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## Melioidosis is an opportunistic infectious disease: the 30-year Darwin Prospective Melioidosis Study --Manuscript Draft--

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<b>Manuscript Region of Origin:</b>	AUSTRALIA
<b>Abstract:</b>	<p>Background: The global distribution of melioidosis is under considerable scrutiny, with both unmasking of endemic disease in African and Pacific nations and evidence of more recent dispersal in the Americas. Tropical northern Australia has high incidence rates and The Darwin Prospective Melioidosis Study commenced in October 1989. We present epidemiology, clinical features, outcomes and bacterial genomics from this 30-year study, highlighting changes in the last decade.</p> <p>Methods: Prospective study of epidemiological, clinical and laboratory data for all culture-confirmed melioidosis cases from the tropical Northern Territory of Australia from October 1<sup>st</sup> 1989 until September 30<sup>th</sup> 2019. Multivariable analysis determined predictors of clinical presentations and outcome. Incidence, survival and cluster analyses were facilitated by population and rainfall data and genotyping of Burkholderia pseudomallei including multilocus sequence typing and whole-genome sequencing.</p>

Findings: There were 1148 individuals with culture-confirmed melioidosis of whom 133 (11.6%) died. Median age was 50 years, 48 were children  $\leq 14$  years old (4.2%), 721 (63%) were male and 600 (52%) Indigenous Australians. All but 186 (16%) had clinical risk factors; diabetes 513 (45%), hazardous alcohol use 455 (40%); only 3/133 (2.3%) fatalities had no identified risk. Pneumonia was the commonest presentation (595; 52%), bacteraemia occurred in 633/1135 (56%), septic shock in 240 (21%) and 180 (16%) required mechanical ventilation. Cases correlated with rainfall with 80% occurring in the “wet” season (November through April). Median annual incidence rate was 20.5/100,000 people; the highest annual incidence in Indigenous Australians was 103.6/100,000 in 2011-2012. Over the 30 years, annual incidences increased as did the proportion with diabetes, while mortality decreased to 17/278 (6%) over the last 5 years. Genotyping of *B. pseudomallei* confirmed case clusters linked to environmental sources and defined evolving and new sequence types.

Interpretation: Melioidosis is an opportunistic infection , but with early diagnosis, specific antimicrobial therapy and state-of-the-art intensive care, mortality can be reduced to below 10%. However, mortality remains much higher in the many endemic regions where health resources remain limited. Genotyping of *B. pseudomallei* informs evolving local and global epidemiology.

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**Title Page**

**Melioidosis is an opportunistic infectious disease: the 30-year Darwin**

**Prospective Melioidosis Study**

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28

## 29 **ABSTRACT**

30 **Background:** The global distribution of melioidosis is under considerable scrutiny,  
31 with both unmasking of endemic disease in African and Pacific nations and evidence  
32 of more recent dispersal in the Americas. Tropical northern Australia has high  
33 incidence rates and The Darwin Prospective Melioidosis Study commenced in  
34 October 1989. We present epidemiology, clinical features, outcomes and bacterial  
35 genomics from this 30-year study, highlighting changes in the last decade.

36

37 **Methods:** Prospective study of epidemiological, clinical and laboratory data for all  
38 culture-confirmed melioidosis cases from the tropical Northern Territory of Australia  
39 from October 1<sup>st</sup> 1989 until September 30<sup>th</sup> 2019. Multivariable analysis determined  
40 predictors of clinical presentations and outcome. Incidence, survival and cluster  
41 analyses were facilitated by population and rainfall data and genotyping of  
42 *Burkholderia pseudomallei* including multilocus sequence typing and whole-genome  
43 sequencing.

44

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46 133 (11.6%) died. Median age was 50 years, 48 were children  $\leq$  14 years old (4.2%),  
47 721 (63%) were male and 600 (52%) Indigenous Australians. All but 186 (16%) had  
48 clinical risk factors; diabetes 513 (45%), hazardous alcohol use 455 (40%); only  
49 3/133 (2.3%) fatalities had no identified risk. Pneumonia was the commonest  
50 presentation (595; 52%), bacteraemia occurred in 633/1135 (56%), septic shock in  
51 240 (21%) and 180 (16%) required mechanical ventilation. Cases correlated with

52 rainfall with 80% occurring in the “wet” season (November through April). Median  
53 annual incidence rate was 20.5/100,000 people; the highest annual incidence in  
54 Indigenous Australians was 103.6/100,000 in 2011-2012. Over the 30 years, annual  
55 incidences increased as did the proportion with diabetes, while mortality decreased to  
56 17/278 (6%) over the last 5 years. Genotyping of *B. pseudomallei* confirmed case  
57 clusters linked to environmental sources and defined evolving and new sequence  
58 types.

59

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61 specific antimicrobial therapy and state-of-the-art intensive care, mortality can be  
62 reduced to below 10%. However, mortality remains much higher in the many endemic  
63 regions where health resources remain limited. Genotyping of *B. pseudomallei*  
64 informs evolving local and global epidemiology.

65

66 **Funding:** The Australian National Health and Medical Research Council.

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68

69     **Research in context**

70     **Evidence before the study**

71     For almost a century after its first description in Myanmar, melioidosis remained an  
72     enigmatic tropical disease from Southeast Asia described in military conflicts and the  
73     occasional returned traveller. Diagnosis requires culture of *Burkholderia*  
74     *pseudomallei* and that necessitates laboratory resources that remain unavailable in  
75     many tropical locations globally. Clinical descriptions were often of chronic  
76     pulmonary infection or of presumptive latency with activation years after infection.  
77     Recognition of the large and increasing burden of severe sepsis from melioidosis in  
78     endemic regions only became evident following studies in northeast Thailand  
79     supported by quality laboratory microbiological capability. Subsequently sequential  
80     randomized trials of antimicrobial therapy from Thailand have defined current  
81     therapy. In parallel to improved laboratory diagnostic facilities unmasking endemic  
82     locations across the globe, genotyping of *B. pseudomallei* has informed the  
83     epidemiology of melioidosis by tracking the source and mode of individual infections  
84     as well as supporting that intercontinental spread of this sapronotic pathogen has  
85     occurred over millennia.

86     **Added value of this study**

87     The 30-year Darwin Prospective Melioidosis Study began in 1989, the year of  
88     publication of the initial trial from Thailand that showed mortality halving with the  
89     introduction of ceftazidime. The 30-year data from 1148 consecutive cases describe  
90     distinct infecting scenarios and clarify the incubation period of acute disease,  
91     proportions with chronic melioidosis and the rarity of activation from latency.  
92     Demographics, risk factors and correlations with outcomes are quantified as is the  
93     diverse clinical spectrum of melioidosis. Genotyping of *B. pseudomallei* from cases

94 and the environment together with weather data support the importance of  
95 inhalational melioidosis during monsoonal events, as well as showing dynamic local  
96 bacterial dispersal which interplays with an evolving global story of dissemination of  
97 melioidosis, most recently in the Americas. With adequate laboratory and intensive  
98 care resources mortality under 10% has been achieved, with deaths only in those with  
99 clinical risk factors; highlighting the recognition of melioidosis as an opportunistic  
100 infectious disease.

101 **Implications of all the available evidence**

102 The continuing high mortality in many melioidosis endemic regions is a stark  
103 reflection of global resource disparities, with basic microbiology laboratory facilities,  
104 access to the required antibiotics and quality intensive care management for severe  
105 sepsis often unavailable. While considerable funding is being directed at *B.*  
106 *pseudomallei* as a potential biothreat agent, recognition of these clear and persisting  
107 inequities is arguably a greater priority.

## 108    **Introduction**

109    In 1912 melioidosis was first described in Burma as a newly recognised glanders-like  
110    disease of humans. Reports from other Southeast Asian countries soon followed.(1, 2)  
111    The genomics era has provided a fascinating but still incomplete picture of the global  
112    presence and spread of *Burkholderia pseudomallei*, the bacterium which resides in  
113    complex ecosystems in the soil and water of endemic regions and causes melioidosis  
114    in exposed humans and animals. Although glanders was described by Hippocrates,  
115    whole genome sequencing has shown that *Burkholderia mallei*, the bacterium causing  
116    glanders, is a derivative clone of *B. pseudomallei*, having evolved through gene loss  
117    and selection to be a horse-adapted pathogen that can no longer survive in the  
118    environment.(3) Glanders has been used in biological warfare and both *B. mallei* and  
119    *B. pseudomallei* are listed as Tier 1 select agents. Rapidly progressive sepsis with  
120    high mortality from aerosol inhalation of *B. pseudomallei* is the concern that funds the  
121    current diagnostic, therapeutic and vaccine initiatives for melioidosis.(4)

122

123    Genomic analyses suggest that *B. pseudomallei* evolved in the environment of  
124    Australia, subsequently spreading to Southeast Asia during the last ice age.(5) From  
125    Asia *B. pseudomallei* spread to Madagascar and Africa(6) and more recently from  
126    Africa to the Americas. Global modelling predicted an estimated 165,000 human  
127    cases annually, with 89,000 deaths.(7) Substantial improvements in surveillance and  
128    laboratory resources are needed to verify these predictions and confirm the burden of  
129    disease in comparison to currently-designated neglected tropical diseases.(8)

130

131    Geospatial modelling predicted many regions globally, including the southern USA to  
132    be receptive to, if not already endemic for, *B. pseudomallei*.(7) Past investigations

133 linked cases of melioidosis in the USA to either travel or prior residence in overseas  
134 endemic countries or probable infection from unidentified contaminated products  
135 imported from Asia.(9) Recent bacterial genotyping of one recent and another  
136 historical autochthonous case supports the hypothesis that melioidosis is now endemic  
137 in Texas, linking to known endemic regions in the Americas.(10)

138

139 The Darwin Prospective Melioidosis Study (DPMS) began in 1989, documenting all  
140 cases of melioidosis in the tropical “Top End” of the Northern Territory of Australia  
141 (Fig. 1). We described 252 cases after 10 years(11) then 540 total cases after 20  
142 years.(12) With an additional 610 cases over the last 10 years, we now present the  
143 findings from 1148 individuals with melioidosis collected prospectively over 30 years  
144 and describe dynamic changes in epidemiology, clinical parameters and bacterial  
145 genomics occurring over the 3 decades.

146

## 147 **Methods**

148 We included and prospectively followed all patients diagnosed with culture-  
149 confirmed melioidosis in the Top End over 30 years from October 1, 1989 until  
150 September 30, 2019. Culture-confirmed melioidosis is a laboratory-notifiable disease  
151 in the Northern Territory and blood, sputum, urine and pus cultures from both primary  
152 care clinics and hospitals are routinely processed to include identification of *B.*  
153 *pseudomallei*. Once *B. pseudomallei* was confirmed, all patients were managed by the  
154 Royal Darwin Hospital Infectious Diseases Department, with the vast majority  
155 admitted to Royal Darwin Hospital for assessment and commencement of intravenous  
156 therapy. Positive serology in symptomatic patients directed clinicians to provide

157 further cultures, with a confirmed diagnosis of melioidosis and inclusion in the study  
158 only for those culture-positive.

159

160 The definitions and descriptions used for clinical and demographic risk factors and for  
161 clinical illness parameters remained constant throughout the 30-years as described  
162 previously, (11, 12), and are summarised in Supplementary Document 1. First  
163 episode melioidosis was categorised as: acute (defined as symptoms present for < 2  
164 months); chronic (symptoms present  $\geq$  2 months); or activation from latency.  
165 Activation of disease from a latent focal infection was a clinician-directed  
166 categorisation based on two scenarios with culture-confirmed new clinical illness  
167 occurring in individuals with no prior culture-confirmed melioidosis; long-standing  
168 positive melioidosis serology with usually a very high titre and no recent exposure  
169 events; or individuals with long-standing pulmonary radiological abnormalities with  
170 new pneumonia specifically involving the area of radiological abnormality.

171

172 Over the 30 years Darwin and international melioidosis treatment guidelines have  
173 been informed by randomized comparative studies of antimicrobial therapy from  
174 Thailand.(13, 14) DPMS treatment includes  $\geq$ 14 days of intravenous ceftazidime or  
175 meropenem (meropenem being mostly restricted to those in intensive care), followed  
176 by eradication therapy with sulfamethoxazole/trimethoprim, usually for 3 months.(15)  
177 Therapy for all patients is directed by the Infectious Diseases Department, based on  
178 the Darwin treatment guidelines, which have evolved over the 30 years as  
179 described.(15) Once discharged from hospital on oral eradication therapy, patients are  
180 seen in clinic initially weekly then monthly, with follow up continued ideally until at  
181 least 6 months following completion of therapy. However, in reality around half of all

182 patients do not complete eradication therapy.(15) For complex clinical scenarios  
183 decisions are by Department consensus, following formal case discussions (moderated  
184 usually by BJC).

