

Border biosecurity interceptions for air passengers – assessing intervention methods and analytic tools

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Short running title: Air passenger biosecurity interventions

1 Abstract

2 At-border interventions are a critical step along the biosecurity continuum, to measure and
3 control the risks associated with the cross-border movement of people and goods. Air pas-
4 sengers are a high-volume pathway for a range of biosecurity risk materials, against which
5 various interventions may be used (e.g., manual searches, detector dogs, x-rays, etc.). Using a
6 large interception database for air passengers entering the southern Australian state of Tasma-
7 nia, this study applies common statistical modelling tools to assess the efficacy of interventions
8 (namely, dog detectors, and bag searches), and to identify pathway risk factors (e.g., flight
9 origin/route). Tasmania is an island state, and its environment and industries have benefited
10 from a low level of invasive pests due to their geographic isolation. Therefore, relatively strict
11 at-border interventions are used to prevent the entry of new pests, including some serious in-
12 vasive pests already present on mainland Australia (e.g., Queensland and Mediterranean fruit
13 fly, *Bactrocera tryoni*, *Ceratitis capitata*). This analysis considered the effects of interventions
14 on both voluntary declarations by passengers and also detections of undeclared risk material
15 on passengers. The analysis also focused on biosecurity risk items generally (e.g., fruits and
16 vegetables, meat products, cut flowers), and items that are specifically considered to be fruit-fly
17 hosts. The results highlight the strong positive effects of detector dogs on the rate of intercep-
18 tions, particularly of items detected on passengers. Conducting bag searches also appears to
19 increase interceptions, both by increasing the rate of items being detected and by encouraging
20 voluntary declarations. Sensitivity analyses then test the robustness of results to modelling im-
21 plementation methods and distributional assumptions. This study demonstrates how statistical
22 modelling can provide robust insights into biosecurity interventions and risk factors along path-
23 ways, and further highlights the value of high-quality interception data resources for informing
24 and improving biosecurity systems.

25
26 **Keywords:** border biosecurity, detector dogs, sniffer dogs, pathway risk analysis, passenger
27 screening, invasive species, fruit fly, *Bactrocera tryoni*, *Ceratitis capitata*.

1 Introduction

Biosecurity border interventions seek to balance the need for the cross-border movement of goods and people against their biosecurity risks. Although intervention policies inherit from a common framework of international agreements (Outhwaite, 2010), there are substantial differences in the policies and implementation of biosecurity interventions across jurisdictions. These differences produce variation in the risk of pest introductions across states, countries and regions (Whattam et al., 2014; Epanchin-Niell et al., 2021). Australia’s biosecurity system tends to have a higher appropriate level of protection and to implement comparatively stricter interventions than many other jurisdictions (Whattam et al., 2014; Black & Bartlett, 2020). This is partially due to the opportunity presented by Australia’s historical isolation, which has made its primary industries relatively pest-free on a global scale. However, contemporary activities such as tourism and trade are reducing this isolation and increasing the risk of pest introductions. Recent studies estimate an aggregated cost of invasive species to Australia to be AU\$389.59 billion since the 1960s (Bradshaw et al., 2021), and a net present value of AU\$314 billion for the national biosecurity system in terms of the assets that it protects (A. Dodd et al., 2020; Stoeckl et al., 2023). This highlights the critical importance of ensuring that border interventions act as effective barriers against biosecurity threats.

This is particularly true for the Australian island state of Tasmania (Fig. 1). Due primarily to their geographic and evolutionary isolation, island ecosystems possess a disproportionate level of the earth’s endemic species and support a large percentage of its biodiversity (Kier et al., 2009; Weigelt & Kreft, 2013). Tasmania has specifically been identified as a major centre of endemism for Australian flora, for example, more than half of the 30 native Eucalyptus species in Tasmania are endemic (Crisp et al., 2001; Potts et al., 2016). This isolation is an asset to agricultural producers who benefit from a relatively low-pest environment, but this also creates a biosecurity challenge because islands can be particularly vulnerable to the impacts of invasive pests and diseases (Keitt et al., 2011; Fraser, 2016; Brettell et al., 2021). For example, a 2018 incursion of Queensland fruit fly (‘Qfly’, *Bactrocera tryoni*) in northern Tasmania cost millions in direct eradication costs in addition to further indirect costs (e.g., via temporary market access losses; Blake, 2019). Furthermore, globalisation is expected to increase movement

57 across borders in both goods and people and to continue to bridge the geographic barriers that
58 once kept Australia and Tasmania isolated (A. J. Dodd et al., 2015; Seebens et al., 2017, 2021).

59 Air passengers are an important high-volume pathway for pest introductions, with around
60 20 passengers arriving annually in Australia in 2023-24 (BITRE, 2024). Pests may be in-
61 troduced via luggage (e.g., via infested fruit) or by being attached to passenger clothing or
62 belongings (e.g., soil on shoes or sports equipment; McNeill et al., 2011; Pace et al., 2022;
63 Robinson & McNeill, 2022). Air passenger pathways may be particularly important for pest
64 insects, including Mediterranean fruit fly ('Medfly', *Ceratitidis capitata*; Liebhold et al., 2006;
65 McCullough et al., 2006). Passengers are also a pathway for animal diseases, for example,
66 measures targeting African swine fever detected a yearly average of 33,684 pork products from
67 2% of screened passengers entering Australia from 2021/22 – 2023/24; (DAFF, 2024). Inter-
68 ventions on these pathways must therefore be able to mitigate a high volume and a diverse
69 range of biosecurity threats.

