanical ventilation 4. Acute renal failure 5. KPC carbapenem. 6. CRE col pre 60d 7. Multisite colon. Pre trans 8. CRE col post 60d 9. Multisite colon post trans | 75.1 (71.3-78.9) | 16.1 (6.6-25.6) | 14.8% |
| Reviewer suggested model: 1. Reintervention 2. Mechanical ventilation 3. Acute renal failure 4. CRE col pre 60d | 69.2 (65.2-73.2) | 17.4 (7.6-27.2) | 8.21% |

However, we agree with the reviewer that a simpler model may be possible. Therefore, using a base model of CRE colonization within 60 days prior to transplant, we examined the degree to which including each variable from our original model improved model performance. Our analysis, in part, supported the reviewer's suggestion that ATG, KPC carbapenemase, and multisite colonization prior to transplantation could be removed from model without marked changes in prediction accuracy.

| Addition of predictor to base model (CRE col pre 60d) | C-Statistic, AUC (95%) | Improvements in prediction accuracy vs. null model (CRE col pre 60d) (higher percentages are better) |
|---|------------------------|--|
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| CRE colonisation post-transplant | 57.6 (54.1-61.0) | 1.71% |
| Multisite colon post-transplant | 62.6 (58.7-66.5) | 6.85% |

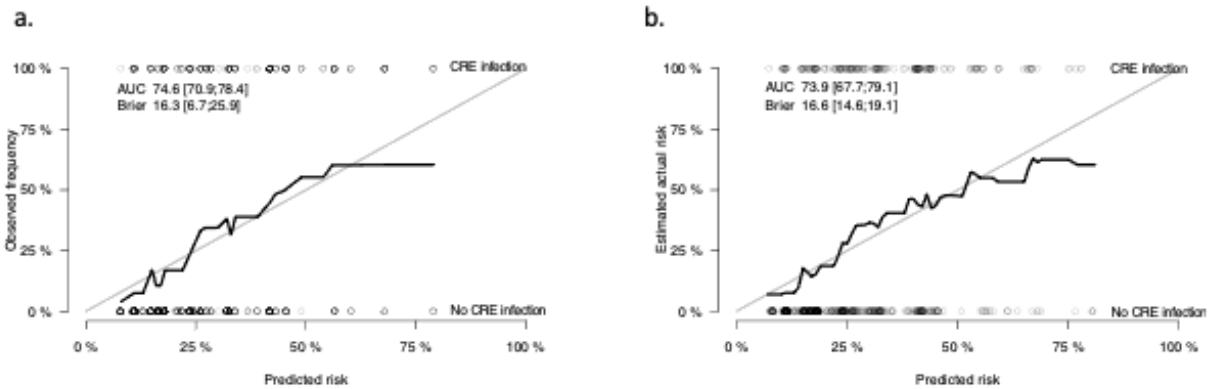
However, colonization documented post-transplant and colonization at additional sites (an indicator of higher inoculum even though not systematically assessed in all patients) were both associated with improved model performance. Therefore, we have propose a revised model that addresses the reviewer’s suggestion as best as possible:

Table 3. Final multivariable analysis for predicting risk of CRE infection after liver transplantation

| Variable | Subhazard ratio (95% CI) | β -coefficient | P value |
|--|--------------------------|----------------------|---------|
| Reintervention | 1.37 (1.04-1.80) | 0.30 (0.02-0.57) | 0.02 |
| Prolonged mechanical ventilation | 2.01 (1.49-2.71) | 0.70 (0.40-0.99) | <0.0001 |
| Acute renal injury | 1.69 (1.26-2.27) | 0.52 (0.23-0.82) | 0.001 |
| CRE colonisation 60 days prior to transplant | 2.17 (1.33-3.53) | 0.78 (0.29-1.26) | 0.002 |
| CRE colonization within 60 days after transplant | 1.41 (0.91-2.19) | 0.35 (-0.09-0.78) | 0.12 |
| Multisite colonization within 60 days after transplant | 2.87 (1.98-4.20) | 1.06 (0.68-1.43) | <0.0001 |

Abbreviations: CI confidence interval, CRE carbapenem-resistant Enterobacteriaceae, KPC Klebsiella pneumoniae carbapenemase-producing.

The discrimination and calibration of this revised simpler model (6 variables) is nearly identical the originally proposed model (9 variables) for predicting CRE infection within 60 days post-transplant as shown in the figure below of the model fit for the (a) derivation dataset; and (b) Bootstrapped-resampled dataset.



Therefore, we have revised all of the corresponding tables, figures and nomogram (in the manuscript and online app) to reflect the new simpler model.

Minor comments:

- **Anti-thymoglobulin is used throughout and perhaps should be changed to anti-thymocyte globulin.**

AUTHORS' RESPONSE: We have done as suggested.

- **BLBLI acronym on page 5 should be spelled out on first use.**

AUTHORS' RESPONSE: We have added the spelling.

- **Line 16 on page 7 should perhaps replace "revise" with "review".**

AUTHORS' RESPONSE: We have done as suggested.

- **Page 10, last paragraph: "Considering the competing risks of death or CRE infection..." CRE infection should be removed as it is not acting as a competing event when evaluating the risk of CRE infection.**

AUTHORS' RESPONSE: We have done as suggested.

- **Figure 1 on page 24 has "participating" misspelled.**

AUTHORS' RESPONSE: We corrected the error.

Should you or any of the reviewers wish any further clarification, please, do not hesitate to contact us. We hope that our manuscript may be now found suitable for publication in your prestigious journal.

Looking forward to hearing from you

Best regards,

Maddalena Giannella

1 CID-105646-unmarked

2 **Development of a Risk Prediction Model for Carbapenem-Resistant Enterobacteriaceae Infection**
3 **after Liver Transplantation: A Multinational Cohort Study**

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84 **Text word count:** 2996 **Abstract word count:** 249

85 **Running title:** CRE infection risk in liver transplant

86 **Key words:** SOT, liver transplantation, CRE carriage, CRE infection

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99 **40-word summary:** One third of CRE carriers develops infection after LT with dramatic impact on
100 survival. Our prediction tool could enable better targeting of early interventions for CRE infection
101 after LT as opposed to universal prophylaxis or treatment in high-prevalence centers.

102 **ABSTRACT**

103 **Background.** Patients colonized with carbapenem resistant Enterobacteriaceae (CRE) are at higher
104 risk of developing CRE infection after liver transplantation (LT) with associated high morbidity and
105 mortality. Prediction model for CRE infection after LT among carriers could be useful to target
106 preventive strategies.

107 **Methods.** Multinational multicenter cohort study of consecutive adult patients underwent LT and
108 colonized with CRE before or after LT, from January 2010 to December 2017. Risk factors for CRE
109 infection were analyzed by univariate analysis and by Fine-Gray sub-distribution hazard model, with
110 death as competing event. A nomogram to predict 30- and 60-day CRE infection risk was created.

111 **Results.** 840 LT recipients found to be colonized with CRE before (n=203) or after (n=637) LT were
112 enrolled. CRE infection was diagnosed in 250 (29.7%) patients within 19 (IQR 9-42) days after LT. Pre-
113 and post-LT colonization, multisite post-LT colonization, prolonged mechanical ventilation, acute
114 renal injury, and surgical re-intervention were retained in the prediction model. Median 30 and 60-
115 day predicted risk was 15% (IQR 11-24%) and 21% (IQR 15-33%), respectively. Discrimination and
116 prediction accuracy for CRE infection was acceptable on derivation (AUC 74.6, Brier index 16.3) and
117 bootstrapped validation dataset (AUC 73.9, Brier index 16.6). Decision-curve analysis suggested net
118 benefit of model-directed intervention over default strategies (treat all, treat none) when CRE
119 infection probability exceeded 10%. The risk prediction model is freely available as mobile
120 application at <https://idbologna.shinyapps.io/CREPostOLTPredictionModel/>.

121 **Conclusions.** Our clinical prediction tool could enable better targeting interventions for CRE infection
122 after transplant.

123

124 **INTRODUCTION**

125 The emergence and spread of carbapenem resistant *Enterobacteriaceae* (CRE) over the last two
126 decades threatens the safety of patients undergoing life-saving surgical procedures or necessary
127 immunosuppressive treatments [1]. Solid organ transplant (SOT) recipients, in particular those
128 undergoing liver transplantation (LT), are at increased risk of colonization with CRE due to repeated
129 contacts with health care system, exposure to broad spectrum antibiotics, complexity of surgery,
130 and altered immune status [2]. Compared with other settings (e.g., geriatric units, long-term care
131 facilities) the incidence of infection per CRE colonization days is higher in SOT recipients [3], and
132 associated with poorer survival [4]. In a prospective multicentre study of 887 SOT recipients in Italy,
133 mortality was more than 10 times higher in patients with positive cultures for CRE [5].

134 Knowledge of CRE infection risk factors is essential to develop targeted preventive strategies. Risk
135 factors for CRE infection after LT were analysed in two large single-centre cohort studies, from Brazil
136 and Italy, including 386 and 553 patients, respectively [6, 7]. In both studies CRE carriage, either
137 detected before or after transplant, was found to be the strongest predictor of subsequent CRE
138 infection. Based on these findings, some investigators have proposed that preventive strategies
139 should be implemented in all CRE carriers undergoing LT [8]. However, a more accurate prediction of
140 which carriers are at higher risk could help to target preventive measures or timely appropriate
141 antibiotic therapy in patients at lower risk (i.e. < 10%) of CRE infection.

142 With this objective in mind, we performed a multicentre, multinational cohort study of patients
143 undergoing LT colonized with CRE before or after transplantation, with the aim of developing a
144 multivariable prediction model for CRE infection within 30 and 60 days of LT.

145

146 **MATERIAL AND METHODS**

147 *Study design*

148 We performed a multicentre multinational retrospective cohort study. The enrolment period was
149 between January 2010 to December 2017, or from the onset of systematic CRE colonization
150 screening in the centre to December 2017 (see supplementary Table 1). Follow-up was of 180 days
151 after LT. Data were accrued from October 2019 to June 2020. Data sources were clinical charts and
152 hospital electronic records, they were de-identified before entry into a standardized electronic case
153 report form (eCRF), and managed using REDCap capture tool hosted by Alma Mater University of
154 Bologna [9]. Collected data were periodically checked for accuracy by an investigator of the
155 coordinating centre (MR). Queries for incongruous or missing data were submitted to investigators
156 to ensure high quality and completeness. The study was first approved by Institutional Review Board
157 (IRB) of the promoting centre (n. 155/2019/Oss/AOUBo on March 20, 2019), then by IRB of all
158 participating centres.

159 *Setting*

160 Fifteen hospitals performing LT participated in the study: seven from Italy (Bologna, Modena, Turin,
161 Padua, Palermo, Milan, and Udine); four from Brazil (2 in São Paulo, 1 in Fortaleza and 1 in Rio de
162 Janeiro); two from Spain (Madrid and Majadahonda); one from United States (Miami); and one from
163 Israel (Petah-Tikva) (see Supplementary Table 1 and Figure 1). An active surveillance screening for
164 CRE colonization was required by the study protocol. All centres performed systematic screening of
165 CRE carriage by rectal swab (RS), at inclusion in waiting list, at LT and weekly after LT until hospital
166 discharge. Screening for CRE colonization at other sites was performed according to clinical
167 judgment and local policy.

168 *Study population*

169 All consecutive adult (≥ 18 years) patients who underwent LT during the study period and were found
170 to be colonized with CRE were enrolled. Patients were included only once at the time of the first CRE
171 carriage detection. CRE was defined as any *Enterobacteriaceae* displaying *in vitro* non-susceptibility
172 to any of the carbapenems according to the criteria (CLSI or EUCAST) adopted at the participating

173 centre during the study period. The colonization status was defined as isolation of CRE from RS or
174 other samples other than blood cultures or sterile fluids (e.g., urine, respiratory samples, superficial
175 skin samples) in absence of symptoms and signs of infection. Multisite colonization was defined
176 when CRE was concomitantly isolated from more than one of such samples. Patients found to be
177 colonized with CRE by samples obtained at inclusion in the waiting list, during the period in waiting
178 list, or at the time of LT were considered as CRE carriers pre-LT, whereas those found to be colonized
179 within 180 days after LT were considered as post-LT CRE carriers.

180 *Predictors of CRE infection after LT*

181 The primary endpoint was the time to development of CRE infection, diagnosed within 180 days
182 after LT, according to Center for Disease Control criteria [10]. The assessment of CRE infection was
183 made by the local investigator and revised by an investigator of the promoting centre (MR), in case
184 of no agreement a third blinded investigator of the promoting centre (MG) was asked to review the
185 case for establishing the final diagnosis. Infection severity was determined using SOFA score and
186 septic shock criteria [11].

187 Candidate prediction variables included: demographic data (age and sex); comorbidities according to
188 Charlson index; underlying liver disease, and severity of liver disease according to Model for End
189 stage Liver Disease (MELD) at inclusion in waiting list and at LT. Complications occurred within 90
190 days before LT were recorded and included: admission to intensive care unit (ICU); grade 3 and/or
191 refractory ascites [12]; hepato-renal syndrome (HRS) [12]; gastrointestinal bleeding; hepatic
192 encephalopathy [13]; development of acute-on-chronic liver failure (ACLF) [14]; any infection,
193 candidemia, and *Clostridioides difficile* infection. For graft characteristics donor age, cold ischemia
194 time, and combined transplant were collected. Intraoperative variables included: antibiotic
195 prophylaxis, biliary anastomosis, bleeding with need of massive transfusion (≥ 40 units of cellular
196 blood products), prolonged intervention (≥ 8 hours). Complications occurred from LT to the
197 diagnosis of CRE infection or death or 180 days (whichever occurred first) were recorded and

198 included: re-intervention, acute kidney injury (AKI) according to KIDGO criteria [15], renal
199 replacement therapy (RRT), prolonged (≥ 48 hours) mechanical ventilation (MV), graft dysfunction
200 (primary or secondary), biopsy-proven rejection, re-transplantation, and CMV DNAemia $>100,000$
201 copies/ml [16]. Management of CRE colonization was recorded according to the following
202 categories: no strategy adopted, anti-CRE surgical prophylaxis, decolonization with non-absorbable
203 antibiotics, faecal microbiota transplantation, pre-emptive strategy (defined as early start of anti-
204 CRE coverage upon to clinical deterioration). For outcome, duration of the hospital stay after LT,
205 number of re-admissions within 180-days after LT, and all-cause 180-days mortality were recorded.

206 *Sample size*

207 Sample size required for the clinical prediction model was estimated for a time-to-event outcome
208 using the pmsampsize package in R (version 4.02, Core Team (2020). **R**: A language and environment
209 for statistical computing. **R** Foundation for Statistical Computing, Vienna, Austria) using the methods
210 proposed by Riley et al. [17]. We estimated that 850 patients with 213 outcome events (event rate
211 0.25 by day 180) would be sufficient to evaluate 20 candidate predictors (>10 events per candidate
212 predictor) with an estimated 60-day event rate of 0.17.

213 *Missing data*

214 A complete-case analysis was performed.

215 *Statistical analysis methods*

216 Categorical variables were expressed as absolute numbers and their relative frequencies. Continuous
217 variables were expressed as mean \pm standard deviation (SD) if normally distributed, or as median
218 and interquartile range (IQR) if non-normally distributed. We used the Fine-Gray sub-distribution
219 hazard model to develop a clinical prognostic index for CRE infection over the first 180 days after LT
220 with death as competing event.

221 Although colonization status has been modelled in previous studies as a time-varying covariate [18],
222 inclusion of the time-dependent covariates limits the ability of the Fine-Gray sub-distribution hazard
223 model to make inferences about the effects of covariates on the cumulative incidence function (CIF)
224 in the presence of competing risks, which is an underlying motivation for our model. Furthermore,
225 simulation studies have shown that a sub-distribution approach for time-dependent covariates often
226 produces misleading results [19], and is not currently recommended [20]. Therefore, we focused on
227 colonization status as a risk factor only in the 60-days prior to and 60 days following OLT to minimize
228 time-dependent effects.

229 Using a base model of CRE colonization within 60 days prior to transplant, we examined the degree
230 to which including each other variable from our original model improved its performance. Additional
231 candidate variables were selected *a priori* based on statistical significance in univariate analysis,
232 published data and clinical experience and whether they improved prediction accuracy of the
233 baseline model. Only variables available for assessment during the risk period were included. We
234 assumed that a predictive model for CRE infection would have the most clinical utility for accurate
235 prediction of CRE infection in the first 30 and 60 days post OLT, as the risk appeared to plateau after
236 this period. Therefore we used C-index (AUC) to assess the discrimination of the model with
237 censored CRE infection status, and calibration plots of day-60 observed versus expected absolute
238 risks with the Brier score to assess the accuracy of predictions. To control for centre effect, the
239 analysis was also performed for each participating centre. Model discrimination and calibration
240 measures were calculated using 100-fold cross validation bootstrap resampling to avoid bias. A
241 decision curve was prepared to examine potential model utility across a range of CRE infection
242 probabilities (0-60%) at 60 days post LT. All analyses were performed using R version 4.02 with
243 riskRegression, prodlim, ggplot2, survival, and rmda packages. The final regression model was
244 visualized by preparing a nomogram to predict 30-day and 60-day risk using methods described by
245 Zhang et al [21] and the regplot package. An online prediction tool (Shiny App) was prepared using
246 the DynNom package in R. Results were reported according to TRIPOD guidelines [22].

247

248 **RESULTS**

249 Overall, 840 LT recipients colonized with CRE before or after transplantation were enrolled, with 250
250 episodes of CRE infection post LT (29.7%) during 104,596 patient days of follow-up. Distribution of
251 patients and of CRE infections across participating centres is shown in Figure 1. The characteristics of
252 study population are shown in Table 1. Median age was 55 (IQR 46-62) years, male sex (65.4%). Viral
253 hepatitis was the primary indication for LT in 376/840 subjects (44.8%), followed by alcohol 207/840
254 (24.6%). Over one-quarter 247/840 (29.4%) of patients had hepatocellular carcinoma (HCC). MELD at
255 inclusion in waiting list and at LT was 19 (IQR 14-25) and 23 (IQR 16-30), respectively. CRE carriage
256 was detected before or at LT in 203 (24.2%) patients, the median time from detection to LT was 6
257 (IQR 0-32) days. The remaining 637 (75.8%) patients were found to be colonized with CRE within a
258 median of 13 (IQR 7-28) days after LT. Comparison of pre- and post-LT carriers showed several
259 differences (see Table 1). Specifically, pre-LT carriers presented with more severe liver disease and
260 complicated clinical course before transplantation. CRE infection rate among pre- and post-LT
261 carriers was 33% and 28.7%, and the median time to infection following LT was 9 (IQR 4-22) and 23
262 (IQR 13-52) days, respectively (see Supplementary Figure 1). The median time from detection of CRE
263 carriage to the diagnosis of CRE infection was shorter for post-LT carriers (8 vs. 19 days, $p<0.001$).
264 Among the 203 pre-OLT carriers, 51 received targeted perioperative prophylaxis, the rate of CRE
265 infection was 31.4% and 33.6% among those exposed and unexposed to targeted prophylaxis
266 ($p=0.90$), respectively. Post-LT CRE infections in pre-LT carriers presented with higher severity
267 according to median SOFA and septic shock rates than in post-LT carriers. However, there was no
268 difference in clinical outcome (all-cause 180 days mortality 56.7% vs. 58.5%, $p=0.46$), data shown in
269 Supplementary Table 2.