185

186 Recurrent melioidosis was defined as re-presentation with *B. pseudomallei* culture-  
187 positive clinical disease occurring after the time designated for treatment completion  
188 (both intravenous and oral phases) for the previous episode, irrespective of whether  
189 the patient was adherent to the therapy or initially lost to follow up. Recurrent  
190 melioidosis was determined to be relapse or new infection based on epidemiology and  
191 comparative genomics using multilocus sequence typing and whole genome  
192 sequencing. Primary study outcome was death from melioidosis, either during initial  
193 or recurrent melioidosis episode. Death from other causes was also recorded up to  
194 December 31, 2019.

195

196 Patient demographic, epidemiological, clinical and laboratory details were stored in  
197 MariaDB v10.2.31 (Oracle, California) and analysed using Stata v15.1 (Stata, Texas).

198 Details of the statistical methods used for patient data analysis, including  
199 multivariable logistic regression analyses to identify associations with clinical  
200 presentations, bacteraemia and a fatal outcome from melioidosis are provided in  
201 Supplementary Document 1. Population numbers by region and ethnicity were  
202 extrapolated from Australian census data. Methods for analysis of incidence rates and  
203 rainfall over the 30 years are in Supplementary Document 1, with incidence rate  
204 trends over time and correlations with season, region, ethnicity and rainfall calculated  
205 using generalized additive models. Estimates of survival at 60 days and 5 years after  
206 melioidosis diagnosis were obtained using Kaplan-Meier analyses in Graph Pad Prism

207 v7.04 (Graph Pad, San Diego, CA, USA). The survival curves with 95% confidence  
208 intervals were compared using the log-rank test. For full 5 years survival data, only  
209 cases of melioidosis until September 30<sup>th</sup>, 2014 were included. Multilocus sequence  
210 typing (MLST) was performed on *B. pseudomallei* isolates as summarised in  
211 Supplementary Document 1, with sequence types (STs) deposited on the *B.*  
212 *pseudomallei* MLST website (<https://pubmlst.org/bpseudomallei/>).(16)

213  
214 This study was approved by the Ethics Committee of the Northern Territory  
215 Department of Health and Menzies School of Health Research (02/38).

#### 216 217 **Role of the funding source**

218 The sponsor of the study had no role in study design, data collection, data analysis,  
219 data interpretation, or writing of the report. All authors have full access to all the data  
220 in the study and the corresponding author had final responsibility for the decision to  
221 submit for publication.

#### 222 223 **Results**

224 Two of the patients included in the 20-year cohort(12) were removed from analysis  
225 because they were from Central Australia and not treated at Royal Darwin Hospital.  
226 The final numbers were 252 cases in the first decade, 286 in the second and 610 in the  
227 third. There were 1212 episodes of melioidosis in 1148 individuals over 30 years,  
228 with 133 (11.6%) deaths attributable to melioidosis, 127 occurring from the initial  
229 infection and 6 following recurrent melioidosis (Fig. 2). Overall, sixty (5.2%)  
230 individuals had one or more recurrences of melioidosis, of which 44 were relapse and  
231 20 new infection (Fig. 2). Of the 1015 individuals who survived melioidosis as of

December 31, 2019, 37 (3.6%) were lost to follow-up and 336 (33%) had died from other causes, predominantly comorbidities.

### ***Epidemiology***

Median age was 50 years (range 7 months-97 years), with only 48 (4.2%) children  $\leq 14$  years old (Table 1). As previously identified, the commonest risk factors in those diagnosed with melioidosis were diabetes (45%) and hazardous alcohol use (40%)(11, 12), but neither were individually predictive of mortality (Table 1). The absence of any clinical risk factor was strongly predictive of survival; of the 186 (16%) patients with no identified clinical risk factor only 3 (1.6%) died from melioidosis. Patients with at least one clinical risk factor were 8.4 times (95% CI: 2.7–26) more likely to die from melioidosis than patients without risk factors.

Occupational exposure to soil/surface-water was documented for 187 (16%) and recreational exposure for 892 (78%). A potential infecting event was documented for 255 (22%) (Supplementary Table 1). For 70 patients with a suspected inoculating event on a known date, incubation period was 1-21 days (median 4 days; IQR 3-7 days). Common scenarios included recreational and occupational activities related to gardening and outdoor maintenance, such as cutaneous exposure through cuts and trauma, and presumptive aerosol exposure from lawn mowing, weed-whacking and high-pressure hosing (Supplementary Table 1).

Yearly case numbers tracked with rainfall (Fig. 3A) and severe weather events were linked to regional clusters.(17) Examples are included in Supplementary Document 1. During the wet season in Darwin, the monthly incidence rate per 100,000 people

257 increased on average 14% (95% CI 8.5–19.0) for every 100mm increase in total  
258 monthly rainfall while accounting for ethnicity and annual trends (see Supplementary  
259 Document 1 for methods).

260

261 Genotyping of *B. pseudomallei* confirmed contamination of unchlorinated water  
262 supplies was associated with one cluster of nine cases (four fatal) in a remote  
263 Aboriginal community(18) and a separate cluster of two cases on a rural property.(19)

264

265 923 first presentations (80%) were in the “wet-season” (November through April)  
266 (Supplementary Fig. 1). Annual incidence rates were 4.8–51.2 (median 20.5)  
267 cases/100,000 people (Fig. 3A). 2011-2012 had the highest overall incidence and for  
268 that 12-month period which included a particularly high-rainfall wet season, the  
269 incidence in Indigenous Australians across the Top End was 103.6/100,000 and for  
270 Indigenous Australians > 14 years of age residing in urban Darwin or surrounds the  
271 incidence was 315.4/100,000 (Supplementary Fig. 2). Accounting for trends over  
272 time, the monthly incidence rate for the Indigenous population in Darwin in the four  
273 high-rainfall months of December through March was on average 4.2 times higher  
274 (95% CI 3.5-5.1) than the non-Indigenous population in Darwin and 2.2 times higher  
275 (95% CI 2.0-2.5) than the Indigenous population in remote Top End regions (Fig 3B)  
276 (see Supplementary Document 1 for methods).

277

### 278 ***Clinical Presentation***

279 Of 1148 primary melioidosis presentations, 1013 (88.2%) were acute, 106 (9.2%)  
280 chronic and 29 (2.5%) were considered activation from latency (Fig. 2). Presentations  
281 with chronic melioidosis were predominantly with sub-acute pulmonary disease, often

282 mimicking tuberculosis (n = 44), or non-healing skin infections (n = 36). In  
283 comparison with those presenting with acute melioidosis, patients with chronic  
284 melioidosis were 6.5 times (95% CI 1.6-26) less likely to die from melioidosis (2/106  
285 died compared with 125/1013).

286

287 Blood cultures were positive for *B. pseudomallei* in 633/1135 (56%), septic shock  
288 occurred in 240 (21%), usually on presentation or within 24 hours, 278 (24%) were  
289 managed in the intensive care unit, 180 (16%) required mechanical ventilation and  
290 100 (9%) required renal replacement therapy.

291

292 Pneumonia was the primary diagnosis in 595 (52%) patients, skin abscesses in 149  
293 (13%), of whom only five were bacteraemic and none died, and genitourinary  
294 infection in 140 (12%), of whom 103 (74%) were males with prostatic abscesses.  
295 Bacteraemia with no evident focus of infection was the presentation in 130 (11%),  
296 commonly immunocompromised patients with acute febrile illness. Children were 5.3  
297 (95% CI 4.0-7.1) times more likely than adults to present with skin abscesses (28/48  
298 compared with 121/1100) and 3.9 (95% CI 2.0-7.8) times less likely than adults to be  
299 bacteraemic (7/48 compared with 626/1087).

300

301 In multivariable analysis, presentation with pneumonia was independently associated  
302 with diabetes, chronic lung disease, rheumatic heart disease or cardiac failure, female  
303 sex, Indigenous ethnicity and presentation in the four high-rainfall months of  
304 December through March (Supplementary Table 2A). Presentation with skin  
305 abscesses reflected younger, healthier people (Supplementary Table 2B). Independent  
306 predictors of bacteraemia were age  $\geq 50$  years, diabetes, hazardous alcohol use,

307 chronic kidney disease, immunosuppression, malignancy, Indigenous ethnicity and  
308 presentation in the four high-rainfall months (Supplementary Table 2C).

309

310 Secondary foci were identified up to 3 weeks after admission: secondary pneumonia  
311 in 107/553 (19%) without primary pneumonia, secondary prostatic abscesses in 40,  
312 osteomyelitis in 36, septic arthritis in 29 and secondary skin lesions usually as  
313 multiple pustules in 21 (19 bacteraemic).

314

315 Since 1995 all patients had imaging for internal abscesses, with abdomen/pelvis CT  
316 scan or abdominal ultrasound. Supplementary Table 3 lists organ abscesses and other  
317 infection foci. Notably 99 patients showed CT-scan evidence of mediastinal  
318 lymphadenopathy/inflammatory-masses and 10 an inflammatory gastrointestinal  
319 mass. Over the 30 years only 2 patients (0.2%) had parotid abscess, both adults.

320

### 321 ***Outcomes***

322 For the 60 patients with one or more recurrence, median time from the date of initial  
323 admission to first relapse was 8.2 (range 3.4-54) months, compared to 49 (range 10-  
324 225) months between initial admission and new infection. Correlates with relapse  
325 included diabetes (25/42 (60%) in relapse cases vs 430/979 (44%) in those with no  
326 relapse;  $p = 0.046$ ), and chronic renal disease (11/42 (26%) in relapse cases vs  
327 108/979 (11%) in those with no relapse;  $p = 0.0027$ ).

328

329 Of the 133 deaths from melioidosis, 127 died during the initial melioidosis episode  
330 (Fig. 2); median time from admission to death of 4 days (range 0-481)  
331 (Supplementary Fig. 3A-C). Eight patients were dead before arrival at hospital and 11

died on the day of admission. Five patients died on relapse and 1 from a new melioidosis infection (Fig. 2). Three children died, all with risk factors.(20) For the remaining 130 deaths, median age was 55 (range 20-93) years, with primary diagnosis including pneumonia 90 (68%), bacteraemic sepsis without focus 18 (14%), genitourinary sepsis 9 (7%) and neurological melioidosis 4 (3%). Three of the 133 deaths occurred in patients without identified clinical risk factors and 97 (73%) had diabetes and/or hazardous alcohol use. In multivariable analysis independent predictors of mortality were age  $\geq 50$  years, rheumatic heart disease or cardiac failure, malignancy and presentation during the four high-rainfall months (Supplementary Table 2D).

### ***The dynamic nature of epidemiology and outcomes over 30 years***

Three major trends were identified: an increase in the incidence of melioidosis (Fig. 3A); a rising proportion of cases from urban Darwin and surrounds (Fig. 3B and Supplementary Fig. 4); and a decrease in mortality (Supplementary Table 4). During the first 5 years mortality from melioidosis was 31% (27/88), but this fell to 6% (17/278) in the last 5 years ( $p < 0.0001$ ). This is unlikely to reflect improved case ascertainment capturing less severe disease. Over the 3 decades, there was no trend to increase in rates of bacteraemia or septic shock (Supplementary Table 4). Furthermore, there was a significant trend over the 3 decades of increase in the proportion aged  $\geq 50$  years, females and the proportions with diabetes, malignancy and immunosuppression, while the proportion with no risk factors significantly decreased (Supplementary Table 4, Supplementary Figure 4). The only significant change in clinical presentations was a trend of decreasing presentation with neurological melioidosis (Supplementary Table 4).

357

358 ***The dynamic nature of *B. pseudomallei* genotypes over 30 years***

359 MLST of the 1108 (97%) available isolates from 1148 primary presentations revealed

360 349 *B. pseudomallei* STs of which 243 were found only in a single patient. *B.*

361 *pseudomallei* STs from DPMS patients from Darwin and surrounding rural hamlets

362 were analysed over the 30 years and contrasted with those STs in rural towns and

363 remote Indigenous Top End communities. There was a large diversity of STs which

364 was especially evident in rural and remote Top End regions (Supplementary Fig. 5).

365 The number of different STs each year increased when rainfall and case numbers

366 increased, but in urban Darwin and surrounds there was a trend over time to fewer ST

367 numbers per case numbers (Supplementary Fig. 6). New STs continued to emerge

368 each year, but at a diminishing rate, with significantly lower ST diversity in years 21-

369 30 in comparison to years 1-10 (Supplementary Fig. 6).