70 A range of risk mitigation tools can be employed at multiple points on the air passen-
71 ger pathway, from pre-departure, in-transit and on-arrival screening/inspection phases of the
72 biosecurity continuum (Whattam et al., 2014; Sequeira & Griffin, 2014). Common at-border
73 interventions for air passengers in Australia include manual examination, dog detector teams,
74 and x-rays (Inspector-General of Biosecurity, 2022).

75 Detector dogs may be particularly valuable in air passenger screening, being able to screen
76 large volumes of passengers and luggage efficiently, and able to be trained to target general
77 biosecurity materials as well as specific pests and diseases (Whattam et al., 2014; Moser et al.,
78 2020). Nonetheless, there are limited studies assessing their efficacy relative to other interven-
79 tion methods.

80 Furthermore, biosecurity interventions are often targeted towards specific flights to max-
81 imise the utility of limited resources, e.g., flight-based-traveller profiles used to target inter-
82 national arrivals into Australia (Inspector-General of Biosecurity, 2019). Therefore, further
83 empirical evidence about the relative efficacy of different interventions may help target inter-
84 vention resources towards the highest-risk arrivals.

85 The analysis of biosecurity interception/surveillance data is prone to some common issues

86 in statistical modelling, including zero-inflation (i.e., where data includes a large proportion
87 of zeros, for example where detections of targeted items are rare), overdispersion (e.g., where
88 variance is much higher than predicted), and censoring (e.g., if data is only recorded where
89 contamination is detected; Kachigunda, 2020; Turner et al., 2020; Trouvé & Robinson, 2021;
90 Kachigunda et al., 2022). Failing to account for overdispersion or zero-inflation in data can lead
91 to biased or inaccurate parameter or error estimates (Harrison, 2014; Feng, 2021; Campbell,
92 2021). Although some studies suggest that the outputs of mixed-effects models can be robust
93 to violations of distributional assumptions (e.g., Schielzeth et al., 2020; Knief & Forstmeier,
94 2021), exploring the potential effects of model design and implementation may be important
95 considerations when using interception data to inform biosecurity decision-making.

96 Focusing on domestic interstate flight arrivals into Tasmania, the goal of this study is to
97 assess the efficacy of passenger interventions and pathway risk factors on biosecurity inter-
98 ceptions. This focuses on both general biosecurity risk material ('BRM') interceptions, and
99 interceptions of BRM material specifically relevant to Qfly and Medfly (collectively referred to
100 as fruit fly, 'FF'). Preventing FF incursions has been a focus of Tasmania's border biosecurity
101 system, particularly following the 2018 incursion. This study uses recent air pathway inter-
102 vention data for Tasmania. These data are rich resources for our remit, as they include records
103 for all commercial arrivals, as well as relevant data on the types/amount of BRM intercepted.
104 This provides a valuable opportunity to apply statistical modelling approaches and assess their
105 sensitivity to implementation methods. The specific aims of this analysis were:

- 106 1. To determine the relative effects of different interventions (namely, dog detector teams
107 and luggage searches) on the rates of BRM and FF host interceptions, including voluntary
108 declarations by passengers and involuntary detections of items by biosecurity officers.
109 We did not make any specific directional predictions about the effects of searches and
110 detector dogs on interceptions.
- 111 2. To identify pathway-risk heterogeneity based on the origin and specific routes of flights.
112 We expected substantial variation in interception rates related to flight origin and route,
113 which may be used to identify high-risk arrivals.
- 114 3. To test whether our results are sensitive to overdispersion and zero-inflation by imple-

115 menting Bayesian mixed models with zero-inflated Poisson and negative binomial dis-
116 tributions. We expected the outputs of models and the estimated effects of intervention
117 methods to be robust to different implementation approaches.

118 2 Methods

119 Data context and overview

120 Tasmania is an island state (see Fig. 1), with a cool temperate climate, unique natural ecosys-
121 tems characterised by high endemism (Potts et al., 2016; Crisp et al., 2001), and a large primary
122 industry sector with and income from agriculture, forestry and fishing industries worth around
123 AU\$3.5 billion in 2022–23, or ~9% of the Gross State Product (ABS, 2023). Local industries,
124 communities, and natural ecosystems benefit from the state’s relative isolation and low levels
125 of pests, including species that are present elsewhere in Australia, such as Qfly, Medfly, tomato
126 potato psyllid (*Bactericera cockerelli*), and grape phylloxera (*Daktulosphaira vitifoliae*; Cook &
127 Fraser, 2015; Florec et al., 2013; Moir et al., 2022; Skinner, 2018).

