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The applicability of commonly used predictive scoring systems in Indigenous Australians with sepsis: an observational study

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| Corresponding Author: | Josh Hanson The Kirby Institute Sydney, NSW AUSTRALIA |
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| Abstract: | <p>Background: Indigenous Australians suffer a disproportionate burden of sepsis, however, the performance of scoring systems that predict mortality in Indigenous patients with critical illness is incompletely defined.</p> <p>Materials and methods: The study was performed at an Australian tertiary-referral hospital between January 2014 and June 2017, and enrolled consecutive Indigenous and non-Indigenous adults admitted to ICU with sepsis. The ability of the ANZROD, APACHE-II, APACHE-III, SAPS-II, SOFA and qSOFA scores to predict death before ICU discharge in the two populations was compared.</p> <p>Results : There were 442 individuals enrolled in the study, 145 (33%) identified as Indigenous. Indigenous patients were younger than non-Indigenous patients (median (interquartile range (IQR) 53 (43-60) versus 65 (52-73) years, $p=0.0001$) and comorbidity was more common (118/145 (81%) versus 204/297 (69%), $p=0.005$). Comorbidities that were more common in the Indigenous patients included diabetes mellitus (84/145 (58%) versus 67/297 (23%), $p<0.0001$), renal disease (56/145 (39%) versus 29/297 (10%), $p<0.0001$) and cardiovascular disease (58/145 (40%) versus 83/297 (28%), $p=0.01$). The use of supportive care (including vasopressors, mechanical ventilation and renal replacement therapy) was similar in Indigenous and non-Indigenous patients, and the two populations had an overall case-fatality rate that was comparable (17/145 (12%) and 38/297 (13%) ($p=0.75$)), although Indigenous patients died at a younger age (median (IQR): 54 (50-60) versus 70 (61-76) years, $p=0.0001$). There was no significant difference in the ability of any the scores to predict mortality in the two populations.</p> <p>Conclusions : Although the crude case-fatality rates of Indigenous and non-Indigenous Australians admitted to ICU with sepsis is comparable, Indigenous patients die at a much younger age. Despite this, the ability of commonly used scoring systems to predict outcome in Indigenous Australians is similar to that of non-Indigenous Australians, supporting their use in ICUs with a significant Indigenous patient population and in clinical trials that enrol Indigenous Australians.</p> |
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Additional data availability information:

1 The applicability of commonly used predictive scoring systems in Indigenous Australians
2 with sepsis: an observational study
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33 **Abstract**

34 **Background:** Indigenous Australians suffer a disproportionate burden of sepsis, however, the
35 performance of scoring systems that predict mortality in Indigenous patients with critical
36 illness is incompletely defined.

37 **Materials and methods:** The study was performed at an Australian tertiary-referral hospital
38 between January 2014 and June 2017, and enrolled consecutive Indigenous and non-
39 Indigenous adults admitted to ICU with sepsis. The ability of the ANZROD, APACHE-II,
40 APACHE-III, SAPS-II, SOFA and qSOFA scores to predict death before ICU discharge in the
41 two populations was compared.

42 **Results:** There were 442 individuals enrolled in the study, 145 (33%) identified as Indigenous.
43 Indigenous patients were younger than non-Indigenous patients (median (interquartile range
44 (IQR) 53 (43-60) versus 65 (52-73) years, $p=0.0001$) and comorbidity was more common
45 (118/145 (81%) versus 204/297 (69%), $p=0.005$). Comorbidities that were more common in
46 the Indigenous patients included diabetes mellitus (84/145 (58%) versus 67/297 (23%),
47 $p<0.0001$), renal disease (56/145 (39%) versus 29/297 (10%), $p<0.0001$) and cardiovascular
48 disease (58/145 (40%) versus 83/297 (28%), $p=0.01$). The use of supportive care (including
49 vasopressors, mechanical ventilation and renal replacement therapy) was similar in Indigenous
50 and non-Indigenous patients, and the two populations had an overall case-fatality rate that was
51 comparable (17/145 (12%) and 38/297 (13%) ($p=0.75$)), although Indigenous patients died at
52 a younger age (median (IQR): 54 (50-60) versus 70 (61-76) years, $p=0.0001$). There was no
53 significant difference in the ability of any the scores to predict mortality in the two populations.