370

371 The relative proportions of the common urban STs 36, 109 and 132 remained constant

372 (Supplementary Fig. 5). ST553 was rare in earlier years but increased to become the

373 commonest ST over the last 5 years. Environmental sampling revealed a suburban

374 hotspot for ST553, with ST553 cases clustering in this area.(21) The first case of

375 melioidosis due to ST562 occurred in urban Darwin in 2005(22), and since then the

376 proportion of cases due to this ST has increased as this presumptively introduced

377 Asian ST has become more widely established.

378

379

380 **Discussion**

381 Northeast Thailand and the Top End of Australia are the two locations where  
382 melioidosis has been consistently documented as a major cause of community-  
383 acquired sepsis and in particular community-acquired pneumonia,(23, 24) both having  
384 annual incidence rates of around 20 cases/100,000, with year-to-year variability  
385 linked to rainfall and with incidence rates increasing over time.(25, 26)

386

387 The Darwin prospective melioidosis study highlights the concept that *B. pseudomallei*  
388 is an opportunistic pathogen.(27) With rapid diagnosis, availability of appropriate  
389 specific antimicrobial therapy (ceftazidime or meropenem) and access to state-of-the-  
390 art intensive care, overall mortality of melioidosis can be <10%, with death a very  
391 rare outcome for healthy hosts. The relative contributions to the decrease in mortality  
392 in the Darwin study remains uncertain but it has been noted that the reduction in  
393 mortality coincided with the introduction of an intensivist-led model of care and the  
394 empiric use of meropenem for critically ill patients.(28) That mortality remains 40%  
395 or higher in many melioidosis-endemic regions reflects delays in presentation, limited  
396 laboratory resources and access to therapy and intensive care.(25, 29, 30) Indigenous  
397 Australians had higher incidences of melioidosis throughout the study, likely  
398 reflecting higher rates of both clinical risk factors and environmental exposure.  
399 Overall mortality was not higher for Indigenous Australians and Indigenous ethnicity  
400 was not an independent risk factor for death from melioidosis.

401

402 Support for recognition of melioidosis as an opportunistic infection is reflected in the  
403 findings from the Darwin study that clinical illness (melioidosis) and severe disease  
404 and death after exposure to *B. pseudomallei* are determined predominantly by host  
405 clinical risk factors. Each wet season weekend thousands of children and healthy

406 adults undertake recreational activities in often muddy sport fields, gardens and  
407 various tropical Darwin environments, where previous studies isolated *B.*  
408 *pseudomallei* from 7/10 popular sport fields.(31) Despite this exposure and the  
409 substantial increase in case numbers over 30 years, the number of children with  
410 melioidosis remains at 0-3 each year. The commonest presentation in children was  
411 skin abscesses, mostly single, without fever and with no bacteraemia or evidence for  
412 infection elsewhere, likely reflecting robust innate and adaptive immunity controlling  
413 the inoculated bacteria at the skin and preventing bacteraemic spread.

414

415 There remain important gaps in understanding the epidemiology of infection with and  
416 disease from *B. pseudomallei*. This is reflected in the paradox that, despite similar  
417 melioidosis incidence rates, seropositivity for *B. pseudomallei* in northeast Thailand is  
418 as high as 50%(32), while in northern Australia seropositivity is under 5%.(33)  
419 Substantial funding of research on melioidosis diagnostics, pathogenesis, therapeutics  
420 and vaccines has been driven by the concern that aerosolized *B. pseudomallei* poses a  
421 serious potential biothreat.(4) While inhalation of aerosols was postulated as the mode  
422 of infection in helicopter crews with melioidosis pneumonia during the Vietnam war,  
423 the relative contributions in endemic areas of inhalation, percutaneous inoculation and  
424 ingestion of *B. pseudomallei* remain unclear, as do the range of clinical scenarios  
425 following each mode of infection. It has been postulated that the far higher rates of  
426 parotid and liver abscesses in Thailand and elsewhere in southeast Asia than seen in  
427 Australia may be attributed to a higher rate of ingestion as the mode of infection in  
428 southeast Asia, where water supplies are often unchlorinated in rural regions. (27, 34)

429

430 While the spectrum of presentations of melioidosis is diverse, over half of cases in  
431 this and other studies present with pneumonia.(23, 24, 35) While the severity and  
432 outcome of melioidosis are predominantly driven by the predisposing host clinical  
433 risk factors, the contribution of mode of infection impacts particularly on melioidosis  
434 pneumonia.(4, 27) Aerosol inhalation of *B. pseudomallei* during severe weather  
435 events is supported by epidemiological analysis, aerosol sampling studies and the  
436 common radiological findings of mediastinal involvement. In Singapore  
437 proportionally more presentations were with pneumonia during heavy monsoonal  
438 seasons(35), consistent with our findings demonstrating strong correlations of high-  
439 rainfall months with pneumonia, bacteraemia and mortality. Air sampling in Taiwan  
440 during typhoons linked detection of *B. pseudomallei* with a surge of melioidosis  
441 downwind in an urban environment.(36) Whole-genome sequencing allowed us to  
442 match *B. pseudomallei* from air sampling to clinical *B. pseudomallei* isolates from a  
443 patient with pneumonia and mediastinal melioidosis.(37)

444

445 The global footprint of environmental *B. pseudomallei* remains unknown but is  
446 informed by improved environmental sampling and melioidosis case detection.  
447 Recent reports from the Americas suggest both unmasking of previously  
448 unrecognized endemic melioidosis in South America and a dynamic situation  
449 potentially with spread of *B. pseudomallei* northwards in the Caribbean and Central  
450 America.(38, 39) The report of two possibly endemic cases of melioidosis in  
451 Texas,(10) necessitates targeted environmental sampling in the southern United States  
452 to establish whether *B. pseudomallei* is endemic. Such confirmation would have  
453 major implications for current biosecurity rulings around the laboratory handling and  
454 transport of *B. pseudomallei* as a listed Tier 1 select agent.

455

456 Genotyping of recent clinical and environmental isolates of *B. pseudomallei* from  
457 Puerto Rico supports possible dispersal through severe weather events such as  
458 hurricanes.(39) Genotyping of 30 years of DPMS isolates shows a large genomic  
459 diversity reflecting the ancient origins of *B. pseudomallei* on the Australian continent,  
460 but also the dynamic nature of *B. pseudomallei* in a high-incidence endemic setting.  
461 While several dominant genotypes have persisted throughout the 30 years, two *B.*  
462 *pseudomallei* genotypes have emerged with each reflecting a different epidemiology.  
463 ST553 has increased to become the dominant genotype in the Darwin region, having  
464 proliferated during a period of intense urban construction. ST562 emerged in 2005 as  
465 a new genotype for urban Darwin. It is phylogenetically clearly an ST of Asian origin,  
466 with ST562 also reported from Hainan Island, China and Taiwan.(22) The lack of  
467 diversity amongst Australian ST562 strains suggests a recent point source  
468 introduction to northern Australia from Asia, but how and when this occurred remains  
469 unknown.

470

471 There are some limitations of this study. Over the 30 years we continued to use the  
472 original definitions for risk factors and septic shock(11), without adding  
473 contemporary pneumonia and sepsis severity scores developed since this prospective  
474 study commenced. Limiting severity analysis to bacteraemia, septic shock and death  
475 precluded a more in-depth analysis of pathogenesis. Phylogeographical analysis of *B.*  
476 *pseudomallei* genotypes is dependent on accurate location information and even with  
477 prospective collection of real-time patient history, attribution of location of infection  
478 will sometimes be incorrect for those who work and travel across the regions of the  
479 Top End. Our assessment of the extent of internal organ infection from disseminated

480 melioidosis is likely to be an underestimate, as routine imaging only commenced for  
481 all patients in 1995 and even with routine imaging less extensive organ involvement  
482 may be missed.

483

484 In conclusion, the 30-year prospective study of melioidosis cases in the tropical  
485 Northern Territory confirms the importance of diabetes as the major risk factor for  
486 melioidosis and defines the spectrum of presentations and disease in a well-resourced  
487 environment. Early diagnosis, specific antimicrobial therapy and state-of-the-art  
488 intensive care therapy have reduced the mortality from melioidosis to under 10%  
489 overall and to zero in healthy individuals. Genotyping of *B. pseudomallei* informs  
490 both local molecular epidemiology and the evolving global melioidosis story.

491

## 492 **Contributors**

493 Conceptualization (BJC); Patient data collection, management and follow up (BJC,  
494 EMM, RWB, RNP, CSM, APR, ES, JD, SHE, SJ, SL, PM, VLK, NMA); Project  
495 administration (BJC, MM); Supervision (BJC, MM, VLK, NMA); Laboratory  
496 procedures (RWB, MM, MK, JRW); Data curation (LMW, MM, JRW); Data  
497 accessed and verified (BJC, LMW, EMM). Formal analysis including genomics and  
498 visualization (LMW, MK, EMM, JRW, CW, PM, MM, BJC); Original draft  
499 preparation (BJC); Review and editing (all); Funding acquisition (BJC, MM, MK,  
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501

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## **Data sharing**

Individual patient data will not be available, but the data dictionary and bacterial genomic data can be provided on request.

## **Conflicts of interest**

We declare no conflicts of interest.

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644 **Tables**

645 **TABLE 1**

646 **EPIDEMIOLOGY, CLINICAL RISK FACTORS AND OUTCOMES FOR 1148**  
647 **PATIENTS WITH MELIOIDOSIS**

		Patients		Died from melioidosis						
		<i>n</i>	% of total	<i>n</i>	Mortality					
<b>Epidemiology</b>	Age <15y	48	4.2%	3	6.3%					
	Age 15-49y	524	46%	45	8.6%					
	Age 50+	576	50%	85	15%					
	Male	721	63%	86	12%					
	Female	427	37%	47	11%					
	Indigenous Australian	600	52%	72	12%					
	Non-Indigenous	548	48%	61	11%					
<b>Regions</b>	Darwin urban	632	55%	81	13%					
	Darwin rural hamlets	154	13%	12	7.8%					
	Regional & remote Top End	349	30%	37	11%					
	Outside Top End	13	1.1%	3	23%					
<b>Clinical Risk factors</b>						<i>p</i> <sup>3</sup>	RR	95% CI		
	Diabetes	513	45%	62	12%	0.72	1.1	0.78	-	1.5
	Hazardous alcohol use	455	40%	56	12%	0.72	1.1	0.80	-	1.5
	Chronic lung disease	312	27%	45	14%	0.13	1.4	0.98	-	1.9
	Chronic renal disease	140	12%	24	17%	0.10	1.6	1.06	-	2.4
	Malignancy	111	9.7%	20	18%	0.10	1.7	1.07	-	2.6
	Immunosuppressive therapy and other immunosuppression <sup>1</sup>	106	9.2%	18	17%	0.13	1.5	0.98	-	2.4
	Rheumatic heart disease and/or congestive cardiac failure	102	8.9%	19	19%	0.10	1.7	1.10	-	2.7
	Kava use	39	3.4%	5	13%	0.80	1.1	0.48	-	2.6
	Other <sup>2</sup>	39	3.4%	7	18%	0.29	1.6	0.79	-	3.2
	No clinical risk factors	186	16%	3	2.3%	<0.0001	0.12	0.04	-	0.37

648 <sup>1</sup>Clinical risk factor parameters as defined in refs 11 and 12 and Supplementary Figure 1

649 <sup>2</sup>Includes the only 4 patients with HIV infection (0.3%); a rate similar to that seen at Royal Darwin  
650 Hospital for sepsis in general.

651 <sup>3</sup>P values adjusted for multiple testing using the False Discovery Rate (FDR) method.