128 The main entry pathway for passengers is by air, with a significant but smaller volume of

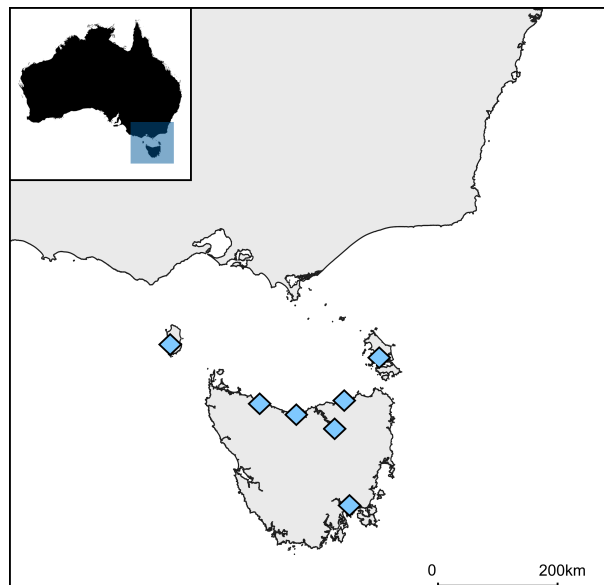


Figure 1: Location of Tasmania in relation to mainland Australia (inset), including the locations of the seven arrival ports for air passengers, namely (clockwise from top left) King Island, Burnie (Wynard), Devonport, Launceston, Bridport, Flinders Island, and Hobart. (Note, airports have been anonymised for the remainder of the analysis and labelled Airport_A, Airport_B. etc.)

129 maritime arrivals (e.g., ferries, cruise vessels, private vessels, etc.). Interceptions from air pas-
130 sengers are recorded in the Biosecurity Activity Database System (referred to as ‘BAS data’),
131 from which data from 1/Jan/2019 – 1/Sept/2023 was available. There are BAS interception
132 records for 59,917 domestic interstate flight arrivals, carrying over 6.5 M passengers (~1.4
133 M/year on average), from which 66,675 BRM interceptions were made.

134 BRM items are generally defined to include fresh produce (i.e., fruits and vegetables), an-
135 imal products including seafood, live animals, plant material (e.g., nursery stock, seeds), and
136 soil attached to sports equipment or clothing. Biosecurity interventions for Tasmania have a
137 particular focus on preventing incursions of FF into Tasmania, and a large subset of BRM in-
138 terceptions (43,803, or approximately 2/3) are of items considered to be FF hosts. For this
139 analysis, FF hosts include 130 taxa listed as Medfly and/or Qfly hosts in the Plant Biosecurity
140 Manual Tasmania 2023 (Biosecurity Tasmania, 2023). For details of BRM and FF host item
141 definitions and of intercepted BRM items, see Supplementary Materials A.

142 **Data processing**

143 Data for 59,917 arrivals was found to be within the scope of analysis, which excludes flights
144 from international origins or within the state, and flights with no data (i.e., cancelled, diverted,
145 missed, or cleared remotely; ~14% of all records). A further subset of 27 records was excluded
146 because of apparent data entry issues, and 25 arrivals into one airport were excluded as no
147 commercial flights arrive at this location, and interceptions for the remaining private arrivals
148 were extremely low, causing computational issues with model implementation. As only a small
149 fraction of actual arrivals are excluded, and missed arrivals do not appear to be targeted/biased
150 towards specific arrivals, we are confident that these exclusions do not reduce the operational
151 relevance of the analyses.

152 Six count variables were used as response variables, namely:

- 153 1. the total number of BRM interceptions per flight (*N.Total*);
- 154 2. the number of BRM declarations by passengers (*N.Declarations*);
- 155 3. the number of undeclared BRM interceptions (*N.Detections*);

- 156 4. the total number of FF host interceptions per flight (N_{Total_FF});
- 157 5. the number of FF host declarations by passengers ($N_{Declarations_FF}$); and,
- 158 6. the number of undeclared FF host interceptions ($N_{Detections_FF}$).

159 Total BRM and FF host interceptions are the sum of their corresponding declared and un-
160 declared detection counts. Both BRM and FF variables were used to explore how interventions
161 perform against both general biosecurity threats as well as high-priority/high-risk biosecurity
162 materials. Both detections and declarations were included to explore how interventions influ-
163 ence both voluntary and involuntary compliance behaviour in passengers (e.g., whether detector
164 dogs primarily increase interception through direct detections, or whether they also encourage
165 voluntary declarations).

166 The number of interceptions was calculated as the sum of each distinct type of BRM/FF
167 host material, separated by the passenger (e.g., if 2 passengers are intercepted each carrying
168 3 types of BRM, $N_{Total} = 6$). The rationale is that each commodity type may represent a
169 distinct biosecurity threat, as may the same kind of commodity being carried by two separate
170 passengers.

171 **Statistical analysis A: Intervention and pathway risk effects**

172 Generalized linear mixed effects ('glm') models with a Poisson distribution were implemented
173 via package 'lme4' (v1.1-33, Bates et al., 2015), in the R statistical environment (v4.2.3, R
174 Core Team, 2013). This was chosen for the primary analysis, as lme4 is an accessible package
175 that can implement models using common distributions, relative to more complex Bayesian
176 implementation methods that may be required for more advanced model types. Therefore, this
177 approach may be more relevant for use by non-academic users such as biosecurity managers.

