

1 Vegetation responses to managed river flow events 2 and regimes

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14 **Keywords**

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16 **Abstract**

17 Vegetation communities are an important part of riverine ecosystems and can be severely impacted by
18 changed flow conditions in regulated waterways, making them a priority for waterway management.
19 Environmental flows are a commonly used tool to mitigate some of the impacts from regulated flow
20 regimes and provide benefits to vegetation communities. Ongoing and effective use of environmental
21 flows requires evidence of vegetation responses to managed flow events and regimes, yet few studies
22 have demonstrated these responses in managed waterways.

23 We conducted vegetation surveys at 44 sites within seven major river systems across Victoria (Aus.)
24 between 2016 and 2023. Vegetation cover and richness were sampled at a range of bank elevations
25 impacted by different flows within each waterway. Surveys before and after flow events over multiple
26 years enabled us to evaluate vegetation responses to flow events and regimes. Generalised linear
27 mixed models were used to evaluate responses of richness and cover for four different plant
28 functional groups (aquatic, emergent, riparian and terrestrial) within delimited flow elevation zones on
29 the bank.

30 Our findings demonstrate that while individual managed and natural flow events may produce modest
31 immediate vegetation responses, their cumulative impact over time can be substantial, increasing
32 plant cover and richness that may support the resilience and stability of riverine systems. The
33 effectiveness of environmental flows is highly influenced by context and a range of flow and non-flow
34 variables. Short duration environmental flows with appropriate timing and duration benefit native
35 aquatic, emergent and riparian species. Terrestrial species can benefit from flows but are the most
36 sensitive to prolonged flow. Managers will also need to apply on-ground actions to enable ecological
37 outcomes from flows in some cases.

38

39 1. Introduction

40 Riverine vegetation communities across the globe are experiencing severe degradation due to factors
41 such as vegetation clearing, livestock grazing, pollution, proliferation of exotic plant species, and
42 alteration of natural flow regimes (Dufour et al. 2019; Greet et al. 2019; Tickner et al. 2020; Vörösmarty
43 et al. 2010). These pressures have led to significant declines in the integrity and functionality of
44 riverine ecosystems. Effective management and restoration of riverine vegetation communities is now
45 recognised as a critical priority for waterway managers worldwide (DEPI 2013; Dudgeon and
46 Arthington 2019; MDBA 2020; Tickner et al. 2020; VEW 2023). Vegetation is a priority not only
47 because of the intrinsic ecological value of these communities and species, and their social and
48 cultural values to humans, but also due to the essential ecosystem services they provide, including
49 erosion control, water quality improvement, flow regulation, habitat provision for terrestrial and
50 aquatic species, pollination support, and microclimate (e.g. light and temperature) stabilisation
51 (Capon and Pettit 2018; Riis et al. 2020).

52 River regulation has profoundly impacted the population dynamics of riverine plant species, affecting
53 processes such as longevity, recruitment and dispersal (Greet et al. 2011; Poff et al. 1997; Stromberg
54 et al. 2007). To counteract these impacts, environmental flows have become a widely used tool in
55 regulated waterways, involving the deliberate release of water from reservoirs to achieve specific
56 ecological outcomes (Arthington et al. 2023; Overton et al. 2014; Wineland et al. 2022). For riverine
57 vegetation, the benefits of individual environmental flow events can include increased native plant
58 growth, the triggering of recruitment events and increased seed dispersal (Greet et al. 2024; Greet et
59 al. 2013, Gomez-Sapiens et al. 2020). Large and long-term benefits are likely to be encountered when
60 the individual events are incorporated into regimes within an adaptive management framework that
61 provide cumulative benefits to the riparian community over time (Foster et al. 2018; Friedman et al.
62 2022; Arthington et al. 2023). Riverine vegetation communities that maintain higher native plant cover
63 and species diversity are more likely to withstand periodic disturbances such as drought or floods
64 (White and Stromberg 2011). Consequently, environmental flow delivery as part of the broader flow
65 regime is likely to enhance long-term resilience.

66 Given the growing competing demands on water resources, particularly in regions experiencing a
67 warming and drying climate, it is becoming increasingly important to evaluate responses to
68 environmental water to continually improve and adapt environmental flow management to maximise
69 its effectiveness (King et al. 2015; Arthington et al. 2023). However, evaluating the effects of
70 environmental flows on riverine vegetation is inherently complex, and it can be particularly difficult to
71 attribute ecological changes to environmental flow management (King et al. 2015; Arthington et al.
72 2023). Evaluation requires the assessment of isolated impacts of specific flow events (short-term
73 responses) as well as the impacts of successive flows (environmental or otherwise) within the broader
74 flow regime (medium- to long-term responses). Data collection needs to be repeated at relatively short
75 intervals to capture the effects of specific flow events, but continued over consecutive years to
76 capture variability in vegetation responses and cumulative impacts to sequences of flow events (King
77 et al 2015). Evaluation should also consider the relative importance of environmental flows in
78 comparison to other key factors that can affect vegetation, such as rainfall, livestock grazing, soil
79 properties, the presence of exotic plant species and historical disturbances (Lester et al. 2020; Nicol
80 et al. 2021; Campbell et al. 2023; Good and Jones 2024).

81 Environmental flows are used within many waterways of southeastern Australia to improve vegetation
82 communities that are degraded via regulation. Objectives for vegetation outcomes vary in detail

83 among waterways and regions but generally aim to improve riverine vegetation communities by
84 increasing propagule dispersal (Greet et al. 2024), stimulating plant germination and recruitment
85 opportunities (Tonkin et al. 2020; Pereira et al. 2021), increasing water resources for plant growth
86 (Tonkin et al. 2020; Deng et al. 2024), and reducing competition from terrestrial plants that are less-
87 tolerant of inundation (Greet et al. 2013; Miller et al. 2013; Vivian et al. 2020). Confirmation of these
88 individual mechanisms in small case studies, ex-situ experiments, and expert-elicited data (e.g. Greet
89 et al. 2023; Main et al. 2023, Nystrand et al. 2024) provides some support for existing management
90 strategies. However, these studies need to be complemented with field data on flow responses and
91 the influence of non-flow factors in both the short- and long-term (King et al. 2015; Arthington et al.
92 2023).

93 In this study, we aim to evaluate vegetation responses to environmental water management and to use
94 this information to inform water management practices for vegetation outcomes. The study aims to
95 address the lack of empirical data on vegetation responses to environmental flows, which are
96 currently used to achieve high-level management objectives. The data come from eight years of
97 monitoring vegetation responses to environmental flows across waterways in south-eastern Australia.
98 Standard vegetation attributes of cover and richness were used to characterise vegetation
99 communities. Our specific objectives were to: 1) describe detailed vegetation trends associated with
100 environmental flows across different seasons, plant position on banks (elevation), and plant
101 functional groups; 2) quantify vegetation responses to broader flow regimes and individual flow events
102 while accounting for factors such as plant origin (native/exotic) and livestock grazing presence; 3)
103 provide evidence-based recommendations for water management in regulated rivers to optimise
104 ecological outcomes; and 4) understand the data and analytical constraints and opportunities
105 provided by the dataset to inform future monitoring and evaluation. We hypothesized that native
106 vegetation cover and species richness would increase in response to environmental water
107 management, with increases being largest for plant groups with greater preference or tolerance of
108 inundation in the bank zones inundated by environmental flows (Wijepala et al. 2026).

109

110 2. Methods

111 2.1. Study area

112 Our study area encompassed seven major river systems in the state of Victoria, Australia (Figure 1).
113 The climate of the study area is varied, ranging from temperate in the south to semi-arid in the north
114 and west. Waterways in the north and west are part of the Murray-Darling Basin (MDB) while the
115 southern systems that flow to the south are part of coastal catchments that drain into Bass Strait.
116 Average annual rainfall ranges from 200-300 mm in the north-west to over 1,000 mm in the east
117 (Bureau of Meteorology 2023). Most environmental water is delivered in waterways located in the drier
118 northern catchments of the study area within the MDB. The seven river systems include most major,
119 lowland regulated systems in Victoria below storage reservoirs from which environmental water is
120 released.

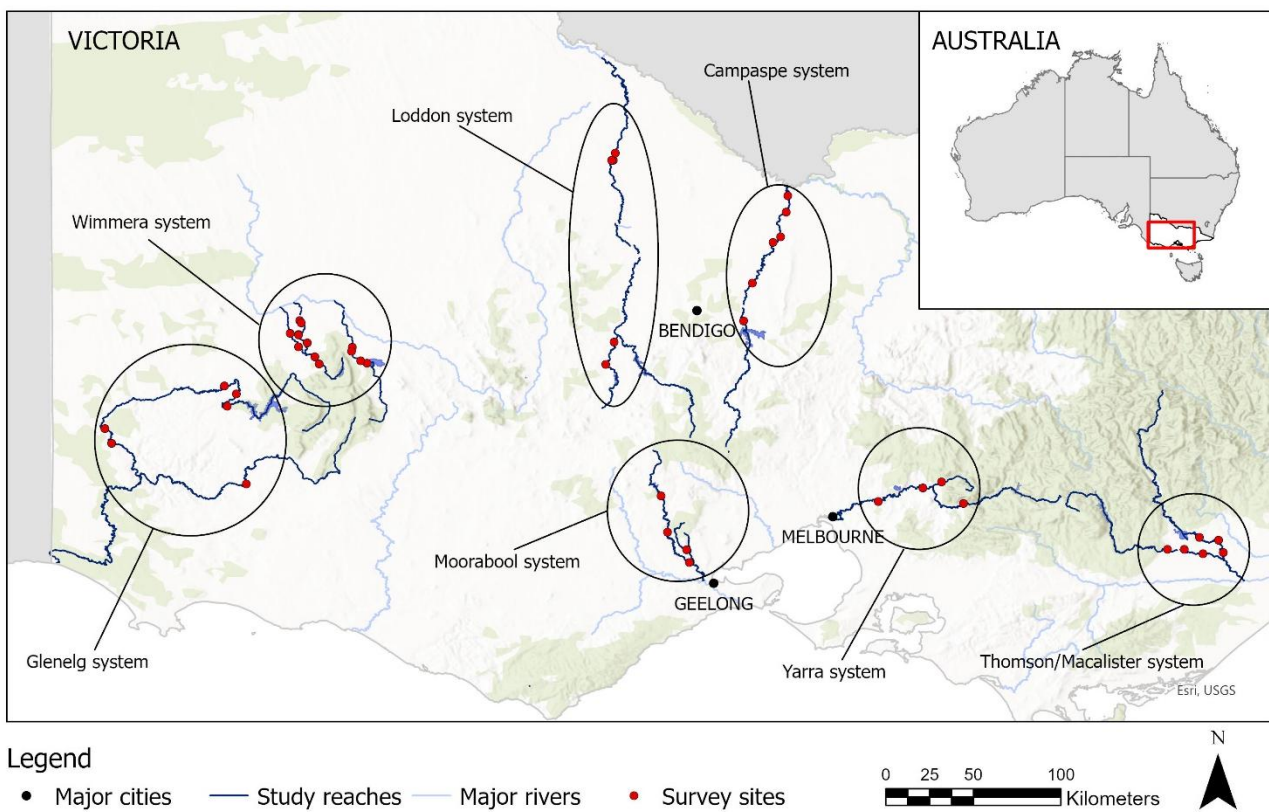
121 Forty-four sites were surveyed across 20 river reaches. Criteria for site inclusion included high priority
122 areas for waterway managers, site accessibility, and a spread of sites across different climatic regions
123 (Tonkin et al., 2020). All sites occur within landscapes largely cleared for agriculture or residential
124 areas but have a narrow strip of remnant native vegetation along the water frontage (e.g. generally 10-
125 50 m). At most sites, the natural vegetation type is floodplain riparian forest, dominated by *Eucalyptus*

126 *camaldulensis* subsp. *camaldulensis* (River Red Gum), one of the subspecies of Australia's most
127 widespread riparian tree.

128 Prior to river regulation, flow regimes across all seven surveyed river systems would have consisted of
129 high flows in winter and low flows in summer. However, contemporary summer flows are artificially
130 elevated due to regulated flow releases from reservoirs for irrigation, while winter flows are reduced
131 due to flow capture and storage. While many of the systems are naturally intermittent, the regulated
132 flow regimes are predominantly perennial except for several reaches in the Wimmera system that have
133 intermittent or ephemeral regulated flow regimes.

134 Surveys were conducted between 2016 and 2023. In late 2022, the wettest spring on record for
135 Victoria resulted in widespread flooding across all the surveyed river systems, with record flood peaks
136 experienced in several of the surveyed rivers (Bureau of Meteorology 2023).

137



138

139 **Figure 1. Study sites and associated river systems within the state of Victoria, Australia.**

140

141 2.2. Survey methods

142 2.2.1. Vegetation surveys

143 At each of the 44 sites, five to ten permanent transects were established perpendicular to the flow of
144 the river running from the low-flow water margin (bank toe) to a bank elevation position known or
145 believed to be above the highest point that managed flows reach. Due to different channel forms (e.g.
146 bank steepness) and stream orders, the lengths of transects varied within and between sites (range: 2–
147 21 m). On each transect, a series of 4 m long sub-transects were positioned every metre,
148 perpendicular to the transect (i.e. parallel to the waterway). Longer transects contained more sub-
149 transects and so the number of sub-transects varied within and between sites (Appendix A, Table A1).

150 For very long transects (>10 m), the sub-transects were spaced at 2 m intervals after the first 5 m.
151 Along each sub-transect, point-intercept samples were taken at 10 cm intervals (i.e. 40 per sub-
152 transect) to record both the substrate type (water, bare ground, log, rock, litter) and the species
153 identity of any plant that occupied the vertical point-intercept space, up to 5 m from the ground.

154 Livestock grazing was recorded as present (n=9) or absent (n=35) at each site. The identity of the
155 livestock (sheep or cattle) was recorded but no data on stocking densities or durations were available.
156 Grazing presence did not change substantially over time at any site.

157 2.2.2. *Survey timing*

158 Surveys were timed to occur before and after specific environmental flow deliveries (e.g. high flows or
159 freshening flow pulses ['freshes']) or natural flow events (e.g. floods) within each system. Specific
160 timing varied but was generally within 3 weeks before an event and approximately 8–12 weeks after an
161 event to allow time for recruitment and growth responses (DELWP 2017; Webb and Morris 2019). In
162 any single year, a site was typically surveyed three times: in early spring before a spring fresh or high
163 flow, in summer between a spring and summer fresh, and in autumn after one or more summer
164 freshes. Between 2016 and 2023, each site was surveyed between three and 16 times, totalling 263
165 site surveys (Appendix A, Table A1). Some sites had surveys undertaken in one year only (i.e. three
166 surveys timed around one spring and one summer fresh) while others had surveys undertaken across
167 multiple years. These different survey frequencies were due to funding variability between years.
168 Additionally, freshes or high flows were not always delivered in all years in all sites; for example, the
169 spring 2022 floods negated the needs for environmental flows in many waterways.

170 2.2.3. *Bank elevation*

171 The elevation of each sub-transect was determined using a high accuracy Trimble Geo7X with post-
172 processing of point aggregates. Elevation surveys were conducted during the first year of vegetation
173 data collection for each system. Bank geomorphology was considered relatively stable throughout the
174 majority of the study until the flooding events at the end of the survey period, which impacted some
175 transects. Horizontal and vertical accuracy was generally <10 cm. Elevation of water levels were also
176 recorded to validate and calibrate elevation data for vegetation and flow levels.

177 2.3. **Hydrological data**

178 Continuous stream flow elevation was used to develop hydrology metrics for individual flow events or
179 regimes. Flow elevations for each site were estimated by various methods depending on the available
180 data and available funding. Data sources used included:

- 181 1. Logger data: Data collected at half of the field sites from Troll loggers for short periods of time
182 (up to two years between 2017 and 2020). Loggers were levelled into Australian Height Datum
183 (AHD).
- 184 2. Water Measurement Information System (WMIS) data: Hydrographic data collected under
185 Victoria's Regional Water Monitoring Partnership (accessible via <http://data.water.vic.gov.au/>).
- 186 3. Water Data Online: Hydrographic data sourced from the Bureau of Meteorology (BOM)
187 (accessible via <http://www.bom.gov.au/waterdata/>).
- 188 4. Melbourne Water: Hydrographic data collected by Melbourne Water (accessible via
189 <https://www.melbournewater.com.au/water/rainfall-and-river-levels#/>).
- 190 5. Southern Rural Water (SRW): Hydrographic data.
- 191 6. AHD and Zero Gauge Height (ZGH) values from sources above or collected directly from the
192 site.

193 Data quality varied among sites due to differences in sources, but the highest quality data for all sites
194 were extracted from existing data sets and then reviewed and quality-checked by a hydrological
195 consultant. For each site surveyed, flow data were compiled, and an output of flow level (elevation in
196 AHD) was produced at regular intervals (interval time depending on data inputs but was at least once
197 daily).

198 **2.4. Aligning hydrological and bank elevation data**

199 High-accuracy Global Positioning System (GPS) points were required for bank elevation survey
200 locations and flow elevation in this study. The data sources differed, so calibration of datasets was
201 required to ensure compatibility. In all cases, the GPS data sources were calibrated to the levels of the
202 vegetation survey bank elevation locations so that flow data and vegetation data align. Not all bank
203 elevation locations were able to be matched to flow elevation data because of a lack of critical flow
204 information at some sites. Consequently, we were unable to evaluate flow hydrology treatments for
205 these sites.

206 **2.5. Data analysis and modelling**

207 *2.5.1. Species trends and patterns*

208 While most of the results have been presented for functional groups to aid generalisation across a
209 large study area containing hundreds of species with differing distributions, we initially present a
210 summary of important vegetation patterns at the species level to demonstrate the data underpinning
211 the models. We summarised patterns in species abundance in relation to bank elevation as well as
212 trends in abundance over time in relation to flow regimes using example species that are widely
213 distributed across the study area and are highly typical species of their respective functional groups.