270 To screen risk factors for CRE infection, we performed univariate analysis of patients with and
271 without CRE infection as shown in Table 2. Considering the competing risks of death, a Fine-Gray

272 model was used to perform regression of risk factors to estimate their association with the
273 cumulative incidence of CRE infection up to 180 days post LT. Baseline cumulative incidence of
274 death versus CRE infection over 180 following OLT is shown in Figure 2. The following variables were
275 retained in the final model: pre-LT CRE colonization, post-LT CRE colonization, multisite post-LT
276 colonization, prolonged MV, AKI, and surgical re-intervention (Table 3). The fitted regression risk
277 model was then rendered as a nomogram (Figure 3). Model-predicted cumulative probabilities of
278 developing CRE infection at day 30 and 60 post LT are shown in Figure 4. The median 30 and 60-day
279 predicted risk was 15% (IQR 11-24%) and 21% (IQR 15-33%), respectively. Nearly one-quarter
280 (23.5%) of patients at day 30, and 3% of patients at day 60 had a cumulative predicted risk of CRE
281 infection $\leq 10\%$. The model showed acceptable 60-day discrimination and prediction accuracy for
282 CRE infection when assessed against the derivation AUC 74.6 (95% 70.9-78.4), Brier index 16.3
283 (95%CI 6.7-25.9) and bootstrapped validation dataset AUC 73.9 (95%CI 67.7-79.1), Brier index 16.6
284 (95%CI 14.6-19.1) (Figure 5). Discrimination and calibration were also evaluated by centre and were
285 consistent with the overall performance, except for higher variability and lower AUCs in centres with
286 lower number of recruited patients (see Supplementary Figure 2). The score was designed to be
287 used in the immediate peri-transplant period, ideally from the day of transplantation up to 2-3
288 weeks after transplantation. During this period, the patient should be frequently evaluated for
289 carriage status, multisite colonization, prolonged MV, AKI, and/or re-intervention. According with
290 the presence of such variables the cumulative risk of CRE infection within 30 and 60 days after LT can
291 be predicted (see examples in Figure 6 and mobile application at
292 <https://idbologna.shinyapps.io/CREPostOLTPredictionModel/>, instructions for using app are further
293 provided as supplementary material).

294 Finally, we explored the potential clinical utility of our model using a decision-curve analysis (DCA) to
295 examine the “net benefit” of applying the prediction model across a range of CRE infection threshold
296 probabilities [23]. A theoretical risk-model guided strategy (i.e. empiric administration of CRE-active
297 antibiotics) is compared against two default strategies- “treat all” and “treat none.” A model is only

298 useful at a given disease threshold if it has a higher net benefit than treat all or treat none. Our
299 analysis suggested that the model-directed interventions for CRE post LT would show net benefit
300 over default strategies when the overall CRE infection threshold probability exceeded 10% (Fig. 7).

301

302 **DISCUSSION**

303 Assessing a large multinational cohort of LT recipients colonized with CRE (n=840) we have found
304 that one third developed CRE infection after LT. This rate was similar among pre- and post-LT CRE
305 carriers. Most infection episodes occurred within 2 months after LT. The time to infection was earlier
306 for pre-LT CRE carriers, and the clinical severity was higher in this group. All-cause 6-month mortality
307 among carriers was as high as 30%, with similar rates between pre- and post-LT CRE carriers.
308 However, it reached 58% among patients who developed CRE infection.

309 We confirmed that the CRE carriage status is strongly associated with the risk of CRE infection after
310 LT, either if detected before or after LT [6, 7]. However, we also confirmed our previous hypothesis
311 that, given a probability of CRE infection higher than 10%, a strategy of treating all CRE carriers could
312 be inferior to a targeted strategy in terms of efficacy, as well as safety, considering potential adverse
313 events of anti-CRE drugs and the risk for further selection pressure. A longitudinal risk stratified
314 approach, could be useful for several reasons. In general, LT is not an elective surgery, thus it could
315 be difficult to implement a strategy before transplantation other than in a minority of patients
316 known to be colonized at the time of inclusion or while on waiting list. Another opportunity for
317 intervention could be surgical prophylaxis. However, data about colonization is frequently available
318 only during the immediate post-operative period. Finally, in most patients CRE colonization becomes
319 evident in the first weeks after LT, typically with a complicated post-operative course, which also
320 increases the risk of infection.

321 Along with pre- and post-LT colonization status, multisite colonization after LT was a predictor of
322 infection. The relationship between multisite colonization with infection development has been
323 already observed in other settings [3]. It is indicative of high colonization burden leading to an
324 increased risk of CRE infection [24]. Indeed, its inclusion in our prognostic model improved model
325 performance. Nevertheless, multisite colonization was not systematically screened in this study.
326 Therefore, we cannot rule out that colonization from different sites may be weighted differently in
327 the model if this risk factor had been systematically screened, as patients with a more complicated
328 course may have been cultured more frequently.

329 The other complications included in our score are widely described as factors involved in the
330 increase risk of infection after LT [25]. AKI has been also associated with microbiome dysbiosis that
331 could favour prolonged CRE carriage and infection [26]. As for MV, it is worth mentioning that we
332 considered such variable, as a qualitative data, from LT to CRE infection, death or end of follow-up. A
333 more detailed information about timing and duration of MV may improve the model performance.

334 Our study has several limitations. First, the retrospective design could have limited identification of
335 all CRE carriers. However, the presence of a local prospective registry of CRE carriers and/or good
336 collaboration with local microbiology laboratories reduced this bias. Heterogeneity in local assays
337 and criteria for identification and determination of CRE could exist. However, obtained data are in
338 line with international literature. The majority of involved centres were high endemic CRE hospitals,
339 with KPC as main mechanism of CRE. Thus, our findings could not apply to transplant centres with
340 low prevalence of CRE, or with KPC alternative main mechanisms among CRE. Finally, we analysed
341 the CRE infection risk only in colonized patients, thus the performance of our prediction tool should
342 be confirmed in unselected LT patients.

343 To conclude, one third of CRE carriers develop infection after LT with a dramatic impact on patient
344 survival. Effective preventive or pre-emptive strategies are needed. Our clinical prediction tool could
345 enable better targeting of early interventions for CRE infection after transplant as opposed to

346 universal prophylaxis or treatment in high-prevalence centres. Further studies are needed to further
347 validate the model and establish which strategy would be more effective, in terms of adverse events
348 and collateral resistance impact and, ultimately patient survival after transplantation.

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350

351 **ACKNOWLEDGMENTS/FUNDING**

352 No funding was received for this project.

353 **CONFLICT OF INTEREST**

354 Authors have no conflict of interest related to the present study.

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439 **Table 1.** Characteristics of study population according to the timing of CRE carriage detection

| | Total N=840 (%) | pre-OLT carrier N=203 (%) | post-OLT carrier N=637 (%) | P |
|--|----------------------------|--|---|----------|
| Demographic data | | | | |
| Age (years) [median (IQR)] | 55 (46-62) | 53 (42-60) | 56 (47-62) | 0.005 |
| Sex, male | 549 (65.4) | 132 (65) | 417 (65.5) | 0.9 |
| Comorbidities | | | | |
| Myocardial infarction | 23 (2.7) | 5 (2.5) | 18 (2.8) | 0.827 |
| Congestive heart failure | 29 (3.5) | 7 (3.4) | 22 (3.5) | 0.9 |
| Peripheral vascular disease | 33 (3.9) | 6 (3) | 27 (4.2) | 0.4 |
| Cerebrovascular disease | 22 (2.6) | 8 (3.9) | 14 (2.2) | 0.18 |
| COPD | 46 (5.5) | 6 (3) | 40 (6.3) | 0.07 |
| Connective tissue disease | 9 (1.1) | 3 (1.5) | 6 (0.9) | 0.38 |
| Peptic ulcer disease | 62 (7.4) | 20 (9.9) | 42 (6.6) | 0.09 |
| Diabetes without organ damage | 147 (17.5) | 38 (18.7) | 109 (17.1) | 0.3 |
| Moderate/severe renal disease | 82 (9.8) | 28 (13.8) | 54 (8.5) | 0.03 |
| Diabetes with organ damage | 76 (9) | 15 (7.4) | 61 (9.6) | 0.2 |
| Any tumor within 5 years | 271 (32.3) | 45 (22.2) | 226 (35.5) | 0.001 |
| Moderate/severe liver disease | 820 (97.6) | 199 (98) | 621 (97.5) | 0.79 |
| Charlson index [median (IQR)] | 5 (4-7) | 5 (3-6) | 5 (4-7) | 0.004 |
| Underlying liver disease | | | | |
| Viral hepatitis | 376 (44.8) | 74 (36.5) | 302 (47.4) | 0.006 |
| Alcohol | 207 (24.6) | 55 (27.1) | 152 (23.9) | 0.2 |
| Metabolic disease | 69 (8.2) | 12 (5.9) | 57 (8.9) | 0.1 |
| Autoimmune disease | 41 (4.9) | 12 (5.9) | 29 (4.6) | 0.45 |
| Fulminant hepatitis | 52 (6.2) | 7 (3.4) | 45 (7.1) | 0.04 |
| Hepatocellular carcinoma | 247 (29.4) | 37 (18.2) | 210 (33) | <0.001 |
| Prior transplant | 39 (4.6) | 20 (9.9) | 19 (3) | <0.001 |
| MELD at waiting list inclusion (median, IQR) | 19 (14-25) | 21 (16-28) | 19 (14-24) | <0.001 |
| MELD at OLT (median, IQR) | 23 (16-30) | 27 (21-32) | 22 (15-29) | <0.001 |
| Variables relative to the last 90 days before OLT | | | | |
| Hepato-renal syndrome | 235 (28) | 88 (43.3) | 147 (23.1) | <0.001 |
| Ascites grade III/refractory | 385 (45.8) | 108 (53.2) | 277 (43.5) | 0.01 |
| GI bleeding | 138 (16.4) | 43 (21.2) | 95 (14.9) | 0.025 |
| Encephalopathy | 343 (40.8) | 96 (47.3) | 247 (38.8) | 0.02 |
| ICU admission | 161 (19.2) | 70 (34.5) | 91 (14.3) | <0.001 |
| ACLF | 98 (11.7) | 44 (21.7) | 54 (8.5) | <0.001 |
| Any infection | 326 (38.8) | 137 (67.5) | 189 (29.7) | <0.001 |
| Intraoperative variables | | | | |
| Combined transplant | 45 (5.4) | 12 (5.9) | 33 (5.2) | 0.7 |
| Donor age | 51 (36-65) | 50 (35-65) | 51 (36-65) | 0.4 |
| Cold ischemia time (hours) (median, IQR) | 7 (6-8) | 7 (6-8) | 7 (6-8) | 0.62 |
| Biliary anastomosis | | | | |
| Duct to duct without Kehr | 382 (45.5) | 84 (41.4) | 298 (46.8) | 0.19 |
| Duct to duct with Kehr | 347 (41.3) | 76 (37.4) | 271 (42.5) | 0.20 |

| | | | | |
|--|-------------|------------|------------|--------|
| Choledochojejunostomy | 111 (13.2) | 43 (21.2) | 68 (10.7) | <0.001 |
| Intraoperative bleeding | 294 (35) | 88 (43.3) | 206 (32.3) | 0.005 |
| Prolonged surgery (>8 hours) | 248 (29.5) | 70 (34.5) | 178 (27.9) | 0.046 |
| Induction regimen | | | | |
| None | 87 (10.4) | 24 (11.8) | 63 (9.9) | 0.51 |
| Bolus of steroids | 701 (83.5) | 171 (84.2) | 530 (83.2) | 0.75 |
| Antithymocyte globulins | 26 (3.1) | 3 (1.5) | 23 (3.6) | 0.09 |
| Basiliximab | 167 (19.9) | 37 (18.2) | 130 (20.4) | 0.55 |
| Rituximab | 14 (1.7) | 0 (0) | 14 (2.2) | 0.02 |
| Maintenance regimen | | | | |
| Steroids | 612 (72.9) | 162 (79.8) | 450 (70.6) | 0.011 |
| Cyclosporin | 42 (5) | 8 (3.9) | 34 (5.3) | 0.28 |
| Tacrolimus | 770 (91.7) | 181 (89.2) | 589 (92.5) | 0.15 |
| Micophenolate mophetil | 287 (34.2) | 70 (34.5) | 217 (34.1) | 0.9 |
| Post-operative complications | | | | |
| Acute renal injury | 462 (55) | 118 (58.1) | 344 (54) | 0.33 |
| Renal replacement therapy | 299 (35.6) | 90 (44.3) | 209 (32.8) | 0.003 |
| Mechanical ventilation ≥48 hours | 266 (31.7) | 64 (31.5) | 202 (31.7) | 0.96 |
| PGNF | 128 (15.2) | 28 (13.8) | 100 (15.7) | 0.58 |
| Re-intervention | 322 (38.3) | 74 (36.5) | 248 (38.9) | 0.56 |
| Re-transplantation | 79 (9.4) | 11 (5.4) | 68 (10.7) | 0.025 |
| Rejection | 152 (18.1) | 30 (14.8) | 122 (19.2) | 0.17 |
| CMV treatment | 225 (26.8) | 46 (22.7) | 179 (28.1) | 0.15 |
| Infections other than CRE | 479 (57) | 106 (52.2) | 373 (58.6) | 0.12 |
| <i>Clostridioides difficile</i> infection | 37 (4.4) | 8 (3.9) | 29 (4.6) | 0.44 |
| Candidemia | 68 (8.1) | 11 (5.4) | 57 (8.9) | 0.14 |
| CRE carriage | | | | |
| Time between OLT and carriage detection (days) (median, IQR) | 9 (0.25-21) | -6 (-32-0) | 13 (7-28) | <0.001 |
| First positive sample | | | | |
| Rectal swab | 719 (85.6) | 189 (93.1) | 530 (83.2) | 0.001 |
| Respiratory sample | 45 (5.4) | 3 (1.5) | 42 (6.6) | 0.002 |
| Urine | 69 (8.2) | 22 (10.8) | 47 (7.4) | 0.14 |
| Multi-site colonization | 113 (13.5) | 26 (12.8) | 87 (13.7) | 0.81 |
| Resistance mechanism | | | | |
| KPC | 600 (71.4) | 149 (73.4) | 451 (70.8) | 0.53 |
| VIM | 7 (0.8) | 3 (1.5) | 4 (0.6) | 0.23 |
| OXA-48 | 17 (2) | 6 (3) | 11 (1.7) | 0.21 |
| Not tested | 115 (13.7) | 19 (9.4) | 96 (15.1) | 0.05 |
| Management of CRE colonization | | | | |
| No strategy adopted | 521 (62) | 85 (41.9) | 436 (68.4) | <0.001 |
| Target perioperative prophylaxis | 51 (6.1) | 51 (25.1) | 0 (0) | <0.001 |
| Decolonization | 16 (1.9) | 4 (2) | 12 (1.9) | 0.57 |
| Pre-emptive strategy ¹ | 21 (2.5) | 6 (3) | 15 (2.4) | 0.4 |
| Outcome | | | | |
| Length of ICU stay after OLT (days) (median, IQR) | 7 (4-15) | 7 (4-14) | 7 (4-15) | 0.38 |
| Length of hospital stay after OLT (days) (median, IQR) | 27 (16-50) | 23 (14-37) | 29 (17-54) | <0.001 |
| All-cause 6-months mortality | 263 (31.3) | 69 (34) | 194 (30.5) | 0.39 |

| | | | | |
|--|-------------|----------------|---------------|--------|
| Time from OLT to death (days) (median, IQR) | 70 (29-145) | 32 (10.5-77.5) | 88.5 (43-162) | <0.001 |
| Hospital re-admissions (median, IQR) | 0 (0-1) | 0 (0-1) | 0 (0-1) | 0.07 |

440 ¹Start of an anti-CRE treatment if complications occur

441 Abbreviations: IQR interquartile range, COPD chronic obstructive pulmonary disease, MELD Model

442 End Stage Liver Disease, OLT orthotopic liver transplant, ICU intensive care unit, ACLF acute on

443 chronic liver failure, PGNF primary graft non-function, CMV cytomegalovirus, CRE carbapenem-

444 resistant *Enterobacteriaceae*, KPC *Klebsiella pneumoniae* carbapenemase-producing, SBP

445 spontaneous bacterial peritonitis.