178 Four fixed effects were included, namely: arrival airport (*Location*), intervention regime
179 (*Regime*), number of bag searches (*BagSearchCount*), and number of passengers per flight
180 (*PassengerCount*). *Regime* includes five combinations of dog detector teams ('DDTs') and
181 biosecurity inspectors ('BIs'), i.e. one BI, two BIs, one DDT, one DDT with one BI, and two
182 DDTs. Both DDTs and BIs have been deployed across all airports. Airports were included as
183 fixed effects, as they may differ both in their interception efficacy and in the underlying rates of

184 contamination on flights arriving at each location. Count predictor variables (i.e., passenger and
185 bag search counts) were square-root transformed and Z-scaled to reduce skewness, to improve
186 both model performance and the interpretability of effect estimates (per Schielzeth, 2010).

187 Models included two random effects, to assess the level of variance associated with the
188 flight's state of origin (*FlightOrigin*) and specific flight route (*FlightNumber*; nested within
189 origin). Flights without a number recorded were categorised as 'Itinerant/Other', with a large
190 majority considered to be private non-commercial arrivals, but also likely to include a small
191 percentage of commercial flights for which their numbers were not entered into the database.
192 For further details of model structure see Supplementary Materials B.

193 Unless otherwise stated, all values in square brackets below represent 95% confidence in-
194 tervals (or credibility intervals for Bayesian models below; '95CI') for the estimated effects.
195 The statistical significance of any fixed effects is inferred from whether their 95CIs include
196 zero. Random effects are assessed based on how much variance is explained in models, and
197 whether 95CIs for any specific random intercept predictions include zero. Where appropriate,
198 parameter/effect estimates below have been converted to percentage changes in the expected
199 number of interceptions for ease of interpretability. Marginal means were extracted from mod-
200 els using the package 'emmeans' (v1.8.7, Lenth, 2023), to estimate expected interception rates
201 under different intervention regimes.

202 **Statistical analysis B: Model sensitivity**

203 Sensitivity to overdispersion and zero-inflation was tested by re-fitting a subset of models in a
204 Bayesian framework via the package 'brms' (v2.19.0, Bürkner, 2017). From the six response
205 variables used in the main analysis, two were selected for sensitivity analyses. These were
206 the total BRM interceptions (*N_Total*, i.e., the most inclusive aggregation of interception data),
207 and the number of FF host declarations (*N_Declarations_FF*, i.e., the most sparse response
208 variable).

209 Four alternative distributions were tested for each response variable, namely Poisson (as in
210 the main analysis but implemented in a Bayesian framework), zero-inflated Poisson, negative
211 binomial, and zero-inflated negative binomial. These were chosen as common alternatives to

212 account for cases with excess zeros and overdispersion in ecology and other fields where count
213 data is common (Campbell, 2021; Lindén & Mäntyniemi, 2011; Pittman et al., 2022). Models
214 used the same fixed and random effects specifications as in the glm models, with default non-
215 informative priors to reflect our lack of prior knowledge for parameter estimates (chains = 3,
216 iterations = 3000, warmup = 1000). The outputs for the fixed effects of intervention regimes
217 and bag searches and the random intercepts associated with flight origins were estimated and
218 qualitatively compared between models.

219 Measures of model fit were also estimated for all models, i.e., Akaike/Watanabe–Akaike
220 information criterion ‘AIC’/‘WAIC’ as a measure of the quality of model fit for the dataset, and
221 marginal and conditional R^2 values as measures of the proportion of variance explained by fixed
222 effects and both fixed and random effects respectively (via package ‘performance’, v0.10.3,
223 Lüdtke et al., 2021; Nakagawa & Schielzeth, 2013). Overdispersion and zero-inflation tests
224 were also conducted (also via ‘performance’).

225 **3 Results**

226 **Intervention and pathway risk effects**

227 Models identified significant effects of biosecurity interventions upon interceptions of both
228 BRM and FF host items. Full results, code, models and outputs are available via Open Science
229 Framework (doi: [access via [review-only link](#)]), and detailed model outputs are available in the
230 Supplementary Materials B. Estimated BRM and FF host interception rates were significantly
231 higher when detector dogs were present. For example, the estimated total BRM interceptions
232 (N_{Total}) per flight with one DDT was 0.88 [0.80, 0.96], compared to 0.50 [0.46, 0.55] for one
233 BI. This effect appeared to primarily be driven by increases in the number of detections, with
234 DDTs having strong positive effects on BRM and FF host item detections (Fig. 2).

235 The number of bag searches conducted (performed both by DDTs and BIs) had a positive
236 effect on the estimated total, declared, and detected BRM counts. Unsurprisingly, the square-
237 root number of bag searches per arrival was associated with a per-unit increase in total BRM
238 interceptions of 24.0% [23.3%, 24.6%]. A similar effect was found for BRM declarations (i.e.,

239 23.8% [22.8%, 24.9%]) and detections (i.e., 24.1% [23.3%, 24.9%]). Similar effects were ob-
 240 served for FF host interceptions in total (23.8% [23.0%, 24.6%], declarations (23.2% [22.1%,
 241 24.4%]), and detections (24.3% [23.2%, 25.4%]). In both cases, the effect was similar for
 242 detections and declarations, suggesting that conducting more bag searches increases the rate
 243 of BRM being detected and encourages more declarations. As expected, increased passenger
 244 counts were also associated with increased interception rates across all response variables. Fi-
 245 nally, there were also some differences between arrival airports in their estimated interception
 246 rates (see Supplementary Materials B).