54 **Conclusions:** Although the crude case-fatality rates of Indigenous and non-Indigenous
55 Australians admitted to ICU with sepsis is comparable, Indigenous patients die at a much
56 younger age. Despite this, the ability of commonly used scoring systems to predict outcome in
57 Indigenous Australians is similar to that of non-Indigenous Australians, supporting their use in

58 ICUs with a significant Indigenous patient population and in clinical trials that enrol Indigenous
59 Australians.

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77 **Introduction**

78 Globally, sepsis is estimated to kill 5.3 million people every year, predominantly in low- and
79 middle income countries (1). Even in the well-resourced Australian health system, sepsis kills
80 over 5000 people annually (2, 3). However, the case-fatality rate is not uniform across Australia
81 (4). This can be partly explained by differences in the complexity of patient care at different
82 sites, but other factors including patient age and demographics, the geographical location of
83 the Intensive Care Unit (ICU), and the local prevalence of different pathogens also contribute
84 (4).

85 In an effort to ensure that all Australians admitted to ICU are receiving high-quality care,
86 predictive scoring systems are used to measure patients' disease severity and determine their
87 expected outcomes (5-9). While these scores have limited utility in the management of
88 individual patients, they can be used by institutions to benchmark ICU performance. They are
89 also important in clinical trials where they can be used to help evaluate interventions by
90 providing a measure of study patients' disease severity. However, these scoring systems are
91 most frequently derived and validated in metropolitan referral centres in high-income settings,
92 their predictive ability may differ in other patient populations (10, 11).

93 Indigenous Australians are disproportionately represented in Australian ICUs and sepsis is one
94 of the most common indications for admission (12). Several studies have examined the
95 characteristics of undifferentiated cohorts of Indigenous Australians admitted to ICU, and have
96 identified that they are younger, have greater comorbidity and live more frequently in remote
97 locations (12-15). While, the overall case-fatality rate of Indigenous Australians is similar to
98 that of non-Indigenous Australians in these series, there has been limited examination of the
99 comparative performance of predictive scoring systems in Indigenous patients (12-15). This is

100 important, because in some parts of the country, Indigenous Australians represent a significant
101 proportion of ICU admissions (13, 14).

102 This study was performed to determine the applicability of commonly used predictive scoring
103 systems in Indigenous Australians admitted to ICU with sepsis. It was hoped that the study
104 might validate the use of these scoring systems in both Aboriginal and Torres Strait Islander
105 Australians, which would support their use in future clinical sepsis trials enrolling Indigenous
106 patients. It would also provide justification for their use in benchmarking the performance of
107 ICUs with a greater proportion of Indigenous patients (16).

108

109 **Materials and methods**

110 This **retrospective study** used data collected in the ICU of Cairns Hospital, the only ICU in the
111 Far North Queensland (FNQ) region. FNQ's population of 279,354 is dispersed across an area
112 of 204,255 km²; approximately 17% of whom identify as Indigenous Australians (17).

113 Consecutive adults (≥ 18 years) admitted between 1 January 2014 and 30 June 2017 to the ICU
114 with a primary admission diagnosis of sepsis (**APACHE III-J** diagnostic codes 501-504) were
115 eligible for the study. Demographic, clinical and laboratory data were collected and correlated
116 with the patients' clinical course. **Comorbidities** were said to be present if documented in the
117 medical record (MetaVision®). Hazardous alcohol consumption was defined as regular
118 consumption of greater than 4 standard drinks/day. The data were collected by 4 members of
119 the research team (SH SS TG JB) who met regularly to confirm uniformity of documentation.
120 If patients were admitted to ICU more than once during the study period, only their first
121 admission was included in the analysis.

122 All individuals receiving care in Queensland's public health system, are asked whether they
123 identify as an Aboriginal Australian, a Torres Strait Islander Australian, both or neither.
124 Commonly used disease severity prediction scores including the Australia and New Zealand
125 Risk Of Death (ANZROD) score (5), the Acute Physiology, Age, Chronic Health Evaluation-
126 II and III (APACHE-II and APACHE-III) scores (18), the Simplified Acute
127 Physiology Score (SAPS) II score (19) and the quick SOFA (qSOFA) score (20) were
128 calculated using variables collected at the time of ICU admission. The Sequential Organ Failure
129 Assessment (SOFA) score was calculated using the worst values recorded in the first 24 hours
130 of the ICU admission (9). The scores' ability to predict both death prior to ICU discharge and
131 at 90 days was determined.

132 **Statistical analysis**

133 Data were de-identified, entered in an electronic database (Microsoft Excel) and analysed using
134 statistical software (Stata version 14.2). Groups were compared using the Kruskal-Wallis, Chi-
135 square test or Fisher's exact test where appropriate. Multivariate analysis was performed using
136 backwards stepwise logistic regression. The Indigenous population was further examined by
137 dividing the Indigenous population into those who identified as Aboriginal Australians and
138 those who identified as Torres Strait Islander Australians. Those who identified as both, were
139 not included in analyses comparing the two Indigenous populations.