446

447 **Table 2.** Comparison of patients with and without CRE infection within 180 days after liver transplant

| | Patients without CRE infection, n=590 (%) | Patients with CRE infection, n=250 (%) | P |
|--|--|---|----------|
| Demographic data | | | |
| Age (years) [median (IQR)] | 55 (46-61) | 56 (47-62) | 0.5 |
| Sex, male | 389 (65.9) | 160 (64) | 0.59 |
| Comorbidities | | | |
| Myocardial infarction | 17 (2.9) | 6 (2.4) | 0.7 |
| Congestive heart failure | 19 (3.2) | 10 (4) | 0.57 |
| Peripheral vascular disease | 27 (4.6) | 6 (2.4) | 0.14 |
| Cerebrovascular disease | 15 (2.5) | 7 (2.8) | 0.83 |
| COPD | 30 (5.1) | 16 (6.4) | 0.51 |
| Connective tissue disease | 6 (1) | 3 (1.2) | 0.81 |
| Peptic ulcer disease | 44 (7.5) | 18 (7.2) | 0.89 |
| Diabetes without organ damage | 103 (17.5) | 44 (17.6) | 0.96 |
| Moderate/severe renal disease | 45 (7.6) | 37 (14.8) | 0.001 |
| Diabetes with organ damage | 55 (9.3) | 21 (8.4) | 0.67 |
| Any tumor within 5 years | 192 (32.5) | 79 (31.6) | 0.79 |
| Moderate/severe liver disease | 576 (97.6) | 244 (97.6) | 0.98 |
| Charlson index [median (IQR)] | 5 (4-7) | 5 (4-7) | 0.56 |
| Underlying liver disease | | | |
| Viral hepatitis | 260 (44.1) | 116 (46.4) | 0.53 |
| Alcohol | 157 (26.6) | 50 (20) | 0.04 |
| Metabolic disease | 48 (8.1) | 21 (8.4) | 0.89 |
| Autoimmune disease | 32 (5.4) | 9 (3.6) | 0.26 |
| Fulminant hepatitis | 36 (6.1) | 16 (6.4) | 0.87 |
| Hepatocellular carcinoma | 174 (29.5) | 73 (29.2) | 0.93 |
| Prior transplant | 24 (4.1) | 15 (6) | 0.22 |
| MELD at waiting list inclusion (median, IQR) | 19 (14-24) | 19 (14-26) | 0.73 |
| MELD at OLT (median, IQR) | 22.5 (16-29) | 25 (16-33) | 0.02 |
| Events prior 90 days to OLT | | | |
| Hepato-renal syndrome | 150 (25.4) | 85 (34) | 0.01 |
| Ascites grade III/refractory | 278 (47.1) | 107 (42.8) | 0.25 |
| Bleeding | 89 (15.1) | 49 (19.6) | 0.11 |
| Encephalopathy | 245 (41.5) | 98 (39.2) | 0.53 |
| ICU admission | 110 (18.6) | 51 (20.4) | 0.55 |
| ACLF | 66 (11.2) | 32 (12.8) | 0.5 |
| Infections | 223 (37.8) | 103 (41.2) | 0.36 |
| Intraoperative variables | | | |
| Combined transplant | 25 (4.2) | 20 (8) | 0.03 |
| Donor age | 51 (35-65) | 51 (36-67) | 0.6 |
| Cold ischemia time (hours) (median, IQR) | 7 (6-8) | 7 (5:50-8) | 0.36 |
| Choledochojejunostomy | 71 (12) | 40 (16) | 0.22 |
| Intraoperative bleeding | 196 (33.2) | 98 (39.2) | 0.1 |
| Prolonged surgery | 162 (27.5) | 86 (34.4) | 0.04 |
| Induction regimen | | | |
| Bolus of steroids | 476 (80.7) | 225 (90) | 0.001 |

| | | | |
|---|------------|------------|--------|
| Antithymocyte globulins | 12 (2) | 14 (5.6) | 0.006 |
| Basiliximab/Daclizumab | 113 (19.2) | 54 (21.6) | 0.42 |
| Rituximab | 6 (1) | 8 (3.2) | 0.02 |
| Maintenance regimen | | | |
| Steroids | 414 (70.2) | 198 (79.2) | 0.007 |
| Cyclosporin | 31 (5.3) | 11 (4.4) | 0.6 |
| Tacrolimus | 549 (93.1) | 221 (88.4) | 0.03 |
| Micophenolate mophetil | 215 (36.4) | 72 (28.8) | 0.03 |
| Post-operative complications | | | |
| Acute renal injury | 286 (48.5) | 176 (70.4) | <0.001 |
| Renal replacement therapy | 173 (29.3) | 126 (50.4) | <0.001 |
| Mechanical ventilation >48 hours | 135 (22.9) | 131 (52.4) | <0.001 |
| PGNF | 74 (12.5) | 54 (21.6) | 0.001 |
| Re-intervention | 200 (33.9) | 122 (48.8) | <0.001 |
| Re-transplantation | 43 (7.3) | 36 (14.4) | 0.001 |
| Rejection | 101 (17.1) | 51 (20.4) | 0.26 |
| CMV DNAemia>100,000 copies/mL | 156 (26.4) | 69 (27.6) | 0.73 |
| Infections other than CRE | 333 (56.4) | 146 (58.4) | 0.6 |
| <i>Clostridioides difficile</i> infection | 26 (4.4) | 11 (4.4) | 0.99 |
| Candidemia | 36 (6.1) | 32 (12.8) | 0.001 |
| CRE carriage | | | |
| Pre-OLT carriage | 136 (23.1) | 67 (26.8) | 0.25 |
| Post-OLT carriage | 454 (76.9) | 183 (73.2) | 0.25 |
| Rectal swab | 517 (87.6) | 202 (80.8) | 0.01 |
| Respiratory sample | 21 (3.6) | 24 (9.6) | <0.001 |
| Urine sample | 50 (8.5) | 19 (7.6) | 0.67 |
| Multisite colonization | 54 (9.2) | 59 (23.6) | <0.001 |
| pre-OLT | 17 (2.9) | 9 (3.6) | <0.001 |
| post-OLT | 37 (6.3) | 50 (20) | 0.66 |
| Genotype of the colonizing strain | | | |
| KPC | 418 (70.8) | 182 (72.8) | 0.61 |
| VIM | 4 (0.7) | 3 (1.2) | 0.45 |
| OXA-48 | 11 (1.9) | 6 (2.4) | 0.6 |
| Not tested | 76 (12.9) | 39 (15.6) | 0.3 |
| Management of CRE colonization | | | |
| Target perioperative prophylaxis | 35 (5.9) | 17 (6.8) | 0.6 |
| Decolonization | 12 (2) | 4 (1.6) | 0.7 |
| Pre-emptive strategy ¹ | 11 (1.9) | 10 (4) | 0.07 |
| Outcome | | | |
| Length of ICU stay after OLT (days) (median, IQR) | 6 (3-10) | 14 (5-29) | <0.001 |
| Length of hospital stay after OLT (days) (median, IQR) | 22 (15-40) | 43 (24-76) | <0.001 |
| All-cause 6-months mortality | 118 (20) | 145 (58) | <0.001 |
| Hospital re-admissions (median, IQR) | 0 (0-1) | 0 (0-1) | <0.001 |

448 ¹Start of an anti-CRE treatment if complications occur.

449 Abbreviations: IQR interquartile range, COPD chronic obstructive pulmonary disease, MELD Model

450 End Stage Liver Disease, OLT orthotopic liver transplant, ICU intensive care unit, SBP spontaneous

451 bacterial peritonitis, ACLF acute on chronic liver failure, PGNF primary graft non-function, CMV
452 cytomegalovirus, CRE carbapenem-resistant *Enterobacteriaceae*, KPC *Klebsiella pneumoniae*
453 carbapenemase-producing.

454 **Table 3.** Final multivariable analysis for predicting risk of CRE infection after liver transplantation

| Variable | Subhazard ratio (95% CI) | β-coefficient | P value |
|--|---------------------------------|---------------------------------------|----------------|
| Reintervention | 1.37 (1.04-1.80) | 0.30 (0.02-0.57) | 0.02 |
| Prolonged mechanical ventilation | 2.01 (1.49-2.71) | 0.70 (0.40-0.99) | <0.0001 |
| Acute renal injury | 1.69 (1.26-2.27) | 0.52 (0.23-0.82) | 0.001 |
| CRE colonisation 60 days prior to transplant | 2.17 (1.33-3.53) | 0.78 (0.29-1.26) | 0.002 |
| CRE colonization within 60 days after transplant | 1.41 (0.91-2.19) | 0.35 (-0.09-0.78) | 0.12 |
| Multisite colonization within 60 days after transplant | 2.87 (1.98-4.20) | 1.06 (0.68-1.43) | <0.0001 |

455

456 Abbreviations: CI confidence interval, CRE carbapenem-resistant Enterobacteriaceae.

457

458 Figure legends:

459 **Figure 1.** Study flow chart with detailed data about involved centers.

460 **Figure 2.** Cumulative incidence of CRE infection and death post-transplant. The cumulative incidence
461 was derived from the estimated sub-distribution hazard function following regression with a Fine-
462 Gray model.

463 **Figure 3.** Model-based nomogram for predicting 30 and 60-day cumulative risk of developing CRE
464 infection. Points are summed for each risk factor.

465 **Figure 4.** Distribution of model-predicted (a) 30-day and (b) 60-day risk for CRE infection following
466 LT.

467 **Figure 5.** Calibration curves for the (a) derivation dataset and (b) Bootstrap resampled validation
468 dataset. The C-index (AUC) and Brier score are expressed as the point estimates and 95% CI. An AUC
469 > 0.8 is considered to give good discriminatory accuracy for a clinical prediction model. The Brier
470 score is a measure of the accuracy of probabilistic predictions. The lower the Brier score is for a set
471 of predictions, the better the predictions are calibrated. An ideal model has pairs of observed and
472 predicted probabilities that lie on the 45-degree angle line. Dotplots shown the distribution of
473 predicted risk for patients who developed (top) and did not develop (bottom) CRE infection.

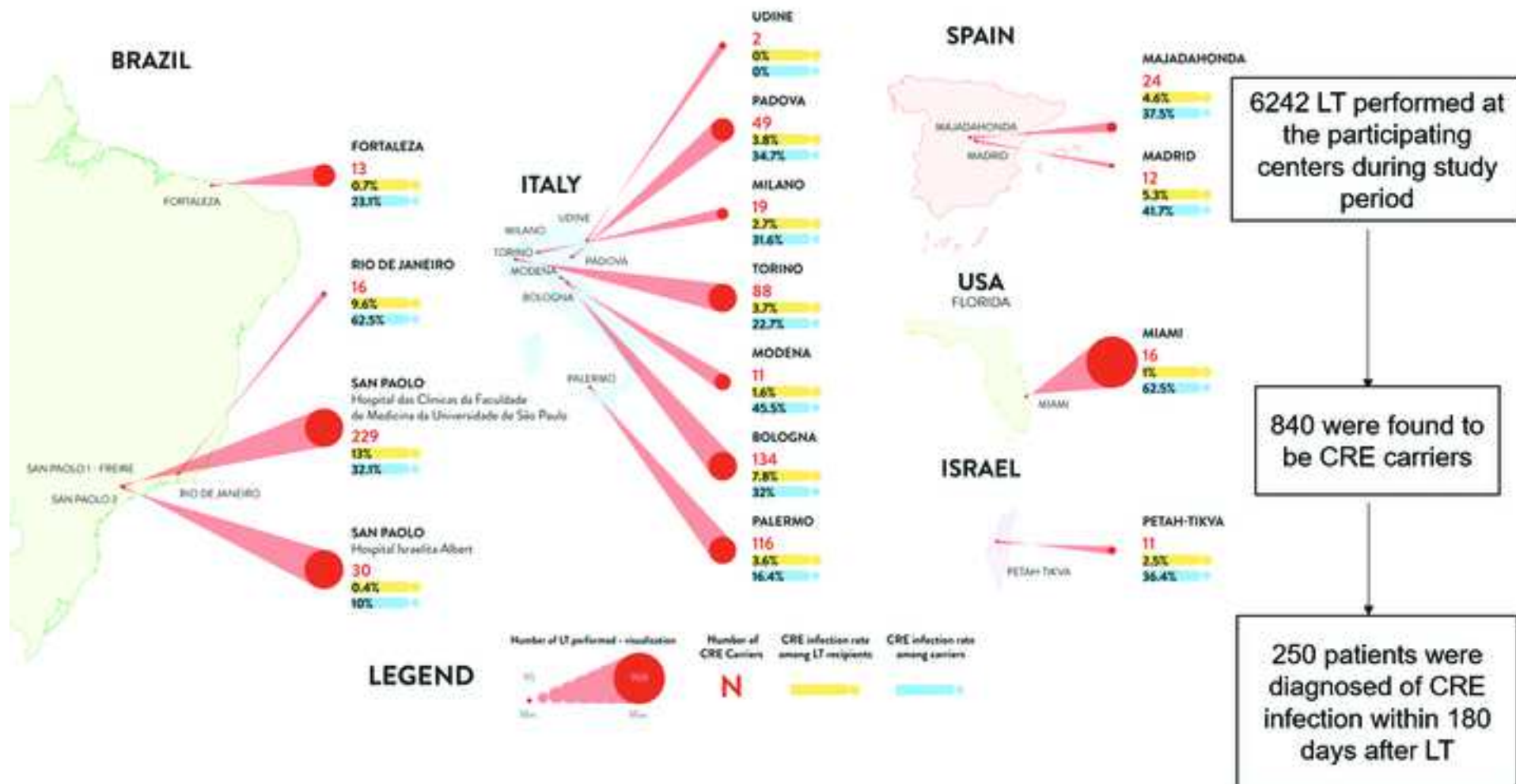
474 **Figure 6.** Examples of the cumulative incidence of CRE infection and nomogram-derived prediction
475 for a low-risk (panels a, b); and a high-risk (panels c, d) LT patient. The blue lines with shading in the
476 nomogram represent the distribution of individual risk factors (0=no, 1=yes) and the distribution of
477 total assigned points on the bottom scale in the original derivation data set. Red points on the upper
478 scale indicate assigned points per risk factor. The red arrows at the bottom of the nomogram indicate
479 the total calculated points for the patient and corresponding 30-day and 60 day predicted
480 cumulative risk with 95% CI.

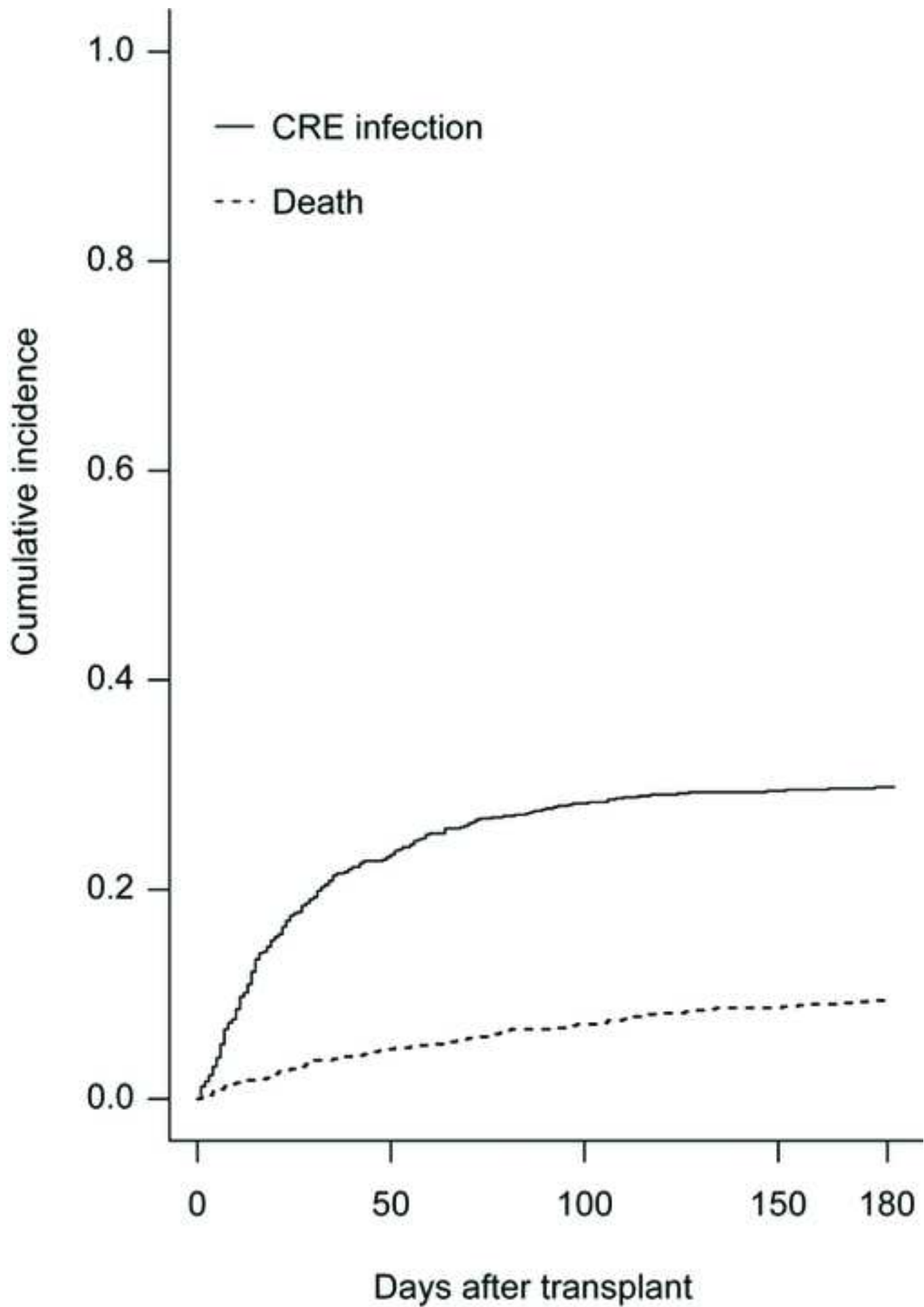
481 **Figure 7.** Decision-curve analysis of model-directed anti CRE therapy. Net benefit at 60 days
482 represents true cases of CRE infection that would be treated with a CRE-active antibiotic regimen-
483 patients without CRE infection unnecessarily receiving CRE-active therapy. The curves indicate that a
484 model-directed antibiotic strategy should have a higher net benefit than default strategies (all
485 patients receive CRE-active therapy or no patients receive CRE-active therapy) at CRE infection
486 probabilities greater than 10-50 %.