652

[Click here to access/download;Figure;FIGURE 1.tif](#) 

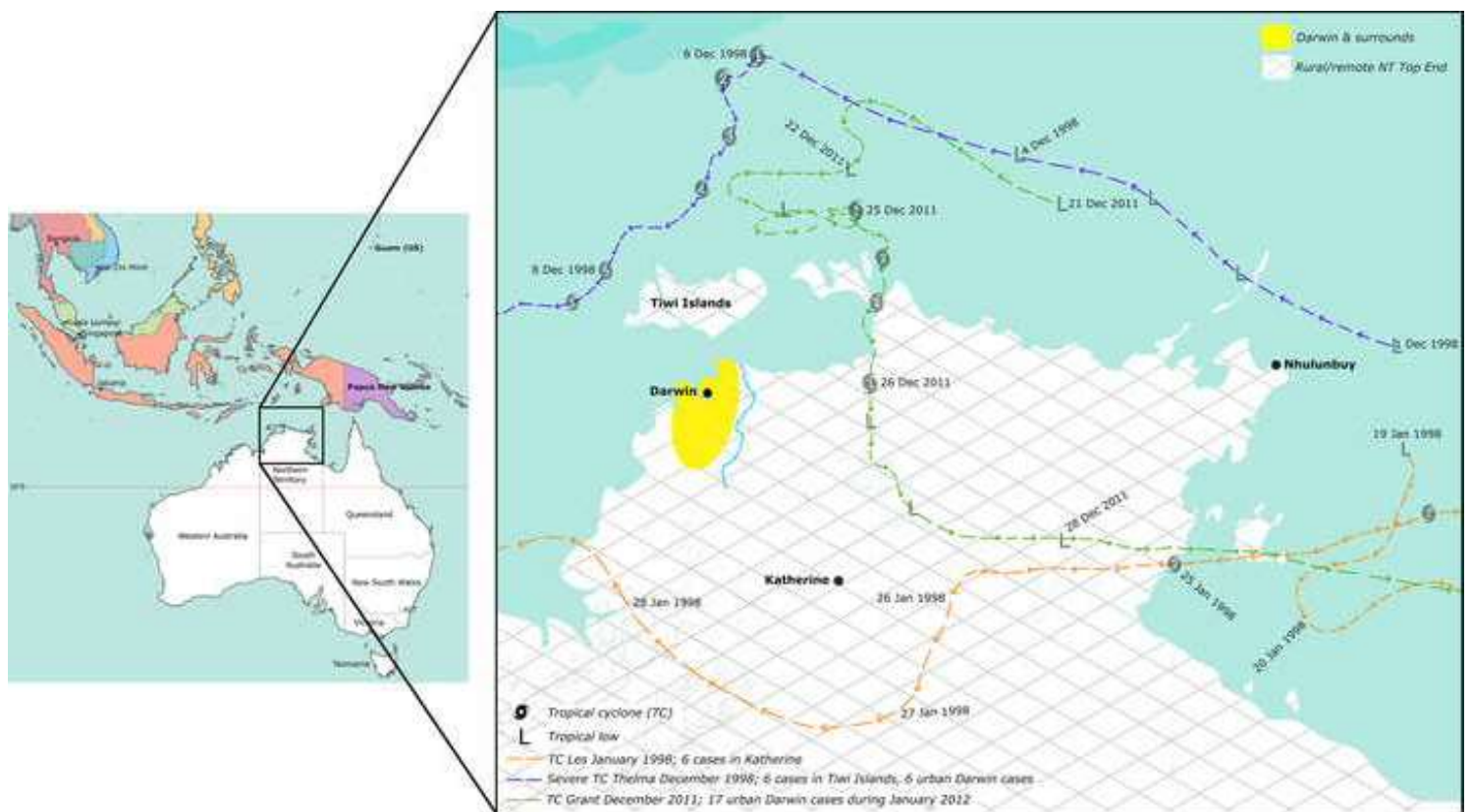


Figure 2

[Click here to access/download;Figure;FIGURE 2.tiff](#)

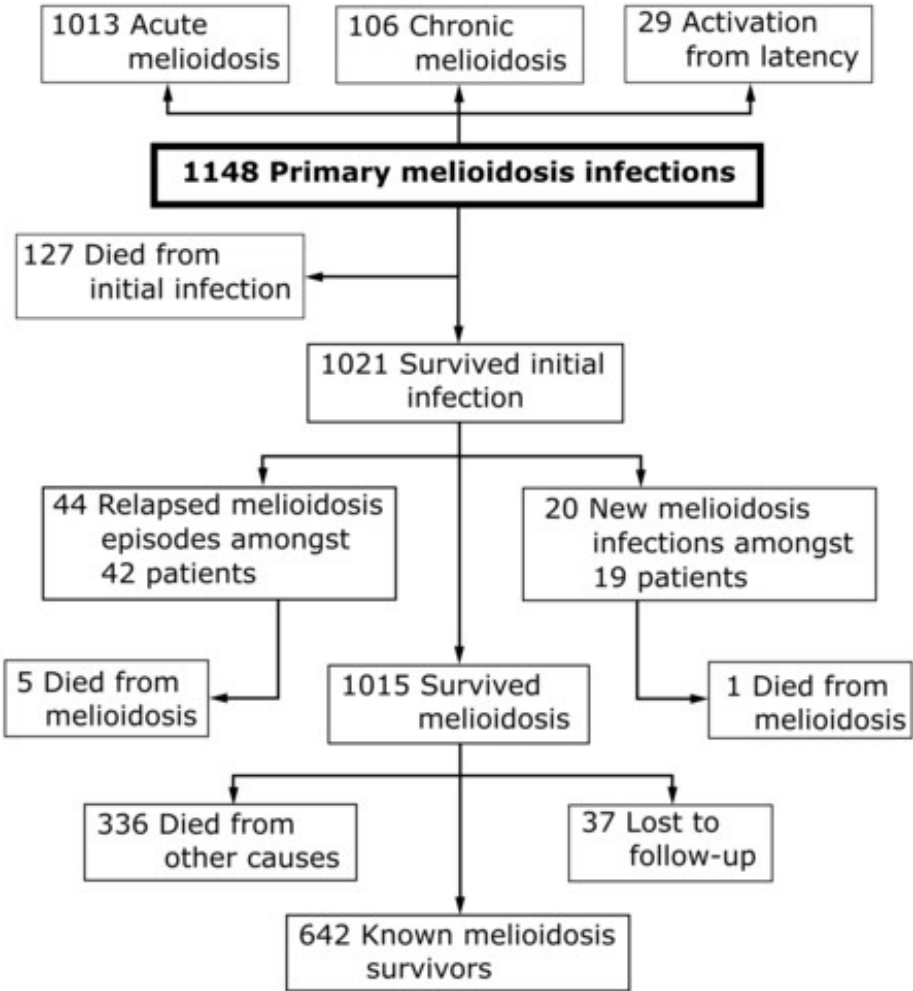


Figure 3A [Click here to access/download;Figure;FIGURE 3A.tif](#)

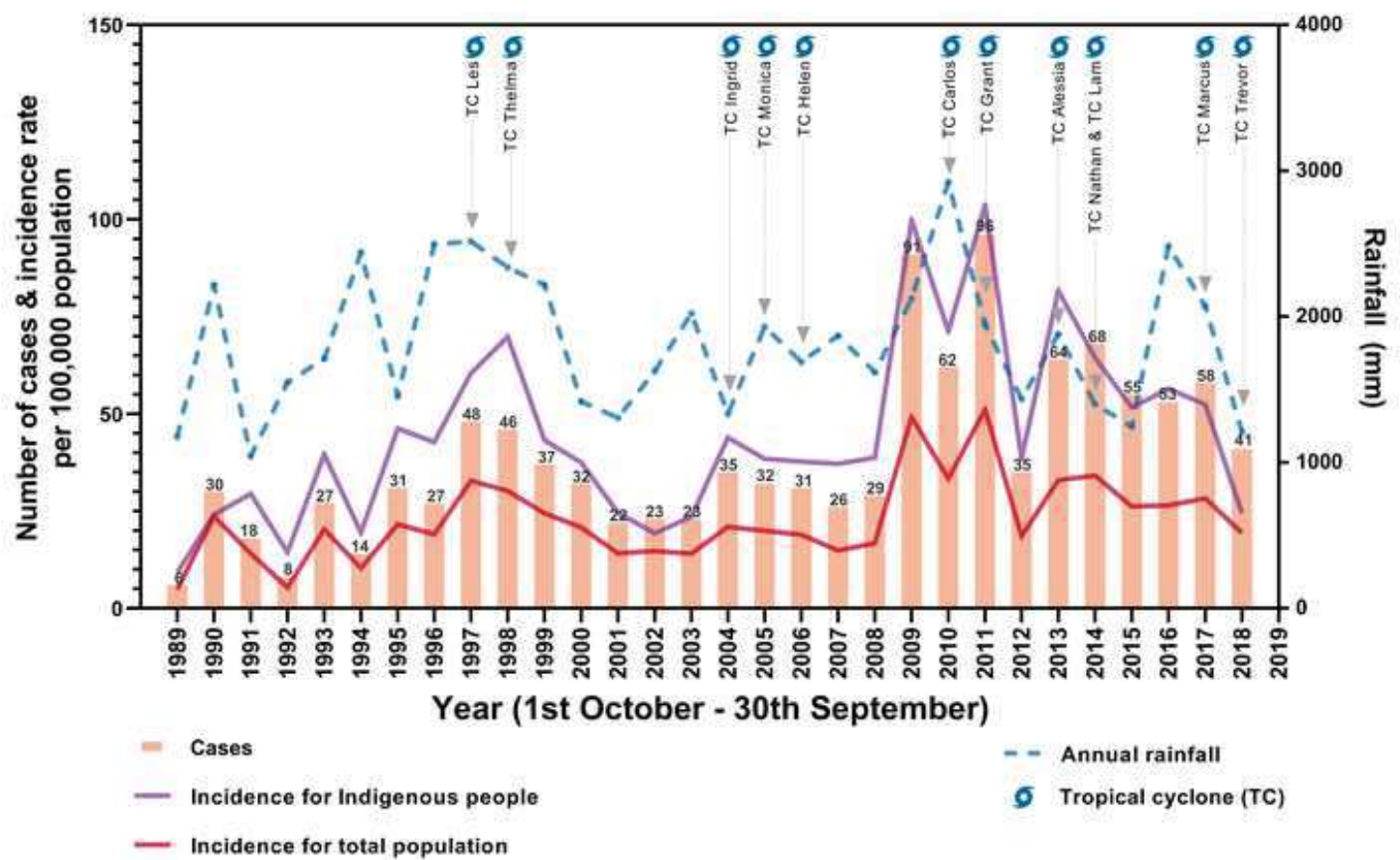
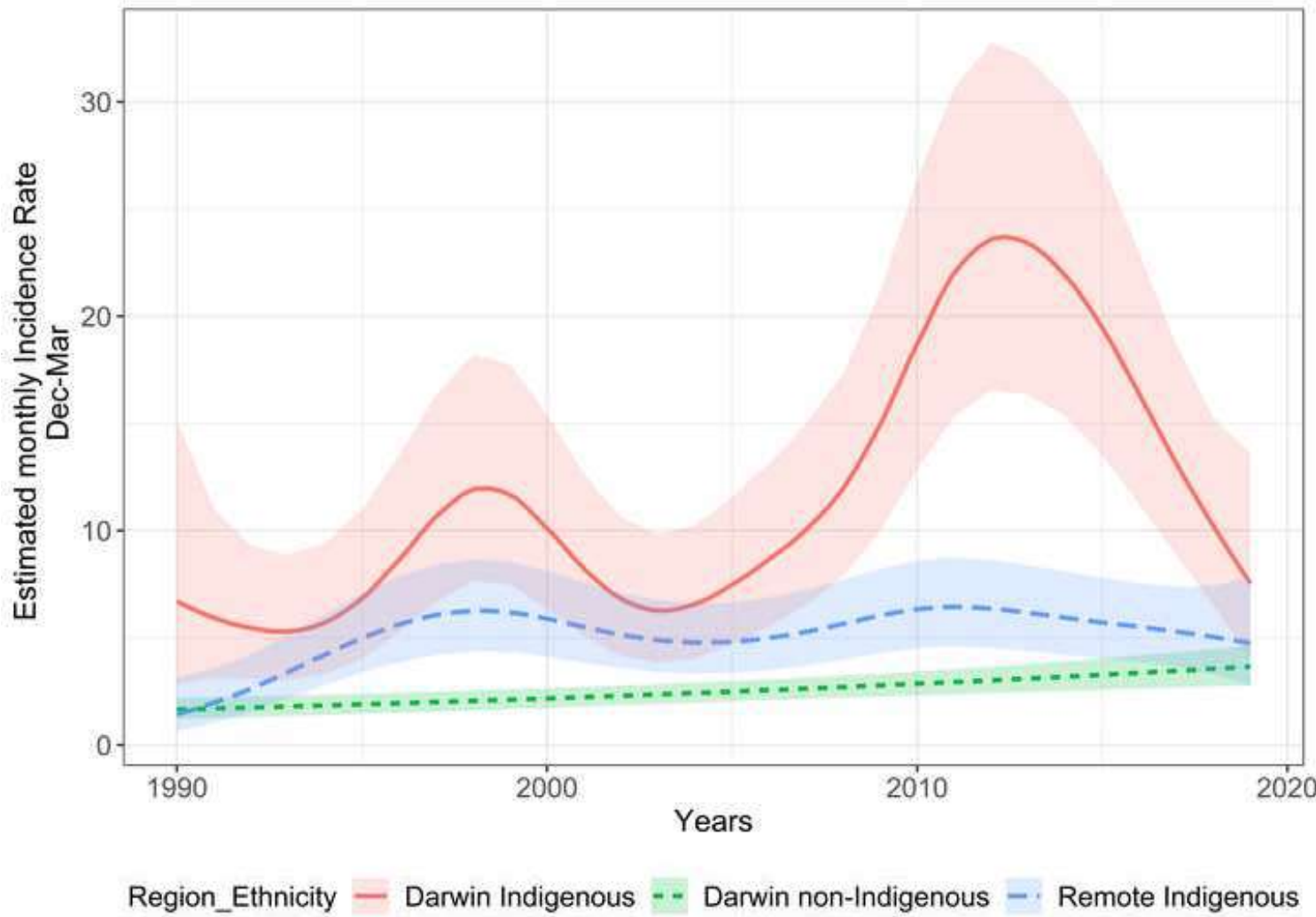