140 The ability of the scores to predict death were determined by measuring the area under receiver
141 operator characteristic (AUROC) curves, the optimal cut-off for the tests for the different
142 populations were determined using Liu's method (21).

143

144

145 **Ethics approval**

146 Ethics approval was obtained from the Far North Queensland Human Research Ethics
147 Committee (HREC/17/QCH/93/AMO2). As the data were retrospective and de-identified, the
148 Committee waived the requirement for informed consent.

149

150 **Results**

151 There were 442 individuals admitted to ICU for sepsis, on 500 occasions, during the study
152 period. Their median (interquartile range (IQR)) age at their first presentation was 59 (48-70)
153 years, 238 (54%) were male. Of the 442 patients, 145 (33%) identified as Indigenous
154 Australians, 94 (65%) identified as Aboriginal, 36 (25%) identified as Torres Strait Islanders,
155 while 15 (10%) identified as both. Among the 442 patients, 416 (94%) were FNQ residents;
156 Indigenous patients were more likely to reside in a rural or remote location than non-Indigenous
157 patients (88/144 (61%) versus 128/272 (47%), $p=0.006$).

158 **Patient characteristics**

159 Indigenous patients were younger than non-Indigenous patients (median (IQR) 53 (43-60)
160 versus 65 (52-73) years, $p=0.0001$) and were more likely to have a significant comorbidity
161 (118/145 (81%) versus 204/297 (69%), $p=0.005$).

162 Indigenous patients were more likely to have diabetes mellitus than non-Indigenous patients
163 (odds ratio (OR): 4.7, 95% confidence interval (CI): 3.1-7.2), which contributed to a
164 significantly greater burden of cardiovascular disease and renal disease (table 1). Diabetes was
165 more common in Torres Strait Islanders than Aboriginal Australians (27/36 (75%) versus 48/94
166 (51%), $p=0.01$). No fewer than 32/145 (22%) Indigenous patients had a history of sepsis (prior
167 hospitalisation with APACHE III-J diagnostic codes 501-504).

168 While Indigenous patients, as a group, were more likely to have a history of hazardous alcohol
169 or tobacco use, hazardous alcohol use was more common in the Aboriginal patients than in
170 Torres Strait Islanders (49/94 (52%) versus 7/36 (9%) (p=0.001). The difference in smoking
171 rates between Aboriginal and Torres Strait Islanders failed to reach statistical significance
172 (63/94 (67%) versus 18/36 (50%), p=0.07). Other demographic characteristics of the cohort
173 and their comorbidities - stratified by Indigenous status - are presented in table 1.

174 **Nature of the sepsis**

175 The commonest presumed source of sepsis was the respiratory tract, although skin and soft
176 tissue infections (SSTI) were also common and occurred more frequently in Indigenous
177 patients, particularly those with diabetes (31/41 (76%) Indigenous patients with a SSTI were
178 diabetic, compared with 53/104 (51%) with another source, p=0.007). *Staphylococcus aureus*
179 was the pathogen isolated most commonly and was responsible for 66 (15%) admissions; 13
180 (18%) of the isolates were methicillin resistant. In general, however, drug resistant pathogens
181 and “tropical” pathogens were relatively uncommon (table 2).

182 **Clinical and laboratory findings at presentation**

183 Renal impairment and metabolic acidosis were more common among Indigenous patients, but
184 other laboratory findings in the Indigenous and non-Indigenous patients were similar (table 3).



185 **Supportive care**


186 The supportive care provided to the Indigenous and non-Indigenous patients was very similar.
187 While a greater proportion of Indigenous patients required renal replacement therapy (RRT),
188 this did not reach statistical significance (19/145 (13%) versus 27/297 (9%), p=0.20) (table 4).

189

190 **Case-fatality rate**

191 There were 55 deaths in the cohort prior to ICU discharge: 17/145 (12%) Indigenous patients
192 versus 38/297 (13%) non-Indigenous patients, $p=0.75$. There were 14 (15%) deaths among the
193 94 Aboriginal Australians and 2 (6%) among the 36 Torres Strait Islander Australians ($p=0.23$).
194 There was one (7%) death among the 15 who identified as both Aboriginal and Torres Strait
195 Islander Australians.

196 Patients that died before ICU discharge were older than those  that survived (median (IQR): 62
197 (54-73) versus 59 (46-69) years), $p=0.01$) and more likely to have a significant comorbidity
198 (47/55 (85%) versus 275/387 (71%), $p=0.03$). Among the 442 FNQ residents,  29/50 (58%)
199 living in a rural or remote location died compared with 187/366 (51%), with an urban address
200 ($p=0.36$). In a multivariate analysis model which included age, Indigenous status, significant
201 comorbidity and remote or rural residence, only age had a statistically significant association
202 with death (OR: 1.03 (95%CI): 1.01-1.05), $p=0.007$).