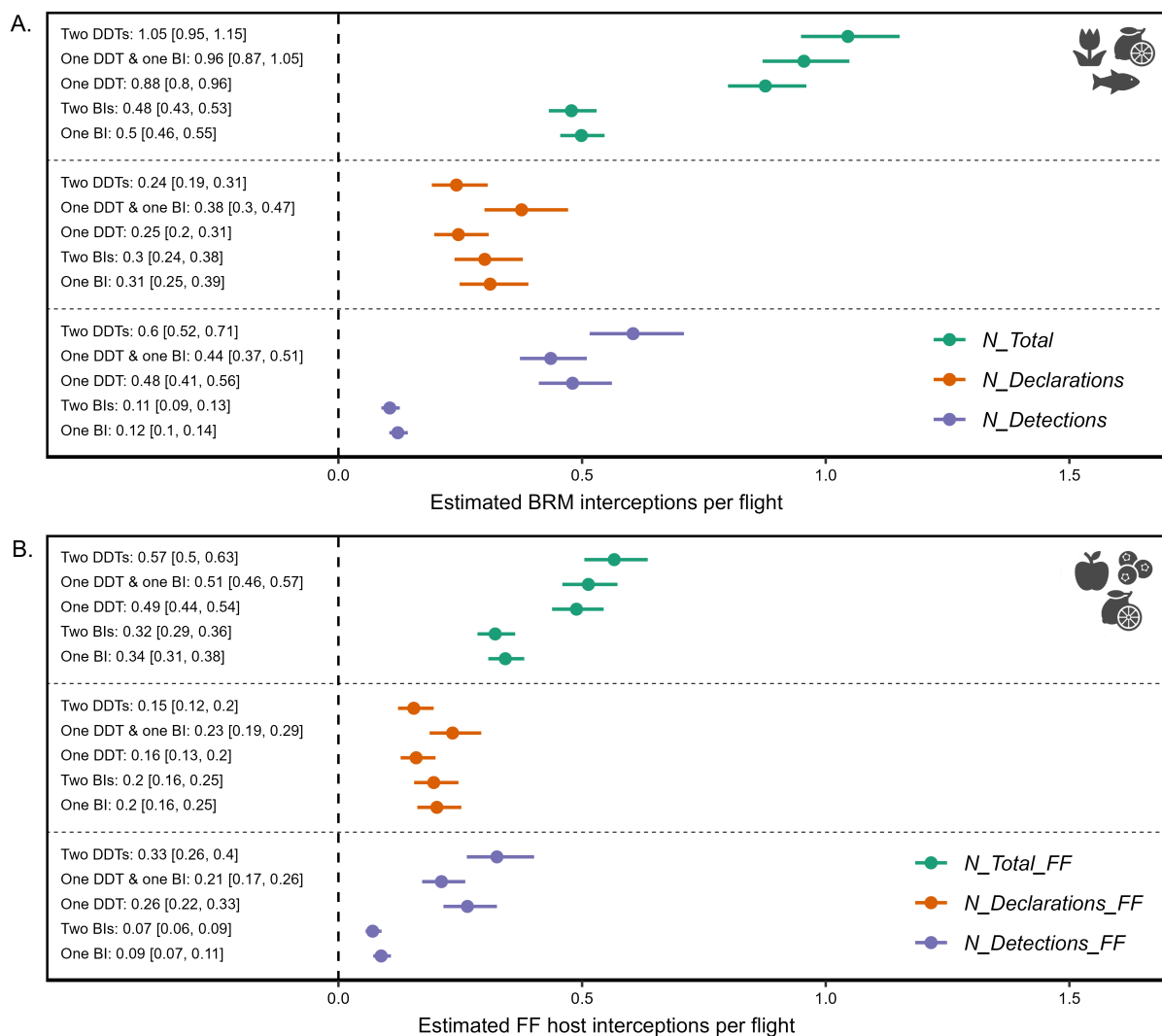


Figure 2: Estimated interception rates for air passenger interception regimes, for (A) BRM interceptions and (B) FF host items. Note, that estimated rates are the predicted number of interceptions per flight, and are independent of other factors included in the models (i.e., are estimated based on a flight with a mean number of passengers, and a mean number of bag searches, and averaged across arrival airports). Total, declared and detected rates come from distinct models, so estimates are not expected to be additive.

247 Random factors flight number and flight origin both explained some variance in interception
 248 rates (e.g., for N_Total , $V_{FlightOrigin} = 0.004$, $V_{FlightNumber} = 0.042$), although the random effects
 249 only explained a small proportion of variation relative to fixed effects (i.e., for N_Total , $R^2_{marginal}$
 250 $= 0.543$; $R^2_{conditional} = 0.570$, proportional $V_{FlightOrigin} = 0.002$, and proportional $V_{FlightNumber} =$
 251 0.025). It should also be noted that the overdispersion may lead to overestimates of R^2 values
 252 (e.g., Harrison, 2014), so these values should be interpreted cautiously. Nonetheless, random
 253 intercept predictions for flight origin show how pathway factors may be used to identify and
 254 target interceptions towards higher risk arrivals (Fig. 3). Similar pathway heterogeneity can
 255 also be identified for flight number (see Supplementary Figs. B.1– B.2).