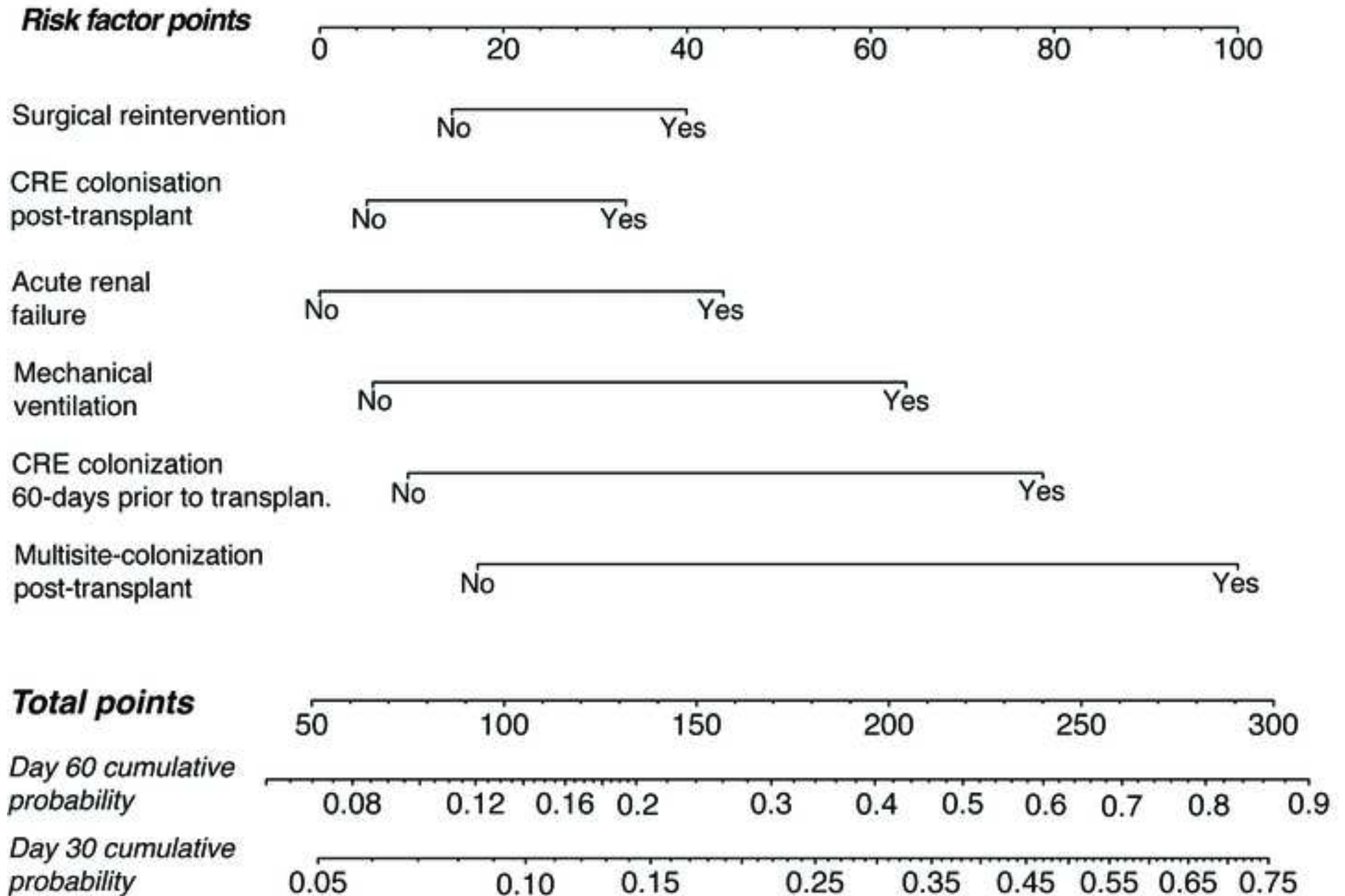
487

Figure 1

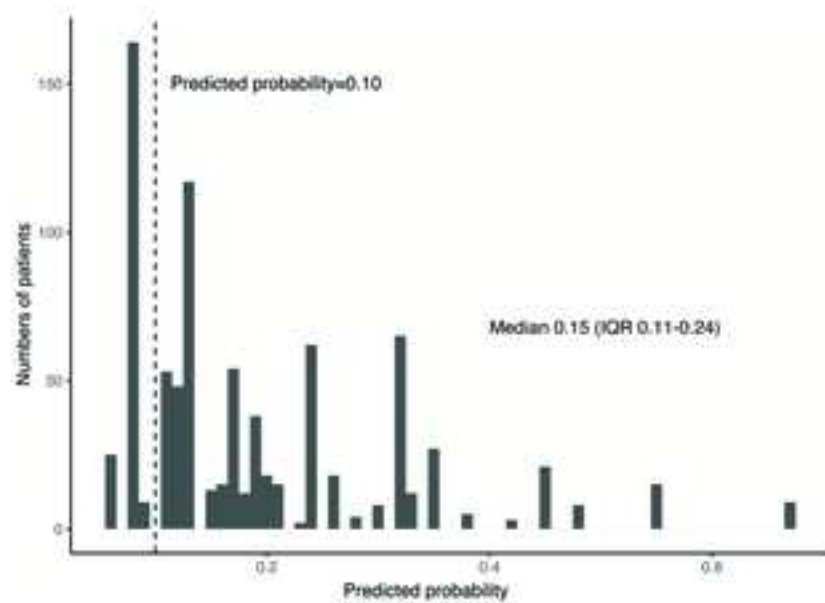
[Click here to access/download;Figure \(.tif and .eps files only\);Figure 1.tif](#)



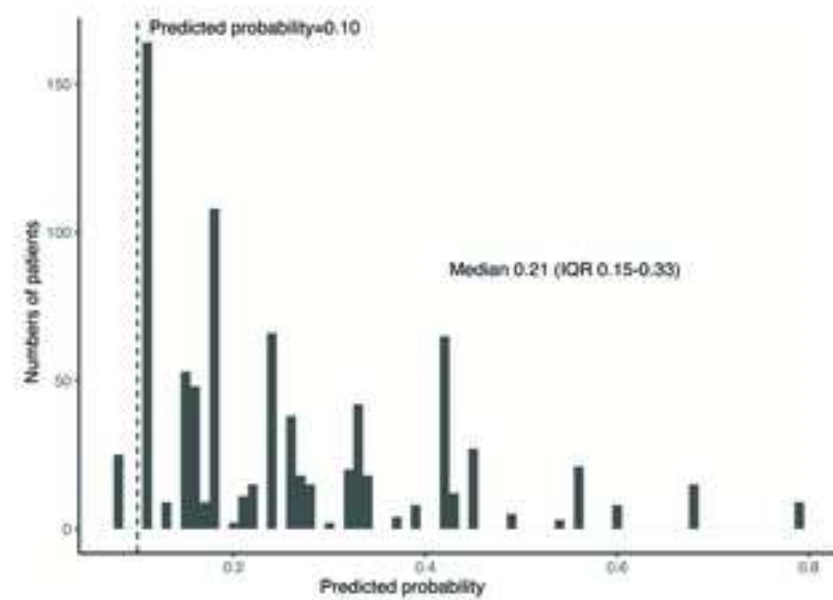


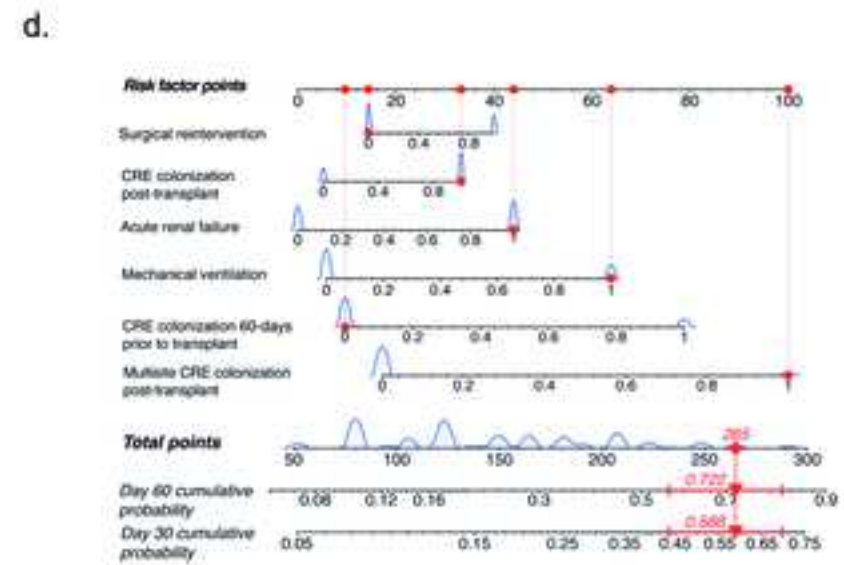
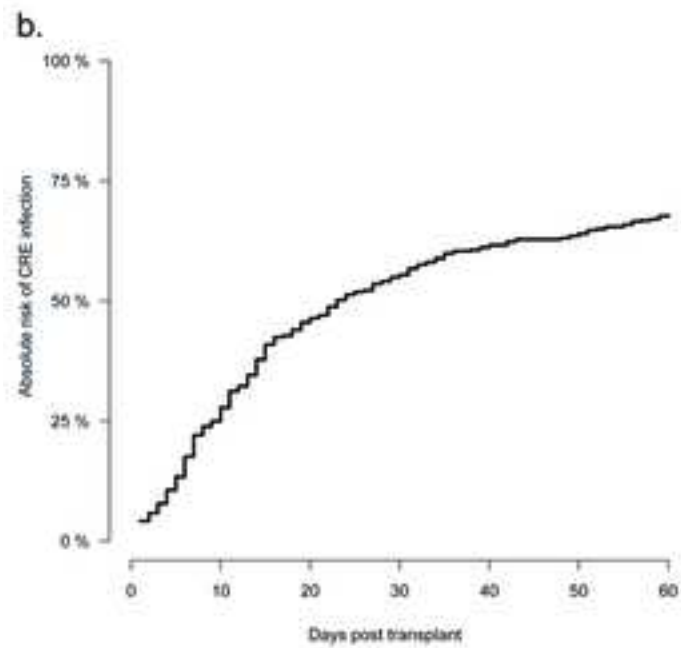
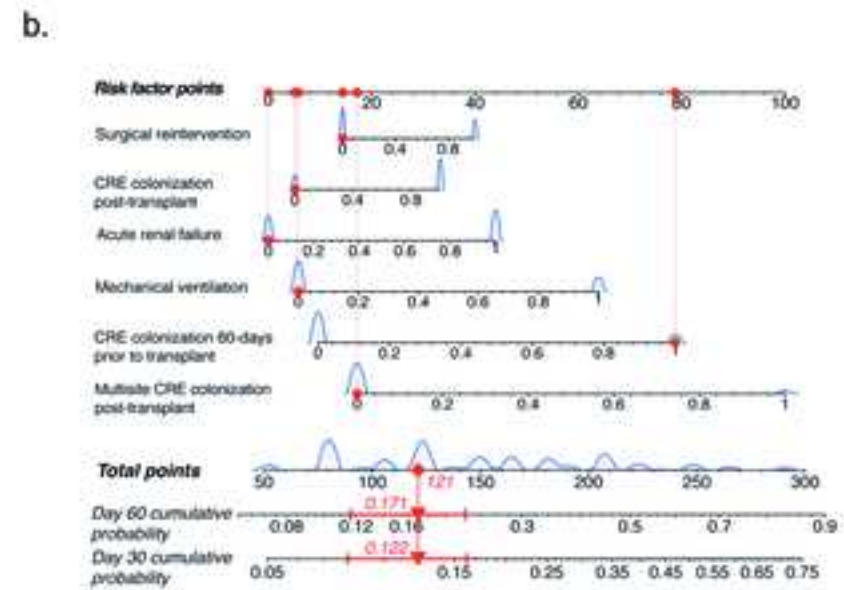
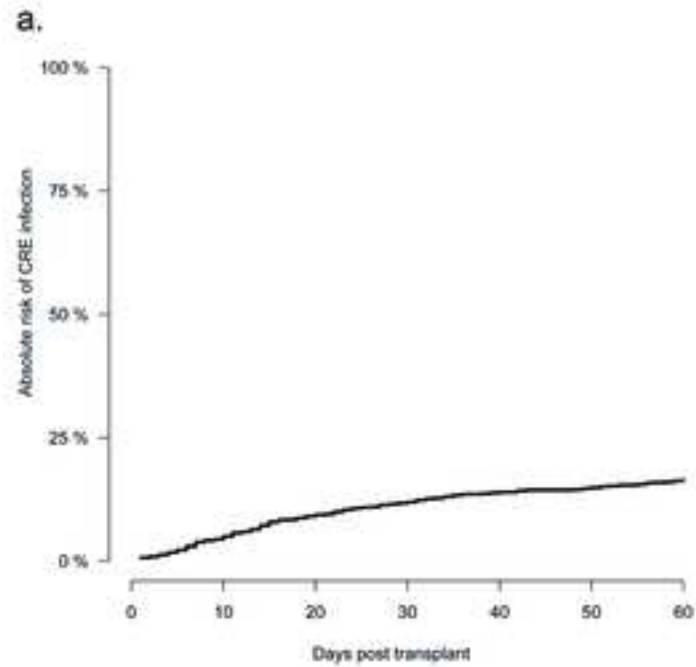


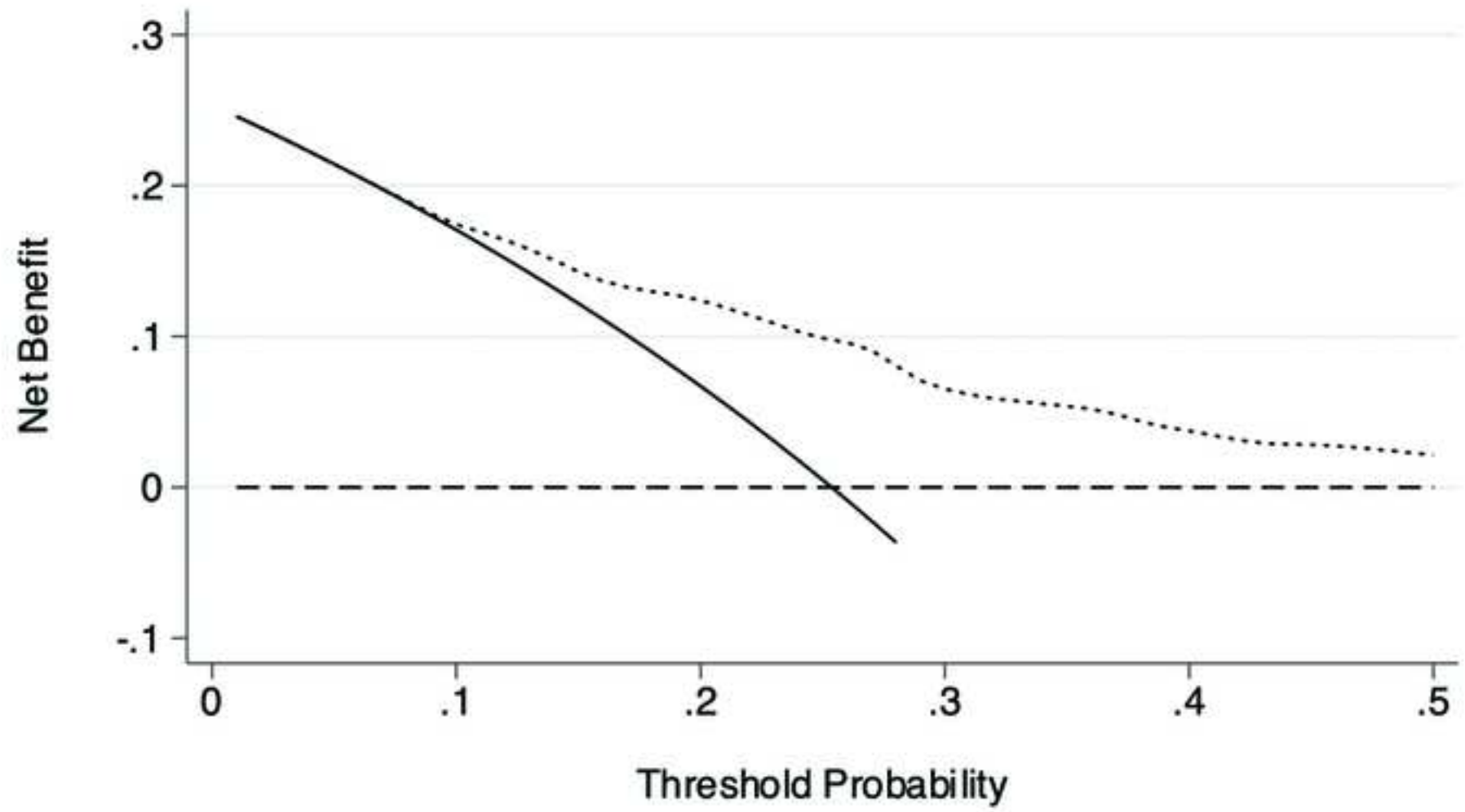
a.



b.