203  At 90 days, 93/442 (21%) had died. Death at 90 days was linked to age ($p=0.0001$), but not
204 to Indigenous status ($p=0.26$) or rural/remote residence ($p=0.29$). Indigenous patients,
205 however, died at a younger age than non-Indigenous patients (median (IQR): 56 (52-62) versus
206 68 (61-76) years, $p=0.0001$).

207 **Disease severity scores and their ability to predict death**

208 The severity scores of the Indigenous and non-Indigenous patients were similar (table 5). There
209 was no statistically significant difference in the ability of the various severity scores to predict
210 death in Indigenous and non-Indigenous patients (tables 6 and 7). Among Indigenous patients,
211 the ANZROD and APACHE-III scores had the highest AUROC curve (0.85 (95% CI: 0.77-
212 0.92 and 0.84 (95% CI: 0.76-0.92), although these values were only statistically superior to
213 that of the qSOFA scores (tables 6 and 7). The very low case-fatality rate in the Torres Strait

214 Islanders and relatively small sample size precluded meaningful comparison of the relative
215 performance of the prediction scores in Aboriginal and Torres Strait Islanders.

216 **Discussion**

217 In this cohort of ICU patients with sepsis there were significant differences in the age, burden
218 of comorbidities and source of infection between Indigenous and non-Indigenous individuals.
219 However, the ability of commonly used disease severity scoring systems to predict mortality
220 in the two populations was similar.



221 FNQ, in tropical Australia, shares a border with Papua New Guinea and has a unique blend of
222 infectious diseases; it has the country's highest incidence of leptospirosis and an increasing
223 incidence of melioidosis and rickettsial disease (22-24). The rates of methicillin-resistance in
224 *S. aureus* isolates are among the highest reported in the country, while antibiotic resistance in
225 other common pathogens is also increasing (25, 26). The population is widely dispersed across
226 a large geographical area and there are very limited specialist services outside the
227 administrative hub of Cairns. It is a region which contains 3 of the 10 most socio-economically
228 disadvantaged local government areas in the country, all 3 are communities with a
229 predominantly Indigenous population (27). It is the only part of Australia which has the
230 homelands of both Aboriginal and Torres Strait Islander Australians, peoples that are
231 frequently conflated but who are ethnologically quite distinct. All these factors might be
232 expected to have implications for local patterns of sepsis, its presentation and its outcomes (16,
233 28-30). However, while Indigenous patients were younger, more likely to be live in a
234 rural/remote location and have a significant comorbidity than non-Indigenous patients, the
235 severity of their sepsis – as determined by commonly used predictive scoring systems was
236 similar to that of non-Indigenous patients.


237 Although Indigenous Australians bear a greater burden of sepsis, there are surprisingly few
238 studies that examine sepsis in the Indigenous population systematically. The demography and
239 comorbidities in our series are very similar to that of a prospective study of sepsis in the
240 Northern Territory, which also included patients that were not admitted to the ICU (16). That
241 study identified that Indigenous Australians were over-represented in the cohort, suffered
242 disproportionately from diabetes and renal disease, and reported higher rates of smoking and
243 hazardous alcohol use. Although APACHE-II and SOFA scores were collected in the study,
244 the comparability of their performance in Indigenous and non-Indigenous populations was not
245 presented.



246 While there are relatively few published data examining the ICU care of Australian Indigenous
247 patients with sepsis specifically, there are several studies that have examined the clinical
248 characteristics and outcomes of Indigenous patients admitted to ICU (12-15). These studies are
249 strikingly similar and show that Indigenous patients with critical illness are younger, have
250 greater comorbidity and are more frequently admitted from remote locations than non-
251 Indigenous patients. These studies also universally show that there is no difference in the
252 proportion of Indigenous and non-Indigenous patients who die in the ICU. Whilst some authors
253 have suggested that Indigenous deaths in ICU are therefore “healthcare preventable” (14), this
254 represents a narrow view of healthcare, de-emphasising the primary holistic care that should
255 prevent the hospitalisation in the first place. Whilst other authors have noted that there is “no
256 mortality gap” between Indigenous and non-Indigenous Australians admitted to ICU (12), this
257 overlooks the fact the Indigenous patients who are admitted to ICU are younger and are dying
258 at a younger age. In an ICU series from the Northern Territory, 7% of both Indigenous and
259 non-Indigenous patients admitted to the ICU died, but the Indigenous patients who died were
260 much younger than their non-Indigenous counterparts (mean age of 45 versus 56) (13). This
261 pattern was seen in our series: although the proportion of Indigenous patients and non-

262 Indigenous patients dying before ICU discharge was similar, the median age of death of
263 Indigenous patients (56 years) was 12 years lower than that of the non-Indigenous patients. To
264 close this gap, we need evidence-based strategies to address the unique challenges faced by
265 Indigenous Australians in the country's health system. Our study shows that predictive scoring
266 systems are a valid way of measuring disease severity among Indigenous Australians with
267 sepsis and can therefore be used to compare the efficacy of interventions with confidence.