Figure 3B

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**SUPPLEMENTARY TABLE 1 PRESUMPTIVE INFECTION EVENTS IN 255 PATIENTS**

<b>Nature of Event</b>	<b>Number of cases</b>	<b>Event examples</b>
<b>Cut skin documented</b>	93 (36%)	15 gardening, 8 stick injury, 6 hunting, 5 fishing, 5 playing sports, 4 whipper-snipper/weed-wacker injury, 3 splinter, 2 swimming, 1 puncture wound, 1 pitchfork in duck yard
<b>Injury/accident with soil/water exposure</b>	14 (5%)	5 motor vehicle accident, 2 ingrown toenail resection, 1 crush injury, 1 fall from bike, 1 fall from flying fox, 1 fall from ladder, 1 fall from tree, 1 fall onto bitumen/asphalt, 1 potting mix in mouth post dental work
<b>Flood water exposure</b>	22 (9%)	17 cut skin
<b>Rainstorm exposure</b>	35 (14%)	6 cyclone exposures, 6 exposed under shelter, 1 fishing
<b>Gardening</b>	18 (7%)	4 mowing lawn, 2 mulching
<b>High pressure hosing</b>	8 (3%)	
<b>Ingestion</b>	2 (1%)	1 eating muddy sandwich, 1 mastitis breast milk ingestion
<b>Bite/sting</b>	8 (3%)	4 dog, 2 insect, 1 crocodile, 1 jellyfish
<b>Burns</b>	4 (2%)	
<b>Children playing in muddy yard</b>	5 (2%)	
<b>Other surface water/soil/mud exposure - non-occupational</b>	21 (8%)	3 aerosolized soil exposures, 3 car washing/cleaning, 2 walking in yard, 2 digging trench, 1 hunting, 1 clearing drains
<b>Other surface water/soil/mud exposure - occupational</b>	18 (7%)	4 earthmovings, 3 digging trench, 3 drilling, 1 fencing, 1 jackhammering, 1 cleaning pumps, 1 brick laying, 1 clearing drains, 1 commando crawling, 1 cleaning
<b>Swimming in river or waterhole</b>	7 (3%)	

**SUPPLEMENTARY TABLE 2A ASSOCIATIONS WITH PRESENTATION WITH PNEUMONIA**

	Patients			Prim Pneu			Bivariate			Multivariable			
	<i>n</i>	%		<i>n</i>	%		<i>p</i> <sup>1</sup>	OR		<i>p</i>	OR	OR 95% CI	
Age ≥50													
No	572	50%		274	48%		0.012			0.065			
Yes	576	50%		321	56%			1.4			1.3	0.98	-
Diabetes													
No	635	55%		286	45%		<0.0001			0.0004			
Yes	513	45%		309	60%			1.8			1.6	1.2	-
Hazardous alcohol use													
No	693	60%		338	49%		0.013			0.089			
Yes	455	40%		257	57%			1.4			1.3	0.96	-
Chronic lung disease													
No	836	73%		382	46%		<0.0001			<0.0001			
Yes	312	27%		213	68%			2.6			2.4	1.8	-
RHD/CCF													
No	1046	91%		524	50%		0.0003			0.023			
Yes	102	8.9%		71	70%			2.3			1.7	1.1	-
Season													
Apr-Nov	381	33%		146	38%		<0.0001			<0.0001			
Dec-Mar	767	67%		449	59%			2.3			2.2	1.7	-
Indigenous													
No	548	48%		244	45%		<0.0001			0.0001			
Yes	600	52%		351	59%			1.8			1.8	1.4	-
Sex													
Male	720	63%		340	47%		0.0001			0.011			
Female	428	37%		255	60%			1.6			1.4	1.1	-
Region													
Darwin Urban	785	68%		424	54%		0.026			0.0011			
Remote Top End	363	32%		171	47%			0.8			0.6	0.4	-
Kava													
No	1109	97%		584	53%		0.0043			0.038			
Yes	39	3.4%		11	28%			0.4			0.4	0.2	-

<sup>1</sup>P values adjusted for multiple testing using the False Discovery Rate (FDR) method.

**SUPPLEMENTARY TABLE 2B ASSOCIATIONS WITH PRESENTATION WITH SKIN ABSCESS(ES)**

	Patients			Primary Skin		Bivariate			Multivariable				
	<i>n</i>	%		<i>n</i>	%	<i>p</i> <sup>1</sup>	OR		<i>p</i>	OR	OR 95% CI		
Age ≥50													
No	572	50%		101	18%	<0.0001			0.0021				
Yes	576	50%		48	8.3%		0.4			0.5	0.3		0.8
Diabetes													
No	635	55%		127	20%	<0.0001			<0.0001				
Yes	513	45%		22	4.3%		0.2			0.2	0.1		0.4
Hazardous alcohol use													
No	693	60%		111	16%	0.0002			0.0004				
Yes	455	40%		38	8.4%		0.5			0.5	0.3		0.7
Chronic lung disease													
No	836	73%		140	17%	<0.0001			<0.0001				
Yes	312	27%		9	2.9%		0.1			0.2	0.1		0.4
RHD/CCF													
No	1046	91%		146	14%	0.0006			<i>p</i> ≥0.100				
Yes	102	8.9%		3	2.9%		0.2						
Season													
Apr-Nov	381	33%		73	19%	<0.0001			<0.0001				
Dec-Mar	767	67%		76	10%		0.5			0.5	0.4		0.8
Indigenous													
No	548	48%		114	21%	<0.0001			<0.0001				
Yes	600	52%		35	5.8%		0.2			0.3	0.2		0.4
Immunosuppression													
No	1042	91%		147	14%	0.0001			0.0074				
Yes	106	9.2%		2	1.9%		0.1			0.1	0.03		0.6
Malignancy													
No	1037	90%		146	14%	0.0002			0.011				
Yes	111	10%		3	2.7%		0.2			0.2	0.1		0.7
Chronic kidney disease													
No	1008	88%		147	15%	<0.0001			0.018				
Yes	140	12%		2	1.4%		0.1			0.2	0.04		0.8
Kava consumption													
No	1109	97%		149	13%	0.0065	(skin <i>n</i> =0)						
Yes	39	3.5%		0	0%								

<sup>1</sup>P values adjusted for multiple testing using the False Discovery Rate (FDR) method.

**SUPPLEMENTARY TABLE 2C ASSOCIATIONS WITH BACTERAEemia ON PRESENTATION**

	Patients			Bacteremic			Bivariate			Multivariable			
	<i>n</i>	%		<i>n</i>	%		<i>p</i> <sup>1</sup>	OR		<i>p</i>	OR	OR 95% CI	
Indigenous													
No	548	48%		246	45%		<0.0001			<0.0001			
Yes	600	52%		387	65%			2.2			2.3	1.8	-
Age ≥50													
No	572	50%		276	48%		<0.0001			<0.0001			
Yes	576	50%		357	62%			1.7			2.0	1.5	-
Diabetes													
No	635	55%		289	46%		<0.0001			<0.0001			
Yes	513	45%		344	67%			2.4			2.1	1.6	-
Hazardous alcohol use													
No	693	60%		358	52%		0.0041			0.0001			
Yes	455	40%		275	60%			1.4			1.7	1.3	-
Chronic kidney disease													
No	1008	88%		525	52%		<0.0001			0.0003			
Yes	140	12%		108	77%			3.1			2.2	1.4	-
Malignancy													
No	1037	90%		552	53%		0.0001			0.0027			
Yes	111	10%		81	73%			2.4			2.1	1.3	-
Immunosuppression													
No	1042	91%		547	53%		<0.0001			<0.0001			
Yes	106	9.2%		86	81%			3.9			4.4	2.6	-
RHD_CCF													
No	1046	91%		565	54%		0.016			<i>p</i> ≥0.100			
Yes	102	8.9%		68	67%			1.7					
Season													
Apr-Nov	381	33%		175	46%		<0.0001			0.0004			
Dec-Mar	767	67%		458	60%			1.7			1.6	1.2	-

<sup>1</sup>P values adjusted for multiple testing using the False Discovery Rate (FDR) method.

**SUPPLEMENTARY TABLE 2D ASSOCIATIONS WITH MORTALITY**

	Patients			Died			Bivariate			Multivariable				
	<i>n</i>	%		<i>n</i>	%		<i>p</i> <sup>1</sup>	OR		<i>p</i>	OR	OR 95% CI		
Age ≥50														
No	572	50%		48	8.4%		0.0034			0.0027				
Yes	576	50%		85	15%			1.9			1.8	1.2	-	2.7
Chronic lung disease														
No	836	73%		88	11%		0.0788			<i>p</i> ≥0.100				
Yes	312	27%		45	14%			1.4						
RHD/CCF														
No	1046	91%		114	11%		0.0543			0.012				
Yes	102	9%		19	19%			1.9			2.0	1.2	-	3.6
Immunosuppression														
No	1042	91%		115	11%		0.0788			<i>p</i> ≥0.100				
Yes	106	9.2%		18	17%			1.6						
Malignancy														
No	1037	90%		113	11%		0.0543			0.0198				
Yes	111	10%		20	18%			1.8			2.0	1.1	-	3.4
Chronic kidney disease														
No	1008	88%		109	11%		0.0543			0.058				
Yes	140	12%		24	17%			1.9			1.6	0.98	-	2.7
Season														
Apr-Nov	381	33%		31	8.1%		0.0285			0.033				
Dec-Mar	767	67%		102	13%			1.7			1.6	1.04	-	2.5
Decade														
1989-1998	252	22%		51	20%		0.0001							
1999-2008	286	25%		28	9.8%			0.43		0.0005	0.41	0.24	-	0.67
2009-2018	610	53%		54	8.9%			0.38		<0.0001	0.32	0.21	-	0.50

<sup>1</sup>P values adjusted for multiple testing using the False Discovery Rate (FDR) method.

Note: diabetes and hazardous alcohol use each had *p*>0.100 on bivariate analysis and therefore were not included in the model.

**SUPPLEMENTARY TABLE 3 INTERNAL ORGAN ABSCESES AND OTHER  
FOCI OF INFECTION FROM 1148 CASES OF MELIOIDOSIS**

Site	Number (%)
<b>Prostatic abscess(es)</b>	143 (20%) <sup>1</sup>
<b>Mediastinal lymphadenopathy/mass</b>	99 (9%)
<b>Splenic abscess(es)</b>	72 (6%)
<b>Liver abscess(es)</b>	47 (3%)
<b>Kidney abscess(es)</b>	37 (3%)
<b>Muscle abscess(es)</b> <sup>2</sup>	37 (3%)
<b>Lymphadenitis</b>	24 (2%)
<b>Pericarditis</b>	11 (<1%)
<b>Para-intestinal mass</b>	10 (<1%)
<b>Brain abscess</b>	7 (<1%)
<b>Subphrenic abscess</b>	7 (<1%)
<b>Mycotic (pseudo)aneurysm</b>	6 (<1%)
<b>Epididymo-orchitis</b>	6 (<1%)
<b>Adrenal abscess</b>	5 (<1%)
<b>Mastitis/breast abscess</b>	5 (<1%)
<b>Extradural abscess</b>	3 (<1%)
<b>Parotid abscess</b>	2 (<1%)