- Net Benefit: All patients receive CRE treatment
- - - Net Benefit: No patients receive CRE treatment
- Model identifies patients for CRE treatment

Supplementary Table 1: Details about participating centers

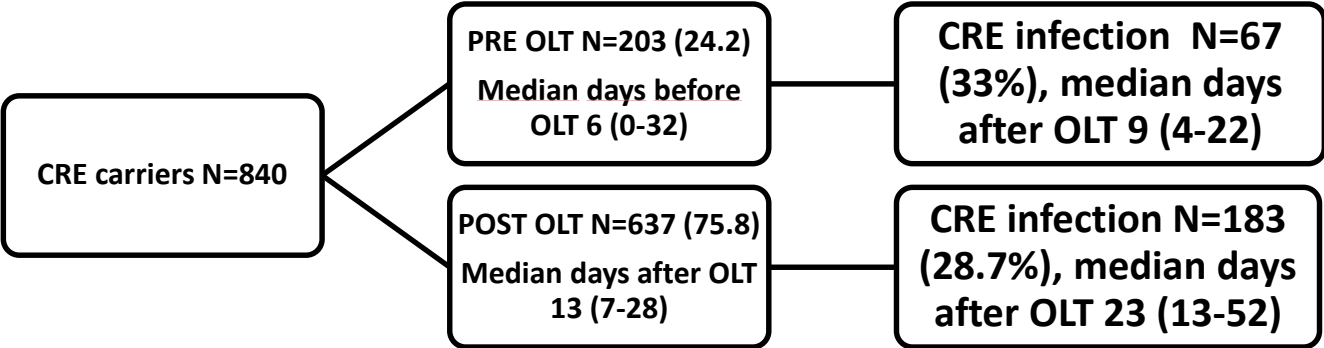
| Center | Year of implementation systematic screening for CRE carriage before and after LT | Microbiology methods for CRE carriage screening | Standard regimen of antibiotic prophylaxis and its duration | Local policy for management of CRE carriers at LT | Local policy for management of CRE carriers after LT |
|---|---|--|--|--|---|
| Policlinico S. Orsola - Bologna | 2010 | Selective culture - molecular test | Amoxicillin/clavulanate 24 hours | No strategy | Antibiotic coverage for CRE only upon to the suspicion of infection |
| IRCCS ISMETT - Palermo | 2013 | Selective culture | Ampicillin + Cefotaxime 48 hours | No strategy | Antibiotic coverage for CRE only upon to the suspicion of infection |
| AOU Città della Salute e della Scienza - Torino | 2014 | Immune assay | Piperacillin/tazobactam 72 hours | Targeted anti-CRE prophylaxis in selected cases | Antibiotic coverage for CRE if complications occur |
| Multivisceral Transplant Unit - Padua | 2013 | Selective culture | III generation cephalosporin 24 hours | No strategy | Antibiotic coverage for CRE only upon to the suspicion of infection |
| Ca' Granda Ospedale Maggiore | 2014 | Selective culture | Piperacillin/tazobactam 24 hours | No strategy | Antibiotic coverage for CRE only upon to the suspicion of infection |

| | | | | | |
|---|------|-------------------|---|---|--|
| Policlinico - Milano | | | | | |
| Azienda Ospedaliero- Universitaria - Modena | 2012 | Molecular test | Ampicillin+Cefotaxime 72 hours | Targeted anti-CRE prophylaxis IN selected cases | Antibiotic coverage for CRE in absence of symptoms if complications occur |
| Presidio Ospedaliero Universitario S. Maria della Misericordia - Udine | 2012 | Selective culture | Piperacillin/tazobactam > 72 hours | Targeted anti-CRE prophylaxis | Antibiotic coverage for CRE if complications occur |
| Hospital das Clinicas da Faculdade de Medicina da Universidade - São Paulo | 2009 | Selective culture | Ampicillin+Cefotaxime 48 hours | Targeted anti-CRE prophylaxis | No strategy |
| Hospital Israelita Albert - São Paulo | 2010 | Selective culture | Cefotaxime + Amoxicillin/clavulanate 48 hours | Targeted anti-CRE prophylaxis | No strategy |

| | | | | | |
|--|------|-------------------|---|----------------------------------|---|
| Quinta D'or Hospital – Rio de Janeiro | 2010 | Selective culture | Ampicillin/sulbactam 24 hours | Targeted anti-CRE prophylaxis | Antibiotic coverage for CRE only upon to the suspicion of infection |
| Hospital Universitario Walter Cantidio - Fortaleza | 2015 | Selective culture | Cefotaxime + Ampicillin 24 hours | Targeted anti-CRE prophylaxis | No strategy |
| Hospital Universitario Puerta de Hierro - Majadahonda | 2009 | Selective culture | Cefotaxime + Amoxicillin/clavulanate 48 hours | No strategy | No strategy |
| Hospital General Universitario Gregorio Maranon - Madrid | 2016 | Selective culture | Cefazolin 48 hours | Targeted anti-CRE prophylaxis | Antibiotic coverage for CRE only upon to the suspicion of infection |
| Jackson Memorial Institute - Miami | 2010 | Selective culture | Ampicillin/sulbactam + Fluconazole 24 hours | Targeted anti-CRE prophylaxis | No strategy |

| | | | | | |
|--|------|----------------|-----------------------|-------------|-------------|
| Rabin Medical Center, Beilinson Hospital – Petah-Tikva | 2013 | Molecular test | Cefazolin 12 hours | No strategy | No strategy |
|--|------|----------------|-----------------------|-------------|-------------|

Supplementary Figure 1. Distribution of CRE infections according to CRE carriage acquisition before or after OLT



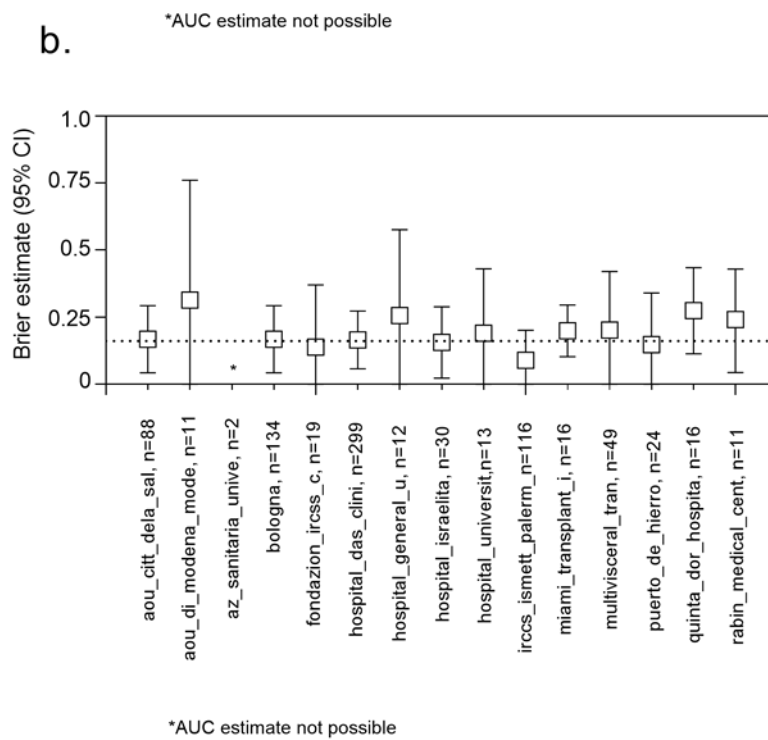
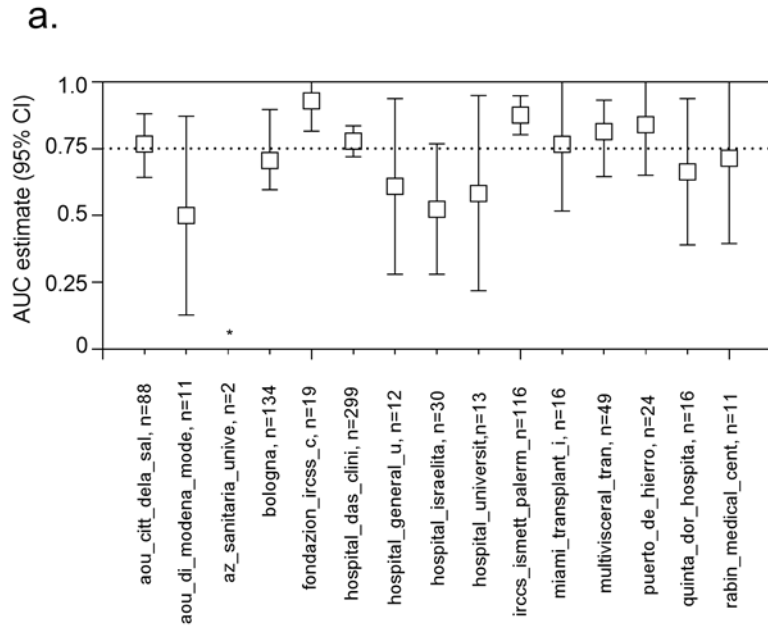
Supplementary Table 2: Characteristics of CRE infections according to time of CRE carriage acquisition.

| | Total N=250 (%) | CRE pre N=67 (%) | CRE post N=183 (%) | p |
|--|----------------------------|-----------------------------|-------------------------------|----------|
| Time to infection | | | | |
| from OLT (days) (median, IQR) | 19 (9-42) | 9 (4-22) | 23 (13-52) | <0.001 |
| from carriage detection (days) (median, IQR) | 10 (3-25) | 19 (8-56) | 8 (3-18) | <0.001 |
| Clinical severity | | | | |
| SOFA score at infection onset (median, IQR) | 8 (5-13) | 11 (5-15) | 6 (4-12) | 0.005 |
| Septic shock | 96 (38.4) | 35 (52.2) | 61 (33.3) | 0.005 |
| Type of infection | | | | |
| BSI | 89 (35.6) | 25 (37.3) | 64 (35) | 0.42 |
| Primary | 34/89 (38.2) | 10 (40) | 24 (37.5) | 0.8 |
| Secondary | 37/89 (41.6) | 11 (44) | 26 (40.6) | 0.77 |
| Catheter-related | 18/89 (20.2) | 4 (16) | 14 (21.9) | 0.54 |
| Pneumonia | 60 (24) | 16 (23.9) | 44 (24) | 0.97 |
| IAI | 56 (22.4) | 9 (13.4) | 47 (25.7) | 0.026 |
| UTI | 22 (8.8) | 4 (6) | 18 (9.8) | 0.25 |
| SSI | 65 (26) | 23 (34.3) | 42 (23) | 0.07 |
| Management of CRE infection | | | | |
| Appropriate empiric therapy | 149 (59.6) | 45 (67.2) | 104 (56.8) | 0.09 |
| Appropriate targeted therapy | 216 (86.4) | 59 (88.1) | 157 (85.8) | 0.41 |
| Treatment duration (days) (median, IQR) | 14 (7-20) | 12 (5-22) | 14 (7-19) | 0.5 |
| Source control | 134 (53.6) | 28 (68.3) | 63 (67.7) | 0.95 |
| Time from CRE infection to source | 2 (0-7) | 3.5 (1-10) | 1 (0-6) | 0.028 |

| | | | | |
|--|------------|------------|------------|-------|
| control (days) (median, IQR) | | | | |
| Outcome | | | | |
| Length of ICU stay (days) (median, IQR) | 14 (5-29) | 14 (6-24) | 13 (5-30) | 0.98 |
| Length of hospital stay after OLT (days) (median, IQR) | 43 (24-76) | 33 (19-49) | 46 (26-85) | 0.002 |
| All-cause 6-months mortality | 145 (58) | 38 (56.7) | 107 (58.5) | 0.46 |
| Hospital re-admissions (median, IQR) | 0 (0-1) | 0 (0-1) | 0 (0-1) | 0.49 |

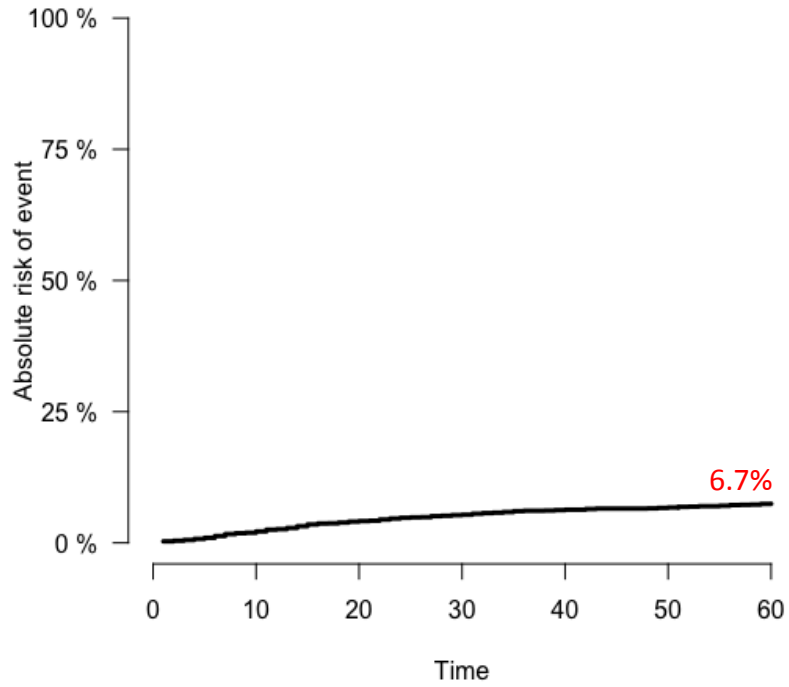
Abbreviations: IQR interquartile range, OLT orthotopic liver transplant, SOFA sequential organ failure assessment, BSI bloodstream infection, IAI intra-abdominal infection, UTI urinary tract infection, SSI surgical site infection, CRE carbapenem-resistant *Enterobacteriaceae*, ICU intensive care unit.

Supplemental Fig 2. Study-site specific CRE risk model (a) discrimination; and (b) prediction accuracy as indicated by c-index (AUC) and the Breir score. AUC> 0.8 indicate good discriminatory accuracy of the mode. are indicative of better model descimination. Lower Brier scores (range 0-1) are indicative of better prediction accuracy

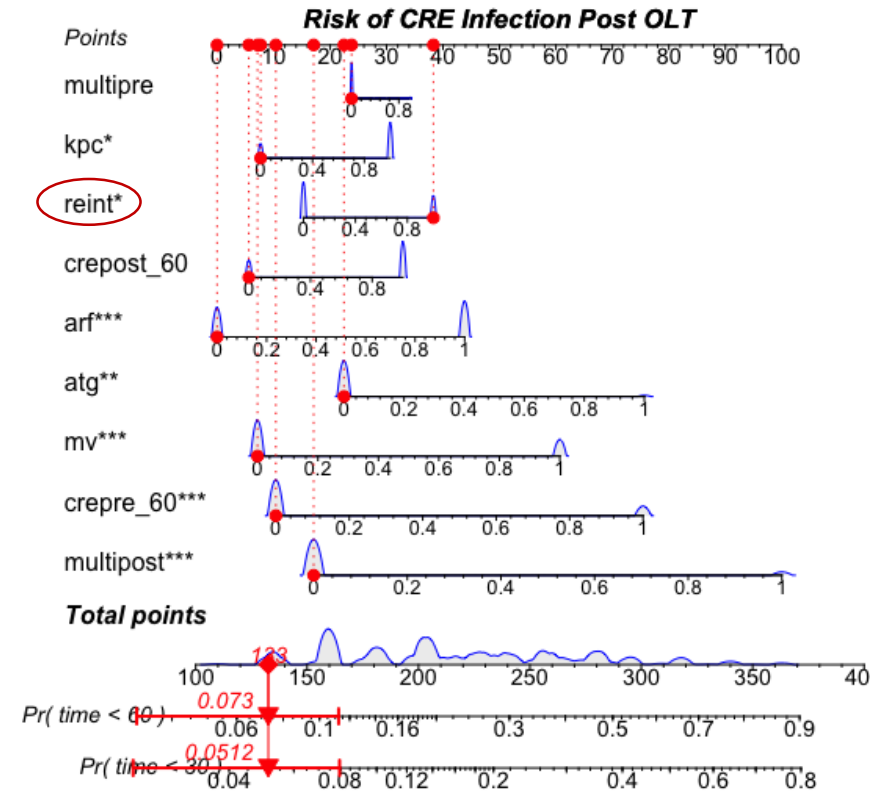


Patient ID #22

Original Fine-gray model



Weighted Cox-regression



Type "0" when risk factor absent; type "1" when risk factor present

atg
0

reint
1

Risk factors
Are entered in
As "0" or "1"

arf
0

mv
0

kpc
0

crepre_60
0

multiple
0

crepost_60
0

multipost
0

Predicted Survival at this Follow Up:

time
60

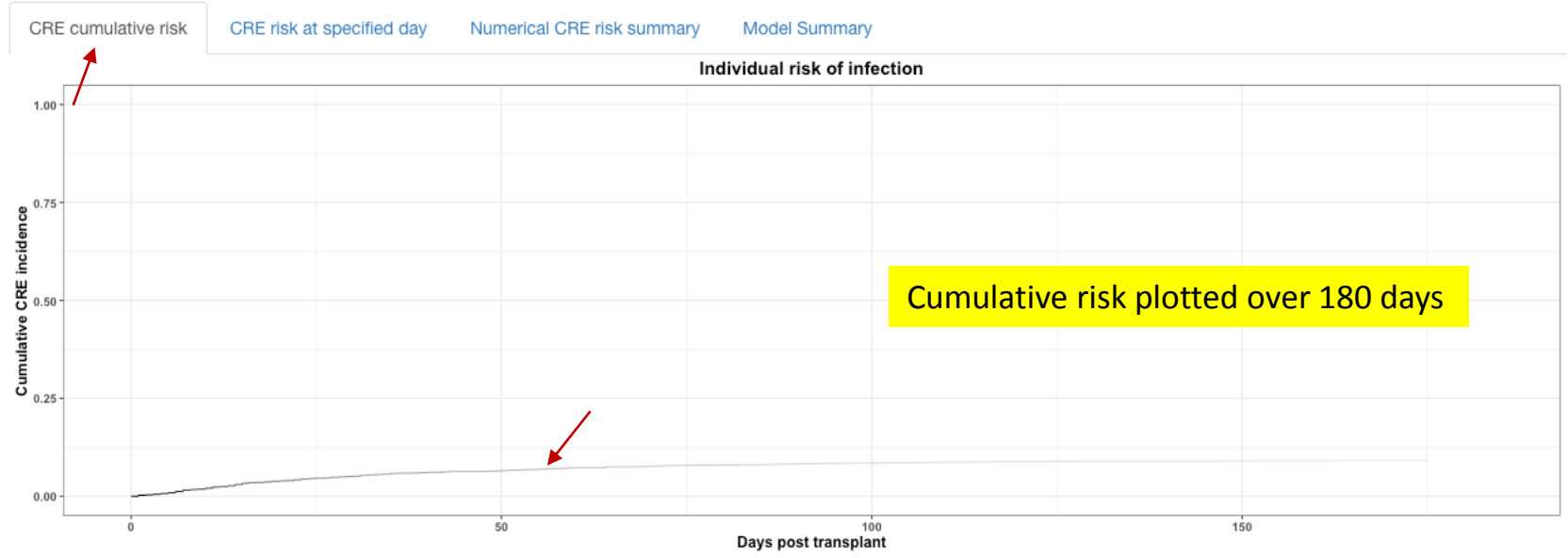
Enter time for specific prediction

Alpha blending (transparency)

Predict

Press Quit to exit the application

Quit



Type "0" when risk factor absent; type "1" when risk factor present

atg

reint

arf

mv

kpc

crepre_60

multiple

crepost_60

multiplast

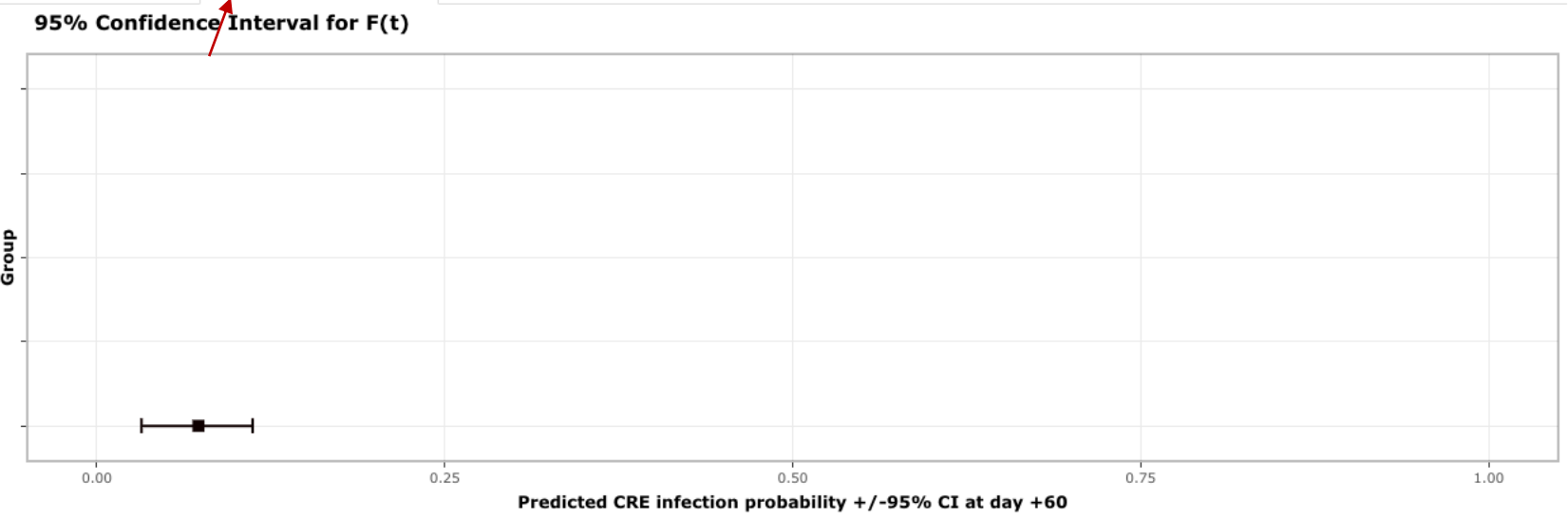
Predicted Survival at this Follow Up:

time

Alpha blending (transparency)