268 Our study has many limitations. As a single centre  trial, it reflects only the experience of a
269 unique part of Australia; the applicability of the results to the broader Indigenous patient
270 population requires validation, although the similarity between the Indigenous population seen
271 in this study and those in other locations is striking  (12-16). The study was retrospective and
272 examined only patients admitted to ICU and will therefore underestimate the true sepsis burden
273 (31).

274 Clearly there is still much to do to address the disparity in health outcomes between Indigenous 
275 and non-Indigenous Australians (30). Once Indigenous Australians with sepsis enter ICU, they
276 receive high quality care, but they are still dying at a much younger age than their non-
277 Indigenous counterparts. Interventions at a community level including efforts to facilitate
278 access to care, reduce crowding, enhance health literacy, improve sanitation and ensure
279 appropriate nutrition are likely to be most helpful (32). At a primary health level optimising
280 management of conditions like diabetes that predispose to sepsis and ensuring comprehensive
281 vaccination, particularly for those at high risk are also likely to assist. Expanded programmes
282 to assist with smoking cessation and encourage alcohol moderation are also essential (33, 34).
283 However, even with optimal preventative interventions, patients will still require ICU care for
284 sepsis. This study suggests that standard predictive scores predict outcomes in Indigenous

285 patients as well as they do in non-Indigenous patients and therefore may be used in future
286 studies to examine strategies to enhance the care of all Australians

287

288

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291 of the patients.

292

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406 **Table 1 Demographic characteristics and comorbidities of the cohort, stratified by**
 407 **Indigenous status**
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| Variable | Indigenous n=145 | Non-Indigenous n=297 | p |
|---|---------------------|-------------------------|---------|
| Age | 53 (43-60) | 65 (52-73) | 0.0001 |
| Male gender | 68 (47%) | 170 (57%) | 0.04 |
| Residence in a remote location ^a | 88/144 (61%) | 128/272 (47%) | 0.006 |
| Inter-hospital transfer | 56 (39%) | 115 (39%) | 0.98 |
| Admitted from Emergency Department | 44 (30%) | 95 (32%) | 0.73 |
| Admitted from a nursing home | 0 | 0 | - |
| Planned admission after surgery | 3 (2%) | 2 (1%) | 0.34 |
| Admitted from hospital ward | 42 (29%) | 83 (28%) | 0.82 |
| Admission at night | 94 (65%) | 182 (61%) | 0.47 |
| Hazardous alcohol consumption | 58 (40%) | 53 (18%) | <0.0001 |
| Cigarette smoker | 88 (61%) | 134 (45%) | 0.002 |
| History of sepsis | 32 (22%) | 30 (10%) | 0.001 |
| History of cardiovascular disease | 58 (40%) | 83 (28%) | 0.01 |
| History of respiratory disease | 31 (21%) | 49 (17%) | 0.21 |
| History of renal disease | 56 (39%) | 29 (10%) | <0.0001 |
| History of haematological disease or malignancy | 7 (5%) | 29 (10%) | 0.10 |
| History of liver disease | 18 (12%) | 22 (7%) | 0.09 |
| History of diabetes mellitus | 84 (58%) | 67 (23%) | <0.0001 |
| History of metastatic cancer | 6 (4%) | 29 (10%) | 0.04 |
| Immunocompromised | 9 (6%) | 52 (18%) | 0.001 |
| Significant comorbidity | 118 (81%) | 204 (69%) | 0.005 |
| Body mass index | 27 (22-33) | 28 (24-33) | 0.07 |

409 Absolute numbers (%) or median (interquartile range) are presented.

410 ^a Only includes the 416 Far North Queensland residents.