<sup>1</sup> Calculated for males

<sup>2</sup> Psoas, calf, thigh most common

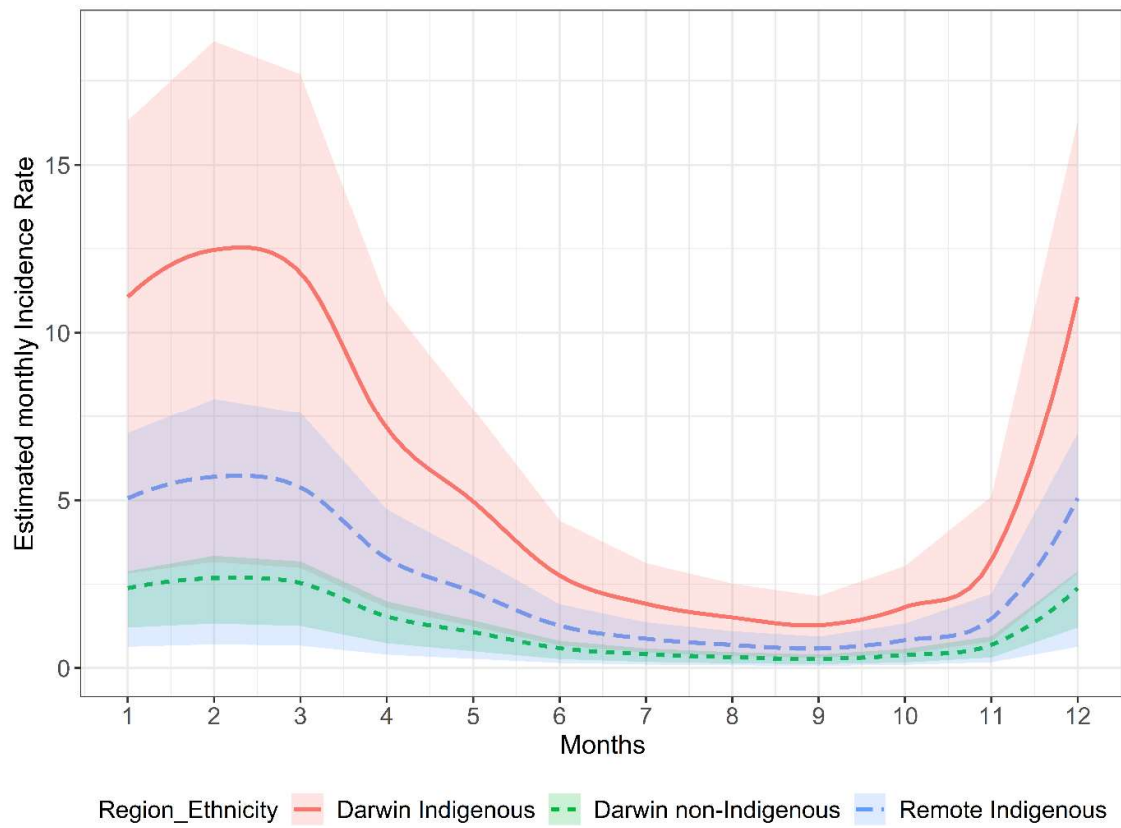
Note: Routine abdominal imaging on all cases commenced in 1995. Prior to 1995 abdominal imaging was performed at clinician discretion.

**SUPPLEMENTARY TABLE 4 EPIDEMIOLOGY, CLINICAL RISK FACTORS,  
CLINICAL PRESENTATIONS AND OUTCOMES BY DECADE**

		1989/1990- 1998/1999 <i>n</i> =252		1999/2000- 2008/2009 <i>n</i> =286		2009/2010- 2018/2019 <i>n</i> =610		<i>p</i> (trend)
		<i>n</i>	%	<i>n</i>	%	<i>n</i>	%	
Epidemiology	Male	190	75%	180	63%	351	58%	<0.0001
	Indigenous Australian	130	52%	148	52%	322	53%	-
	Age <15y	9	3.6%	15	5.2%	24	3.9%	-
	Age 15-49y	129	51%	141	49%	254	42%	
	Age 50+	114	45%	130	46%	332	54%	0.0051
	Urban Darwin and surrounds	152	60%	179	63%	455	75%	<0.0001
Clinical Risk Factors	Diabetes	93	37%	117	41%	303	50%	0.0002
	Hazardous alcohol use	95	38%	115	40%	245	40%	-
	Chronic lung disease	64	25%	77	27%	171	28%	-
	Chronic renal disease	24	10%	40	14%	76	13%	-
	Malignancy	11	4.4%	21	7.3%	79	13%	<0.0001
	Immunosuppressive therapy and other immunosuppression	14	5.6%	17	5.9%	75	12%	0.0004
	Rheumatic heart disease and/or congestive cardiac failure	17	6.7%	23	8.0%	62	10%	0.091
	Kava use	20	7.9%	7	2.4%	12	2.0%	<0.0001
	No clinical risk factors	54	21%	52	18%	80	13%	0.0015
Clinical Presentations  and Outcomes	Pneumonia	127	50%	149	52%	319	52%	-
	Genitourinary	37	15%	38	13%	65	11%	0.081
	Bacteraemia no evident focus	23	9.1%	37	13%	70	12%	-
	Skin infection	31	12%	38	13%	80	13%	-
	Soft tissue abscess(es)	13	5.2%	6	2.1%	27	4.4%	-
	Neurological	9	3.6%	5	1.7%	5	0.8%	0.0045
	Osteomyelitis	3	1.2%	4	1.4%	8	1.3%	-
	Septic arthritis	6	2.4%	7	2.4%	16	2.6%	-
	Other diagnosis	3	1.2%	2	0.7%	20	3.3%	-
	Bacteraemic	118	47%	180	63%	335	55%	0.15
	Septic shock	49	19%	72	25%	119	20%	-
	ICU admission	39	16%	92	32%	147	24%	0.069
	Mechanical ventilation	38	15%	66	23%	76	12%	-
	Died from melioidosis	51	20%	28	10%	54	9%	<0.0001

## SUPPLEMENTARY FIGURE 1

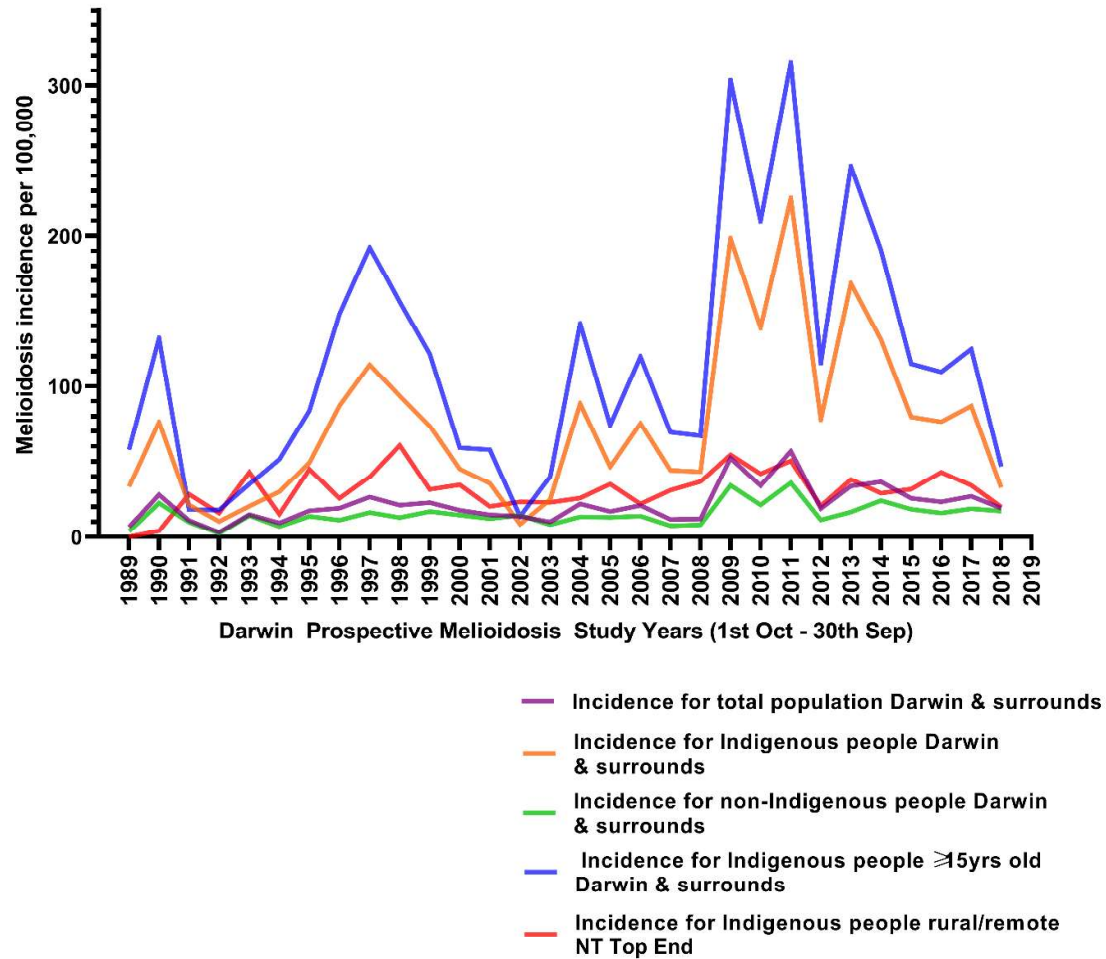
### ESTIMATED INCIDENCE RATES BY MONTH, ETHNICITY AND REGION FOR THE 30 YEARS



See Supplementary Document 1 for generalized additive model (GAM) methods. Shaded colour represents 95% CI for the estimated monthly incidence rates for each of the three groups.