214 *2.5.2. Response variables*

215 Native plant cover and species richness were used to characterise vegetation communities. These
216 variables are the most commonly used for describing plant responses and they align with the broad
217 objectives to sustain or increase plant communities. For the bulk of the analysis and modelling, plant
218 species were classified by functional group to enable pooling data across sites and generalisation of
219 outcomes. Our classification consisted of four groups which we hereafter refer to as functional
220 groups: aquatic (including submerged and floating species), emergent, riparian, and terrestrial, which
221 matches terminologies in waterway management frameworks used in southeast Australia (e.g. VEWH
222 2021) and other similar studies used to evaluate vegetation responses to environmental flows (e.g.
223 Tonkin et al. 2020; Jones et al. 2025). This plant grouping based on commonly used and clearly defined
224 attributes enables an assessment of how plant communities are responding to environmental flows as
225 specified in objectives. A full list of species and their classifications is provided in the data and code
226 repository, see Appendix A, Table A2 for a summary of the most common species within each group.

227 Several aggregations of cover were tested during a Pilot Study. Initially, we assessed individual species
228 cover. Projected foliage cover (percentage) was estimated for each species in each sub-transect as
229 the percentage of hits per 40 points. Subsequently, cover within groups was assessed by summing the
230 hits by species and estimating cover as a percentage of hits per 40 points. The summed cover allows
231 cover values within a group to exceed 100% where plants overlap each other (separated into native
232 and exotic species groups). Species richness within functional groups was estimated as the number of
233 unique species per functional group (separated into native and exotic species) recorded at a sub-
234 transect.

235 *2.5.3. Predictor variables*

236 Analyses included a set of predictor variables intended to characterise flow regimes, flow events, and
237 other factors influencing spatial and temporal variation in plant species cover or occurrence. Flow
238 events and bank zones are clearly defined in our study systems because managed flows are delivered
239 relatively consistently between years. We classified flow events based on their depth, duration, and
240 timing, but did not distinguish natural from managed events because they were expected to have
241 similar responses. The predictor variables used were:

- 242 • Flow regimes: characterised by the number of days per year (averaged over the preceding three
243 water years) where discharge exceeded baseflow or spring-fresh levels (z-scaled prior to
244 analysis to facilitate comparisons between sites), resulting in two regime metrics (days above
245 baseflow, and days above spring fresh per year).
- 246 • Flow events: the ‘sampling period’ defined by the survey timing in relation to an event (before a
247 spring fresh, between a spring and summer fresh, after a summer fresh) that enable a before-
248 after comparison for each event type.
- 249 • Zone: bank position in which a sub-transect occurred (below baseflow level, between
250 baseflow and spring-fresh level, above spring-fresh level).
- 251 • Functional group: categorising preferred inundation tolerances (aquatic, emergent, riparian,
252 terrestrial)
- 253 • Origin: differentiating indigenous from introduced species (native or exotic)
- 254 • Grazing: site grazing status by livestock (grazed or ungrazed)
- 255 • System: names for connected sets of waterbodies (e.g. tributaries) and waterways.

256 The identity of sites and transects (nested within sites), sub-transect height (metres above baseflow),
257 survey year, and system/waterbody (to group spatially connected sites) were included in analyses to
258 account for otherwise unexplained spatial and temporal variation in plant cover and species richness.
259 See Appendix B Table B1 for a complete list of model variables and their description, units, modelling
260 variable type and grouping.

261 2.5.4. *Adaptive preregistration and model development process*

262 The analysis structure and workflow were developed as part of a study of adaptive pre-registration in
263 ecology (Gould et al. 2025). Adaptive preregistration is an open-science approach to minimise
264 questionable research practices and ensure reproducibility and robustness of research findings, while
265 enabling flexibility given the highly variable nature of ecological data (Powers & Hampton 2018, Wood
266 et al. 2020) and the iterative nature of the model development process (Augusiak et al. 2014, Jakeman
267 et al. 2006). The workflow is described by a series of proposed analyses and decision points, with
268 explicit decision criteria to guide choices at each stage. We used the ecological modelling
269 preregistration template (Gould et al. 2024) to preregister the analysis and modelling for this study¹.
270 The study design is underpinned by a thorough conceptual model of ecological processes that are
271 used to justify the study purpose, design, target response and predictor variables, derivation of
272 variables, and constraints as a core element of the preregistration.

273 Adaptive Preregistration of model development and analysis occurred across three distinct stages
274 (Table B2, Figure B1), with two interim preregistrations (Version 1 and Version 2) preceding the final
275 preregistration (Version 3). Each preregistration version corresponded to distinct phases in the model
276 development process. Two candidate models were initially preregistered (Version 1, Table B2, Figure
277 B1), one for each vegetation response variable, however there was uncertainty around whether the
278 data could support the desired models due to low sample sizes relative to the complexity and spatio-

¹ The final version of the preregistration is available through GitHub (sign in required) and located at
https://github.com/egouldo/VEFMAP_VEG_Stage6/blob/master/analysis/preregistration_template.md

279 temporal patchiness of the data. Based on preregistered exploratory analysis (Version 1), we therefore
280 preregistered and conducted a pilot study (described below, but see also Table B2, Figure B1) to
281 resolve critical uncertainties in model specification and determine the feasibility of different model
282 structures. From the pilot study, we derived and preregistered a set of preferred candidate models for
283 fitting to the full dataset, as well as a candidate set of simplified models and decision-trees for
284 triggering their fitting and acceptance should the preferred model specifications fail to converge or
285 provide appropriate parameter estimates. Results from the pilot analysis informed subsequent
286 preregistration of the Main Analysis on the full dataset (Version 3).

287 The process shifted a lot of the thinking and trouble shooting from later parts of the analysis to prior to
288 the analysis, ensuring that all initial designs and processes were tied directly to the project objectives
289 and followed a logical and transparent workflow. Preregistration of the Pilot Study and then the final
290 analysis allowed flexibility within the design process that maintained transparency while enabling
291 clear decision points based on unknown outcomes of the pilot analysis – given the large dataset and
292 complex model structure.

293 2.5.5. *Pilot study*

294 The final data set comprised 42,356 non-zero observations for cover and 31,084 non-zero
295 observations for species richness, across approx. 12,000 replicate sub-transect surveys. In addition,
296 exploratory data analysis (preregistration version 1, Table B2) revealed considerable zero-inflation and
297 potential over-dispersion of the response variables, particularly vegetation cover. To address this, we
298 defined a maximal model structure and tested this structure with a pilot data set (all observations
299 from the Campaspe River from 2017–2020) to identify feasible model structures and likely decision
300 points. We developed separate analyses for vegetation cover and species richness, with the same
301 maximal model structure for both. The maximal model structure was (in high-level, R mixed-model
302 notation):

```
303 response (cover or species richness [multiple aggregations tested]) ~ cover in previous survey +  
304     functional group × origin × flow metrics +  
305     functional group × origin × zone × sampling period +  
306     grazing +  
307     (1 | site / transect) + (1 | sub-transect height) + (1 | survey year) + (1 | system) +  
308     offset(number of points).
```

309 Most variables are described above but see Appendix B for full descriptions. Vegetation cover in the
310 previous survey was included only for analyses of cover and meant that models of vegetation cover
311 examined how flow influenced the change in cover through time, rather than absolute vegetation cover
312 (akin to models of population growth rate used for the long-term modelling of fish populations in
313 Victorian rivers (Tonkin et al. 2024)). Flow regime was included with both linear and quadratic
314 associations. System was excluded from the pilot study given the use of data from a single waterway
315 (the Campaspe River). Results of the pilot study informed subsequent analyses.

316 2.5.6. *Main Analysis*

317 Based on the pilot study results, we preregistered the following modelling strategy (Version 3, Table B2,
318 Figure B2):

- 319 1. Refit the maximal models for each response variable as described above in 2.5.5.
- 320 2. Following pre-specified decision-trees, should maximal models fail to converge or provide
321 appropriate parameter estimates, fit model from a candidate set of simplified models
322 (preregistration pp.20-21).

- 323 3. Simplified models separately capture different flow components for each response variable.
324 For vegetation cover, two versions of the flow events model were specified (Flow event models
325 A & B, Figure B1).
326 4. A second flexible strategy for determining model functional form (Figure B2) and to account for
327 potential over-dispersion was preregistered based on the pilot study finding that the data
328 distributions in the full dataset did not match expectations informed by the pilot analysis.

329 Maximal models failed to converge on the full dataset, consequently we fitted the preregistered
330 candidate simplified models. Candidate simplified models of vegetation cover and species richness
331 focused on functional groups as opposed to individual species because these supported model
332 structures closest to the maximal structure. We systematically re-assessed interactions in the
333 maximal models within the candidate simplified models based on the expectations that additional
334 data would either support more-complex model structures, or else introduce additional heterogeneity
335 (and, therefore, support less-complex model structures) than in the pilot study. This unregistered
336 analysis did not include re-assessment of the functional group classifications. The complexity of the
337 fitted model and data precluded interactions between species origin or grazing and functional groups
338 or zone.

339 Posterior predictive checks indicated that a zero-inflated Poisson distribution reliably captured the
340 distribution of observed vegetation cover within functional groups and a Poisson distribution (without
341 zero-inflation) reliably captured observed distributions of species richness (results not shown, see
342 Appendix C for checking of full models). Models of cover had excess over-dispersion, but a zero-
343 inflated negative binomial distribution did not substantially reduce unmodelled over-dispersion and
344 resulted in worse characterisation of zero-inflation and the observed distribution of cover data (results
345 not shown). Therefore, subsequent models of vegetation cover and species richness assumed that
346 data were derived from a Poisson distribution, with additional parameters for zero-inflation specific to
347 each functional group in the case of vegetation cover.

348 2.5.7. *Analysis of vegetation cover*

349 The full data set supported a slightly more complex model than identified in the pilot study, with this
350 model including the three-way interaction between zone, sampling period, and functional group.
351 However, separate models were still required for flow metrics and categorical flow terms (zone and
352 sampling period). We defined the model with flow metrics as the flow-regime model, which examines
353 how realised flow conditions over the preceding three years influence vegetation cover within each
354 functional group and had the following structure:

```
355 functional group cover ~ cover in previous survey +  
356     functional group × (days above baseflow + days above spring fresh) +  
357     functional group × (days above baseflow2 + days above spring fresh2) +  
358     origin +  
359     (1 | site / transect) + (1 | sub-transect height) + (1 | survey year) + (1 | system) +  
360     offset(number of points)
```

361 Similarly, we defined the model with categorical flow terms as the flow-event model, which examines
362 how vegetation cover in each zone changes before and after key flow events (spring and summer
363 freshes) and had the following structure:

```
364 Functional group cover ~ cover in previous survey +  
365     functional group × zone × sampling period +  
366     origin + grazing +  
367     (1 | site / transect) + (1 | sub-transect height) + (1 | survey year) + (1 | system) +
```

368 offset(number of points)

369 2.5.8. *Analysis of vegetation species richness*

370 As for cover, the full data set supported a more complex model than the pilot study, with all predictors
371 included in a single model and a three-way interaction between zone, sampling period, and functional
372 group. Quadratic associations with flow metrics were unable to be fitted at the functional-group level
373 and were instead included as a single term for all functional groups. The final model structure was:

374 Functional group species richness ~ functional group × (days above baseflow + days above spring fresh)+
375 days above baseflow² + days above spring fresh² +
376 functional group × zone × sampling period +
377 origin + grazing +
378 (1 | site / transect) + (1 | sub-transect height) + (1 | survey year) + (1 | waterbody) +
379 offset(number of points)

380 2.5.9. *Computational details and model performance*

381 All models were fitted using the glmmTMB R package version 1.1.8 (Brooks et al. 2017) in R 4.3.1 (R
382 Core Team 2023). Comparisons were made between model outputs from the Pilot study (data from the
383 one system with most data) and the full model across all systems. Model performance was evaluated
384 using goodness of fit tests, indicated by Pearson's r^2 and posterior predictive checks. Thresholds for
385 interpreting the model fit statistics were defined a priori in the preregistration of the study and are
386 summarised as: < 0.25 is poor, 0.25 - 0.5 is moderate, 0.5 - 0.75 is good, 0.75 - 1.0 is excellent, but
387 probably indicates overfitting. These thresholds are broadly consistent with many ecological studies,
388 e.g., Thomas et al. (2019) and Hale et al. (2023), that indicate values over ~0.5 provide adequate to
389 strong support. Posterior predictive checks were implemented with the performance R package
390 version 0.10.8 (Lüdecke et al. 2021) and modelled associations were extracted with the effects R
391 package version 4.2-2 (Fox 2003). The full set of packages used in the analysis for this study is
392 described in Appendix D Model code, including version history and associated pre-registration
393 documents is recorded on a private GitHub repository and will be made available alongside manuscript
394 submission.

395 3. Results

396 3.1. **Model evaluation**

397 Models for vegetation cover and species richness had moderate goodness-of-fit, with r^2 values (based
398 on Pearson's r) of 0.54 for the flow-regime model of vegetation cover, 0.55 for the flow-event model of
399 vegetation cover, and 0.32 for the model of vegetation species richness. Posterior predictive checks
400 indicated that models were sufficiently capturing the distribution of observed data, including the high
401 proportion of zero values in cover data, although with some evidence of mismatch for low and
402 intermediate (but not zero) values of cover (Figures C1–C3 in Appendix C). Alternative model
403 distributions did not improve model fit (results not shown), indicating high levels of heterogeneity in
404 vegetation cover data.

405 3.2. **General vegetation patterns**

406 Species cover and richness varied among waterways and sites. Waterways with more sites generally
407 had greater species richness totals (Appendix E, Table E1) due to species-area relationships. Overall,
408 aquatic, emergent and riparian vegetation groups were dominated by native species, while the
409 terrestrial group was dominated by exotic species (Appendix E, Tables E1 and E2). Waterways in higher
410 rainfall regions (e.g. Macalister, Thomson, Yarra Rivers) had a higher prevalence of exotic species

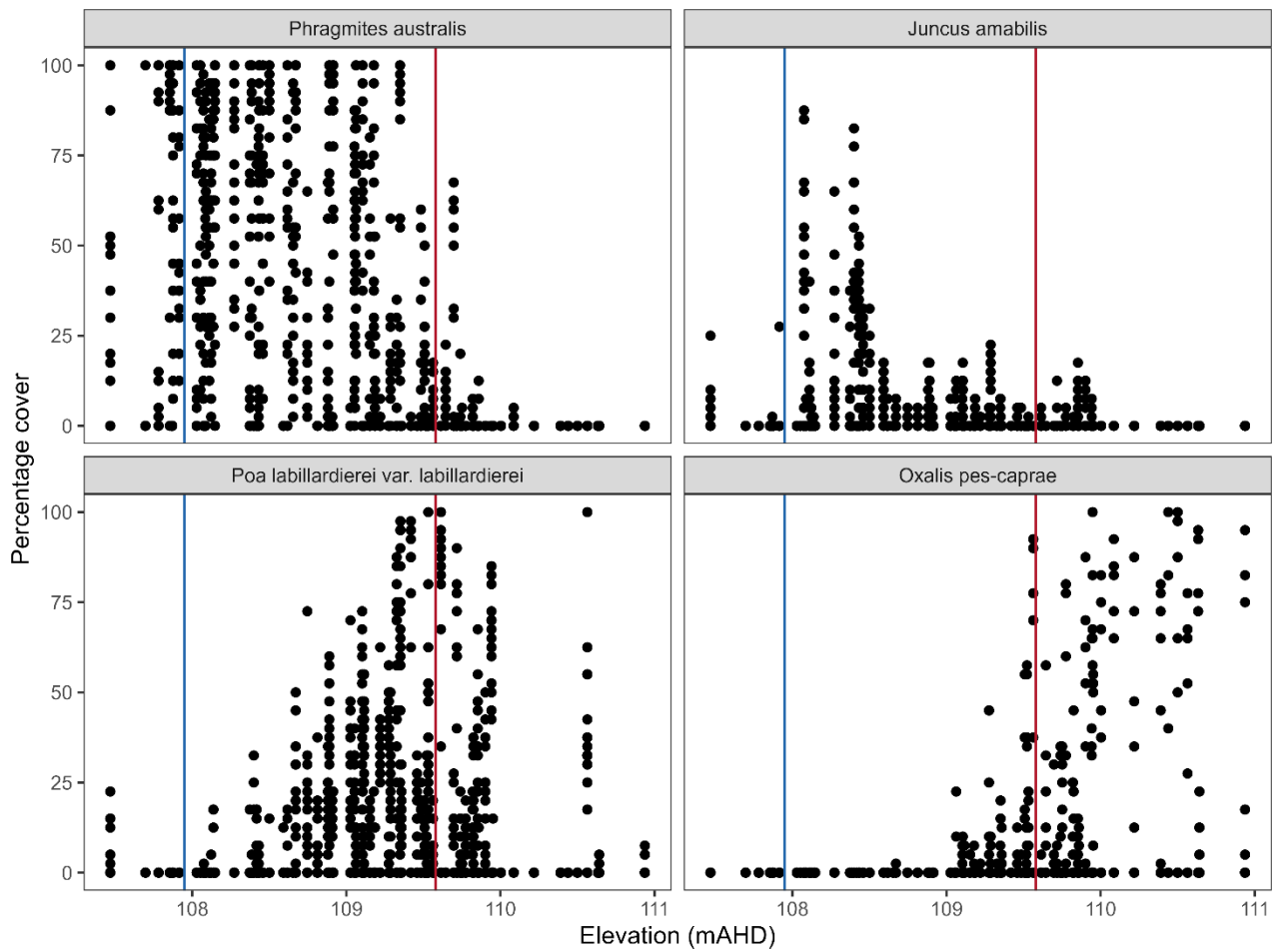
411 compared to natives, and lower abundance of aquatic and emergent species than waterways in lower
412 rainfall regions (e.g. Loddon, Wimmera, Campaspe Rivers). The correlation with high rainfall could be a
413 result of the increased water resources or factors at the system level, including plant traits, levels of
414 disturbance, and neighbouring land use.