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Table 2 Source and aetiology of the sepsis, stratified by Indigenous status

| Variable | Indigenous n=145 | Non- Indigenous n=297 | p |
|--|------------------|-----------------------|--------|
| Respiratory source | 51 (35%) | 97 (33%) | 0.60 |
| Genitourinary source | 32 (22%) | 64 (22%) | 0.90 |
| Bone/joint source | 7 (5%) | 6 (2%) | 0.13 |
| Central nervous system source | 0 (0%) | 6 (2%) | 0.18 |
| Skin/soft tissue source | 41 (28%) | 43 (14%) | 0.0001 |
| Abdominal source | 13 (9%) | 43 (14%) | 0.10 |
| Other source | 10 (7%) | 47 (16%) | 0.01 |
| Bacterial infection | 103(71%) | 194 (65%) | 0.23 |
| Gram negative bacteria | 63 (43%) | 131 (44%) | 0.90 |
| Gram positive bacteria | 55 (38%) | 84 (28%) | 0.04 |
| Fungal infection | 7 (5%) | 9 (3%) | 0.42 |
| Viral infection | 10 (7%) | 22 (7%) | 0.85 |
| Drug resistant organism | 5 (3%) | 11 (4%) | 1.0 |
| Bacteraemia | 56 (39%) | 128 (43%) | 0.37 |
| Polymicrobial infection | 34 (23%) | 46 (15%) | 0.04 |
| <i>Escherichia coli</i> | 15 (10%) | 44 (15%) | 0.20 |
| <i>Staphylococcus aureus</i> | 26 (18%) | 40 (13%) | 0.22 |
| Methicillin-resistant <i>S. aureus</i> | 7 (5%) | 6 (2%) | 0.13 |
| <i>Pseudomonas aeruginosa</i> | 15 (10%) | 28 (9%) | 0.76 |
| <i>Klebsiella pneumoniae</i> | 9 (6%) | 15 (5%) | 0.66 |
| Influenza A | 5 (3%) | 13 (4%) | 0.80 |
| <i>Burkholderia pseudomallei</i> | 8 (6%) | 8 (3%) | 0.17 |
| <i>Streptococcus pyogenes</i> | 16 (11%) | 9 (3%) | 0.001 |
| Leptospirosis | 0 | 11 (4%) | 0.02 |
| <i>Streptococcus pneumoniae</i> | 6 (4%) | 10 (3%) | 0.79 |
| <i>Pneumocystis jirovecii</i> | 2 (1%) | 4 (1%) | 1.0 |
| <i>Mycobacterium tuberculosis</i> | 0 | 1 (0.3%) | 1 |
| <i>Plasmodium falciparum</i> | 0 | 1 (0.3%) | 1 |
| <i>Rickettsia australis</i> | 1 (0.7%) | 0 | 0.33 |
| <i>Vibrio vulnificus</i> | 1 (0.7%) | 0 | 0.33 |

423 Absolute numbers (%) presented.

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Table 3 Clinical and laboratory findings at presentation, stratified by Indigenous status