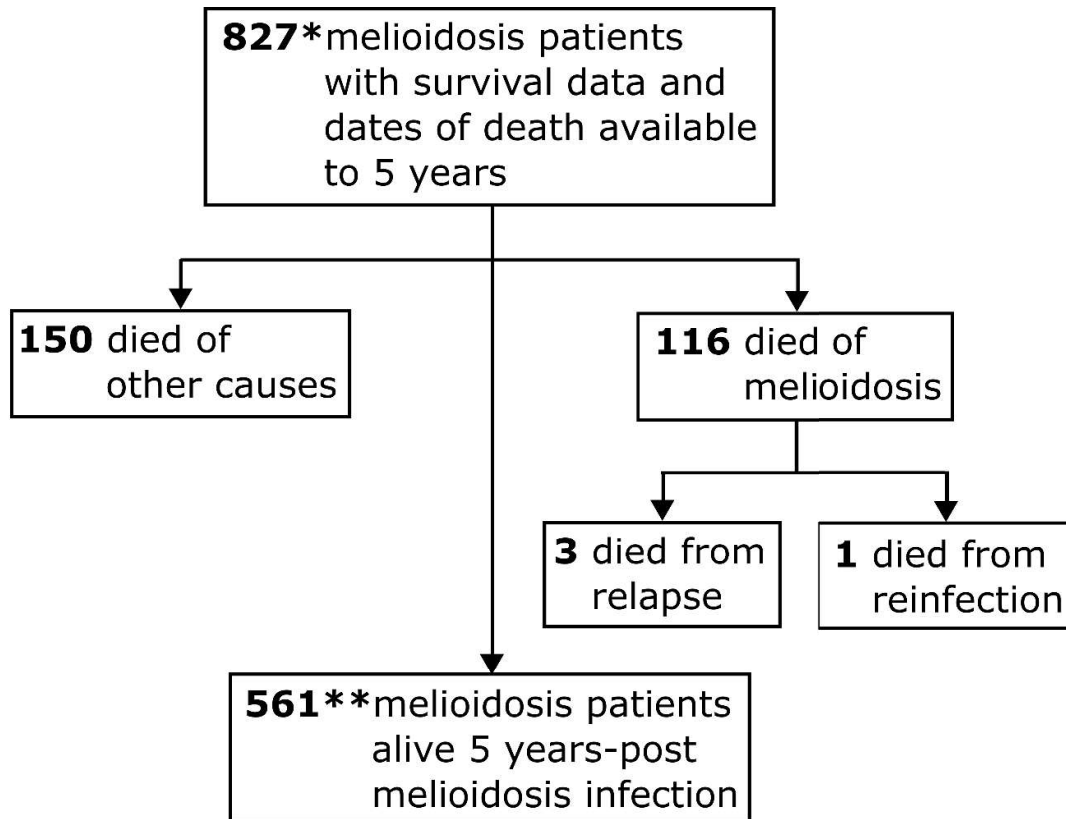
**SUPPLEMENTARY FIGURE 2**

**INCIDENCE RATES BY REGION AND ETHNICITY OVER 30 YEARS**



### SUPPLEMENTARY FIGURE 3A

#### KAPLAN-MEIER SURVIVAL ANALYSIS DATA

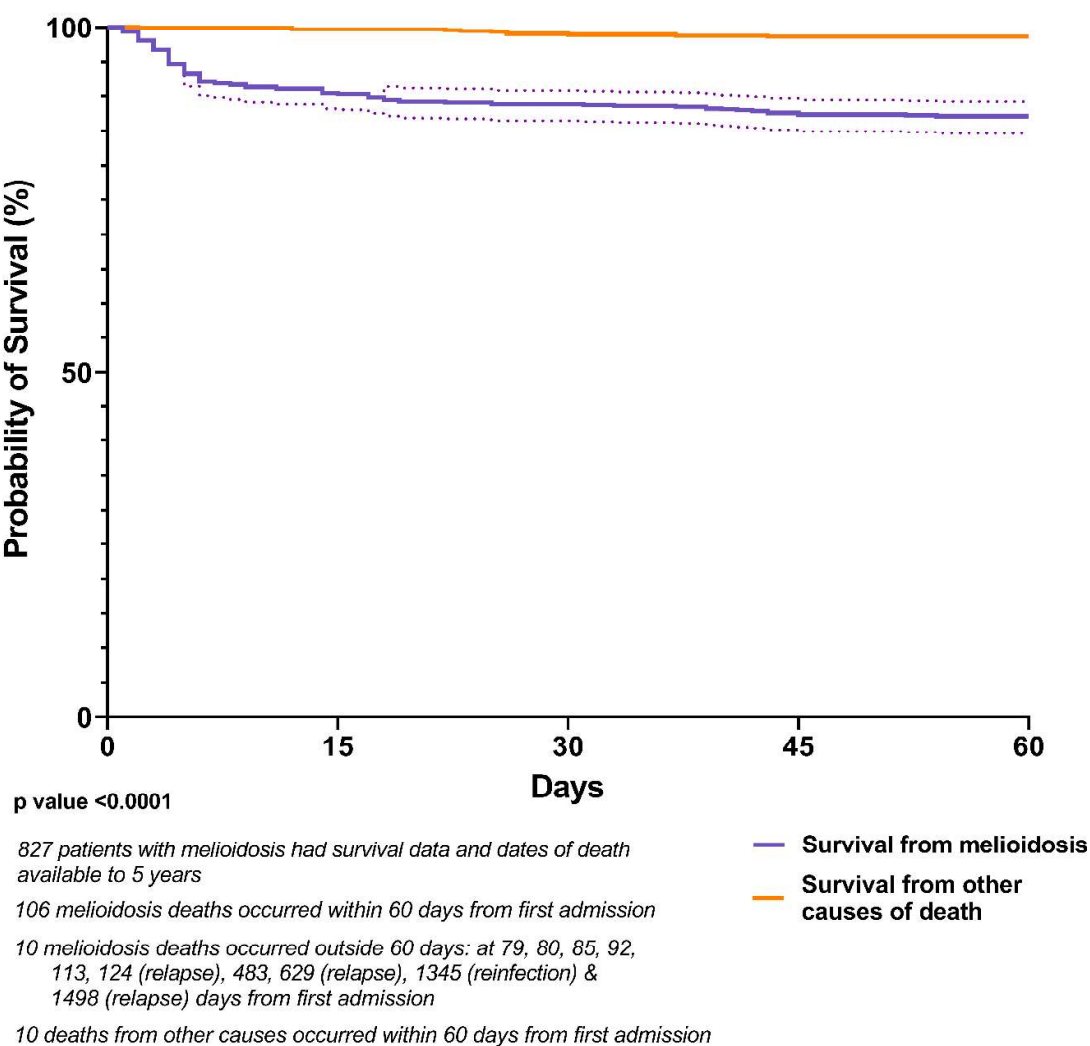


\* 321 DPMS patients were not included in the Kaplan-Meier survival analysis because either their admission date was after September 30th 2014 or they were lost to follow-up or their date of death was not recorded

\*\* An additional 142 deaths occurred from other causes amongst these 561 patients over subsequent years but there have been no further deaths from melioidosis

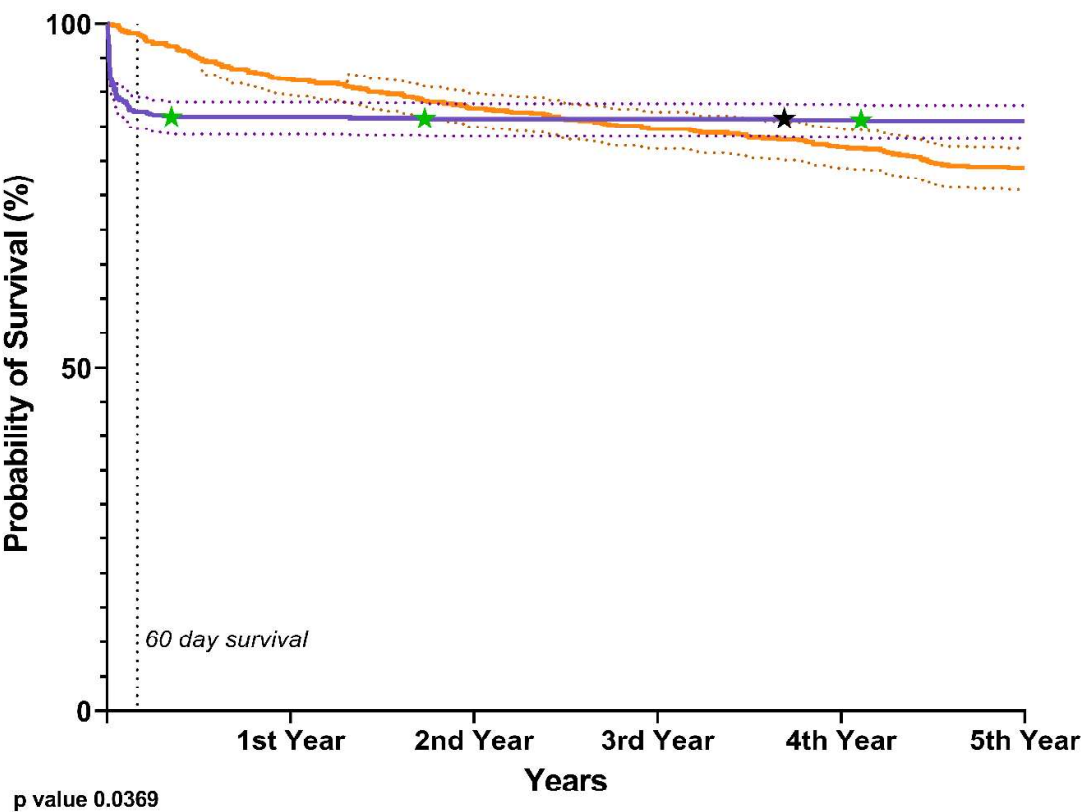
**SUPPLEMENTARY FIGURE 3B**

**KAPLAN-MEIER 60 DAY SURVIVAL ANALYSIS**



95% confidence intervals were used for error bars.

**SUPPLEMENTARY FIGURE 3C**  
**KAPLAN-MEIER 5 YEAR SURVIVAL ANALYSIS**

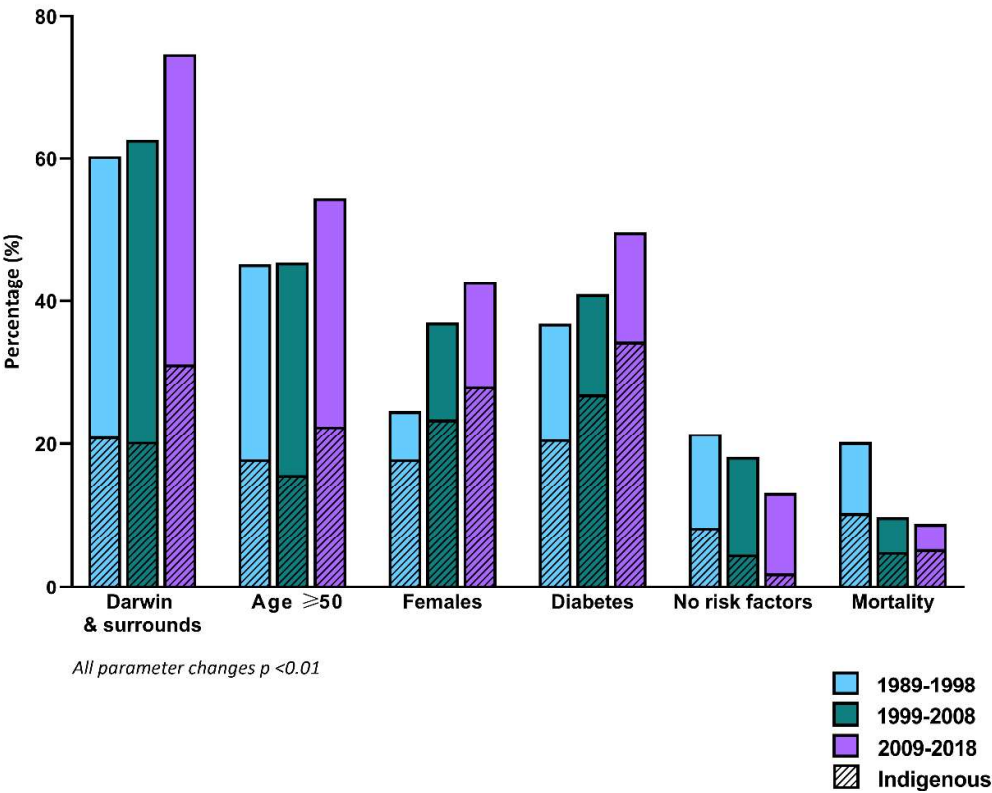


827 patients with melioidosis had survival data and dates of death available to 5 years  
 116 (all) melioidosis deaths occurred within 5 years from first admission  
 3 melioidosis relapse deaths occurred at day 124, 629 & 1498 from first admission  
 1 melioidosis reinfection death occurred at day 1345 from first admission  
 150 deaths from other causes occurred within 5 years from first admission

95% confidence intervals were used for error bars.

**SUPPLEMENTARY FIGURE 4**

**SIGNIFICANT PARAMETER CHANGES OVER THE 30 YEARS OF THE DARWIN PROSPECTIVE MELIOIDOSIS STUDY BY DECADE**



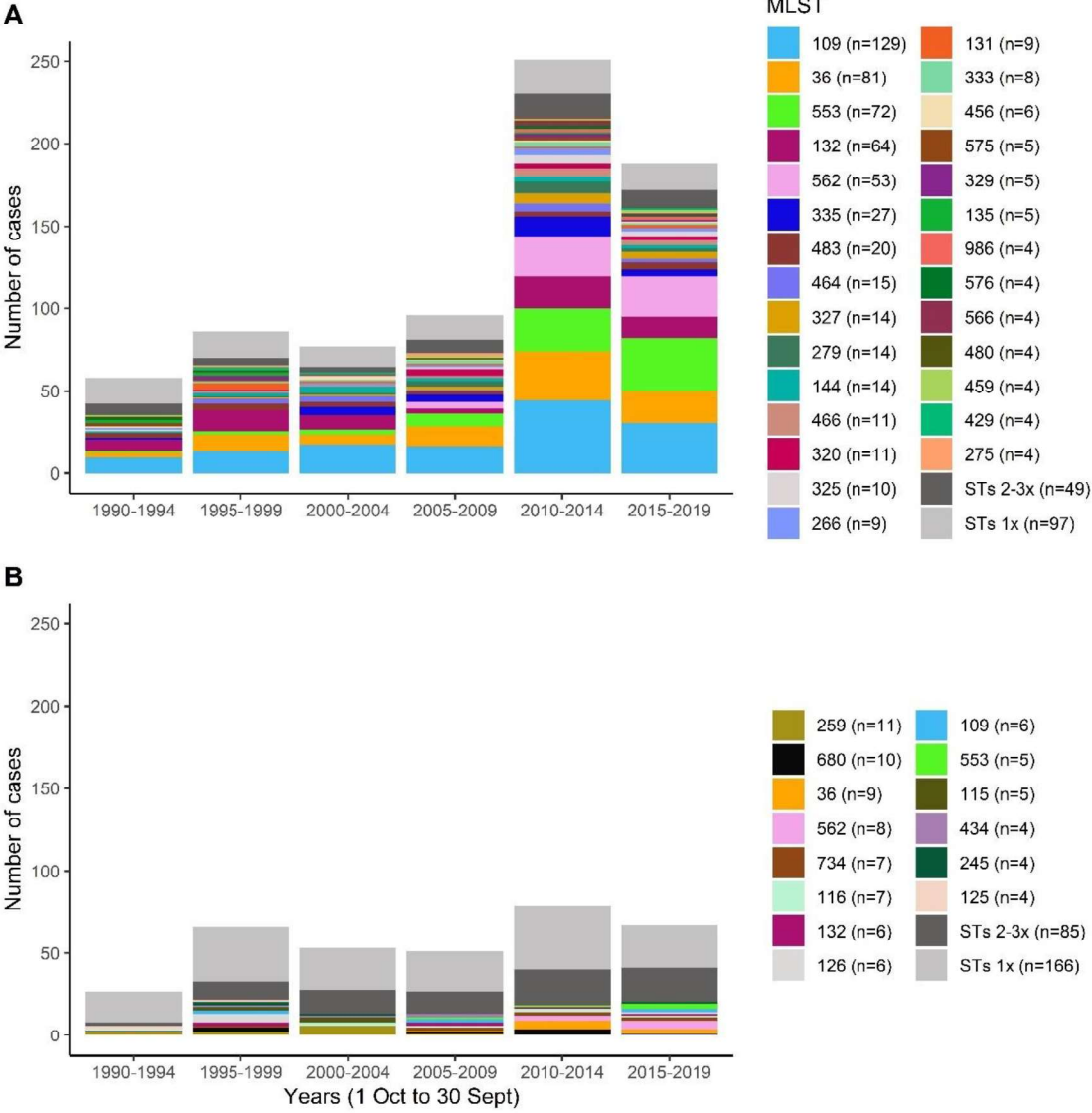
See Supplementary Table 4 for data on trends by decade.