Press Quit to exit the application

CRE cumulative risk CRE risk at specified day Numerical CRE risk summary Model Summary



Risk estimate plotted for Day 60
Note... the confidence intervals are affected by the number of patients with this combination of risk factor, thus the model provides the user an indication of uncertainty in prediction based on the specific combination of risk factors

Type "0" when risk factor absent; type "1" when risk factor present

atg
0

reint
1

arf
0

mv
0

kpc
0

crepre_60
0

multipre
0

crepost_60
0

multiplist
0

Predicted Survival at this Follow Up:

time
60

Alpha blending (transparency)

Predict

Press Quit to exit the application

Quit

CRE cumulative risk CRE risk at specified day **Numerical CRE risk summary** Model Summary

```

=====
time atg reint arf mv kpc crepre_60 multipre crepost_60 multiplist Prediction Lower.bound Upper.bound
-----
1 60 0 1 0 0 0 0 0 0 0.073 0.032 0.112
=====

```

Numeric Risk prediction with 95% CI

Type "0" when risk factor absent; type "1" when risk factor present

atg

reint

arf

mv

kpc

crepre_60

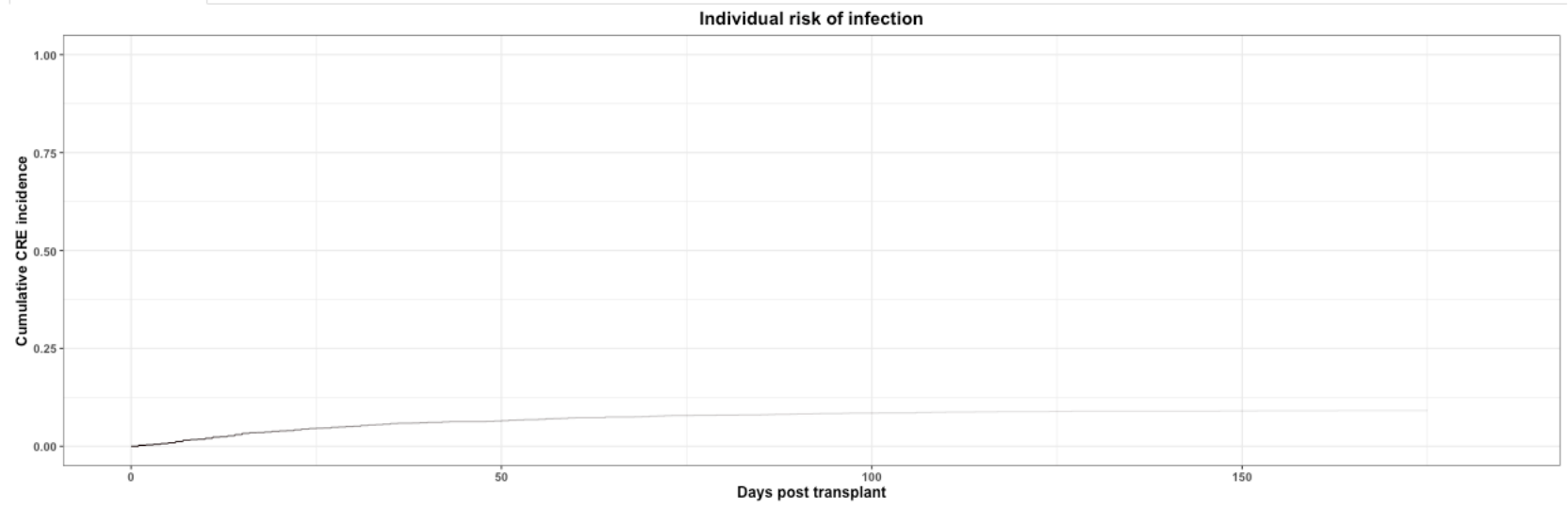
multiple

crepost_60

multiplast

Risk factors
Are entered in
As "0" or "1"

CRE cumulative risk CRE risk at specified day Numerical CRE risk summary Model Summary



Predicted Survival at this Follow Up:

time

You can go back and change the time frame for prediction...

Alpha blending (transparency)

Predict

Press Quit to exit the application

Quit

Type "0" when risk factor absent; type "1" when risk factor present

atg

reint

arf

mv

kpc

crepre_60

multiple

crepost_60

multiplast

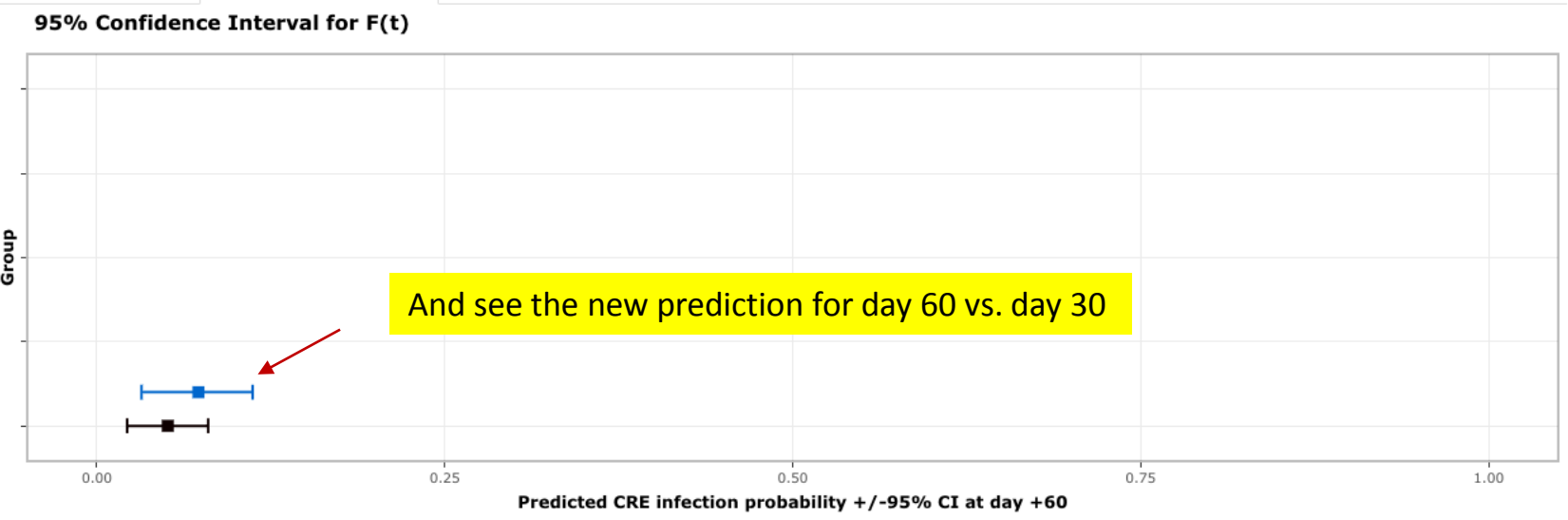
Predicted Survival at this Follow Up:

time

Alpha blending (transparency)

Press Quit to exit the application

[CRE cumulative risk](#) [CRE risk at specified day](#) [Numerical CRE risk summary](#) [Model Summary](#)



Type "0" when risk factor absent; type "1" when risk factor present

atg

reint

arf

mv

kpc

crepre_60

multiple

crepost_60

multiplast

Predicted Survival at this Follow Up:

time

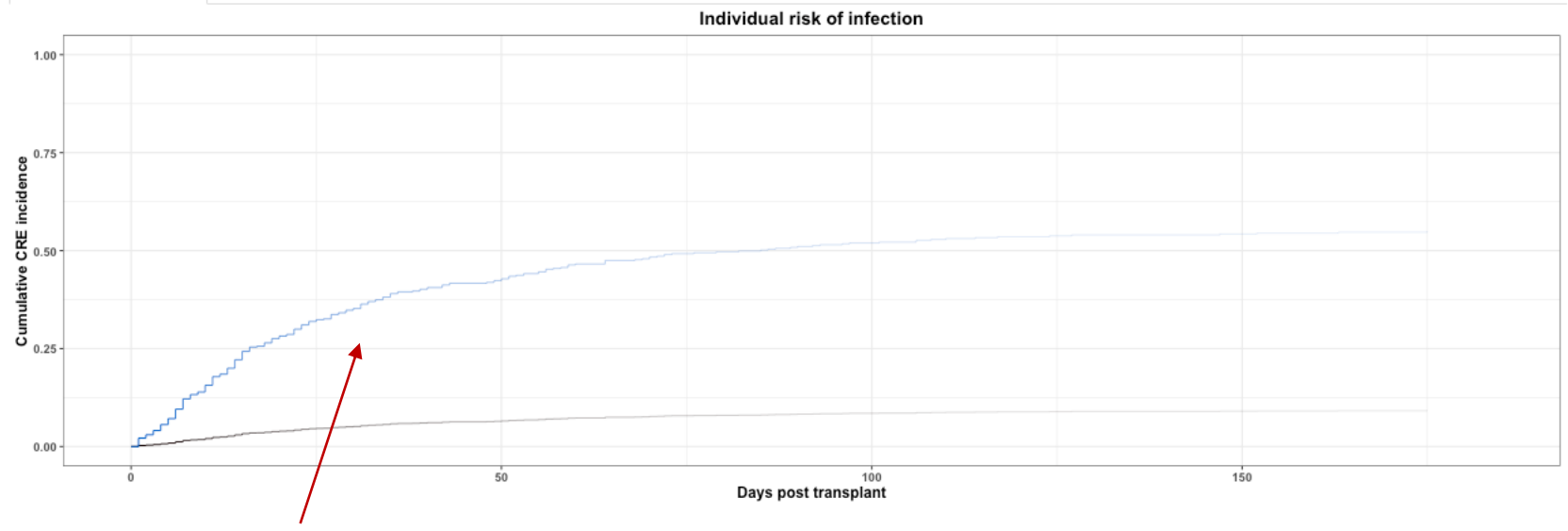
Alpha blending (transparency)

Predict

Press Quit to exit the application

Quit

CRE cumulative risk CRE risk at specified day Numerical CRE risk summary Model Summary



You can also add new risk factors, to see how it affects the prediction...
i..e this »low-risk patient« develops, ARF, goes on mechanical ventilation, and Becomes colonised post OLT with KPC

Type "0" when risk factor absent; type "1" when risk factor present

atg

reint

arf

mv

kpc

crepre_60

multiple

crepost_60

multiplast

Predicted Survival at this Follow Up:

time

Alpha blending (transparency)

Press Quit to exit the application

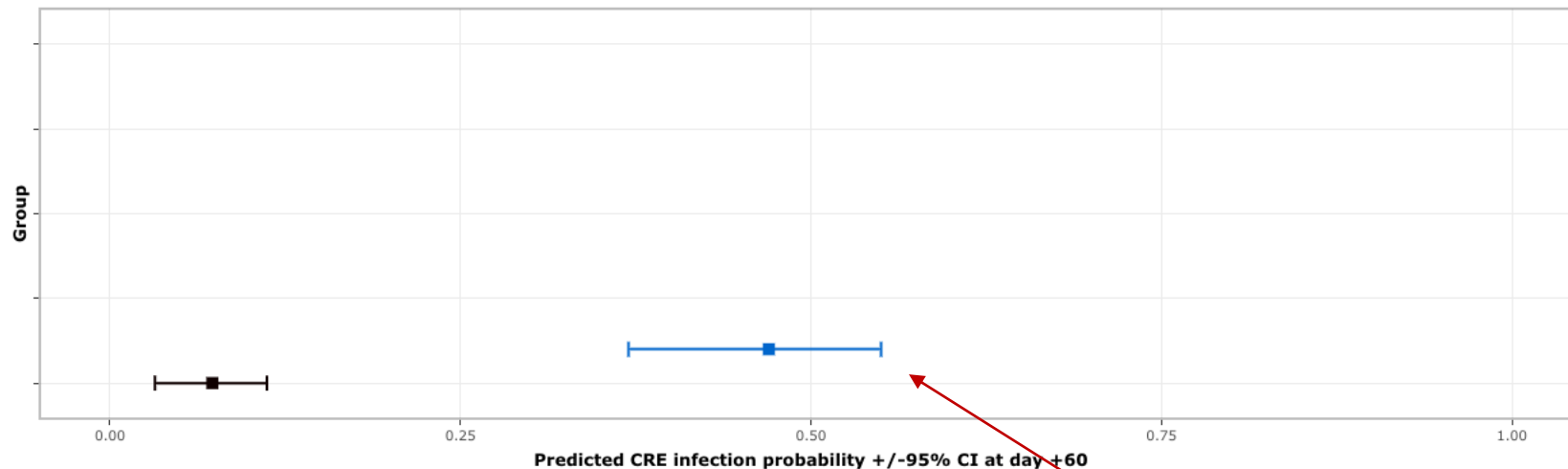
CRE cumulative risk

CRE risk at specified day

Numerical CRE risk summary

Model Summary

95% Confidence Interval for F(t)



Comparing Day CRE prediction from previous single risk factor, and now with additional risk factors (note wider CI as fewer patients in the dataset had this combination of variables)

https://idbologna.shinyapps.io/CRE_POST_OLT/

1 CID-105646-marked

2 **Development of a Risk Prediction Model for Carbapenem-Resistant Enterobacteriaceae Infection**
3 **after Liver Transplantation: A Multinational Cohort Study**

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6 Alberto Ferrarese¹⁰, Patrizia Burra¹⁰, Ruan Fernandes¹¹, Luis Fernando Aranha Camargo¹¹, Angel
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84 **Text word count:** 2996 **Abstract word count:** 249

85 **Running title:** CRE infection risk in liver transplant

86 **Key words:** SOT, liver transplantation, CRE carriage, CRE infection

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98

99 **40-word summary:** One third of CRE carriers develops infection after LT with dramatic impact on
100 survival. Our prediction tool could enable better targeting of early interventions for CRE infection
101 after LT as opposed to universal prophylaxis or treatment in high-prevalence centers.

102 **ABSTRACT**

103 **Background.** Patients colonized with carbapenem resistant Enterobacteriaceae (CRE) are at higher
104 risk of developing CRE infection after liver transplantation (LT) with associated high morbidity and
105 mortality. Prediction model for CRE infection after LT among carriers could be useful to target
106 preventive strategies.

107 **Methods.** Multinational multicenter cohort study of consecutive adult patients underwent LT and
108 colonized with CRE before or after LT, from January 2010 to December 2017. Risk factors for CRE
109 infection were analyzed by univariate analysis and by Fine-Gray sub-distribution hazard model, with
110 death as competing event. A nomogram to predict 30- and 60-day CRE infection risk was created.

111 **Results.** 840 LT recipients found to be colonized with CRE before (n=203) or after (n=637) LT were
112 enrolled. CRE infection was diagnosed in 250 (29.7%) patients within 19 (IQR 9-42) days after LT. **Pre-**
113 **and post-LT colonization, multisite post-LT colonization, prolonged mechanical ventilation, acute**
114 **renal injury, and surgical re-intervention** were retained in the prediction model. Median 30 and 60-
115 day predicted risk was **15%** (IQR **11-24%**) and **21%** (IQR **15-33%**), respectively. Discrimination and
116 prediction accuracy for CRE infection was acceptable on derivation (AUC **74.6**, Brier index **16.3**) and
117 bootstrapped validation dataset (AUC **73.9**, Brier index **16.6**). Decision-curve analysis suggested net
118 benefit of model-directed intervention over default strategies (treat all, treat none) when CRE
119 infection probability exceeded 10%. The risk prediction model is freely available as mobile
120 application at <https://idbologna.shinyapps.io/CREPostOLTPredictionModel/>.

121 **Conclusions.** Our clinical prediction tool could enable better targeting interventions for CRE infection
122 after transplant.

123

124 **INTRODUCTION**

125 The emergence and spread of carbapenem resistant *Enterobacteriaceae* (CRE) over the last two
126 decades threatens the safety of patients undergoing life-saving surgical procedures or necessary
127 immunosuppressive treatments [1]. Solid organ transplant (SOT) recipients, in particular those
128 undergoing liver transplantation (LT), are at increased risk of colonization with CRE due to repeated
129 contacts with health care system, exposure to broad spectrum antibiotics, complexity of surgery,
130 and altered immune status [2]. Compared with other settings (e.g., geriatric units, long-term care
131 facilities) the incidence of infection per CRE colonization days is higher in SOT recipients [3], and
132 associated with poorer survival [4]. In a prospective multicentre study of 887 SOT recipients in Italy,
133 mortality was more than 10 times higher in patients with positive cultures for CRE [5].

134 Knowledge of CRE infection risk factors is essential to develop targeted preventive strategies. Risk
135 factors for CRE infection after LT were analysed in two large single-centre cohort studies, from Brazil
136 and Italy, including 386 and 553 patients, respectively [6, 7]. In both studies CRE carriage, either
137 detected before or after transplant, was found to be the strongest predictor of subsequent CRE
138 infection. Based on these findings, some investigators have proposed that preventive strategies
139 should be implemented in all CRE carriers undergoing LT [8]. However, a more accurate prediction of
140 which carriers are at higher risk could help to target preventive measures or timely appropriate
141 antibiotic therapy in patients at lower risk (i.e. < 10%) of CRE infection.

142 With this objective in mind, we performed a multicentre, multinational cohort study of patients
143 undergoing LT colonized with CRE before or after transplantation, with the aim of developing a
144 multivariable prediction model for CRE infection within 30 and 60 days of LT.