256 Model sensitivity

257 Tests showed that models used in the main analysis for N_Total and $N_Declarations_FF$ both
 258 had probable zero-inflation (ratio of predicted to observed zeros: 0.85 and 0.92, respectively),
 259 and overdispersion was present in both cases (N_Total : dispersion ratio = 1.634, $\chi^2 = 97786.221$,
 260 $P < 0.001$; $N_Declarations_FF$: dispersion ratio = 1.709, $\chi^2 = 102303.640$, $P < 0.001$). No-

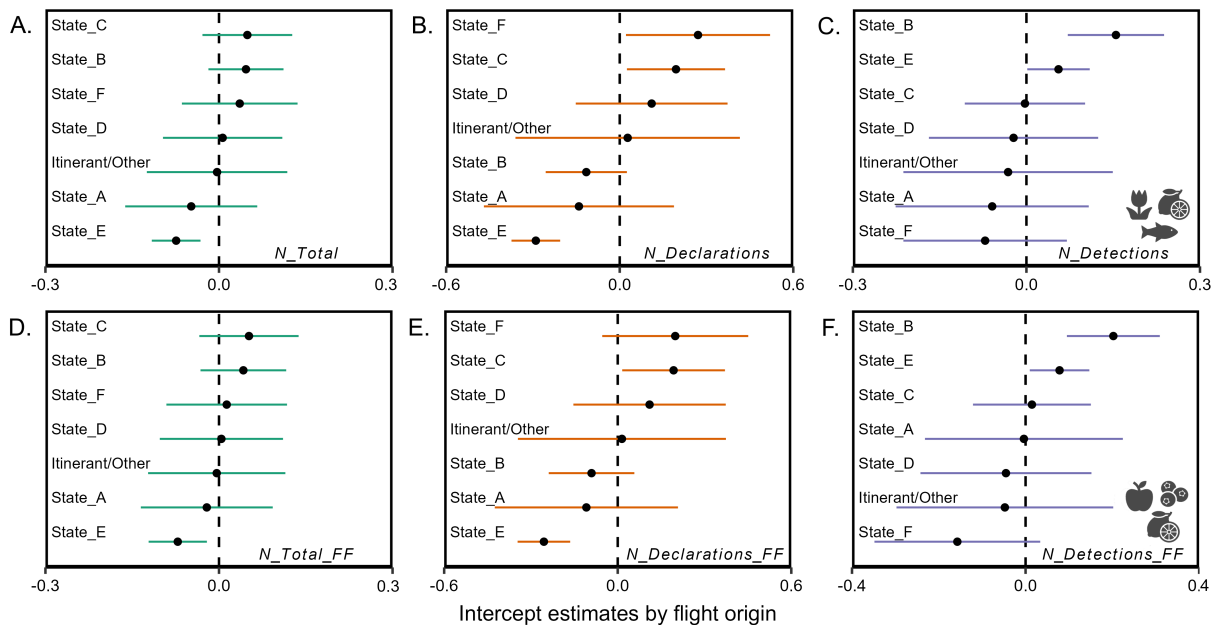


Figure 3: Predicted random intercepts by flight origin for (A-C) BRM interceptions, declarations, and detections; and (B) FF host interceptions, declarations, and detections. Error bars represent 95CIs, and intervals that do not include zero are considered to have significantly higher or lower levels of BRM interceptions than an average flight. Intercept estimates are in the modelled unit, i.e., the log of the proportional difference between the group and the overall expected BRM/FF host count.

261 tably, overdispersion can be a common consequence of zero-inflation, in which case a zero-
 262 inflated Poisson approach may be sufficient to account for both issues (see Yang et al., 2009).
 263 Measures of model fit also showed that all models accounting for zero-inflation had lower
 264 WAIC scores than those that did not, whereas negative binomial models had lower scores com-
 265 pared to Poisson models (see Supplementary Table B.1).

266 Nonetheless, sensitivity analysis showed that the outputs were relatively robust to imple-
 267 mentation methods, with the patterns identified qualitatively similar between implementation
 268 types, but with some variation in the magnitude and uncertainty of effects. Estimated intercep-
 269 tion rates under differing regimes showed similar patterns when using a Bayesian implementa-
 270 tion, although with slightly greater uncertainty (e.g., the estimated rate with one BI was 0.50
 271 [0.44, 0.57] compared to 0.50 [0.46, 0.55] in the main model; see Fig. 4). Incorporating zero-
 272 inflation into Poisson models led to higher rate estimates (see also Supplementary Fig. B.3).
 273 Whereas, negative binomial models produced lower estimates of BRM interceptions, while also