**SUPPLEMENTARY FIGURE 5**

**BURKHOLDERIA PSEUDOMALLEI MULTILOCUS SEQUENCE TYPES  
DIVERSITY DYNAMICS OVER 30 YEARS**

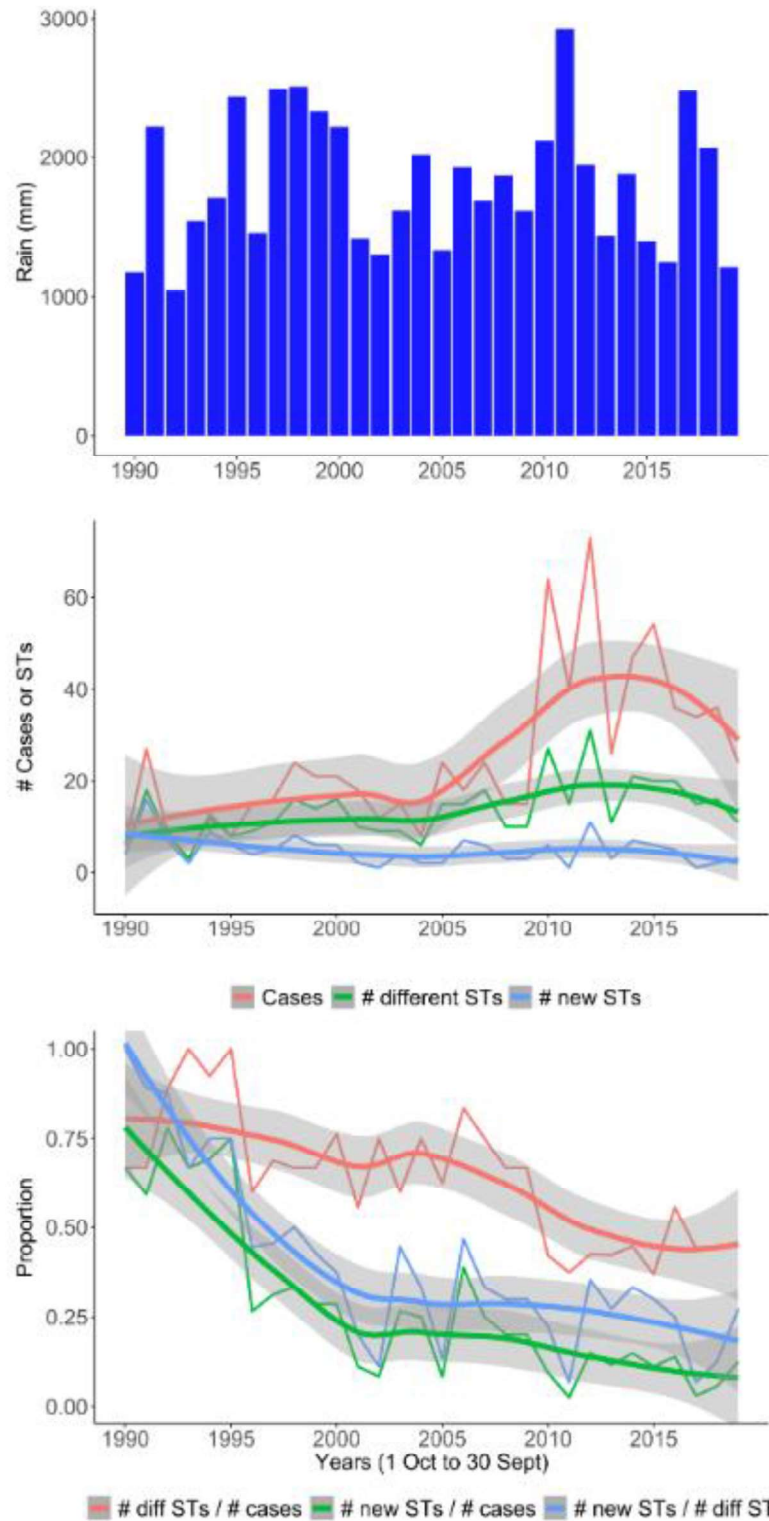
A. DARWIN AND SURROUNDS

B. RURAL AND REMOTE TOP END



# SUPPLEMENTARY FIGURE 6

## *BURKHOLDERIA PSEUDOMALLEI* MULTILOCUS SEQUENCE TYPES DIVERSITY DYNAMICS IN DARWIN AND SURROUNDS OVER 30 YEARS



# **1    Supplementary Document 1**

## **2    Supplementary Methods**

### **3    Setting**

4    The Top End of the Northern Territory encompasses around 500,000 km<sup>2</sup> (Fig. 1). Darwin  
5    (12°S latitude) with surrounding rural hamlets has a population around 150,000. The  
6    remaining population of around 50,000 live in the rural towns of Katherine and Nhulunbuy  
7    and numerous remote Indigenous communities with populations of 50-3000 people.

### **8    Definitions for demographical, clinical risk factors and clinical illness parameters**

9    Patient location was based on residence or a known likely location when infection occurred.  
10    Patients were each assigned to one of urban Darwin, Darwin rural hamlets, rural towns and  
11    remote Indigenous Top End communities or from outside the Top End (Fig. 1).

12    Clinical risk factors and clinical illness parameters used constant definitions over the 30  
13    years. Variables recorded were age, sex, ethnicity (Indigenous Australian Aboriginal or  
14    other), and the previously identified clinical risk factors of diabetes, hazardous alcohol use,  
15    chronic renal disease and chronic lung disease. Hazardous alcohol use was defined as greater  
16    than an average daily consumption of six standard drinks (60 g alcohol total) for males and  
17    four (40g alcohol total) for females. Chronic renal disease was defined as a creatinine of >  
18    150 umol/L (N. R. <90 umol/L) before the admission with melioidosis, or after completion of  
19    therapy if not previously documented. Chronic lung disease was defined as a documented  
20    diagnosis of chronic obstructive airways disease or bronchiectasis. Recent or current  
21    malignancy, immunosuppressive illness or immunosuppressive therapy, confirmed rheumatic  
22    heart disease or congestive cardiac failure and a history of recent kava ingestion were also

23 documented. “No clinical risk factors” referred to any patient with none of the above  
24 presumptive clinical risk factors.

25 Each patient was assigned to a single primary clinical diagnosis on presentation, representing  
26 the dominant organ involvement on clinical assessment by the Infectious Diseases team;  
27 pneumonia, skin infection without systemic symptoms, genitourinary melioidosis,  
28 bacteraemia with no evident focus, soft tissue abscess(es) either subcutaneous and/or lymph  
29 node, septic arthritis, osteomyelitis, neurological melioidosis and other. Presence or absence  
30 of bacteraemia was recorded. Septic shock was defined as the presence of hypotension not  
31 responsive to fluid replacement, together with hypoperfusion abnormalities manifest as end  
32 organ dysfunction (American College of Chest Physicians/Society of Critical Care Medicine  
33 Consensus Conference: definitions for sepsis and organ failure and guidelines for the use of  
34 innovative therapies in sepsis. Crit Care Med 1992;20: 864–874). This historical definition of  
35 septic shock was used throughout the 30 years and for this analysis contemporary pneumonia  
36 and sepsis severity scores such as the pneumonia severity index (PSI) and the APACHE score  
37 were not used, having not been created when this prospective study commenced.

### 38 **Statistical and graphical methods**

39 Patient demographic, epidemiological, clinical and laboratory details were stored in MariaDB  
40 v10.2.31 (Oracle, California) and analysed using Stata v15.1 (Stata, Texas). Relative risks  
41 were calculated to assess the risks of clinical risk factors among melioidosis patients on  
42 binary outcomes such as mortality and 2-sided Fisher’s exact test and 95% confidence  
43 intervals were computed. P values of bivariate analyses were adjusted for multiple testing  
44 using the False Discovery Rate (FDR) method. A multivariable logistic regression model was  
45 constructed to identify demographic, clinical risk factor, seasonal and regional associations  
46 with various clinical presentations (pneumonia, skin abscess), bacteraemia and a fatal

47 outcome from melioidosis. P values of bivariate analyses to select parameters for inclusion in  
48 the models were adjusted for multiple testing using the False Discovery Rate (FDR) method.  
49 All variables with  $p < 0.100$  in bivariate analyses were included in the initial models. Variable  
50 selection for the final multivariable models was through backward stepwise elimination  
51 ( $p < 0.100$  threshold). Of note, for the outcome of mortality, diabetes and hazardous alcohol  
52 use each had  $p > 0.100$  on bivariate analysis and therefore were not included in the model.

53 Incidence rate (IR) trends over time were estimated with the package “mgcv” in R (version  
54 4.0.2, R Development Core Team 2018) using a generalized additive model (GAM) with  
55 outcome being monthly melioidosis cases and parameters being region (Darwin vs remote),  
56 ethnicity (Indigenous vs non-Indigenous), season (dry vs wet season) and smoothing terms  
57 months (cyclic cubic regression spline) and years (1989-2019). The latter was estimated by  
58 region and ethnicity using thin-plate regression splines. In order to get monthly IR per  
59 100,000 population, an offset (log) population and a negative binomial family (log link) were  
60 applied. To account for temporal autocorrelation, a first-order autoregressive (AR1) error  
61 model with estimated rho 0.08 was used. Nested models were compared using the Akaike's  
62 information criterion (AIC) and model residuals were checked for no temporal  
63 autocorrelation and meeting distributional assumptions.

64 The association between IR trends over time and rainfall (monthly rainfall, Australian Bureau  
65 of Meteorology Darwin Airport weather station) were estimated in a separate model for the  
66 Darwin region and wet season (Nov to April) only. The same GAM model structure was used  
67 (estimated rho 0.23) with outcome being monthly cases in Darwin in the wet season, offset  
68 (log) population, parameters being ethnicity, rainfall and smoothing term years by ethnicity.

69 **Multilocus sequence typing**

70 The multilocus sequence typing (MLST) scheme for *B. pseudomallei* was first developed in  
71 2003 and targets the genetic sequence of seven housekeeping loci: ace (acetoacetyl coenzyme  
72 A reductase), gltB (glutamate synthase), gmhD (ADP-L-glycero-D-manno-heptose 6-  
73 epimerase), lepA (GTPbinding elongation factor), lipA (lipoic acid synthetase), narK (nitrite  
74 extrusion protein) and ndh (NADH dehydrogenase). For each housekeeping locus, the  
75 different sequences obtained from the *B. pseudomallei* isolates are assigned as distinct alleles.  
76 Each isolate is then defined by a string of seven integers (the allelic profile), which  
77 correspond to the allele numbers at the seven loci. Each unique allelic profile is assigned a  
78 sequence type (ST), which can then be assigned to any *B. pseudomallei* strain having that  
79 specific allelic profile. New allelic profiles not on the database are assigned the next available  
80 ST number by JRW, the curator of the global *B. pseudomallei* MLST database (previously  
81 housed at <http://bpseudomallei.mlst.net>; now at <http://pubmlst.org/bpseudomallei/>). In  
82 addition to conventional MLST with sequencing of each of the seven alleles, *in silico* MLST  
83 is now possible using allelic sequence data derived from whole genome sequencing to  
84 determine the ST.(16)

85

## 86 **Supplementary Results**

### 87 **Melioidosis and severe weather events**

88 The flooding and evacuation of the regional town of Katherine in January 1998, which  
89 followed extensive inland movement of tropical cyclone Les was associated with 6 cases of  
90 melioidosis (Fig. 1). The Category 5 severe tropical cyclone Thelma in December 1998 (Fig.  
91 1), resulted in 6 cases from the remote Tiwi Islands north of Darwin and 6 mainland cases  
92 from the Darwin region (Fig. 1)(17). Several severe weather events with high rainfall  
93 particularly in the Darwin region were associated with the two highest 12-monthly total case

94 numbers in the Top End – 91 cases in the 2009-2010 12 months and 96 cases in 2011-2012,  
95 with tropical cyclone Grant causing widespread flooding in December 2011 (Fig. 1, Fig. 3A).