| Variable | Indigenous n=145 | Non- Indigenous n=297 | p |
|--|----------------------|-----------------------|--------|
| Heart rate (beats/min) | 99 (89-118) | 97 (81-114) | 0.28 |
| Systolic blood pressure (mmHg) | 107 (94-118) | 106 (94-122) | 0.42 |
| Mean arterial pressure (mmHg) | 73 (66-82) | 72 (65-83) | 0.50 |
| Temperature (°C) | 36.8 (36.5-37.4) | 36.9 (36.5-37.3) | 0.53 |
| Respiratory rate (breaths/min) | 20 (17-26) | 20 (16-25) | 0.99 |
| Glasgow Coma Score | 15 (14-15) | 15 (14-15) | 0.98 |
| Glucose (mmol/L) | 6.9 (5.0-9.4) | 6.6 (5.4-8.5) | 0.96 |
| Lactate (mmol/L) | 1.8 (1.1-3.0) | 1.7 (1.1-2.8) | 0.29 |
| pH | 7.36 (7.24-7.42) | 7.37 (7.30-7.43) | 0.13 |
| PaO ₂ /FiO ₂ | 291 (194-398) | 269 (158-375) | 0.02 |
| PaCO ₂ (mmHg) | 34 (27-39) | 34 (29-42) | 0.0496 |
| Bicarbonate (mmol/L) | 18 (15-21) | 20 (17-23) | 0.001 |
| Base excess (mmol/L) | -6.2 (-10.9 to -2.8) | -4.5 (-8.4 to -2.2) | 0.004 |
| Anion gap | 9 (7-12) | 9 (7-12) | 0.44 |
| Haemoglobin (g/dL) | 100 (87-122) | 111 (93-124) | 0.007 |
| White cell count (x10 ⁹ /L) | 13.8 (9.3-21.9) | 13.8 (7.9-22.2) | 0.67 |
| Neutrophils (x10 ⁹ /L) | 10.7 (7.1-19.2) | 11.6 (6.2-19.4) | 0.80 |
| Eosinophils (x10 ⁹ /L) | 0 (0-0.1) | 0(0-0) | 0.001 |
| Platelets (x10 ⁹ /L) | 170 (111-257) | 1650(101-236) | 0.17 |
| C-reactive protein (mg/L)) | 165 (67-295) | 172 (85-286) | 0.57 |
| Troponin I (ng/mL) | 0.07 (0.04-0.39) | 0.11 (0.04-0.43) | 0.85 |
| Prothrombin (seconds) | 16 (14-21) | 16 (14-18) | 0.07 |
| APTT (seconds) | 39 (34-47) | 36 (31-40) | 0.0001 |
| INR | 1.5 (1.3-1.9) | 1.4 (1.3-1.6) | 0.03 |
| Fibrinogen (g/L) | 6.2 (4.6-8.3) | 6.6 (4.9-8.2) | 0.35 |
| Total Bilirubin (µmol/L) | 18 (12-31) | 20 (13-31) | 0.31 |
| Conjugated bilirubin (µmol/L) | 8 (4-19) | 7 (4-15) | 0.43 |
| Albumin (g/L) | 24 (20-27) | 25 (22-29) | 0.007 |
| Protein (g/L) | 58 (52-65) | 52 (48-60) | 0.0001 |
| AST (IU/mL) | 45 (21-104) | 52 (27-106) | 0.12 |
| ALT (IU/mL) | 25 (12-42) | 35 (20-65) | 0.0001 |
| GGT (IU/mL) | 41 (22-65) | 51 (26-94) | 0.01 |
| ALP (IU/mL) | 91 (69-134) | 82 (57-119) | 0.01 |
| LDH (IU/mL) | 333 (248-482) | 329 (239-439) | 0.43 |
| Sodium (mmol/L) | 133 (131-137) | 135 (133-138) | 0.0004 |
| Potassium (mmol/L) | 4.1 (3.6-4.8) | 4.0 (3.7-4.5) | 0.31 |
| Chloride (mmol/L) | 104 (99-109) | 104 (100-108) | 0.92 |
| Creatinine (µmol/L) | 162 (87-403) | 118 (75-190) | 0.0003 |
| eGFR (ml/min/1.73m ²) | 29 (11-65) | 49 (24-80) | 0.0002 |
| Calcium (mmol/L) | 2.2 (2.1-2.3) | 2.2 (2.1-2.3) | 0.99 |
| Magnesium (mmol/L) | 0.73 (0.66-0.87) | 0.75 (0.65-0.88) | 0.48 |
| Phosphate (mmol/L) | 1.5 (1.0-2.0) | 1.1 (0.9-1.5) | 0.0001 |

438 Median (Interquartile range) presented. PaO₂/FiO₂: ratio of arterial oxygen partial pressure (PaO₂ in mmHg) to
 439 fractional inspired oxygen. PaCO₂: arterial carbon dioxide partial pressure. APTT: activated partial
 440 thromboplastin time; INR: International normalised ratio; AST: aspartate aminotransferase; ALT: alanine
 441 aminotransferase; GGT: Gamma-glutamyl transferase; ALP: alkaline phosphatase; LDH: lactate dehydrogenase;
 442 eGFR: estimated glomerular filtration rate.

443 **Table 4 Supportive care delivered to the cohort, stratified by Indigenous status**

| Variable | Indigenous n=145 | Non- Indigenous n=297 | p |
|---------------------------------------|------------------|-----------------------|------|
| Vasopressors on admission | 102 (70%) | 204 (69%) | 0.72 |
| Number of vasopressors required | 1 (1-1) | 1 (0-1) | 0.89 |
| Antibiotics administered on admission | 142 (98%) | 288 (97%) | 0.56 |
| Endotracheal intubation | 28 (19%) | 64 (22%) | 0.59 |
| Minute ventilation | 7.3 (5.8 - 8.8) | 7.7 (6.6 – 9.0) | 0.30 |
| PICC line | 40 (28%) | 89 (30%) | 0.61 |
| Central venous line | 61 (42%) | 121 (41%) | 0.79 |
| Arterial line | 122 (84%) | 246 (83%) | 0.73 |
| Nasogastric feeding | 28 (19%) | 63 (21%) | 0.64 |
| Indwelling urinary catheter | 117 (81%) | 252 (85%) | 0.27 |
| Renal replacement therapy | 19 (13%) | 27 (9%) | 0.20 |