415 At the individual site and species level, there were clear patterns of distribution from low to high bank
416 elevation that correspond with flow levels and events (Figure 2). The individual species patterns are
417 informative for investigating trends and patterns that are reflected by the broader functional groups in
418 the models. For example, at a site on the Campaspe River, the emergent *Phragmites australis*
419 (Common Reed) was common from the baseflow to spring fresh level, the riparian *Juncus amabilis*
420 (Hollow Rush) was common above the baseflow margin, the damp-tolerant terrestrial grass *Poa*
421 *labillardierei* (Common Tussock-grass) was distributed around the spring fresh peak level, and the
422 flood-intolerant exotic terrestrial herb *Oxalis pes-caprae* (Soursob) was largely restricted to elevations
423 above the spring fresh. The dominant species varied among waterways (Appendix A, Table A2).

424 Trends over time for the three most abundant functional groups (emergent, riparian and terrestrial)
425 show broadly increasing vegetation cover over the survey period for all three groups, within spring
426 fresh zone targeted by elevated environmental flows (Figure 3). The increasing trends in this zone are
427 indicative of a cumulative response to repeated management over consecutive years. In contrast, there
428 was no apparent trend for any group above the spring fresh level or for riparian and terrestrial groups
429 below the baseflow level. Emergent plants that more commonly occur on the river bed showed an
430 increase in cover over the survey period. Terrestrial and riparian species were generally heavily
431 impacted by the natural floods in late 2022, resulting in dramatic declines in 2023 (survey 16, Figure 3).
432 However, the emergent species were only marginally impacted by the floods. The terrestrial group
433 contained a large proportion of annual species and geophytes, and therefore showed strong seasonal
434 abundance fluctuations (Figure 3).

435 Additionally, there were clear negative relationships between native and exotic species, where one
436 was often able to outcompete the other (Appendix E, Figure E1). Most sites in the study area had
437 terrestrial vegetation communities dominated by exotic species (e.g. invasive herbs and agricultural
438 grasses), and the prevalence of exotic riparian species differed among sites and rainfall regions. This
439 means that responses by native riparian species may be constrained by the increases in exotic
440 species that exclude them.

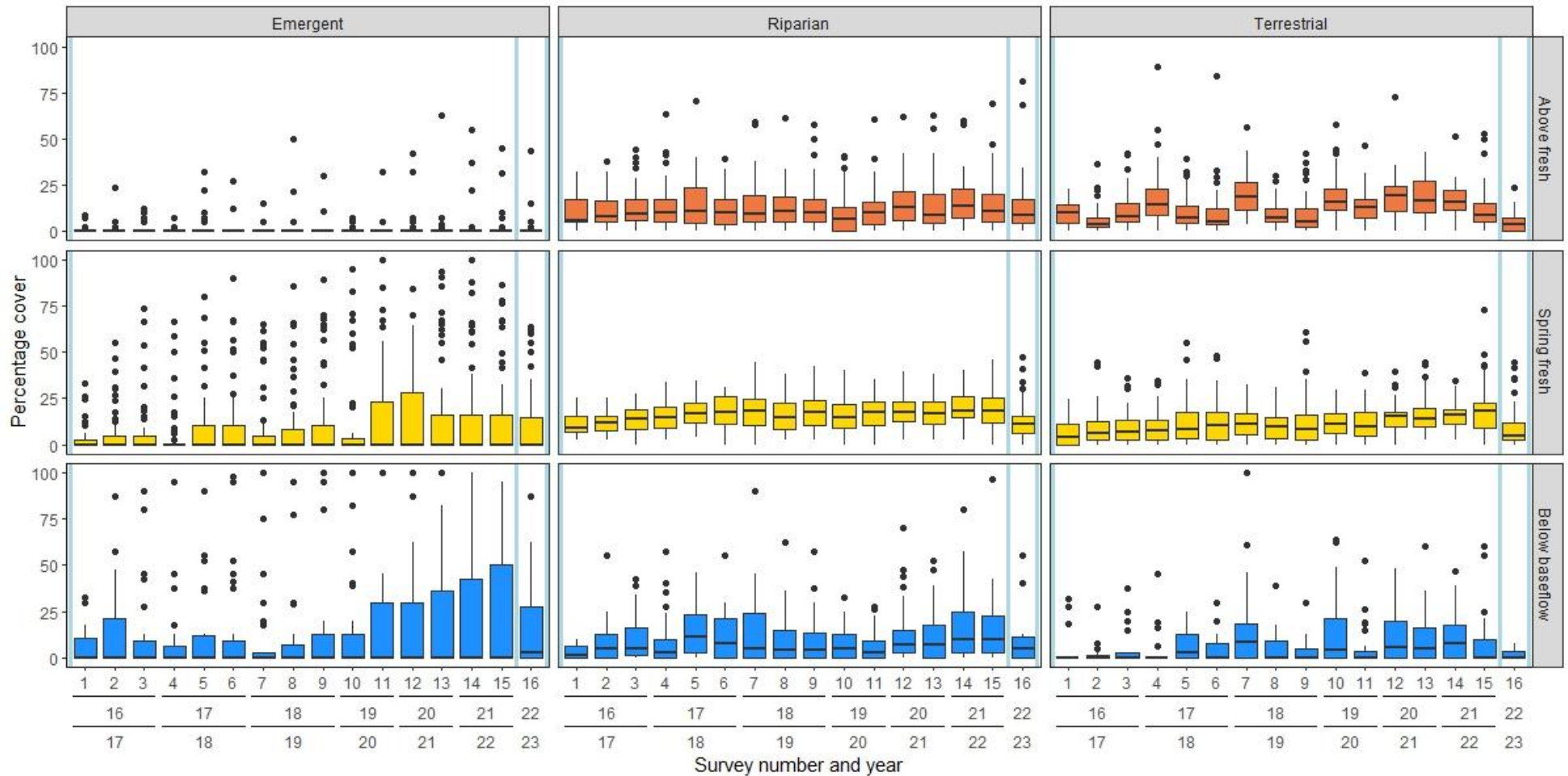
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443 **Figure 2. Four common plant species distributions in relation to elevation and flow events at a single**
 444 **site (Spencer) on the Campaspe River. Blue lines indicate the baseflow peak level, and red lines**
 445 **indicate spring fresh peak levels.**

446



447

448 **Figure 3. The abundance (as percentage cover) of all plants in three functional groups (emergent, riparian and terrestrial) within the three bank zones**
 449 **recorded during 16 surveys over the study period at all sites on the Campaspe River. Vertical lines indicate major floods. Boxplots show the median and**
 450 **the range between the first and third quartiles of the data, the whiskers extend up to 1.5 x the inter-quartile range.**

451 **3.3. Flow regime influences on plant cover and richness**

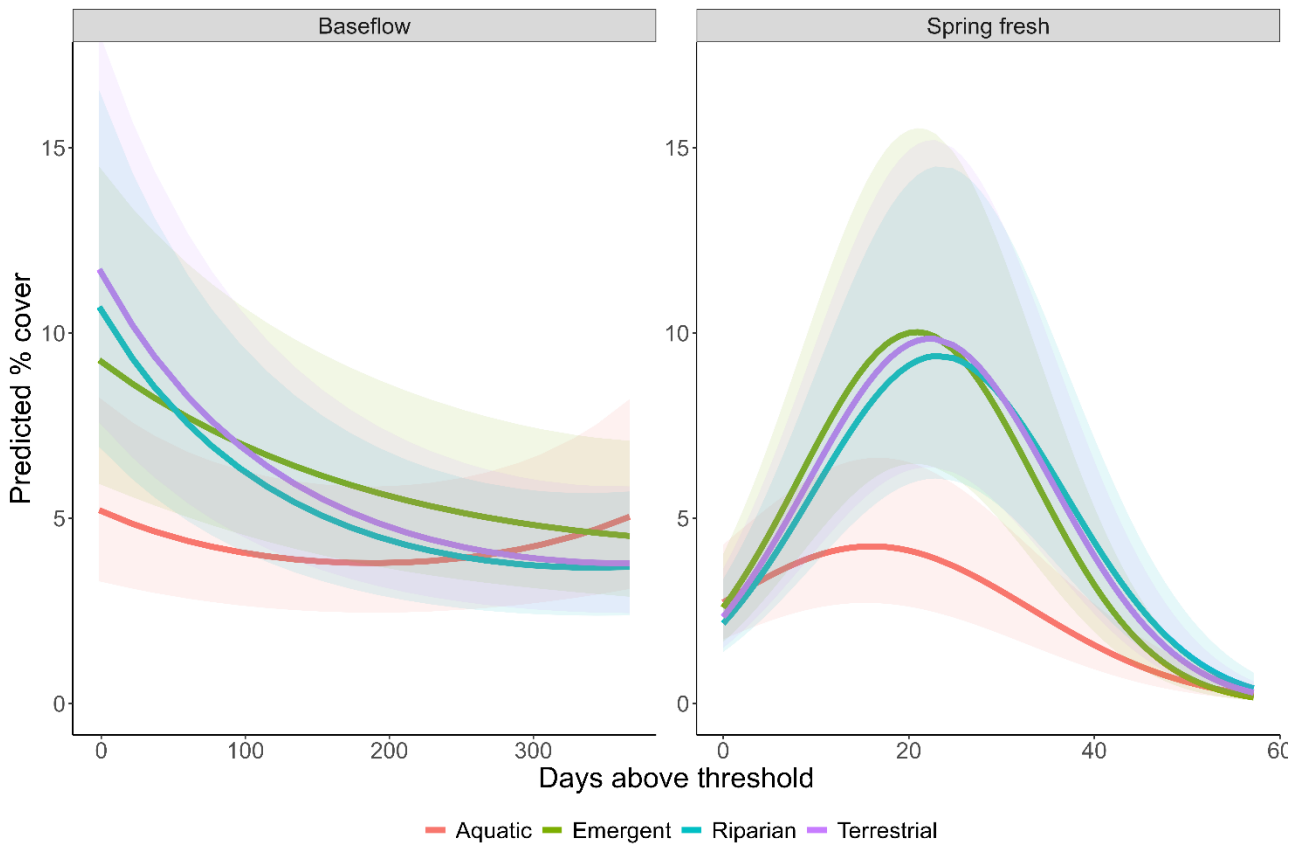
452 Associations of plant cover with flow regimes (days above baseflow or spring-fresh levels over the prior
453 three years) generally reflected negative effects of sustained baseflows, positive effects of short
454 freshes (15–25 days), and negative effects of sustained, elevated flows (> 25 days per year above
455 spring-fresh level) (Figure 4). These results implicitly consider the cumulative influence of flows and
456 rainfall in prior years as opposed to event-based responses. Aquatic species differed most from the
457 other functional groups, with evidence of positive associations with sustained baseflows (Figure 4a)
458 and weaker associations with sustained, elevated flows (Figure 4b). The elevated cover of aquatic
459 plants with few days above baseflow was a common outcome in sites where persistent shallow water
460 provided ideal conditions for aquatic plants. Few sites in this study have long cease to flow periods, so
461 the low flow systems typically had high cover of aquatic plants.

462 Species richness of functional groups showed peaked associations between species richness and the
463 number of days above spring-fresh level, particularly for riparian and terrestrial species, indicating
464 positive effects of elevated flows (above spring-fresh levels) for 10–30 days per year but negative
465 effects when elevated flows were sustained for >30 days per year (Figure 5). Terrestrial species
466 richness peaked at the lowest number of elevated flow days (~25), followed by riparian plants (~34)
467 and then emergent (~38). Richness flow regime models identified negative associations between the
468 number of riparian and terrestrial species and the number of days with discharge above baseflow
469 level, with no clear evidence of associations between days above baseflow and the number of aquatic
470 or emergent species (Figure 5).

471 Conditional on associations with flow regimes, variation in vegetation cover occurred primarily among
472 sites, with less variation among systems, transects, sub-transect heights, and years (Appendix F,
473 Figure F1). Residual variation among systems (i.e., variation at system level, but not explained by flow
474 regimes) indicated generally higher-than-expected vegetation cover in the Moorabool and West
475 Gippsland systems and average or slightly-below-average cover elsewhere (Appendix F, Figure F2).
476 These two systems are both southern systems, but the Moorabool has relatively low local rainfall and
477 flow compared to high rainfall and flow in West Gippsland. Cover of vegetation functional groups did
478 not differ considerably between native or exotic species (Appendix F, Figure F3).

479 Variation in vegetation species richness for regime and event models was similar among systems,
480 sites, transects, and sub-transect heights, with less variation among years (Appendix F, Figure F8).
481 Residual variation among systems (i.e., variation at system level, but not explained by flow regimes)
482 indicated generally higher-than-expected species richness in the Moorabool and (to a lesser extent)
483 Yarra systems, average species richness in the Campaspe, Glenelg, and Wimmera systems, and
484 lower-than-expected species richness in the Loddon and West Gippsland systems (Appendix F, Figure
485 F9). Again, these system patterns do not follow a clear pattern of rainfall or climate, and the patterns of
486 low or high richness can be driven by multiple factors such as high productivity leading to high diversity
487 or competitive exclusion; with low productivity leading to lack of resources or reduced competition.

488

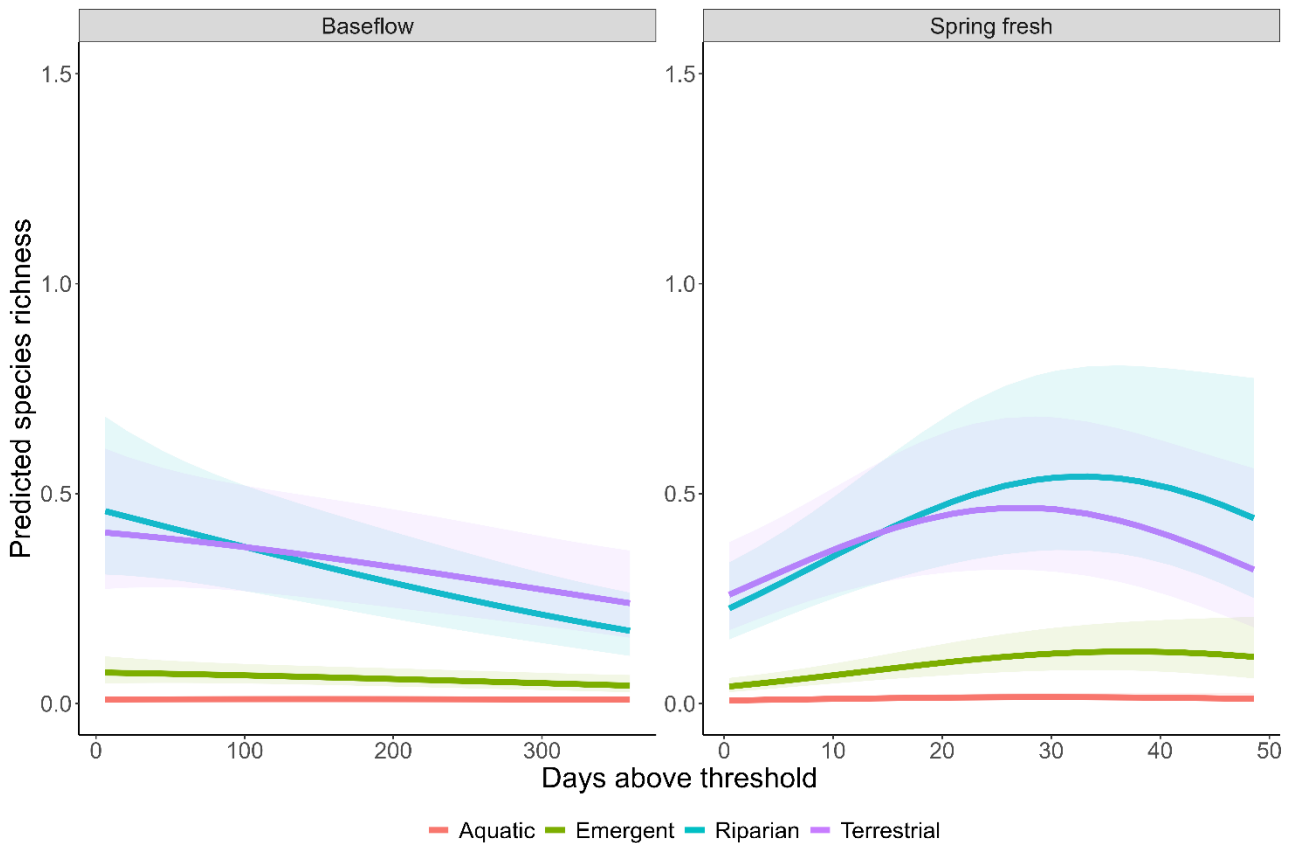


489

490 **Figure 4. Predicted associations between vegetation cover and the number of days with discharge**
 491 **above (a) baseflow levels and (b) spring fresh levels, and in the three years preceding a survey.**

492 Solid lines are mean estimates of cover for each functional group and shaded regions bound 95% confidence intervals.
 493 Note that the y-axis is truncated to a maximum value of 15%, which reflects the relatively low values of average cover
 494 given zero-inflation in the dataset.

495



496

497 **Figure 5. Predicted associations between vegetation species richness and the number of days with**
 498 **discharge above spring-fresh and baseflow levels in the three years preceding a survey.**

499 Solid lines are mean estimates of species richness for each functional group and shaded regions bound 95% confidence
 500 intervals. Note that the y-axis is truncated to a maximum value of 1.00 species, which reflects the relatively low values of
 501 average species richness at the sub-transect level on average over all bank elevations.