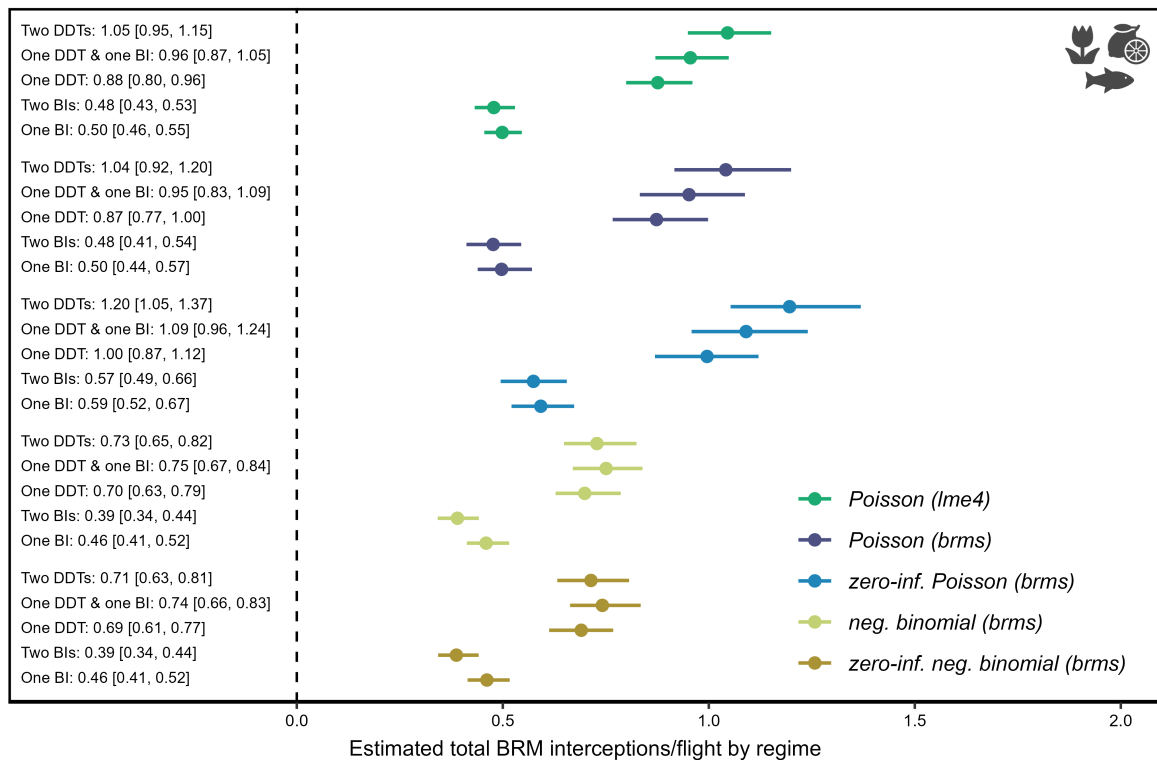


Figure 4: Estimated BRM interception rates for air passenger intervention regimes using different model implementations. Estimates are for five different model implementations, based on the modelled distribution and on a frequentist (lme4) vs Bayesian (brms) framework (see further details under Supplementary Table B.1).

274 showing similar differences between regimes.

275 Random intercept predictions also appeared to show qualitatively similar patterns for FF
276 host detections (Fig. 5) and BRM interceptions (Supplementary Fig. B.4). Bayesian approaches
277 produced greater uncertainty in the mean intercept estimates/predictions when directly com-
278 paring the Poisson implementation in lme4 and brms (Fig. 4., 5), although the means were
279 relatively consistent.

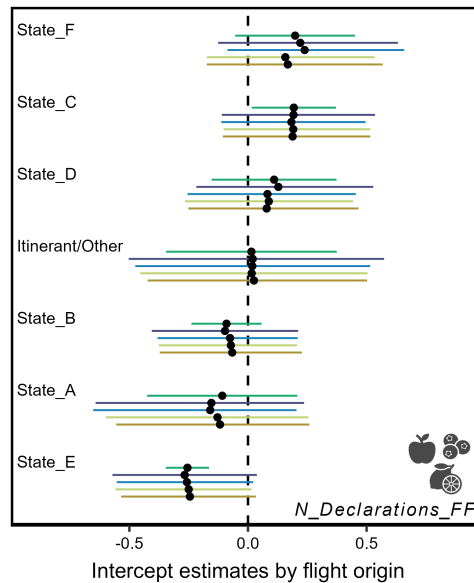


Figure 5: Predicted random intercepts for FF host item detections, by flight origin. Estimates are included from five different model implementations, which from the top include the following; (green) Poisson-lme4; (purple) Poisson-brms; (blue) zero-inflated Poisson-brms; (gold) negative binomial-brms; and, (orange) zero-inflated negative binomial-brms. Intercept estimates are in the modelled unit, i.e., the log of the proportional difference between the group and the overall expected FF host count.

280 4 Discussion

281 Biosecurity interceptions were strongly influenced by the methods used, for example, the num-
282 ber of bag searches conducted increased both detections and declarations from passengers. This
283 suggests that increased effort in active at-border surveillance by officers will increase the ef-
284 ficacy of interventions by promoting both voluntary compliance by passengers and detections
285 of undeclared risk items that may otherwise have been missed. Similarly, dog detector teams
286 are increasingly deployed for border interventions along high-volume phytosanitary risk path-
287 ways, both in Australia and many other countries (Whattam et al., 2014; Inspector-General of

288 Biosecurity, 2022). Few studies have quantitatively assessed the efficacy of detector dogs, al-
289 though a recent study from Williams & Sharp (2023) showed how the presence of a dog versus
290 an officer alone can alter passenger behaviour including eye contact, gestures or interactions
291 with the officer/dog. Our study provides further insights into the effects dogs may have on
292 passenger behaviour and the rate of interceptions at airports. The increase also appears to be
293 largely driven by detections of undeclared items instead of voluntary declarations, suggesting
294 that dogs may be particularly useful for capturing a component of the biosecurity risk material
295 that may otherwise not be found through more passive, voluntary compliance-based methods
296 (e.g., public awareness and education campaigns, biosecurity signage and announcements).