444 Absolute numbers (%) and median (interquartile range) presented. PICC: Peripherally inserted central catheter

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473 **Table 5 Severity score values on admission to ICU, stratified by Indigenous status**

| Variable | number | Indigenous n=145 | Non- Indigenous n=297 | p |
|------------|--------|------------------|-----------------------|------|
| qSOFA | 348 | 1 (1-2) | 1 (1-2) | 0.83 |
| SOFA | 378 | 9 (6-11) | 8 (6-11) | 0.20 |
| ANZROD | 430 | 0.24 (0.10-0.44) | 0.23 (0.10-0.46) | 0.86 |
| APACHE-II | 430 | 21 (15-26) | 20 (15-26) | 0.65 |
| APACHE-III | 430 | 70 (52-87) | 69 (53-87) | 0.87 |
| SAPS-II | 413 | 35 (25-51) | 38 (28-58) | 0.12 |

474 Median (interquartile range) presented. qSOFA: quick SOFA score; SOFA: Sequential Organ Failure Assessment
 475 score; ANZROD: Australia and New Zealand Risk Of Death score; APACHE-II: Acute Physiology, Age, Chronic
 476 Health Evaluation-II score; and APACHE-III: Acute Physiology, Age, Chronic Health Evaluation-III score;
 477 SAPS-II: Simplified Acute Physiology Score.

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Table 6 Performance of the severity scores in predicting death before ICU discharge, stratified by Indigenous status

| Severity score | Number of Indigenous patients in whom the score could be calculated | AUROC (95% CI) | Optimal cut-off in Indigenous patients | Number of Non- Indigenous patients in whom the score could be calculated | AUROC (95% CI) | Optimal cut-off in Non-Indigenous patients | p |
|----------------|---|------------------|--|--|------------------|--|------|
| qSOFA | 116 | 0.71 (0.57-0.84) | 2 | 232 | 0.58 (0.46-0.70) | 2 | 0.17 |
| SOFA | 120 | 0.75 (0.64-0.87) | 10 | 258 | 0.72 (0.63-0.80) | 10 | 0.62 |
| ANZROD | 138 | 0.85 (0.77-0.92) | 0.33 | 292 | 0.79 (0.71-0.87) | 0.35 | 0.32 |
| APACHE-II | 138 | 0.78 (0.64-0.91) | 30 | 292 | 0.76 (0.67-0.84) | 27 | 0.61 |
| APACHE-III | 138 | 0.84 (0.76-0.92) | 82 | 292 | 0.79 (0.71-0.87) | 88 | 0.37 |
| SAPS-II | 136 | 0.80 (0.70-0.90) | 46 | 277 | 0.85 (0.78-0.92) | 66 | 0.40 |

qSOFA: quick SOFA score; SOFA: Sequential Organ Failure Assessment score; ANZROD: Australia and New Zealand Risk Of Death score; APACHE-II: Acute Physiology, Age, Chronic Health Evaluation-II score; and APACHE-III: Acute Physiology, Age, Chronic Health Evaluation-III score; SAPS-II: Simplified Acute Physiology Score.

Table 7 Performance of the severity scores in predicting 90-day mortality, stratified by Indigenous status

| Severity score | Number of Indigenous patients in whom the score could be calculated | AUROC (95% CI) | Optimal cut-off in Indigenous patients | Number of Non-Indigenous patients in whom the score could be calculated | Non-Indigenous n=297 | Optimal cut-off in Non-Indigenous patients | p |
|----------------|---|------------------|--|---|----------------------|--|------|
| qSOFA | 116 | 0.66 (0.53-0.79) | 2 | 232 | 0.59 (0.50-0.68) | 2 | 0.42 |
| SOFA | 120 | 0.75 (0.64-0.85) | 11 | 258 | 0.66 (0.58-0.74) | 9 | 0.22 |
| ANZROD | 138 | 0.85 (0.79-0.92) | 0.33 | 292 | 0.78 (0.71-0.84) | 0.29 | 0.09 |
| APACHE-II | 138 | 0.74 (0.62-0.85) | 30 | 292 | 0.72 (0.65-0.79) | 21 | 0.78 |
| APACHE-III | 138 | 0.85 (0.78-0.92) | 73 | 292 | 0.77 (0.71-0.84) | 75 | 0.13 |
| SAPS-II | 136 | 0.78 (0.68-0.88) | 36 | 277 | 0.75 (0.69-0.82) | 39 | 0.66 |

qSOFA: quick SOFA score; SOFA: Sequential Organ Failure Assessment score; ANZROD: Australia and New Zealand Risk Of Death score; APACHE-II: Acute Physiology, Age, Chronic Health Evaluation-II score; and APACHE-III: Acute Physiology, Age, Chronic Health Evaluation-III score; SAPS-II: Simplified Acute Physiology Score.



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Supporting Information

Final dataset for publication.xlsx