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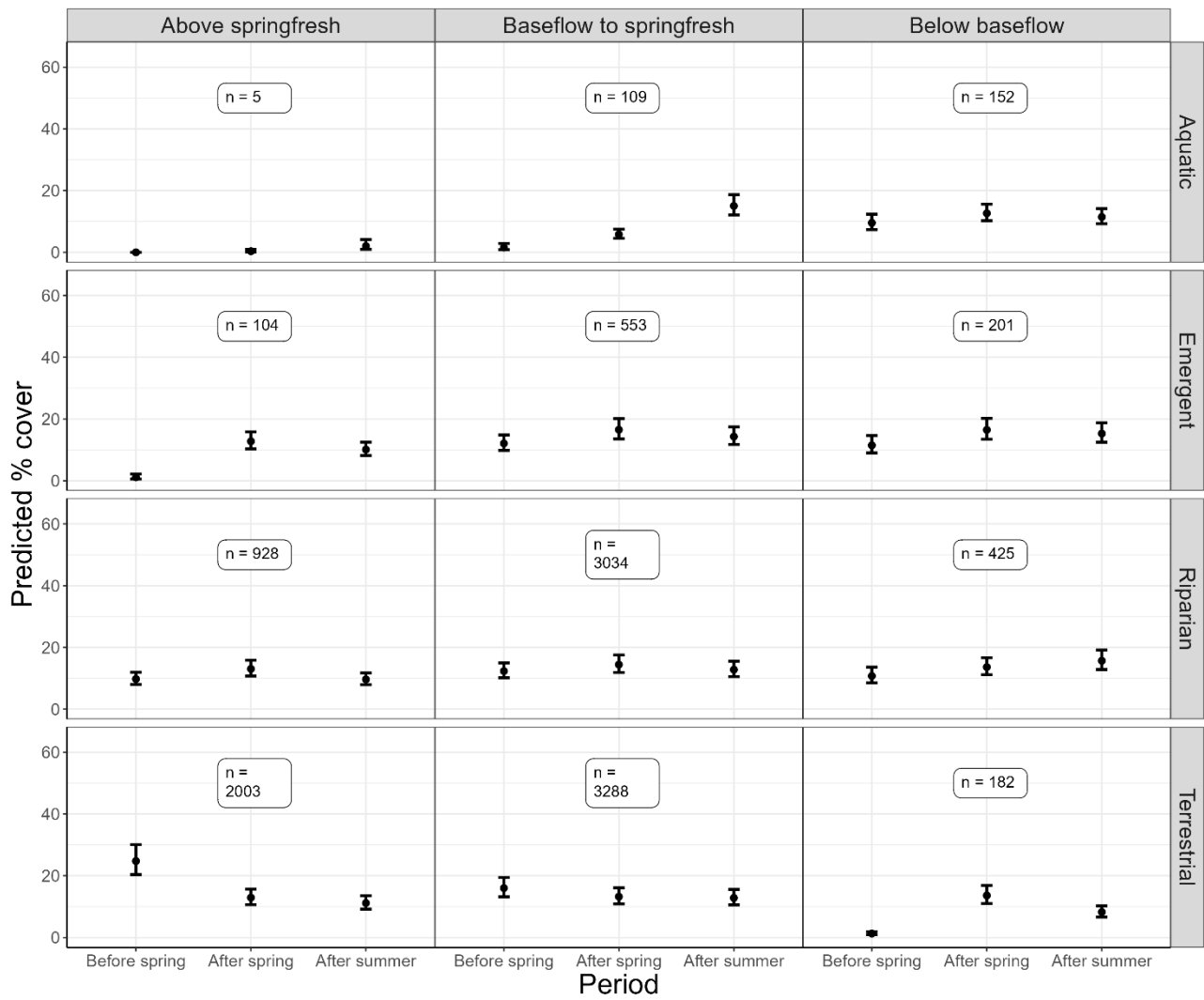
504 **3.4. Flow event influences on plant cover and richness**

505 Associations of plant cover with flow events (zone and sampling period) suggested that vegetation
 506 responses to flow events (spring and summer freshes) differed among zones (bank location relative to
 507 baseflow and spring-fresh levels) and functional groups (Figure 6). The summed cover of aquatic
 508 species above the spring-fresh level or below the baseflow level did not change markedly following
 509 spring or summer freshes because of the very low abundance at that high bank elevation (Figure 6). By
 510 contrast, there was a clear, near-linear increase in cover of aquatic species between the baseflow and
 511 spring-fresh levels following both spring and summer fresh events which suggests a positive response
 512 for these plants in the zone targeted by the freshes (Figure 6). The summed cover of emergent species
 513 increased following spring freshes, with increases apparent in all zones but clearest above the spring-
 514 fresh level where cover was lowest at the start of the growth season (Figure 6). Cover of emergent
 515 species decreased slightly following summer freshes in all zones due to seasonal senescence but
 516 remained elevated above their pre-spring fresh level (Figure 6). Riparian species had relatively weak
 517 associations with flow events in all zones. Terrestrial species had clear, but opposite associations
 518 above the spring-fresh level and below the baseflow level (Figure 6). Over the spring fresh period, there
 519 was a decline in cover above the fresh level and an increase below the baseflow level, but this was
 520 largely driven by the different pre-fresh cover levels. Before the spring fresh, terrestrial plants have high
 521 abundance at high bank elevations following autumn recruitment from rainfall, but they have low cover

522 below baseflow level when the winter baseflows are higher and prevent terrestrial recruitment.
523 Summer freshes were associated with minor declines in terrestrial cover in inundated zones.

524 Models of flow events (spring and summer freshes) highlighted relatively low intra-annual variation in
525 vegetation species richness, with slight increases in the number of riparian species following spring
526 freshes in all zones and slight decreases in the number of terrestrial species above baseflow and
527 spring-fresh levels following spring and summer freshes (Figure 7). Riparian and terrestrial plants
528 showed clearly divergent preferences for wetter and drier bank positions respectively. Aquatic
529 vegetation richness was low and largely restricted to below the baseflow levels, while emergent
530 species followed a similar but more slightly more abundant pattern.

531 Conditional on associations with flow events, variation in vegetation cover occurred primarily among
532 sites, transects, and sub-transect heights, with slightly less variation among systems and much less
533 variation among years (Appendix F, Figure F4). High levels of variation among transects and sub-
534 transect heights were not observed in the flow-regime model, which indicates that flow regimes may
535 partially explain variation in vegetation cover among transects and sub-transect heights (compare
536 Appendix F, Figures F1 and F4). Residual variation among systems (i.e., variation at system level but
537 not explained by flow events) indicated generally higher-than-expected vegetation cover in the
538 Moorabool, West Gippsland, and Yarra systems, average cover in the Loddon and Wimmera systems,
539 and lower-than-expected cover in the Campaspe and Glenelg systems (Appendix F, Figure F5). These
540 patterns do not follow a clear rainfall pattern, neither was there a consistent trend of livestock grazing
541 influence. Cover of vegetation functional groups differed little between native and exotic species
542 (Appendix F, Figure F6) and was negatively associated with the occurrence of grazing (Appendix F,
543 Figure F7).

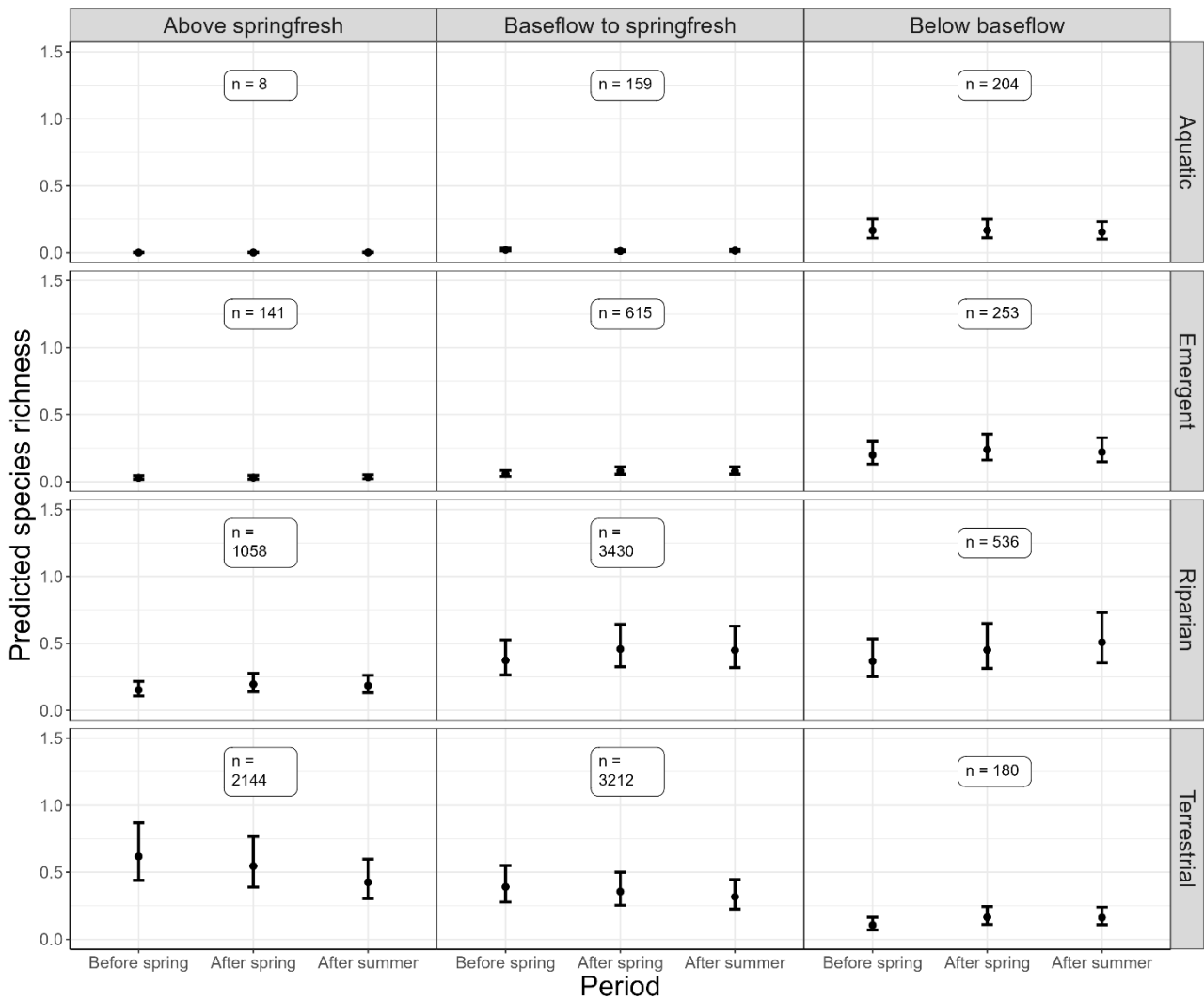


544

545 **Figure 6. Predicted associations between vegetation cover and flow events (spring and summer**
 546 **freshes) in each zone and sampling period.**

547 Solid points are mean estimates of cover for each functional group and error bars bound 95% confidence intervals. Note
 548 that the y-axis is truncated to a maximum value of 60%, which reflects the relatively low values of average cover given
 549 many zeros in the data set.

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Figure 7. Predicted associations between vegetation species richness and flow events (spring and summer freshes) in each zone and sampling period.

Solid points are mean estimates of species richness for each functional group and error bars bound 95% confidence intervals. Note that the y-axis varies between functional groups is truncated to a maximum value of 0.8, which reflects the relatively low values of average species richness at the sub-transect level.

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4. Discussion

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This study highlights the complex and multifaceted nature of riverine vegetation dynamics in regulated river systems, emphasising the critical role of environmental flows in maintaining and restoring these modified ecosystems. Our findings demonstrate that while individual flow events may produce modest immediate vegetation responses, their cumulative impact over time is substantial, which is expected to increase the resilience of riverine systems to disturbances. However, the effectiveness of environmental flows is not uniform across all vegetation groups or sites; it is highly context-dependent, influenced by local environmental conditions, species-specific traits, and the interplay of non-flow factors such as grazing, soil properties, and the presence of exotic species.

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The variability in vegetation responses reveals the necessity for tailored, adaptive management approaches that consider the unique characteristics of each river system and the site-specific ecological objectives. Short-duration freshes are beneficial for emergent and riparian species, while prolonged elevated flows can be detrimental to riparian and terrestrial species in particular if not

571 carefully managed. Moreover, the influence of non-flow factors, specifically the dominance of exotic
572 species and the impacts of livestock grazing, suggests that environmental flows alone will be
573 insufficient to achieve desired ecological outcomes without additional livestock or weed
574 management. Integrated management strategies that address both hydrological and non-hydrological
575 pressures are essential for the long-term health and functionality of riverine ecosystems.

576 **4.1. Vegetation patterns and drivers**

577 The distribution of species along bank elevations at the site scale provides a clear demonstration of
578 the impact of flow events and regimes at different bank zones, with species such as *Phragmites*
579 *australis* occupying lower elevations close to baseflow, while flood-intolerant species like *Oxalis pes-*
580 *caprae* were restricted to higher elevations above the spring-fresh level. This zonation illustrates the
581 importance of hydrological gradients in structuring riverine vegetation communities. However, the
582 variation in dominant species among different waterways and the lack of uniform patterns across sites
583 complicate efforts to generalise vegetation responses to flow events and regimes. These varying
584 species assemblages and traits among waterways and regions mean that no single management
585 approach will apply to all waterways, however there are generalisable categories that can help inform
586 management actions given the conditions and constraints of a system (e.g. Good and Jones 2026).

587 Long term data trends that were available at a small number of sites (6/44) highlight the rates of
588 change in vegetation under relatively consistent flow regimes. These clearly show increasing trends in
589 plant cover for emergent, riparian and terrestrial species in the spring fresh zone, with no trends above
590 the fresh zone (too dry) or for riparian and terrestrial plants below the baseflow level (too wet). The
591 trends suggest a cumulative benefit of regular (annual) delivery of environmental flows for emergent
592 species in the baseflow zone and most species in the fresh zone. The slow rate of change in relatively
593 low rainfall regions provides important context to management expectations, objectives, and the need
594 for long-term data to detect important vegetation change. However, the rates of change are likely to be
595 muted by the competitive influence of exotic riparian species that will reduce the magnitude of change
596 in native riparian plants at many sites (Catford et al. 2011), particularly for sites in more productive
597 regions that have higher exotic plant abundance. These rates will also be important for vegetation
598 communities to recover following flood or drought disturbance and to build resilience prior to future
599 disturbances, which are predicted to increase in frequency and intensity in the future within the study
600 region (Zhang et al. 2020).

601 **4.2. Vegetation responses to flows**

602 Clear short term benefits to aquatic and emergent species following freshes (or natural flows with
603 equivalent flow attributes) are expected due to the high preference and tolerance to inundation (Vivian
604 et al. 2025). Emergent species benefits above the spring-fresh level likely result from belowground
605 water resources via lateral bank recharge (Deng et al. 2024), that diminish in autumn with seasonal
606 plant senescence. The high variability in plant responses to specific flow events is likely to be largely
607 influenced by the relatively small short term changes as a result of brief managed flows compared to
608 the variability among years, seasons and sites. In contrast, the long-term effects of flow regimes
609 showed clear positive associations with managed flows that closely corresponded with vegetation
610 response patterns derived from recent experimental studies. Maintaining beneficial flow regimes in
611 the long term is likely to be important for riparian plant communities to recover following disturbances
612 and build resilience to threats.

613 Sustained baseflows were generally detrimental to riparian and terrestrial species that are intolerant
614 of extended inundation (Vivian et al. 2020; Main et al. 2023). However, aquatic species benefited from

615 these conditions, as sustained baseflows provided consistent water availability, supporting their
616 growth and persistence. Regimes of short-duration freshes (15–25 days) were associated with the
617 highest cover in all vegetation groups within the fresh zone, indicating that regimes of freshes
618 approximately 10-30 days long in late-winter/spring flow events can stimulate vegetation growth
619 without causing the adverse effects associated with prolonged inundation found in controlled nursery
620 studies (Vivian et al. 2020; Kitanovic et al. 2023; Main et al. 2023; Wijepala et al. 2026). Inundation
621 duration beyond ~30 days resulted in declines in all species, matching the tolerance thresholds
622 observed in controlled conditions. Negative impacts on species richness occurred following longer
623 inundation durations (~26, 35, and 38 days for terrestrial, riparian and emergent plants respectively)
624 because richness only declines when the plants die, while declines in health and biomass occur
625 earlier (Vivian et al. 2020; Kitanovic et al. 2023; Main et al. 2023; Wijepala et al. 2026). Aquatic plants
626 showed no significant indication of mortality from extended flows, as expected.

627 The variation in functional group cover and species richness among different systems, sites, and
628 transects further highlights the importance of local context in shaping vegetation responses to flow
629 regimes. For example, the Moorabool and West Gippsland systems exhibited higher-than-expected
630 overall cover, suggesting that other flow or site-specific factors, such as soil properties, water quality,
631 land use, or historical disturbances, may be playing a significant role in these areas. Plant responses
632 to inundation can vary substantially between species even within plant functional groups, making
633 generalisation difficult (Van Eck et al. 2006; Main et al. 2022). Additional factors could be added to
634 future versions of this assessment, but the statistical models are already very complex and there are
635 limits to what can be done without also increasing the vegetation survey data volume at all sites to
636 support model complexity. The lack of significant differences in cover between native and exotic
637 species across functional groups suggests that they have similar responses and flow regimes alone
638 may not be sufficient to control the spread of exotic species, highlighting the need for integrated
639 management approaches to achieve native vegetation objectives that vary among waterways or
640 regions (Good and Jones 2026).

641 While the regimes recorded within this study were influenced by environmental flows and unregulated
642 natural flows in different years, the regimes were indicative of the environmental flow regulated system
643 for each waterway. The spatial influences were also restricted to areas most heavily influenced by
644 consistently applied environmental flows. It is impossible to create a monitoring design in the field
645 where the effects of environmental flows and unregulated natural flows can be fully disentangled from
646 one another. The insights from this analysis relate directly to real-world management scenarios that
647 are shaped by environmental flows.

648 **4.3. The influence of non-flow factors**

649 The role of non-flow factors in shaping vegetation patterns and responses cannot be overstated
650 (Campbell et al. 2023; Good and Jones 2024; Nystrand et al. 2025). In many of the study sites, the
651 presence of exotic species, particularly in agricultural landscapes, was a dominant factor influencing
652 vegetation dynamics. Exotic species, such as invasive herbs and agricultural grasses, have similar
653 responses to flows as native species and compete with native vegetation, leading to reduced native
654 species richness and cover where they are abundant (Appendix E1). The dominant functional group of
655 exotic plants varied among waterways and therefore had a different relative influence depending on
656 the starting species composition. In drier landscapes, exotic plants were primarily terrestrial and so
657 had limited impact on the native riparian plants on the lower bank of waterways when flows were
658 present. While in wetter regions, the exotic plants were commonly flood-tolerant and were more often
659 found to fully exclude native riparian plants. Exotic plant influences will continue to be a major threat

660 to native plant populations in regulated rivers in the study region and will need to be actively managed,
661 particularly where they are primarily flood-tolerant species (Catford et al. 2011; Greet et al. 2013;
662 Tonkin et al 2020).