145

146 **MATERIAL AND METHODS**

147 *Study design*

148 We performed a multicentre multinational retrospective cohort study. The enrolment period was
149 between January 2010 to December 2017, or from the onset of systematic CRE colonization
150 screening in the centre to December 2017 (see supplementary Table 1). Follow-up was of 180 days
151 after LT. Data were accrued from October 2019 to June 2020. Data sources were clinical charts and
152 hospital electronic records, they were de-identified before entry into a standardized electronic case
153 report form (eCRF), and managed using REDCap capture tool hosted by Alma Mater University of
154 Bologna [9]. Collected data were periodically checked for accuracy by an investigator of the
155 coordinating centre (MR). Queries for incongruous or missing data were submitted to investigators
156 to ensure high quality and completeness. The study was first approved by Institutional Review Board
157 (IRB) of the promoting centre (n. 155/2019/Oss/AOUBo on March 20, 2019), then by IRB of all
158 participating centres.

159 *Setting*

160 Fifteen hospitals performing LT participated in the study: seven from Italy (Bologna, Modena, Turin,
161 Padua, Palermo, Milan, and Udine); four from Brazil (2 in São Paulo, 1 in Fortaleza and 1 in Rio de
162 Janeiro); two from Spain (Madrid and Majadahonda); one from United States (Miami); and one from
163 Israel (Petah-Tikva) (see Supplementary Table 1 and Figure 1). An active surveillance screening for
164 CRE colonization was required by the study protocol. All centres performed systematic screening of
165 CRE carriage by rectal swab (RS), at inclusion in waiting list, at LT and weekly after LT until hospital
166 discharge. Screening for CRE colonization at other sites was performed according to clinical
167 judgment and local policy.

168 *Study population*

169 All consecutive adult (≥ 18 years) patients who underwent LT during the study period and were found
170 to be colonized with CRE were enrolled. Patients were included only once at the time of the first CRE
171 carriage detection. CRE was defined as any *Enterobacteriaceae* displaying *in vitro* non-susceptibility
172 to any of the carbapenems according to the criteria (CLSI or EUCAST) adopted at the participating

173 centre during the study period. The colonization status was defined as isolation of CRE from RS or
174 other samples other than blood cultures or sterile fluids (e.g., urine, respiratory samples, superficial
175 skin samples) in absence of symptoms and signs of infection. Multisite colonization was defined
176 when CRE was concomitantly isolated from more than one of such samples. Patients found to be
177 colonized with CRE by samples obtained at inclusion in the waiting list, during the period in waiting
178 list, or at the time of LT were considered as CRE carriers pre-LT, whereas those found to be colonized
179 within 180 days after LT were considered as post-LT CRE carriers.

180 *Predictors of CRE infection after LT*

181 The primary endpoint was the time to development of CRE infection, diagnosed within 180 days
182 after LT, according to Center for Disease Control criteria [10]. The assessment of CRE infection was
183 made by the local investigator and revised by an investigator of the promoting centre (MR), in case
184 of no agreement a third blinded investigator of the promoting centre (MG) was asked to review the
185 case for establishing the final diagnosis. Infection severity was determined using SOFA score and
186 septic shock criteria [11].

187 Candidate prediction variables included: demographic data (age and sex); comorbidities according to
188 Charlson index; underlying liver disease, and severity of liver disease according to Model for End
189 stage Liver Disease (MELD) at inclusion in waiting list and at LT. Complications occurred within 90
190 days before LT were recorded and included: admission to intensive care unit (ICU); grade 3 and/or
191 refractory ascites [12]; hepato-renal syndrome (HRS) [12]; gastrointestinal bleeding; hepatic
192 encephalopathy [13]; development of acute-on-chronic liver failure (ACLF) [14]; any infection,
193 candidemia, and *Clostridioides difficile* infection. For graft characteristics donor age, cold ischemia
194 time, and combined transplant were collected. Intraoperative variables included: antibiotic
195 prophylaxis, biliary anastomosis, bleeding with need of massive transfusion (≥ 40 units of cellular
196 blood products), prolonged intervention (≥ 8 hours). Complications occurred from LT to the
197 diagnosis of CRE infection or death or 180 days (whichever occurred first) were recorded and

198 included: re-intervention, acute kidney injury (AKI) according to KIDGO criteria [15], renal
199 replacement therapy (RRT), prolonged (≥ 48 hours) mechanical ventilation (MV), graft dysfunction
200 (primary or secondary), biopsy-proven rejection, re-transplantation, and CMV DNAemia $>100,000$
201 copies/ml [16]. Management of CRE colonization was recorded according to the following
202 categories: no strategy adopted, anti-CRE surgical prophylaxis, decolonization with non-absorbable
203 antibiotics, faecal microbiota transplantation, pre-emptive strategy (defined as early start of anti-
204 CRE coverage upon to clinical deterioration). For outcome, duration of the hospital stay after LT,
205 number of re-admissions within 180-days after LT, and all-cause 180-days mortality were recorded.

206 *Sample size*

207 Sample size required for the clinical prediction model was estimated for a time-to-event outcome
208 using the pmsampsize package in R (version 4.02, Core Team (2020). **R**: A language and environment
209 for statistical computing. **R** Foundation for Statistical Computing, Vienna, Austria) using the methods
210 proposed by Riley et al. [17]. We estimated that 850 patients with 213 outcome events (event rate
211 0.25 by day 180) would be sufficient to evaluate 20 candidate predictors (>10 events per candidate
212 predictor) with an estimated 60-day event rate of 0.17.

213 *Missing data*

214 A complete-case analysis was performed.

215 *Statistical analysis methods*

216 Categorical variables were expressed as absolute numbers and their relative frequencies. Continuous
217 variables were expressed as mean \pm standard deviation (SD) if normally distributed, or as median
218 and interquartile range (IQR) if non-normally distributed. We used the Fine-Gray sub-distribution
219 hazard model to develop a clinical prognostic index for CRE infection over the first 180 days after LT
220 with death as competing event.

221 Although colonization status has been modelled in previous studies as a time-varying covariate [18],
222 inclusion of the time-dependent covariates limits the ability of the Fine-Gray sub-distribution hazard
223 model to make inferences about the effects of covariates on the cumulative incidence function (CIF)
224 in the presence of competing risks, which is an underlying motivation for our model. Furthermore,
225 simulation studies have shown that a sub-distribution approach for time-dependent covariates often
226 produces misleading results [19], and is not currently recommended [20]. Therefore, we focused on
227 colonization status as a risk factor only in the 60-days prior to and 60 days following OLT to minimize
228 time-dependent effects.

229 Using a base model of CRE colonization within 60 days prior to transplant, we examined the degree
230 to which including each other variable from our original model improved its performance. Additional
231 candidate variables were selected *a priori* based on statistical significance in univariate analysis,
232 published data and clinical experience and whether they improved prediction accuracy of the
233 baseline model. Only variables available for assessment during the risk period were included. We
234 assumed that a predictive model for CRE infection would have the most clinical utility for accurate
235 prediction of CRE infection in the first 30 and 60 days post OLT, as the risk appeared to plateau after
236 this period. Therefore we used C-index (AUC) to assess the discrimination of the model with
237 censored CRE infection status, and calibration plots of day-60 observed versus expected absolute
238 risks with the Brier score to assess the accuracy of predictions. To control for centre effect, the
239 analysis was also performed for each participating centre. Model discrimination and calibration
240 measures were calculated using 100-fold cross validation bootstrap resampling to avoid bias. A
241 decision curve was prepared to examine potential model utility across a range of CRE infection
242 probabilities (0-60%) at 60 days post LT. All analyses were performed using R version 4.02 with
243 riskRegression, prodlim, ggplot2, survival, and rmda packages. The final regression model was
244 visualized by preparing a nomogram to predict 30-day and 60-day risk using methods described by
245 Zhang et al [21] and the regplot package. An online prediction tool (Shiny App) was prepared using
246 the DynNom package in R. Results were reported according to TRIPOD guidelines [22].

247

248 **RESULTS**

249 Overall, 840 LT recipients colonized with CRE before or after transplantation were enrolled, with 250
250 episodes of CRE infection post LT (29.7%) during 104,596 patient days of follow-up. Distribution of
251 patients and of CRE infections across participating centres is shown in Figure 1. The characteristics of
252 study population are shown in Table 1. Median age was 55 (IQR 46-62) years, male sex (65.4%). Viral
253 hepatitis was the primary indication for LT in 376/840 subjects (44.8%), followed by alcohol 207/840
254 (24.6%). Over one-quarter 247/840 (29.4%) of patients had hepatocellular carcinoma (HCC). MELD at
255 inclusion in waiting list and at LT was 19 (IQR 14-25) and 23 (IQR 16-30), respectively. CRE carriage
256 was detected before or at LT in 203 (24.2%) patients, the median time from detection to LT was 6
257 (IQR 0-32) days. The remaining 637 (75.8%) patients were found to be colonized with CRE within a
258 median of 13 (IQR 7-28) days after LT. Comparison of pre- and post-LT carriers showed several
259 differences (see Table 1). Specifically, pre-LT carriers presented with more severe liver disease and
260 complicated clinical course before transplantation. CRE infection rate among pre- and post-LT
261 carriers was 33% and 28.7%, and the median time to infection following LT was 9 (IQR 4-22) and 23
262 (IQR 13-52) days, respectively (see Supplementary Figure 1). The median time from detection of CRE
263 carriage to the diagnosis of CRE infection was shorter for post-LT carriers (8 vs. 19 days, $p<0.001$).
264 Among the 203 pre-OLT carriers, 51 received targeted perioperative prophylaxis, the rate of CRE
265 infection was 31.4% and 33.6% among those exposed and unexposed to targeted prophylaxis
266 ($p=0.90$), respectively. Post-LT CRE infections in pre-LT carriers presented with higher severity
267 according to median SOFA and septic shock rates than in post-LT carriers. However, there was no
268 difference in clinical outcome (all-cause 180 days mortality 56.7% vs. 58.5%, $p=0.46$), data shown in
269 Supplementary Table 2.

270 To screen risk factors for CRE infection, we performed univariate analysis of patients with and
271 without CRE infection as shown in Table 2. Considering the competing risks of death, a Fine-Gray

272 model was used to perform regression of risk factors to estimate their association with the
273 cumulative incidence of CRE infection up to 180 days post LT. Baseline cumulative incidence of
274 death versus CRE infection over 180 following OLT is shown in Figure 2. The following variables were
275 retained in the final model: pre-LT CRE colonization, post-LT CRE colonization, multisite post-LT
276 colonization, prolonged MV, AKI, and surgical re-intervention (Table 3). The fitted regression risk
277 model was then rendered as a nomogram (Figure 3). Model-predicted cumulative probabilities of
278 developing CRE infection at day 30 and 60 post LT are shown in Figure 4. The median 30 and 60-day
279 predicted risk was 15% (IQR 11-24%) and 21% (IQR 15-33%), respectively. Nearly one-quarter
280 (23.5%) of patients at day 30, and 3% of patients at day 60 had a cumulative predicted risk of CRE
281 infection $\leq 10\%$. The model showed acceptable 60-day discrimination and prediction accuracy for
282 CRE infection when assessed against the derivation AUC 74.6 (95% 70.9-78.4), Brier index 16.3
283 (95%CI 6.7-25.9) and bootstrapped validation dataset AUC 73.9 (95%CI 67.7-79.1), Brier index 16.6
284 (95%CI 14.6-19.1) (Figure 5). Discrimination and calibration were also evaluated by centre and were
285 consistent with the overall performance, except for higher variability and lower AUCs in centres with
286 lower number of recruited patients (see Supplementary Figure 2). The score was designed to be
287 used in the immediate peri-transplant period, ideally from the day of transplantation up to 2-3
288 weeks after transplantation. During this period, the patient should be frequently evaluated for
289 carriage status, multisite colonization, prolonged MV, AKI, and/or re-intervention. According with
290 the presence of such variables the cumulative risk of CRE infection within 30 and 60 days after LT can
291 be predicted (see examples in Figure 6 and mobile application at
292 <https://idbologna.shinyapps.io/CREPostOLTPredictionModel/>, instructions for using app are further
293 provided as supplementary material).

294 Finally, we explored the potential clinical utility of our model using a decision-curve analysis (DCA) to
295 examine the “net benefit” of applying the prediction model across a range of CRE infection threshold
296 probabilities [23]. A theoretical risk-model guided strategy (i.e. empiric administration of CRE-active
297 antibiotics) is compared against two default strategies- “treat all” and “treat none.” A model is only

298 useful at a given disease threshold if it has a higher net benefit than treat all or treat none. Our
299 analysis suggested that the model-directed interventions for CRE post LT would show net benefit
300 over default strategies when the overall CRE infection threshold probability exceeded 10% (Fig. 7).

301

302 **DISCUSSION**

303 Assessing a large multinational cohort of LT recipients colonized with CRE (n=840) we have found
304 that one third developed CRE infection after LT. This rate was similar among pre- and post-LT CRE
305 carriers. Most infection episodes occurred within 2 months after LT. The time to infection was earlier
306 for pre-LT CRE carriers, and the clinical severity was higher in this group. All-cause 6-month mortality
307 among carriers was as high as 30%, with similar rates between pre- and post-LT CRE carriers.
308 However, it reached 58% among patients who developed CRE infection.

309 We confirmed that the CRE carriage status is strongly associated with the risk of CRE infection after
310 LT, either if detected before or after LT [6, 7]. However, we also confirmed our previous hypothesis
311 that, given a probability of CRE infection higher than 10%, a strategy of treating all CRE carriers could
312 be inferior to a targeted strategy in terms of efficacy, as well as safety, considering potential adverse
313 events of anti-CRE drugs and the risk for further selection pressure. A longitudinal risk stratified
314 approach, could be useful for several reasons. In general, LT is not an elective surgery, thus it could
315 be difficult to implement a strategy before transplantation other than in a minority of patients
316 known to be colonized at the time of inclusion or while on waiting list. Another opportunity for
317 intervention could be surgical prophylaxis. However, data about colonization is frequently available
318 only during the immediate post-operative period. Finally, in most patients CRE colonization becomes
319 evident in the first weeks after LT, typically with a complicated post-operative course, which also
320 increases the risk of infection.

321 Along with pre- and post-LT colonization status, multisite colonization after LT was a predictor of
322 infection. The relationship between multisite colonization with infection development has been
323 already observed in other settings [3]. It is indicative of high colonization burden leading to an
324 increased risk of CRE infection [24]. Indeed, its inclusion in our prognostic model improved model
325 performance. Nevertheless, multisite colonization was not systematically screened in this study.
326 Therefore, we cannot rule out that colonization from different sites may be weighted differently in
327 the model if this risk factor had been systematically screened, as patients with a more complicated
328 course may have been cultured more frequently.

329 The other complications included in our score are widely described as factors involved in the
330 increase risk of infection after LT [25]. AKI has been also associated with microbiome dysbiosis that
331 could favour prolonged CRE carriage and infection [26]. As for MV, it is worth mentioning that we
332 considered such variable, as a qualitative data, from LT to CRE infection, death or end of follow-up. A
333 more detailed information about timing and duration of MV may improve the model performance.

334 Our study has several limitations. First, the retrospective design could have limited identification of
335 all CRE carriers. However, the presence of a local prospective registry of CRE carriers and/or good
336 collaboration with local microbiology laboratories reduced this bias. Heterogeneity in local assays
337 and criteria for identification and determination of CRE could exist. However, obtained data are in
338 line with international literature. The majority of involved centres were high endemic CRE hospitals,
339 with KPC as main mechanism of CRE. Thus, our findings could not apply to transplant centres with
340 low prevalence of CRE, or with KPC alternative main mechanisms among CRE. Finally, we analysed
341 the CRE infection risk only in colonized patients, thus the performance of our prediction tool should
342 be confirmed in unselected LT patients.

343 To conclude, one third of CRE carriers develop infection after LT with a dramatic impact on patient
344 survival. Effective preventive or pre-emptive strategies are needed. Our clinical prediction tool could
345 enable better targeting of early interventions for CRE infection after transplant as opposed to

346 universal prophylaxis or treatment in high-prevalence centres. Further studies are needed to further
347 validate the model and establish which strategy would be more effective, in terms of adverse events
348 and collateral resistance impact and, ultimately patient survival after transplantation.