297 The ability to analyse pathway risk heterogeneity is limited by the type of data collected
298 on pathway risk factors. Despite the relatively limited set of pathway factors included in mod-
299 els and the relatively small proportion of total variance explained by these factors, models
300 were able to identify specific flight origins and routes as potentially high- and low-risk ar-
301 rivals. This shows how interception data may be useful for supporting risk-based approaches to
302 interventions (e.g., Australia's flight-based-traveller profiles; Inspector-General of Biosecurity,
303 2019) by identifying higher and lower risk arrivals to allocate limited resources to these arrivals
304 (Trouvé et al., 2024). These analyses can provide important quantitative evidence supporting
305 targeted resource allocations at the border, particularly when combined with further contex-
306 tual information such as pre-border pest prevalence data or estimates of potential post-border
307 impacts. While Tasmania currently targets 100% of air arrivals, risk-based approaches could in-
308 clude decisions about where and when to allocate their most effective methods (e.g., DDTs), or
309 to potentially identify a subset of low-risk arrivals that can be met with less resource-demanding
310 methods (e.g., passive interventions, signage, amnesty bins).

311 While this data is valuable in identifying how interventions or pathway risk influence the
312 actual interception rates, many unknowns remain that limit our ability to fully quantify the risk
313 of incursions along this pathway. For example, the risk of FF establishing through this pathway
314 would require us to estimate the actual volume of BRM on flights and the proportion of those
315 items infested with FF (i.e., contamination/infestation rates), the proportion of risk material
316 missed (i.e., leakage), or the viability of any FF individuals or larvae that may infest any of

317 the risk material. These parameters may be estimated using complementary methods, such as
318 endpoint surveys that target a subset of passengers as manual or X-ray searches to estimate
319 BRM contamination rates (Mannix et al., 2024). Samples of intercepted BRM may also be
320 further tested to measure their pest contamination/infestation rates. In many cases, particularly
321 in biosecurity, empirical data is lacking. Structured expert judgement may then be used to
322 elicit unknown parameters from relevant experts (e.g., biosecurity managers, entomologists,
323 academics, etc.), using advanced methods to directly elicit uncertainty in parameter estimates
324 and incorporate this uncertainty into the decision-making process (Hemming et al., 2018; Bau et
325 al., 2024). Therefore, although this study highlights the value of interception data for informing
326 biosecurity practices, additional knowledge is required to more completely assess and quantify
327 risk across a biosecurity continuum.

328 This study used a simple and common approach for count data (i.e., regression modelling
329 based on a Poisson distribution), performed with modelling tools that are accessible, and rela-
330 tively easy to implement. Sensitivity analysis suggested that the outputs of this approach were
331 qualitatively similar to approaches using more advanced tools (i.e., Bayesian modelling meth-
332 ods), or distributions (e.g., that account for zero-inflation). Although there were some notable
333 differences in outputs, for example, interception rates estimated from negative binomial mod-
334 els tended to be lower, suggesting that failing to account for over-dispersion may lead to slight
335 overestimates of predicted rates. Also, Bayesian methods tended to lead to higher uncertainty
336 estimates around fixed- and random-effect parameters, so may represent a more conservative
337 approach to modelling pathway risk factors. Therefore, while our conclusions were generally
338 robust to implementation methods, sensitivity analysis may also be a valuable step for provid-
339 ing additional information for decision-makers about the robustness of any conclusions drawn
340 from modelling. In this case, however, the operational interpretation of the simpler models was
341 borne out by the more complex models.

342 Finally, effective at-border interventions are a key step in the biosecurity continuum. As
343 at-border interventions become more sophisticated and widely implemented, large interception
344 data sets will inevitably become more available to researchers and biosecurity decision-makers.
345 This study highlights how this data can be a valuable resource for informing management

346 decisions for Tasmania and can provide empirical evidence to support the implementation of
347 risk-based approaches or the use of specific methods such as detector dogs, which can improve
348 resource allocations and lead to more effective interventions at borders.

349 **Data availability statement**

350 For privacy and operational purposes, all identifying information has been anonymised from
351 datasets, including airport names, flight numbers, etc. Fully anonymised datasets, analysis
352 code, models and outputs are all available at the Open Science Framework (doi: [access via
353 [review-only link](https://osf.io/78tv9/?view_only=f2128031528348599c165b644d2c776f), https://osf.io/78tv9/?view_only=f2128031528348599c165b644d2c776f]).

354 **Supplementary information**

355 Additional supporting materials include the following:

356 A – Composition of BRM and FF host interceptions

357 B – Supplementary model outputs

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369 **Competing interests**

370 The authors declare that they have no conflict of interest.

371 **Authors' contributions (CRediT taxonomy)**

372 NPM: Conceptualization, Data curation, Formal analysis, Investigation, Methodology, Soft-
373 ware, Resources, Validation, Visualization, Writing – original draft.

374 AMH: Conceptualization, Methodology, Supervision, Writing – review & editing.

375 APR: Conceptualization, Methodology, Funding acquisition, Supervision, Writing – review &
376 editing.

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