663 Grazing pressure was another critical factor, with sites subjected to livestock grazing showing
664 significant reductions in vegetation cover and species richness, particularly among native species.
665 This finding aligns with previous studies that have demonstrated the negative impacts of grazing on
666 riverine vegetation, particularly in terms of trampling, soil compaction, and the preferential
667 consumption of palatable native species (Jones and Vesk 2016; Jones et al. 2022; Nystrand et al.
668 2025). However, the impact of grazing was often inconsistent within a site and therefore the use of a
669 whole site variable for grazing presence is likely to have been less informative than assessing grazing
670 influence at the transect (sub-site) scale. Other non-flow factors that can have an important influence
671 on vegetation, including soil properties (Hale et al. 2014, Fagundes et al. 2019) and historical
672 disturbances (Capon et al. 2016), were not assessed in this study. These interacting influences
673 reinforce the need for a holistic approach to vegetation management—one that considers the full
674 range of environmental influences on riverine ecosystems (Campbell et al. 2023; Good and Jones
675 2026; Nystrand et al. 2025).

676 **4.4. Implications for management and monitoring**

677 Environmental flows have clear benefits to native aquatic, emergent and riparian plants in regulated
678 waterways that are cumulative over several years. These benefits were larger for cover than richness,
679 but a small cumulative increase in richness may be adequate for ecosystem health and function. This
680 positive outcome is also significant considering the high occurrence of exotic species that are present
681 within some sites and respond positively to flows and outcompete native species (Catford et al. 2011;
682 Tonkin et al. 2020). Importantly, given that native and exotic species respond in similar ways to flows,
683 riparian or flood tolerant exotic species cannot be controlled with flows and would need a co-
684 ordinated management strategy of flow and non-flow actions to reduce their populations (Good and
685 Jones 2024).

686 While it is important for managers to consider their local context when applying management actions,
687 this study supports the development of clear management ranges that will best support native
688 vegetation in riparian zones. Freshes of around 15-20 days (range 10-25) are optimal for most riparian
689 and emergent plant cover, but flows should be restricted to <25 days in spring and <10 days in summer
690 (Vivian et al. 2020; Main et al. 2023; Wijepala et al. 2025) to avoid negative impacts. Species richness
691 may be optimised by longer flow durations (~35 days), but this may come at a risk of maintaining plant
692 abundance, health and function. Larger and longer duration freshes (within the bounds above) that
693 occur annually are likely to provide the most benefit to desirable plant functional groups while
694 disadvantaging terrestrial plants, but low flow periods of up to 12-14 weeks after floods (Gower et al.
695 2024) and >4 weeks between freshes should be used to enable growth and recovery.

696 Given the influence of non-flow factors observed in this study and others, integrated management
697 approaches that address grazing pressures, the spread of exotic species, and other land-use impacts
698 are essential for achieving long-term ecological health (Good and Jones 2024; Nystrand et al. 2025).
699 Moreover, the variability in vegetation responses among different systems and sites highlights the
700 importance of local context in informing management decisions. This includes variability at the site
701 level caused by historical and future climate, soils, historical disturbances, and threats. By integrating
702 site-specific knowledge with broader ecological principles, waterway managers can develop more
703 effective strategies for maintaining and restoring the health of riverine vegetation (Good and Jones

704 2024). Future studies could build on the results presented here to address data shortfalls, limited data
705 coverage (spatially and temporally), and the influences of integrated flow and non-flow management.

706

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729

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Appendix A

Table A1. Site and survey information.

Catchment	Waterway	Number of sites	Number of transects	Number of sub-transects	Number of site surveys	Number of visits to waterbody	Survey years
Campaspe	Campaspe	6	61	565	96	16	2016–2023
Glenelg	Glenelg	6	54	267	27	7	2018-19, 2022-23
Glenelg	Wannon	1	5	24	3	3	2018-2019
Loddon	Loddon	2	10	55	6	3	2017-2018
Loddon	Tullaroop	2	20	109	6	3	2017-2018
Loddon	Twelve Mile	2	10	56	6	3	2017-2018
Moorabool	Moorabool	5	46	227	20	6	2017-18, 2022-23
Moorabool	Sutherland	1	10	41	3	3	2017-2018
Thomson	Macalister	4	21	102	10	3	2018-2019
Thomson	Thomson	3	21	106	9	3	2018-2019
Wimmera	Burnt	4	20	89	22	6	2017-2019
Wimmera	McKenzie	4	19	88	24	6	2017-2019
Wimmera	Mt William	4	22	93	18	6	2017-2019
Yarra	Watts	1	5	24	3	3	2018-2019
Yarra	Yarra	3	34	188	10	4	2018-2019

Table A2. The most common species within each vegetation group in each waterway.

Waterway	Aquatic (exotic)	Aquatic (native)	Emergent (exotic)	Emergent (native)	Riparian (exotic)	Riparian (native)	Terrestrial (exotic)	Terrestrial (native)
Campaspe	NA	<i>Cycnogeton procerum</i> , <i>Characeae</i> spp., <i>Myriophyllum</i> spp., <i>Vallisneria australis</i> , <i>Potamogeton crispus</i>	NA	<i>Phragmites australis</i> , <i>Bolboschoenus medianus</i> , <i>Typha</i> spp., <i>Typha domingensis</i> , <i>Bolboschoenus</i> spp.	<i>Paspalum distichum</i> , <i>Cyperus eragrostis</i> , <i>Juncus acutus</i> subsp. <i>acutus</i> , <i>Juncus articulatus</i> subsp. <i>articulatus</i> , <i>Agrostis stolonifera</i>	<i>Lachnagrostis filiformis</i> s.l., <i>Juncus usitatus</i> , <i>Carex bichenoviana</i> , <i>Persicaria prostrata</i> , <i>Juncus amabilis</i>	<i>Cynodon dactylon</i> var. <i>dactylon</i> , <i>Panicum coloratum</i> , <i>Paspalum dilatatum</i> , <i>Lolium</i> spp., <i>Bromus diandrus</i>	<i>Poa labillardierei</i> var. <i>labillardierei</i> , <i>Paspalidium jubiflorum</i> , <i>Hemarthria uncinata</i> var. <i>uncinata</i> , <i>Themeda triandra</i> , <i>Rytidosperma</i> spp.
Glenelg	<i>Elodea canadensis</i>	<i>Cycnogeton procerum</i> , <i>Ranunculus amphitrichus</i> , <i>Cycnogeton alcockiae</i> , <i>Characeae</i> spp., <i>Potamogeton ochreatus</i>	<i>Typha latifolia</i>	<i>Phragmites australis</i> , <i>Baumea juncea</i> , <i>Typha</i> spp., <i>Baumea</i> spp., <i>Baumea arthropphylla</i>	<i>Paspalum distichum</i> , <i>Agrostis stolonifera</i> , <i>Juncus articulatus</i> subsp. <i>articulatus</i> , <i>Polypogon monspeliensis</i> , <i>Cotula coronopifolia</i>	<i>Juncus amabilis</i> , <i>Eragrostis infecunda</i> , <i>Alternanthera denticulata</i> , <i>Selliera radicans</i> , <i>Lachnagrostis filiformis</i> s.l.	<i>Lolium</i> spp., <i>Holcus lanatus</i> , <i>Phalaris aquatica</i> , <i>Vulpia</i> spp., <i>Romulea rosea</i>	<i>Poa labillardierei</i> var. <i>labillardierei</i> , <i>Distichlis distichophylla</i> , <i>Rytidosperma</i> spp., <i>Crassula</i> spp., <i>Hemarthria uncinata</i> var. <i>uncinata</i>
Wannon	<i>Elodea canadensis</i>	<i>Cycnogeton procerum</i> , <i>Ornduffia reniformis</i> , <i>Characeae</i> spp., <i>Cycnogeton alcockiae</i> , <i>Nitella</i> sp.	<i>Typha latifolia</i>	<i>Phragmites australis</i> , <i>Eleocharis acuta</i> , <i>Baumea juncea</i> , <i>Baumea arthropphylla</i> , <i>Baumea</i> spp.	<i>Agrostis stolonifera</i> , <i>Cotula coronopifolia</i> , <i>Cynosurus echinatus</i> , <i>Cyperus eragrostis</i> , <i>Isolepis hystrix</i>	<i>Lachnagrostis filiformis</i> s.l., <i>Carex appressa</i> , <i>Selliera radicans</i> , <i>Isolepis inundata</i> , <i>Glyceria australis</i>	<i>Anthoxanthum odoratum</i> , <i>Leontodon saxatilis</i> , <i>Hypochaeris radicata</i> , <i>Holcus lanatus</i> , <i>Phalaris aquatica</i>	<i>Poa labillardierei</i> var. <i>labillardierei</i> , <i>Lobelia anceps</i> , <i>Senecio biserratus</i> , <i>Lomandra longifolia</i> , <i>Senecio minimus</i>
Loddon	NA	<i>Cycnogeton procerum</i> , <i>Myriophyllum</i> spp., <i>Ranunculus inundatus</i>	NA	<i>Bolboschoenus medianus</i> , <i>Eleocharis acuta</i> , <i>Phragmites australis</i> ,	<i>Cyperus eragrostis</i> , <i>Agrostis stolonifera</i> , <i>Mentha pulegium</i> ,	<i>Cyperus exaltatus</i> , <i>Carex tereticaulis</i> , <i>Alternanthera denticulata</i> ,	<i>Lolium rigidum</i> , <i>Lolium perenne</i> , <i>Helminthotheca echioides</i> , <i>Symphytotrichum</i>	<i>Poa fordeana</i> , <i>Asperula gemella</i> , <i>Paspalidium jubiflorum</i> , <i>Walwhalleya</i>

Waterway	Aquatic (exotic)	Aquatic (native)	Emergent (exotic)	Emergent (native)	Riparian (exotic)	Riparian (native)	Terrestrial (exotic)	Terrestrial (native)
				<i>Schoenoplectus tabernaemontani</i>	<i>Paspalum distichum</i> , <i>Polypogon monspeliensis</i>	<i>Juncus aridicola</i> , <i>Juncus amabilis</i>	<i>subulatum</i> , <i>Cirsium vulgare</i>	<i>proluta</i> , <i>Asperula conferta</i>
Tullaroop	NA	<i>Cycnogeton procerum</i> , <i>Myriophyllum</i> spp., <i>Ranunculus inundatus</i>	NA	<i>Phragmites australis</i> , <i>Bolboschoenus medianus</i> , <i>Schoenoplectus tabernaemontani</i> , <i>Eleocharis acuta</i>	<i>Cyperus eragrostis</i> , <i>Paspalum distichum</i> , <i>Agrostis stolonifera</i> , <i>Polypogon monspeliensis</i> , <i>Mentha pulegium</i>	<i>Juncus usitatus</i> , <i>Lachnagrostis filiformis</i> s.l., <i>Carex bichenoviana</i> , <i>Juncus amabilis</i> , <i>Carex tereticaulis</i>	<i>Cynodon dactylon</i> var. <i>dactylon</i> , <i>Cirsium vulgare</i> , <i>Phalaris aquatica</i> , <i>Oxalis pes-caprae</i> , <i>Bromus catharticus</i>	<i>Hemarthria uncinata</i> var. <i>uncinata</i> , <i>Lythrum hyssopifolia</i> , <i>Oxalis perennans</i> , <i>Rumex brownii</i> , <i>Euchiton involucratus</i> s.l.
Twelve Mile	NA	<i>Cycnogeton procerum</i> , <i>Ranunculus inundatus</i> , <i>Myriophyllum</i> spp.	NA	<i>Schoenoplectus tabernaemontani</i> , <i>Bolboschoenus medianus</i> , <i>Eleocharis acuta</i> , <i>Phragmites australis</i>	<i>Cyperus eragrostis</i> , <i>Mentha pulegium</i> , <i>Paspalum distichum</i> , <i>Ranunculus sceleratus</i> subsp. <i>sceleratus</i> , <i>Agrostis stolonifera</i>	<i>Eragrostis infecunda</i> , <i>Juncus aridicola</i> , <i>Alternanthera denticulata</i> , <i>Cyperus exaltatus</i> , <i>Juncus amabilis</i>	<i>Lolium rigidum</i> , <i>Helminthotheca echinoides</i> , <i>Symphotrichum subulatum</i> , <i>Polygonum aviculare</i> , <i>Cirsium vulgare</i>	<i>Calotis scapigera</i> , <i>Oxalis perennans</i> , <i>Rumex tenax</i> , <i>Lythrum hyssopifolia</i> , <i>Rytidosperma</i> spp.
Moorabool	<i>Elodea canadensis</i>	<i>Cycnogeton procerum</i> , <i>Potamogeton crispus</i> , <i>Ceratophyllum demersum</i> , <i>Myriophyllum</i> spp., <i>Lemna disperma</i>	NA	<i>Phragmites australis</i> , <i>Schoenoplectus tabernaemontani</i> , <i>Eleocharis acuta</i> , <i>Bolboschoenus medianus</i> , <i>Typha domingensis</i>	<i>Agrostis stolonifera</i> , <i>Juncus acutus</i> subsp. <i>acutus</i> , <i>Paspalum distichum</i> , <i>Juncus articulatus</i> subsp. <i>articulatus</i> , <i>Cyperus eragrostis</i>	<i>Ficinia nodosa</i> , <i>Lachnagrostis filiformis</i> s.l., <i>Carex appressa</i> , <i>Persicaria decipiens</i> , <i>Isolepis inundata</i>	<i>Phalaris aquatica</i> , <i>Ehrharta erecta</i> var. <i>erecta</i> , <i>Anthoxanthum odoratum</i> , <i>Bromus diandrus</i> , <i>Dactylis glomerata</i>	<i>Poa labillardierei</i> var. <i>labillardierei</i> , <i>Microlaena stipoides</i> var. <i>stipoides</i> , <i>Pteridium esculentum</i> , <i>Lomandra longifolia</i> , <i>Senecio quadridentatus</i>
Sutherland	<i>Elodea canadensis</i>	<i>Cycnogeton procerum</i> ,	NA	<i>Phragmites australis</i> ,	<i>Juncus acutus</i> subsp. <i>acutus</i> ,	<i>Ficinia nodosa</i> , <i>Lachnagrostis</i>	<i>Phalaris aquatica</i> ,	<i>Oxalis perennans</i> ,

Waterway	Aquatic (exotic)	Aquatic (native)	Emergent (exotic)	Emergent (native)	Riparian (exotic)	Riparian (native)	Terrestrial (exotic)	Terrestrial (native)
		<i>Ceratophyllum demersum</i> , <i>Characeae</i> spp., <i>Lemna disperma</i> , <i>Myriophyllum</i> spp.		<i>Bolboschoenus medianus</i> , <i>Typha domingensis</i> , <i>Eleocharis acuta</i> , <i>Schoenoplectus tabernaemontani</i>	<i>Polypogon monspeliensis</i> , <i>Paspalum distichum</i> , <i>Agrostis stolonifera</i> , <i>Cyperus eragrostis</i>	<i>filiformis</i> s.l., <i>Montia australasica</i> , <i>Isolepis cernua</i> , <i>Alternanthera denticulata</i>	<i>Ehrharta erecta</i> var. <i>erecta</i> , <i>Cynodon dactylon</i> var. <i>dactylon</i> , <i>Bromus diandrus</i> , <i>Galenia pubescens</i>	<i>Lobelia anceps</i> , <i>Rumex brownii</i> , <i>Rytidosperma</i> spp., <i>Acaena echinata</i>
Macalister	NA	<i>Cycnogeton procerum</i> , <i>Potamogeton crispus</i> , <i>Potamogeton</i> spp.	NA	<i>Phragmites australis</i> , <i>Schoenoplectus tabernaemontani</i> , <i>Baumea</i> spp., <i>Bolboschoenus medianus</i> , <i>Bolboschoenus</i> spp.	<i>Phalaris arundinacea</i> , <i>Cyperus eragrostis</i> , <i>Paspalum distichum</i> , <i>Glyceria maxima</i> , <i>Agrostis stolonifera</i>	<i>Isachne globosa</i> , <i>Persicaria praetermissa</i> , <i>Persicaria subsessilis</i> , <i>Carex appressa</i> , <i>Persicaria decipiens</i>	<i>Cynodon dactylon</i> var. <i>dactylon</i> , <i>Cenchrus clandestinus</i> , <i>Tradescantia fluminensis</i> , <i>Dactylis glomerata</i> , <i>Hypochaeris radicata</i>	<i>Microlaena stipoides</i> var. <i>stipoides</i> , <i>Hemarthria uncinata</i> var. <i>uncinata</i> , <i>Rumex brownii</i> , <i>Lobelia anceps</i> , <i>Lythrum hyssopifolia</i>
Thomson	NA	<i>Potamogeton</i> spp., <i>Potamogeton crispus</i> , <i>Cycnogeton procerum</i>	NA	<i>Schoenoplectus tabernaemontani</i> , <i>Baumea</i> spp., <i>Schoenoplectus</i> spp., <i>Bolboschoenus medianus</i> , <i>Bolboschoenus</i> spp.	<i>Ranunculus repens</i> , <i>Phalaris arundinacea</i> , <i>Agrostis stolonifera</i> , <i>Cyperus eragrostis</i> , <i>Paspalum distichum</i>	<i>Persicaria hydropiper</i> , <i>Persicaria praetermissa</i> , <i>Persicaria subsessilis</i> , <i>Juncus amabilis</i> , <i>Isolepis inundata</i>	<i>Tradescantia fluminensis</i> , <i>Vinca major</i> , <i>Dactylis glomerata</i> , <i>Cynodon dactylon</i> var. <i>dactylon</i> , Monocot grass	<i>Microlaena stipoides</i> var. <i>stipoides</i> , <i>Cardamine</i> spp., <i>Dichondra repens</i> , <i>Lythrum hyssopifolia</i> , <i>Centella cordifolia</i>
Burnt	<i>Callitriche brutia</i> var. <i>brutia</i> , <i>Callitriche stagnalis</i>	<i>Cycnogeton procerum</i> , <i>Myriophyllum simulans</i> , <i>Ottelia ovalifolia</i> subsp. <i>ovalifolia</i> , <i>Gratiola peruviana</i> , <i>Potamogeton cheesemaniae</i>	NA	<i>Eleocharis acuta</i> , <i>Baumea arthropphylla</i> , <i>Baumea articulata</i> , <i>Eleocharis sphacelata</i> , <i>Phragmites australis</i>	<i>Paspalum distichum</i> , <i>Juncus articulatus</i> subsp. <i>articulatus</i> , <i>Polypogon monspeliensis</i> , <i>Cyperus eragrostis</i> , <i>Agrostis stolonifera</i>	<i>Carex tereticaulis</i> , <i>Juncus amabilis</i> , <i>Carex gaudichaudiana</i> , <i>Ficinia nodosa</i> , <i>Baloskion tetraphyllum</i> subsp. <i>tetraphyllum</i>	<i>Ehrharta calycina</i> , <i>Paspalum dilatatum</i> , <i>Lolium rigidum</i> , <i>Romulea rosea</i> , <i>Holcus lanatus</i>	<i>Rytidosperma</i> spp., <i>Lythrum hyssopifolia</i> , <i>Eragrostis brownii</i> , <i>Microlaena stipoides</i> var. <i>stipoides</i> , <i>Goodenia humilis</i>