349

350

351 **ACKNOWLEDGMENTS/FUNDING**

352 No funding was received for this project.

353 **CONFLICT OF INTEREST**

354 Authors have no conflict of interest related to the present study.

355

356

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438

439 **Table 1.** Characteristics of study population according to the timing of CRE carriage detection

| | Total N=840 (%) | pre-OLT carrier N=203 (%) | post-OLT carrier N=637 (%) | P |
|--|----------------------------|--|---|----------|
| Demographic data | | | | |
| Age (years) [median (IQR)] | 55 (46-62) | 53 (42-60) | 56 (47-62) | 0.005 |
| Sex, male | 549 (65.4) | 132 (65) | 417 (65.5) | 0.9 |
| Comorbidities | | | | |
| Myocardial infarction | 23 (2.7) | 5 (2.5) | 18 (2.8) | 0.827 |
| Congestive heart failure | 29 (3.5) | 7 (3.4) | 22 (3.5) | 0.9 |
| Peripheral vascular disease | 33 (3.9) | 6 (3) | 27 (4.2) | 0.4 |
| Cerebrovascular disease | 22 (2.6) | 8 (3.9) | 14 (2.2) | 0.18 |
| COPD | 46 (5.5) | 6 (3) | 40 (6.3) | 0.07 |
| Connective tissue disease | 9 (1.1) | 3 (1.5) | 6 (0.9) | 0.38 |
| Peptic ulcer disease | 62 (7.4) | 20 (9.9) | 42 (6.6) | 0.09 |
| Diabetes without organ damage | 147 (17.5) | 38 (18.7) | 109 (17.1) | 0.3 |
| Moderate/severe renal disease | 82 (9.8) | 28 (13.8) | 54 (8.5) | 0.03 |
| Diabetes with organ damage | 76 (9) | 15 (7.4) | 61 (9.6) | 0.2 |
| Any tumor within 5 years | 271 (32.3) | 45 (22.2) | 226 (35.5) | 0.001 |
| Moderate/severe liver disease | 820 (97.6) | 199 (98) | 621 (97.5) | 0.79 |
| Charlson index [median (IQR)] | 5 (4-7) | 5 (3-6) | 5 (4-7) | 0.004 |
| Underlying liver disease | | | | |
| Viral hepatitis | 376 (44.8) | 74 (36.5) | 302 (47.4) | 0.006 |
| Alcohol | 207 (24.6) | 55 (27.1) | 152 (23.9) | 0.2 |
| Metabolic disease | 69 (8.2) | 12 (5.9) | 57 (8.9) | 0.1 |
| Autoimmune disease | 41 (4.9) | 12 (5.9) | 29 (4.6) | 0.45 |
| Fulminant hepatitis | 52 (6.2) | 7 (3.4) | 45 (7.1) | 0.04 |
| Hepatocellular carcinoma | 247 (29.4) | 37 (18.2) | 210 (33) | <0.001 |
| Prior transplant | 39 (4.6) | 20 (9.9) | 19 (3) | <0.001 |
| MELD at waiting list inclusion (median, IQR) | 19 (14-25) | 21 (16-28) | 19 (14-24) | <0.001 |
| MELD at OLT (median, IQR) | 23 (16-30) | 27 (21-32) | 22 (15-29) | <0.001 |
| Variables relative to the last 90 days before OLT | | | | |
| Hepato-renal syndrome | 235 (28) | 88 (43.3) | 147 (23.1) | <0.001 |
| Ascites grade III/refractory | 385 (45.8) | 108 (53.2) | 277 (43.5) | 0.01 |
| GI bleeding | 138 (16.4) | 43 (21.2) | 95 (14.9) | 0.025 |
| Encephalopathy | 343 (40.8) | 96 (47.3) | 247 (38.8) | 0.02 |
| ICU admission | 161 (19.2) | 70 (34.5) | 91 (14.3) | <0.001 |
| ACLF | 98 (11.7) | 44 (21.7) | 54 (8.5) | <0.001 |
| Any infection | 326 (38.8) | 137 (67.5) | 189 (29.7) | <0.001 |
| Intraoperative variables | | | | |
| Combined transplant | 45 (5.4) | 12 (5.9) | 33 (5.2) | 0.7 |
| Donor age | 51 (36-65) | 50 (35-65) | 51 (36-65) | 0.4 |
| Cold ischemia time (hours) (median, IQR) | 7 (6-8) | 7 (6-8) | 7 (6-8) | 0.62 |
| Biliary anastomosis | | | | |
| Duct to duct without Kehr | 382 (45.5) | 84 (41.4) | 298 (46.8) | 0.19 |
| Duct to duct with Kehr | 347 (41.3) | 76 (37.4) | 271 (42.5) | 0.20 |

| | | | | |
|--|-------------|------------|------------|--------|
| Choledochojejunostomy | 111 (13.2) | 43 (21.2) | 68 (10.7) | <0.001 |
| Intraoperative bleeding | 294 (35) | 88 (43.3) | 206 (32.3) | 0.005 |
| Prolonged surgery (>8 hours) | 248 (29.5) | 70 (34.5) | 178 (27.9) | 0.046 |
| Induction regimen | | | | |
| None | 87 (10.4) | 24 (11.8) | 63 (9.9) | 0.51 |
| Bolus of steroids | 701 (83.5) | 171 (84.2) | 530 (83.2) | 0.75 |
| Antithymocyte globulins | 26 (3.1) | 3 (1.5) | 23 (3.6) | 0.09 |
| Basiliximab | 167 (19.9) | 37 (18.2) | 130 (20.4) | 0.55 |
| Rituximab | 14 (1.7) | 0 (0) | 14 (2.2) | 0.02 |
| Maintenance regimen | | | | |
| Steroids | 612 (72.9) | 162 (79.8) | 450 (70.6) | 0.011 |
| Cyclosporin | 42 (5) | 8 (3.9) | 34 (5.3) | 0.28 |
| Tacrolimus | 770 (91.7) | 181 (89.2) | 589 (92.5) | 0.15 |
| Micophenolate mophetil | 287 (34.2) | 70 (34.5) | 217 (34.1) | 0.9 |
| Post-operative complications | | | | |
| Acute renal injury | 462 (55) | 118 (58.1) | 344 (54) | 0.33 |
| Renal replacement therapy | 299 (35.6) | 90 (44.3) | 209 (32.8) | 0.003 |
| Mechanical ventilation ≥48 hours | 266 (31.7) | 64 (31.5) | 202 (31.7) | 0.96 |
| PGNF | 128 (15.2) | 28 (13.8) | 100 (15.7) | 0.58 |
| Re-intervention | 322 (38.3) | 74 (36.5) | 248 (38.9) | 0.56 |
| Re-transplantation | 79 (9.4) | 11 (5.4) | 68 (10.7) | 0.025 |
| Rejection | 152 (18.1) | 30 (14.8) | 122 (19.2) | 0.17 |
| CMV treatment | 225 (26.8) | 46 (22.7) | 179 (28.1) | 0.15 |
| Infections other than CRE | 479 (57) | 106 (52.2) | 373 (58.6) | 0.12 |
| <i>Clostridioides difficile</i> infection | 37 (4.4) | 8 (3.9) | 29 (4.6) | 0.44 |
| Candidemia | 68 (8.1) | 11 (5.4) | 57 (8.9) | 0.14 |
| CRE carriage | | | | |
| Time between OLT and carriage detection (days) (median, IQR) | 9 (0.25-21) | -6 (-32-0) | 13 (7-28) | <0.001 |
| First positive sample | | | | |
| Rectal swab | 719 (85.6) | 189 (93.1) | 530 (83.2) | 0.001 |
| Respiratory sample | 45 (5.4) | 3 (1.5) | 42 (6.6) | 0.002 |
| Urine | 69 (8.2) | 22 (10.8) | 47 (7.4) | 0.14 |
| Multi-site colonization | 113 (13.5) | 26 (12.8) | 87 (13.7) | 0.81 |
| Resistance mechanism | | | | |
| KPC | 600 (71.4) | 149 (73.4) | 451 (70.8) | 0.53 |
| VIM | 7 (0.8) | 3 (1.5) | 4 (0.6) | 0.23 |
| OXA-48 | 17 (2) | 6 (3) | 11 (1.7) | 0.21 |
| Not tested | 115 (13.7) | 19 (9.4) | 96 (15.1) | 0.05 |
| Management of CRE colonization | | | | |
| No strategy adopted | 521 (62) | 85 (41.9) | 436 (68.4) | <0.001 |
| Target perioperative prophylaxis | 51 (6.1) | 51 (25.1) | 0 (0) | <0.001 |
| Decolonization | 16 (1.9) | 4 (2) | 12 (1.9) | 0.57 |
| Pre-emptive strategy ¹ | 21 (2.5) | 6 (3) | 15 (2.4) | 0.4 |
| Outcome | | | | |
| Length of ICU stay after OLT (days) (median, IQR) | 7 (4-15) | 7 (4-14) | 7 (4-15) | 0.38 |
| Length of hospital stay after OLT (days) (median, IQR) | 27 (16-50) | 23 (14-37) | 29 (17-54) | <0.001 |
| All-cause 6-months mortality | 263 (31.3) | 69 (34) | 194 (30.5) | 0.39 |

| | | | | |
|--|-------------|----------------|---------------|--------|
| Time from OLT to death (days) (median, IQR) | 70 (29-145) | 32 (10.5-77.5) | 88.5 (43-162) | <0.001 |
| Hospital re-admissions (median, IQR) | 0 (0-1) | 0 (0-1) | 0 (0-1) | 0.07 |

440 ¹Start of an anti-CRE treatment if complications occur

441 Abbreviations: IQR interquartile range, COPD chronic obstructive pulmonary disease, MELD Model

442 End Stage Liver Disease, OLT orthotopic liver transplant, ICU intensive care unit, ACLF acute on

443 chronic liver failure, PGNF primary graft non-function, CMV cytomegalovirus, CRE carbapenem-

444 resistant *Enterobacteriaceae*, KPC *Klebsiella pneumoniae* carbapenemase-producing, SBP

445 spontaneous bacterial peritonitis.

446

447 **Table 2.** Comparison of patients with and without CRE infection within 180 days after liver transplant

| | Patients without CRE infection, n=590 (%) | Patients with CRE infection, n=250 (%) | P |
|--|--|---|----------|
| Demographic data | | | |
| Age (years) [median (IQR)] | 55 (46-61) | 56 (47-62) | 0.5 |
| Sex, male | 389 (65.9) | 160 (64) | 0.59 |
| Comorbidities | | | |
| Myocardial infarction | 17 (2.9) | 6 (2.4) | 0.7 |
| Congestive heart failure | 19 (3.2) | 10 (4) | 0.57 |
| Peripheral vascular disease | 27 (4.6) | 6 (2.4) | 0.14 |
| Cerebrovascular disease | 15 (2.5) | 7 (2.8) | 0.83 |
| COPD | 30 (5.1) | 16 (6.4) | 0.51 |
| Connective tissue disease | 6 (1) | 3 (1.2) | 0.81 |
| Peptic ulcer disease | 44 (7.5) | 18 (7.2) | 0.89 |
| Diabetes without organ damage | 103 (17.5) | 44 (17.6) | 0.96 |
| Moderate/severe renal disease | 45 (7.6) | 37 (14.8) | 0.001 |
| Diabetes with organ damage | 55 (9.3) | 21 (8.4) | 0.67 |
| Any tumor within 5 years | 192 (32.5) | 79 (31.6) | 0.79 |
| Moderate/severe liver disease | 576 (97.6) | 244 (97.6) | 0.98 |
| Charlson index [median (IQR)] | 5 (4-7) | 5 (4-7) | 0.56 |
| Underlying liver disease | | | |
| Viral hepatitis | 260 (44.1) | 116 (46.4) | 0.53 |
| Alcohol | 157 (26.6) | 50 (20) | 0.04 |
| Metabolic disease | 48 (8.1) | 21 (8.4) | 0.89 |
| Autoimmune disease | 32 (5.4) | 9 (3.6) | 0.26 |
| Fulminant hepatitis | 36 (6.1) | 16 (6.4) | 0.87 |
| Hepatocellular carcinoma | 174 (29.5) | 73 (29.2) | 0.93 |
| Prior transplant | 24 (4.1) | 15 (6) | 0.22 |
| MELD at waiting list inclusion (median, IQR) | 19 (14-24) | 19 (14-26) | 0.73 |
| MELD at OLT (median, IQR) | 22.5 (16-29) | 25 (16-33) | 0.02 |
| Events prior 90 days to OLT | | | |
| Hepato-renal syndrome | 150 (25.4) | 85 (34) | 0.01 |
| Ascites grade III/refractory | 278 (47.1) | 107 (42.8) | 0.25 |
| Bleeding | 89 (15.1) | 49 (19.6) | 0.11 |
| Encephalopathy | 245 (41.5) | 98 (39.2) | 0.53 |
| ICU admission | 110 (18.6) | 51 (20.4) | 0.55 |
| ACLF | 66 (11.2) | 32 (12.8) | 0.5 |
| Infections | 223 (37.8) | 103 (41.2) | 0.36 |
| Intraoperative variables | | | |
| Combined transplant | 25 (4.2) | 20 (8) | 0.03 |
| Donor age | 51 (35-65) | 51 (36-67) | 0.6 |
| Cold ischemia time (hours) (median, IQR) | 7 (6-8) | 7 (5:50-8) | 0.36 |
| Choledochojejunostomy | 71 (12) | 40 (16) | 0.22 |
| Intraoperative bleeding | 196 (33.2) | 98 (39.2) | 0.1 |
| Prolonged surgery | 162 (27.5) | 86 (34.4) | 0.04 |
| Induction regimen | | | |
| Bolus of steroids | 476 (80.7) | 225 (90) | 0.001 |

| | | | |
|---|------------|------------|--------|
| Antithymocyte globulins | 12 (2) | 14 (5.6) | 0.006 |
| Basiliximab/Daclizumab | 113 (19.2) | 54 (21.6) | 0.42 |
| Rituximab | 6 (1) | 8 (3.2) | 0.02 |
| Maintenance regimen | | | |
| Steroids | 414 (70.2) | 198 (79.2) | 0.007 |
| Cyclosporin | 31 (5.3) | 11 (4.4) | 0.6 |
| Tacrolimus | 549 (93.1) | 221 (88.4) | 0.03 |
| Micophenolate mophetil | 215 (36.4) | 72 (28.8) | 0.03 |
| Post-operative complications | | | |
| Acute renal injury | 286 (48.5) | 176 (70.4) | <0.001 |
| Renal replacement therapy | 173 (29.3) | 126 (50.4) | <0.001 |
| Mechanical ventilation >48 hours | 135 (22.9) | 131 (52.4) | <0.001 |
| PGNF | 74 (12.5) | 54 (21.6) | 0.001 |
| Re-intervention | 200 (33.9) | 122 (48.8) | <0.001 |
| Re-transplantation | 43 (7.3) | 36 (14.4) | 0.001 |
| Rejection | 101 (17.1) | 51 (20.4) | 0.26 |
| CMV DNAemia>100,000 copies/mL | 156 (26.4) | 69 (27.6) | 0.73 |
| Infections other than CRE | 333 (56.4) | 146 (58.4) | 0.6 |
| <i>Clostridioides difficile</i> infection | 26 (4.4) | 11 (4.4) | 0.99 |
| Candidemia | 36 (6.1) | 32 (12.8) | 0.001 |
| CRE carriage | | | |
| Pre-OLT carriage | 136 (23.1) | 67 (26.8) | 0.25 |
| Post-OLT carriage | 454 (76.9) | 183 (73.2) | 0.25 |
| Rectal swab | 517 (87.6) | 202 (80.8) | 0.01 |
| Respiratory sample | 21 (3.6) | 24 (9.6) | <0.001 |
| Urine sample | 50 (8.5) | 19 (7.6) | 0.67 |
| Multisite colonization | 54 (9.2) | 59 (23.6) | <0.001 |
| pre-OLT | 17 (2.9) | 9 (3.6) | <0.001 |
| post-OLT | 37 (6.3) | 50 (20) | 0.66 |
| Genotype of the colonizing strain | | | |
| KPC | 418 (70.8) | 182 (72.8) | 0.61 |
| VIM | 4 (0.7) | 3 (1.2) | 0.45 |
| OXA-48 | 11 (1.9) | 6 (2.4) | 0.6 |
| Not tested | 76 (12.9) | 39 (15.6) | 0.3 |
| Management of CRE colonization | | | |
| Target perioperative prophylaxis | 35 (5.9) | 17 (6.8) | 0.6 |
| Decolonization | 12 (2) | 4 (1.6) | 0.7 |
| Pre-emptive strategy ¹ | 11 (1.9) | 10 (4) | 0.07 |
| Outcome | | | |
| Length of ICU stay after OLT (days) (median, IQR) | 6 (3-10) | 14 (5-29) | <0.001 |
| Length of hospital stay after OLT (days) (median, IQR) | 22 (15-40) | 43 (24-76) | <0.001 |
| All-cause 6-months mortality | 118 (20) | 145 (58) | <0.001 |
| Hospital re-admissions (median, IQR) | 0 (0-1) | 0 (0-1) | <0.001 |

448 ¹Start of an anti-CRE treatment if complications occur.

449 Abbreviations: IQR interquartile range, COPD chronic obstructive pulmonary disease, MELD Model

450 End Stage Liver Disease, OLT orthotopic liver transplant, ICU intensive care unit, SBP spontaneous

451 bacterial peritonitis, ACLF acute on chronic liver failure, PGNF primary graft non-function, CMV
452 cytomegalovirus, CRE carbapenem-resistant *Enterobacteriaceae*, KPC *Klebsiella pneumoniae*
453 carbapenemase-producing.

454 **Table 3.** Final multivariable analysis for predicting risk of CRE infection after liver transplantation

| Variable | Subhazard ratio (95% CI) | β -coefficient | P value |
|--|--------------------------|----------------------|---------|
| Reintervention | 1.37 (1.04-1.80) | 0.30 (0.02-0.57) | 0.02 |
| Prolonged mechanical ventilation | 2.01 (1.49-2.71) | 0.70 (0.40-0.99) | <0.0001 |
| Acute renal injury | 1.69 (1.26-2.27) | 0.52 (0.23-0.82) | 0.001 |
| CRE colonisation 60 days prior to transplant | 2.17 (1.33-3.53) | 0.78 (0.29-1.26) | 0.002 |
| CRE colonization within 60 days after transplant | 1.41 (0.91-2.19) | 0.35 (-0.09-0.78) | 0.12 |
| Multisite colonization within 60 days after transplant | 2.87 (1.98-4.20) | 1.06 (0.68-1.43) | <0.0001 |

455

456 Abbreviations: CI confidence interval, CRE carbapenem-resistant Enterobacteriaceae.

457

458 Figure legends:

459 **Figure 1.** Study flow chart with detailed data about **involved** centers.

460 **Figure 2.** Cumulative incidence of CRE infection and death post-transplant. The cumulative incidence
461 was derived from the estimated sub-distribution hazard function following regression with a Fine-
462 Gray model.

463 **Figure 3.** Model-based nomogram for predicting 30 and 60-day cumulative risk of developing CRE
464 infection. Points are summed for each risk factor.

465 **Figure 4.** Distribution of model-predicted (a) 30-day and (b) 60-day risk for CRE infection following
466 LT.

467 **Figure 5.** Calibration curves for the (a) derivation dataset and (b) Bootstrap resampled validation
468 dataset. The C-index (AUC) and Brier score are expressed as the point estimates and 95% CI. An AUC
469 > 0.8 is considered to give good discriminatory accuracy for a clinical prediction model. The Brier
470 score is a measure of the accuracy of probabilistic predictions. The lower the Brier score is for a set
471 of predictions, the better the predictions are calibrated. An ideal model has pairs of observed and
472 predicted probabilities that lie on the 45-degree angle line. Dotplots shown the distribution of
473 predicted risk for patients who developed (top) and did not develop (bottom) CRE infection.

474 **Figure 6.** Examples of the cumulative incidence of CRE infection and nomogram-derived prediction
475 for a low-risk (panels a, b); and a high-risk (panels c, d) LT patient. The blue lines with shading in the
476 nomogram represent the distribution of individual risk factors (0=no, 1=yes) and the distribution of
477 total assigned points on the bottom scale in the original derivation data set. Red points on the upper
478 scale indicate assigned points per risk factor. The red arrows at the bottom of the nomogram indicate
479 the total calculated points for the patient and corresponding 30-day and 60 day predicted
480 cumulative risk with 95% CI.

481 **Figure 7.** Decision-curve analysis of model-directed anti CRE therapy. Net benefit at 60 days
482 represents true cases of CRE infection that would be treated with a CRE-active antibiotic regimen-
483 patients without CRE infection unnecessarily receiving CRE-active therapy. The curves indicate that a
484 model-directed antibiotic strategy should have a higher net benefit than default strategies (all
485 patients receive CRE-active therapy or no patients receive CRE-active therapy) at CRE infection
486 probabilities greater than 10-50 %.

487