Waterway	Aquatic (exotic)	Aquatic (native)	Emergent (exotic)	Emergent (native)	Riparian (exotic)	Riparian (native)	Terrestrial (exotic)	Terrestrial (native)
McKenzie	<i>Callitriche brutia</i> var. <i>brutia</i> , <i>Callitriche stagnalis</i>	<i>Cycnogeton procerum</i> , Characeae spp., <i>Gratiola pumilo</i> , <i>Ceratophyllum demersum</i> , <i>Gratiola peruviana</i>	NA	<i>Schoenoplectus tabernaemontani</i> , <i>Baumea arthropphylla</i> , <i>Baumea articulata</i> , <i>Eleocharis acuta</i> , <i>Eleocharis sphacelata</i>	<i>Juncus articulatus</i> subsp. <i>articulatus</i> , <i>Agrostis stolonifera</i> , <i>Cynosurus echinatus</i> , <i>Cyperus eragrostis</i> , <i>Cotula coronopifolia</i>	<i>Juncus amabilis</i> , <i>Carex gaudichaudiana</i> , <i>Lachnagrostis filiformis</i> s.l., <i>Juncus procerus</i> , <i>Chorizandra enodis</i>	<i>Ehrharta calycina</i> , <i>Holcus lanatus</i> , <i>Romulea rosea</i> , <i>Briza maxima</i> , <i>Phalaris aquatica</i>	<i>Hemarthria uncinata</i> var. <i>uncinata</i> , <i>Microlaena stipoides</i> var. <i>stipoides</i> , <i>Rytidosperma</i> spp., <i>Poa labillardierei</i> var. <i>labillardierei</i> , <i>Caesia</i> spp.
Mt William	<i>Callitriche stagnalis</i> , <i>Callitriche brutia</i> var. <i>brutia</i>	<i>Cycnogeton procerum</i> , <i>Ceratophyllum demersum</i> , Characeae spp., <i>Myriophyllum simulans</i> , <i>Gratiola peruviana</i>	NA	<i>Phragmites australis</i> , <i>Eleocharis sphacelata</i> , <i>Baumea articulata</i> , <i>Eleocharis acuta</i> , <i>Baumea arthropphylla</i>	<i>Paspalum distichum</i> , <i>Cynosurus echinatus</i> , <i>Juncus articulatus</i> subsp. <i>articulatus</i> , <i>Cotula coronopifolia</i> , <i>Ranunculus sceleratus</i> subsp. <i>sceleratus</i>	<i>Juncus amabilis</i> , <i>Juncus pallidus</i> , <i>Carex appressa</i> , <i>Lachnagrostis filiformis</i> s.l., <i>Triglochin striata</i>	<i>Holcus lanatus</i> , <i>Bromus diandrus</i> , <i>Lolium rigidum</i> , <i>Anthoxanthum odoratum</i> , Monocot grass	<i>Microlaena stipoides</i> var. <i>stipoides</i> , <i>Lobelia anceps</i> , <i>Rytidosperma</i> spp., <i>Euchiton involucratus</i> s.l., <i>Acaena echinata</i>
Watts	<i>Callitriche stagnalis</i> , <i>Elodea canadensis</i>	<i>Gratiola peruviana</i> , <i>Azolla</i> spp.	NA	<i>Phragmites australis</i>	<i>Glyceria maxima</i> , <i>Ranunculus repens</i> , <i>Cyperus eragrostis</i> , <i>Paspalum distichum</i> , <i>Juncus articulatus</i> subsp. <i>articulatus</i>	<i>Persicaria praetermissa</i> , <i>Carex appressa</i> , <i>Persicaria decipiens</i> , <i>Isolepis inundata</i> , <i>Carex fascicularis</i>	<i>Vinca major</i> , <i>Lonicera japonica</i> , <i>Ehrharta erecta</i> var. <i>erecta</i> , <i>Cenchrus clandestinus</i> , <i>Oxalis incarnata</i>	<i>Tetrarrhena juncea</i> , <i>Blechnum nudum</i> , <i>Oxalis exilis</i> , <i>Lythrum hyssopifolia</i> , <i>Centella cordifolia</i>
Yarra	<i>Elodea canadensis</i> , <i>Callitriche stagnalis</i>	<i>Gratiola peruviana</i> , <i>Azolla</i> spp.	NA	<i>Phragmites australis</i>	<i>Phalaris arundinacea</i> , <i>Paspalum distichum</i> , <i>Cyperus</i>	<i>Persicaria hydropiper</i> , <i>Alternanthera denticulata</i> , <i>Persicaria</i>	<i>Cenchrus clandestinus</i> , <i>Tradescantia fluminensis</i> , <i>Allium</i>	<i>Microlaena stipoides</i> var. <i>stipoides</i> , <i>Poa labillardierei</i> var. <i>labillardierei</i> ,

Waterway	Aquatic (exotic)	Aquatic (native)	Emergent (exotic)	Emergent (native)	Riparian (exotic)	Riparian (native)	Terrestrial (exotic)	Terrestrial (native)
					<i>eragrostis, Ranunculus repens, Glyceria maxima</i>	<i>praetermissa, Juncus amabilis, Carex appressa</i>	<i>triquetrum, Cynodon dactylon var. dactylon, Dactylis glomerata</i>	<i>Pteridium esculentum, Poa ensiformis, Lomandra longifolia</i>

Appendix B

Table B1. Data analysis and modelling dataset

Variable Name	Unit	Definition	Variable Type	Variable Group
Species vegetation cover	Continuous: Cover	Cover per <i>species</i> , per <i>sub-transect</i>	Response	Vegetation Cover
Plant functional group cover		Cover per <i>plant functional group</i> , per <i>sub-transect</i>		
Sub-transect vegetation cover		Total cover (sum all species), per sub-transect		
Species richness	Integer: Counts	Number unique Species Functional Group, per <i>sub-transect</i>	Predictor	Species Richness
Days above baseflow / spring-fresh levels	Integer, Counts: Z- scale	Number of days per year (averaged over preceding 3 water-years) discharge > baseflow or spring-fresh levels, Z-scaled		
Sub-transect zone	Factor	Zone in which a sub-transect occurred: <ul style="list-style-type: none"> • Below baseflow level • Between baseflow and spring-fresh levels • Above spring-fresh level 		Flow Regime
Flow event	Factor	Timing of sampling: <ul style="list-style-type: none"> • Before spring fresh • Between spring and summer fresh • After summer fresh 	Predictor	Flow Event
Site grazing status	Factor	Grazing status of site: <ul style="list-style-type: none"> • Grazed • Ungrazed 		Other factors influencing spatio-temporal variation in

Variable Name	Unit	Definition	Variable Type	Variable Group
Species functional group	Factor	For each <i>Origin</i> group (native or exotic): <ul style="list-style-type: none"> • Aquatic • Emergent • Riparian • Terrestrial 		cover / abundance
Species origin	Factor	Origin of the recorded species: <ul style="list-style-type: none"> • Native • Exotic 	Predictor	
Site identity	Factor	Name / Identity of the site	Random effect	
Transect identity	Factor	Identity of nested within sites		
Sub-transect height	Continuous: Metres above baseflow	Height of the sub-transect along bank		
Survey year	Date: YYYY	Year in which the sample was recorded		
Waterbody	Factor	Waterbody in which the site is located		

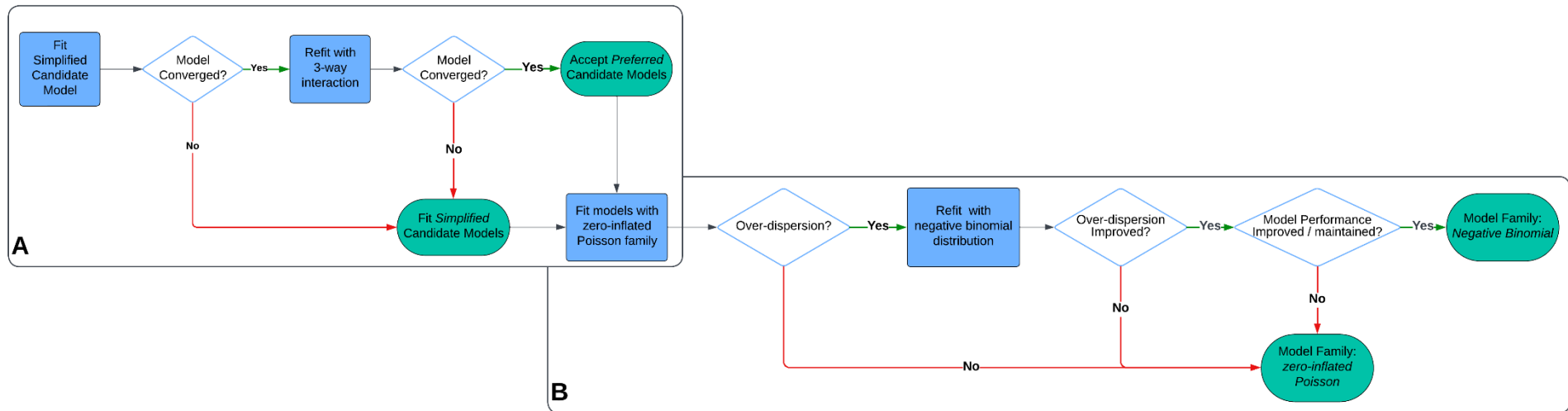


Figure B2. Preregistered decision-tree for selecting model structure (A) and selecting an alternative model functional form over the default zero-inflated *Poisson* (B). The decision-tree was preregistered in version 2 and version 3 of the preregistration (Figure B1, Table B2). Reproduced with permission from Gould et al. (2025).

Initial model fitting assumed a zero-inflated Poisson distribution for both vegetation cover and species richness, with zero-inflated negative binomial distributions designated in cases where over-dispersion was severe (Figure B2). This approach does not introduce an upper bound on estimates of cover (e.g., specifying a maximum cover level of 100%). We used this approach for three reasons. First, overlapping species may generate cover estimates that exceed 100%. Second, parameters of bounded distributions are challenging to estimate when many values occur near the bounds. Third, bounded distributions require link functions (e.g., the logit function) that complicate the specification and interpretation of autoregressive models. We note that the Poisson distribution approximates the bounded binomial distribution in the limit of many trials. The extent of zero-inflation and over-dispersion (and a model's capacity to capture these patterns) was assessed using posterior predictive checks, which use values simulated from the fitted model to determine whether the model structure captures the observed distribution of the response variable (Gelman and Hill, 2007; Gelman et al. 2014). Pilot study models of vegetation cover at the species level did not converge in any instance. Models of vegetation cover aggregated to plant functional groups converged but required separate models of two-way interactions instead of the three- and four-way interactions included in the maximal model structure. Models of vegetation cover aggregated to functional groups (aquatic, emergent, riparian, terrestrial) converged but required removal of interactions with species origin and separate analyses of flow metrics and for each of the categorical flow terms (zone and sampling period). Models of vegetation species richness within functional groups converged but required separate models of two-way interactions instead of the three- and four-way interactions included

in the maximal model structure. Models of vegetation species richness within functional groups converged but required removal of interactions with species origin and separate models for the two categorical flow variables (zone and sampling period).

Table B2. Key stages of the modelling and Adaptive Preregistration process (reproduced with permission from Gould et al. 2025). Version numbers in parentheses are GitHub tags, marking snapshots of the repository for each Preregistration Version.

Preregistration Version	Preregistration Process	Modelling and Analysis Process
Version 1: (v0.10 – v0.8.1)	Establish problem context, study background and aims, describe data collection and data cleaning process, articulate modelling objectives, and specify candidate models.	Data cleaning and preparation undertaken concurrently by lead and supporting researchers due to limited resourcing and short timelines for project deliverables. Candidate models addressing different components of the research question were preregistered. Data structural properties and potential consequences for model fitting were identified
Version 2: (v0.9.0 – v.0.12.5)	Describe Pilot Analysis on data subset and specify registered flexibility in the form of decision heuristics.	Critical uncertainties in modelling decisions were articulated (e.g. how do we classify best flow regime based on inundation data?). In addition, data partitioning, exploratory data analysis, refined candidate models and model checking procedure preregistered
Version 3: (v0.13.0 – v.0.20.1)	Describe Main Analysis, including specification of decision tree for model selection process based on results of pilot analysis.	Conducted modelling and analysis as outlined in preregistration. Reported deviations and rationale as necessary.

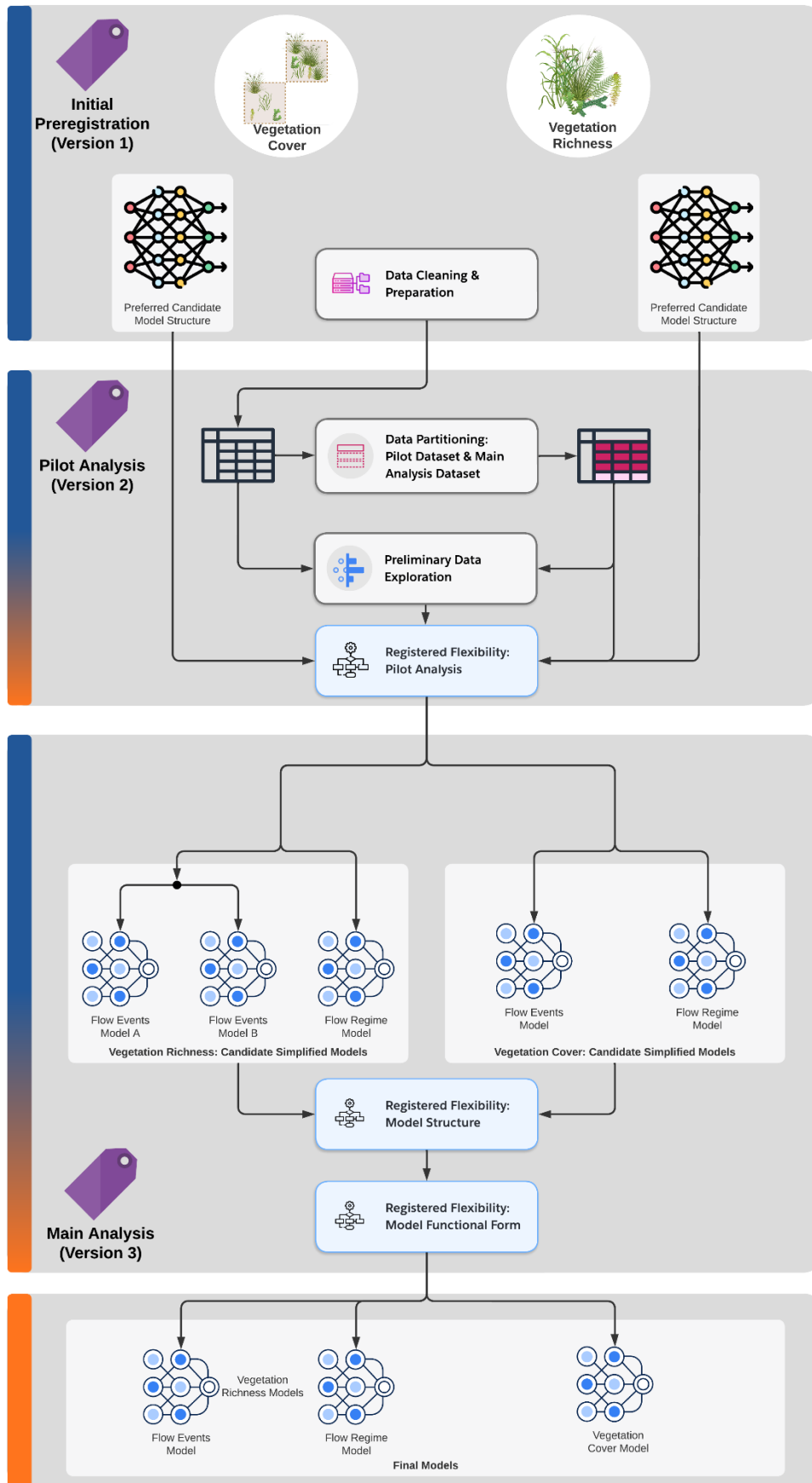


Figure B1. Model development and adaptive preregistration process (reproduced with permission from Gould et al. 2025). Three versions of the model analysis plan were preregistered. Each preregistration version and stage of the modelling process is briefly described in Table B2. The two registered flexibility steps in the Main Analysis (v3) (boxes shown in blue) are shown.

Appendix C

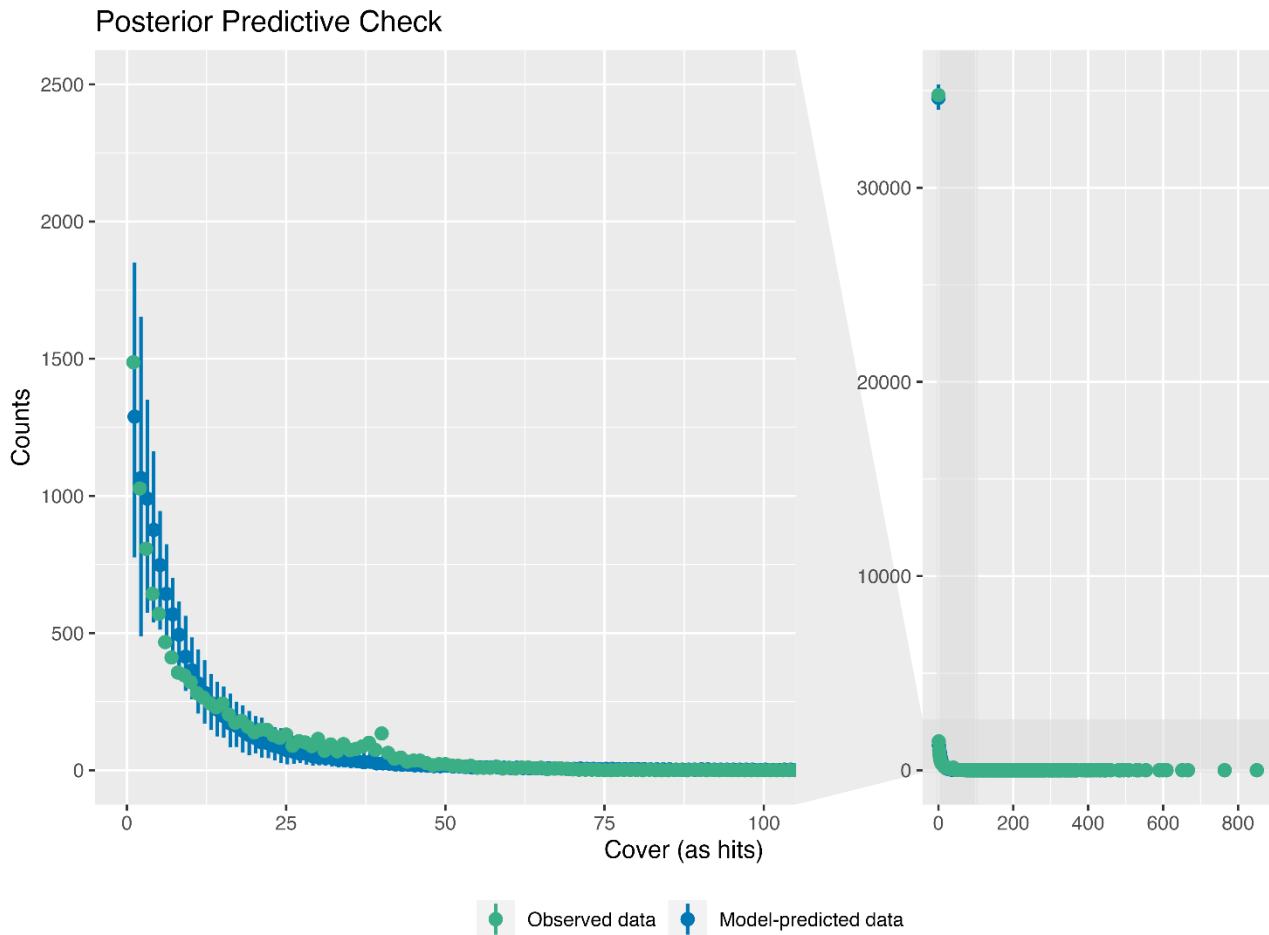


Figure C1. Posterior predictive check for the flow-regime model of vegetation cover. Blue points display the median predicted density of cover and green points display the observed density of vegetation cover. Error bars (not visible) bound 95% confidence intervals.

A model that reliably captures variation in a given response variable is indicated by close agreement between the blue and green points. This figure illustrates close agreement for zero observations (i.e., the model characterises zero-inflation well) but under-prediction at very low levels of cover and over-prediction at intermediate levels of cover. Assessments of alternative model structures (e.g., negative binomial distributions) did not improve alignment between modelled and observed values.

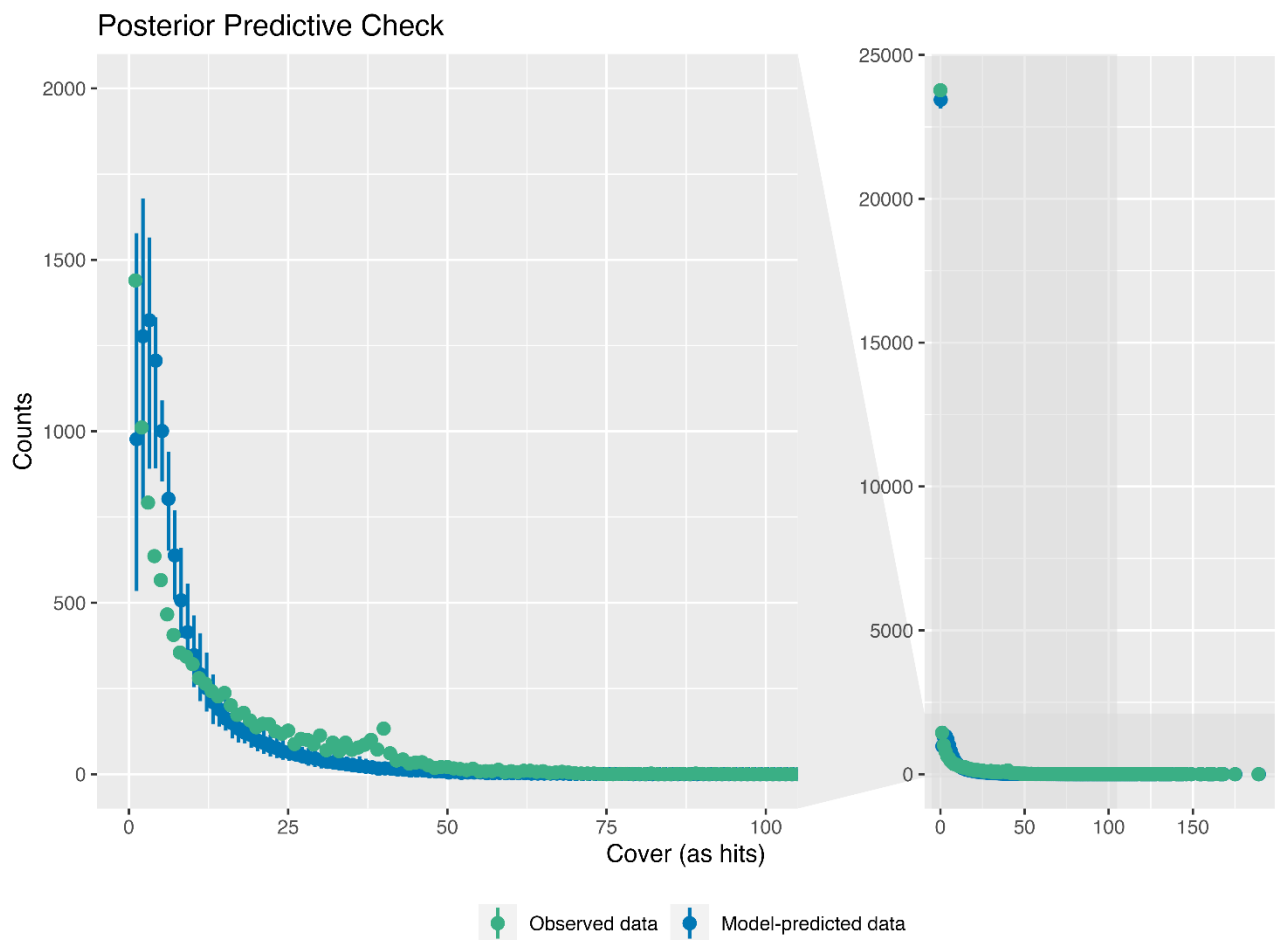


Figure C2. Posterior predictive check for the flow-event model of vegetation cover. Blue points display the median predicted density of cover and green points display the observed density of vegetation cover. Error bars (not visible) bound 95% confidence intervals.

A model that reliably captures variation in a given response variable is indicated by close agreement between the blue and green points. This figure illustrates close agreement for zero observations (i.e., the model characterises zero-inflation well) but under-prediction at very low levels of cover and over-prediction at intermediate levels of cover. Assessments of alternative model structures (e.g., negative binomial distributions) did not improve alignment between modelled and observed values.

Posterior Predictive Check

Model-predicted intervals should include observed data points

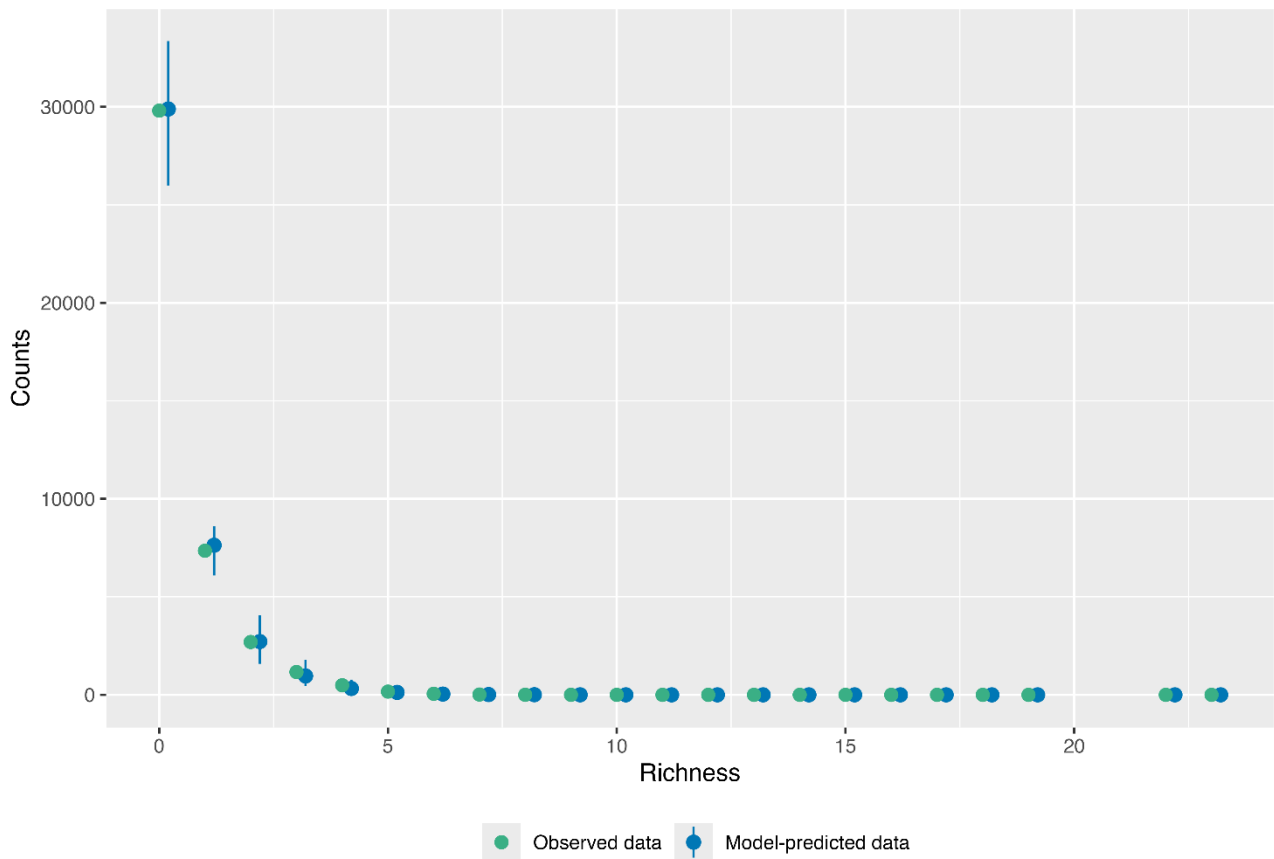


Figure C3. Posterior predictive check for the model of vegetation species richness. Blue points display the median predicted density of species richness and green points display the observed density of vegetation species richness. Error bars (visible for modelled but not observed values) bound 95% confidence intervals.

A model that reliably captures variation in a given response variable is indicated by close agreement between the blue and green points. This figure illustrates close agreement between modelled and observed species richness over the observed range of values.

Table C1. Assessment of overdispersion for all models.

Overdispersion was assessed using the *check_overdispersion()* function from the *performance* package (Ludecke, et al. 2021). For richness models (which do not include a zero-inflation parameter), overdispersion tests were based on the sum of squared Pearson residuals. The ratio of this sum to the residual degrees of freedom (assuming one model degree of freedom for each variance or covariance parameter) should approximately follow a χ^2 distribution (from which a p-value can be calculated)². For cover models, this test is based on simulated residuals rather than Pearson residuals, and this approach does not return an exact χ^2 estimate or residual degrees of freedom³. Overdispersion is likely when the dispersion ratio is greater than 1 and when the *p* value for the dispersion ratio estimate is <0.05. If the *p* value is greater than 0.05 but the dispersion ratio is 1, then overdispersion is unlikely. If the dispersion ratio is < 1 then underdispersion is likely.

Model	χ^2	Dispersion Ratio	Residual <i>df</i>	<i>p</i> -value	Overdispersion
Richness	46,360.75	1.11	41,682	<2e-16	Overdispersion detected
Cover: flow regime	NA	0.45	NA	0.496	No overdispersion detected
Cover: flow event	NA	1.72	NA	0.064	No overdispersion detected

Table C2. Assessment of zero-inflation for all models.

Zero-inflation was assessed using the *check_zeroinflation()* function from the *performance* R package (Ludecke, et al. 2021). For the richness model, which did not contain a zero-inflation parameter, zero-inflation was assessed by simulating from the fitted model and comparing the number of predicted zeros with the number present in the observed data. Zero-inflation occurs when the number of observed zeros exceeds the number of predicted zeros⁴. For the cover models, the test of zero-dispersion is based on simulated residuals⁵. When the ratio of observed versus predicted zeros is less than a tolerance threshold (default of 1 +/-0.05 for the richness model and 1 +/-0.1 for the cover models), the model is said to be underfitting zeros, while if the ratio is greater than the upper tolerance threshold, then the model is said to be overfitting zeros. When the ratio falls within the upper and lower limits of tolerance, no zero-inflation or underfitting of zeros is detected.

Model	Number predicted zeros	Observed zeros	Dispersion ratio	Tolerance	<i>P</i> value	Zero-inflation
Richness	29,033	29,804	0.9741310	0.05	NA	No zero-inflation detected
Cover: flow regime	35,011	35,154	0.9959188	0.10	0.640	No zero-inflation detected
Cover: flow event	23,436	23,773	0.9858253	0.10	0.096	No zero-inflation detected

² See https://easystats.github.io/performance/reference/check_overdispersion.html for details.

³ See https://easystats.github.io/performance/reference/simulate_residuals.html

⁴ See https://easystats.github.io/performance/reference/check_zeroinflation.html?q=zeroinflation#null for details.

⁵ See https://easystats.github.io/performance/reference/simulate_residuals.html

Appendix D

Table D1: R packages, versions and citations used directly in this analysis

Package	Version	Reference
aae.hydro	0.0.1.9003	Yen (2020)
assertthat	0.2.1	Wickham (2019)
base	4.4.0	R Core Team (2024)
brms	2.21.0	Bürkner (2017), Bürkner (2018), Bürkner (2021)
cowplot	1.1.3	Wilke (2024)
effects	4.2.2	Fox and Weisberg (2019), Fox and Weisberg (2018), Fox (2003), Fox and Hong (2009)
ggforce	0.4.2	Pedersen (2024a)
ggtext	0.1.2	Wilke and Wiernik (2022)
glmmTMB	1.1.9	Brooks et al. (2017)
glue	1.7.0	Hester and Bryan (2024)
here	1.0.1	Müller (2020)
interactions	1.1.5	Long (2019)
jtools	2.2.2	Long (2022)
lme4	1.1.35.3	Bates et al. (2015)
mgcv	1.9.1	S. N. Wood (2011), S. N. Wood et al. (2016), S. N. Wood (2004), S. N. Wood (2017), S. N. Wood (2003)
mixedup	0.4.0	Clark (2024)
parameters	0.21.7	Lüdecke et al. (2020)
patchwork	1.2.0	Pedersen (2024b)
performance	0.11.0	Lüdecke, Ben-Shachar, et al. (2021)
pointblank	0.12.1	Iannone, Vargas, and Choe (2024)
qs	0.26.3	Ching (2024)
R.utils	2.12.3	Bengtsson (2023)
remotes	2.5.0	Csárdi et al. (2024)
renv	0.17.3	Ushey (2023)
rmarkdown	2.27	Allaire et al. (2024), Xie, Allaire, and Golemund (2018), Xie, Dervieux, and Riederer (2020)
rstanarm	2.32.1	Goodrich et al. (2024), Brilleman et al. (2018)
see	0.8.4	Lüdecke, Patil, et al. (2021)
sessioninfo	1.2.2	Wickham et al. (2021)
tidyverse	2.0.0	Wickham et al. (2019)
withr	3.0.0	Hester et al. (2024)

Table D2: R Session Information

```
— Session info —  
setting value
```

```

version R version 4.4.0 (2024-04-24)
os      macOS Ventura 13.6.6
system  aarch64, darwin20
ui      RStudio
language (EN)
collate en_US.UTF-8
ctype   en_US.UTF-8
tz      Australia/Melbourne
date    2024-06-05
rstudio 2024.04.1+748 Chocolate Cosmos (desktop)
pandoc  3.1.12.2 @ /opt/homebrew/bin/pandoc

```

-- Packages

! package	* version	date (UTC)	lib	source
aae.hydro	* 0.0.1.9003	2024-06-04	[1]	Github (aae-stats/aae.hydro@ab38b12)
P bit	4.0.5	2022-11-15	[?]	RSPM (R 4.4.0)
P bit64	4.0.5	2020-08-30	[?]	RSPM (R 4.4.0)
P cachem	1.1.0	2024-05-16	[?]	RSPM (R 4.4.0)
P cellranger	1.1.0	2016-07-27	[?]	RSPM (R 4.4.0)
P cli	3.6.2	2023-12-11	[?]	RSPM (R 4.4.0)
P colorspace	2.1-0	2023-01-23	[?]	RSPM (R 4.4.0)
P commonmark	1.9.1	2024-01-30	[?]	RSPM (R 4.4.0)
P crayon	1.5.2	2022-09-29	[?]	RSPM (R 4.4.0)
P curl	5.2.1	2024-03-01	[?]	RSPM (R 4.4.0)
P datapasta	3.1.0	2020-01-17	[?]	RSPM (R 4.4.0)
P devtools	2.4.5	2022-10-11	[?]	RSPM (R 4.4.0)
P digest	0.6.35	2024-03-11	[?]	RSPM (R 4.4.0)
P dplyr	* 1.1.4	2023-11-17	[?]	RSPM (R 4.4.0)
P ellipsis	0.3.2	2021-04-29	[?]	RSPM (R 4.4.0)
P evaluate	0.2.3	2023-11-01	[?]	RSPM (R 4.4.0)
P fansi	1.0.6	2023-12-08	[?]	RSPM (R 4.4.0)
P farver	2.1.2	2024-05-13	[?]	RSPM (R 4.4.0)
P fastmap	1.2.0	2024-05-15	[?]	RSPM (R 4.4.0)
P forcats	* 1.0.0	2023-01-29	[?]	RSPM (R 4.4.0)
P fs	1.6.4	2024-04-25	[?]	RSPM (R 4.4.0)
P generics	0.1.3	2022-07-05	[?]	RSPM (R 4.4.0)
P ggplot2	* 3.5.1	2024-04-23	[?]	RSPM (R 4.4.0)
P ggtext	0.1.2	2022-09-16	[?]	CRAN (R 4.4.0)
P glue	1.7.0	2024-01-09	[?]	RSPM (R 4.4.0)
P gluedown	1.0.9	2024-03-11	[?]	CRAN (R 4.4.0)
P grateful	0.2.4	2023-10-22	[?]	RSPM (R 4.4.0)
P gridtext	0.1.5	2022-09-16	[?]	CRAN (R 4.4.0)
P gt	* 0.10.1	2024-01-17	[?]	RSPM (R 4.4.0)
P gtable	0.3.5	2024-04-22	[?]	RSPM (R 4.4.0)
P here	* 1.0.1	2020-12-13	[?]	RSPM (R 4.4.0)
P highr	0.11	2024-05-26	[?]	CRAN (R 4.4.0)
P hms	1.1.3	2023-03-21	[?]	RSPM (R 4.4.0)
P htmltools	0.5.8.1	2024-04-04	[?]	RSPM (R 4.4.0)
P htmlwidgets	1.6.4	2023-12-06	[?]	RSPM (R 4.4.0)
P httpuv	1.6.15	2024-03-26	[?]	RSPM (R 4.4.0)
P httr	1.4.7	2023-08-15	[?]	RSPM (R 4.4.0)
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P jsonlite	1.8.8	2023-12-04	[?]	RSPM (R 4.4.0)
P knitr	1.47	2024-05-29	[?]	CRAN (R 4.4.0)
P labeling	0.4.3	2023-08-29	[?]	RSPM (R 4.4.0)
P later	1.3.2	2023-12-06	[?]	RSPM (R 4.4.0)
P lattice	0.22-6	2024-03-20	[?]	CRAN (R 4.4.0)
P lifecycle	1.0.4	2023-11-07	[?]	RSPM (R 4.4.0)
P lubridate	* 1.9.3	2023-09-27	[?]	RSPM (R 4.4.0)
P magrittr	2.0.3	2022-03-30	[?]	RSPM (R 4.4.0)
P markdown	1.12	2023-12-06	[?]	RSPM (R 4.4.0)
P Matrix	1.7-0	2024-03-22	[?]	CRAN (R 4.4.0)
P memoise	2.0.1	2021-11-26	[?]	RSPM (R 4.4.0)
P mgcv	1.9-1	2023-12-21	[?]	CRAN (R 4.4.0)
P mime	0.12	2021-09-28	[?]	RSPM (R 4.4.0)
P miniUI	0.1.1.1	2018-05-18	[?]	RSPM (R 4.4.0)
P munsell	0.5.1	2024-04-01	[?]	RSPM (R 4.4.0)
P nlme	3.1-164	2023-11-27	[?]	CRAN (R 4.4.0)
P pillar	1.9.0	2023-03-22	[?]	RSPM (R 4.4.0)

P pkgbuild	1.4.4	2024-03-17	[?]	RSPM	(R 4.4.0)
P pkgconfig	2.0.3	2019-09-22	[?]	RSPM	(R 4.4.0)
P pkgload	1.3.4	2024-01-16	[?]	RSPM	(R 4.4.0)
P profvis	0.3.8	2023-05-02	[?]	RSPM	(R 4.4.0)
P promises	1.3.0	2024-04-05	[?]	RSPM	(R 4.4.0)
P purrr	1.0.2	2023-08-10	[?]	RSPM	(R 4.4.0)
P qs	* 0.26.3	2024-05-16	[?]	CRAN	(R 4.4.0)
P R6	2.5.1	2021-08-19	[?]	RSPM	(R 4.4.0)
P ragg	1.3.2	2024-05-15	[?]	RSPM	(R 4.4.0)
P RApiSerialize	0.1.3	2024-05-14	[?]	CRAN	(R 4.4.0)
P rappdirs	0.3.3	2021-01-31	[?]	RSPM	(R 4.4.0)
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P renv	0.17.3	2023-04-06	[?]	CRAN	(R 4.4.0)
P rlang	1.1.3	2024-01-10	[?]	RSPM	(R 4.4.0)
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P scales	1.3.0	2023-11-28	[?]	RSPM	(R 4.4.0)
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P tibble	3.2.1	2023-03-20	[?]	RSPM	(R 4.4.0)
P tidyr	* 1.3.1	2024-01-24	[?]	RSPM	(R 4.4.0)
P tidyselect	1.2.1	2024-03-11	[?]	RSPM	(R 4.4.0)
P timechange	0.3.0	2024-01-18	[?]	RSPM	(R 4.4.0)
P tzdb	0.4.0	2023-05-12	[?]	RSPM	(R 4.4.0)
P urlchecker	1.0.1	2021-11-30	[?]	RSPM	(R 4.4.0)
P usethis	2.2.3	2024-02-19	[?]	RSPM	(R 4.4.0)
P utf8	1.2.4	2023-10-22	[?]	RSPM	(R 4.4.0)
P vctrs	0.6.5	2023-12-01	[?]	RSPM	(R 4.4.0)
P vroom	1.6.5	2023-12-05	[?]	RSPM	(R 4.4.0)
P withr	3.0.0	2024-01-16	[?]	RSPM	(R 4.4.0)
P xfun	0.44	2024-05-15	[?]	RSPM	(R 4.4.0)
P xml2	1.3.6	2023-12-04	[?]	RSPM	(R 4.4.0)
P xtable	1.8-4	2019-04-21	[?]	RSPM	(R 4.4.0)
P yaml	2.3.8	2023-12-11	[?]	RSPM	(R 4.4.0)

[1] /Users/elliottgould/Documents/GitHub/VEFMAP_VEG_Stage6/systemfonts/renv/library/R-4.4/aarch64-apple-darwin20

[2] /Users/elliottgould/Library/Caches/org.R-project.R/R/renv/sandbox/R-4.4/aarch64-apple-darwin20/84ba8b13

P — Loaded and on-disk path mismatch.

Appendix E

Table E1. Number of species recorded in each waterway by vegetation group.

Waterway	Aquatic (exotic)	Aquatic (native)	Emergent (exotic)	Emergent (native)	Riparian (exotic)	Riparian (native)	Terrestrial (exotic)	Terrestrial (native)	Total (native)	Total (exotic)
Campaspe	0	9	0	7	8	26	89	53	95	97
Glenelg	1	9	1	9	13	33	66	39	90	81
Wannon	0	2	0	3	0	14	11	10	29	11
Loddon	0	0	0	0	1	10	15	13	23	16
Tullaroop	0	2	0	4	4	7	40	7	20	44
Twelve Mile	0	2	0	1	4	14	19	10	27	23
Moorabool	1	10	0	6	5	28	63	36	80	69
Sutherland	0	1	0	3	3	4	23	4	12	26
Macalister	0	1	0	5	6	13	40	10	29	46
Thomson	0	2	0	3	8	14	42	4	23	50
Burnt	0	5	0	2	4	25	48	41	73	52
McKenzie	0	3	0	1	4	20	40	37	61	44
Mt William	2	4	0	4	7	18	51	20	46	60
Watts	0	1	0	0	5	10	27	9	20	32
Yarra	2	2	0	1	11	23	72	30	56	85

Table E2. Mean percentage cover for a sub-transect in each waterway by vegetation group.

Waterway	Aquatic (exotic)	Aquatic (native)	Emergent (exotic)	Emergent (native)	Riparian (exotic)	Riparian (native)	Terrestrial (exotic)	Terrestrial (native)
Campaspe	0	0.4	0	5.3	5.3	16.3	24.1	10.9
Glenelg	0	8.5	0.4	6.2	3.5	17.4	24.5	15.7
Wannon	0	4	0	19.8	0	49.1	6.6	37.2
Loddon	0	0	0	0	0	30	3.6	15.6
Tullaroop	0	0.6	0	23.7	3.9	2.9	31.7	0.5
Twelve Mile	0	2.5	0	0	21.2	38.4	8.7	3.6
Moorabool	0	6.1	0	12.9	2.9	10.5	28.4	26.4
Sutherland	0	0.6	0	10.8	3.2	2.3	62.4	0.7
Macalister	0	0.2	0	3.6	18.1	28.3	30.7	3.5
Thomson	0	0.3	0	1.3	32.4	16.2	47.3	4.6
Burnt	0	3.3	0	2.6	2.3	35.3	29	5.9
McKenzie	0	2.6	0	0	1.2	23.6	22.5	19.3
Mt William	0.1	6.7	0	8.8	5.4	27.6	48.8	2.7
Watts	0	0.9	0	0	17.8	17.4	27.4	19.5
Yarra	0	0.5	0	0.3	17	21	46.3	13.2

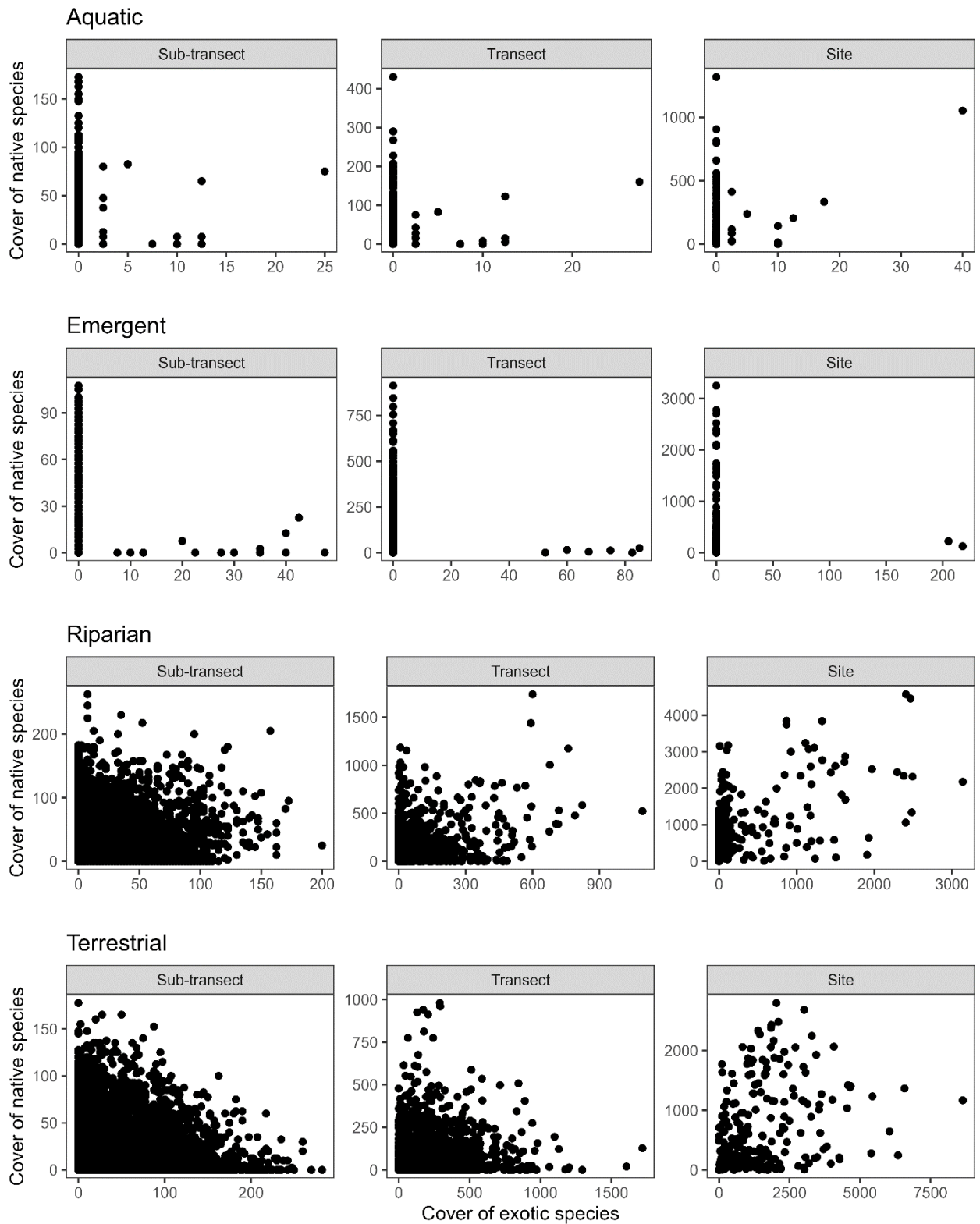


Figure E1. The summed percentage cover of native versus exotic species in the four functional groups within each sub-transect, transect, and site using all survey data.

Appendix F

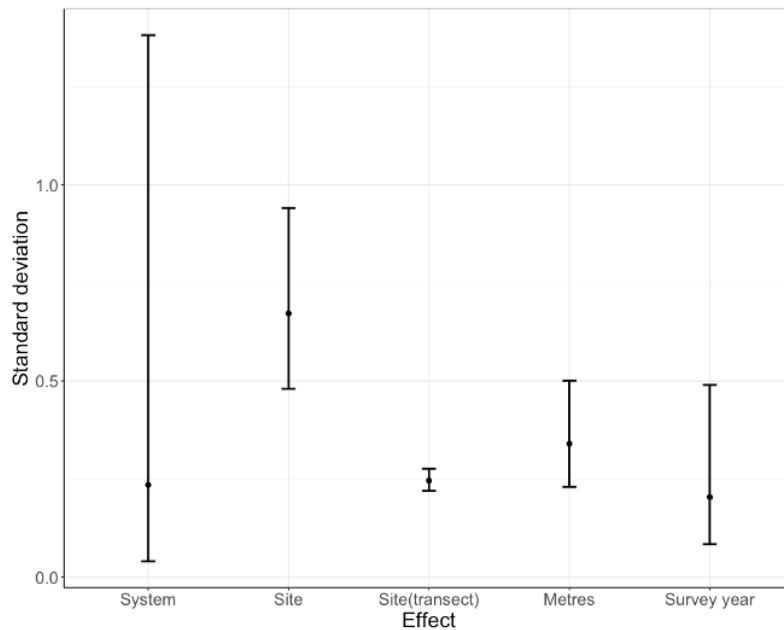


Figure F1. Estimated standard deviations of random intercepts for different factors included in the analysis of vegetation cover as a function of flow regimes (days above baseflow and spring-fresh levels).

Points are mean estimated standard deviations and bars bound 95% confidence intervals. Standard deviations of random effects reflect the amount of variation among levels of each random factor, with high values highlighting the level/s at which variation is most pronounced.

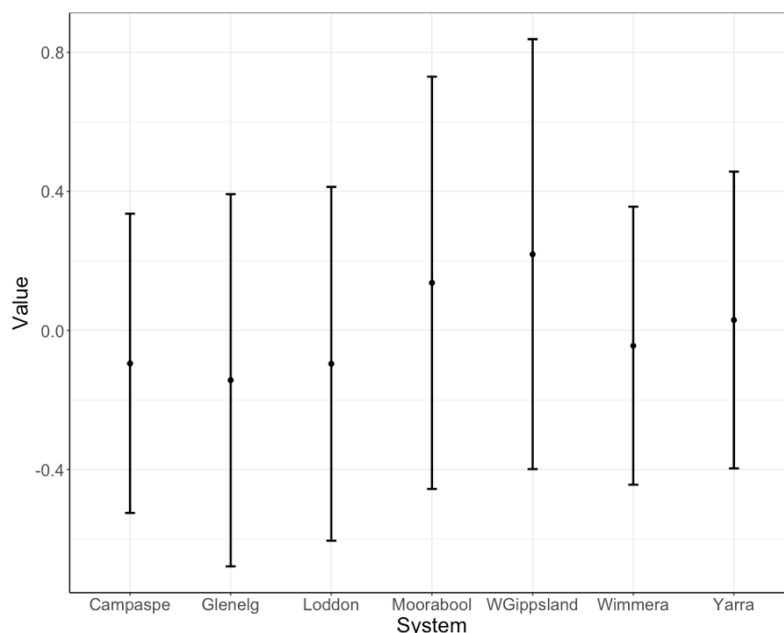


Figure F2. Estimated random intercepts for different systems included in the analysis of vegetation cover as a function of flow regimes (days above baseflow and spring-fresh levels).

Points are mean estimated intercepts on the link scale and bars bound 95% confidence intervals. Random intercepts reflect the deviation of each level (here, different systems) from the average value over all levels. Higher values indicate generally higher-than-average cover and low values the opposite. Systems may include multiple waterways (see main text for sampling details).

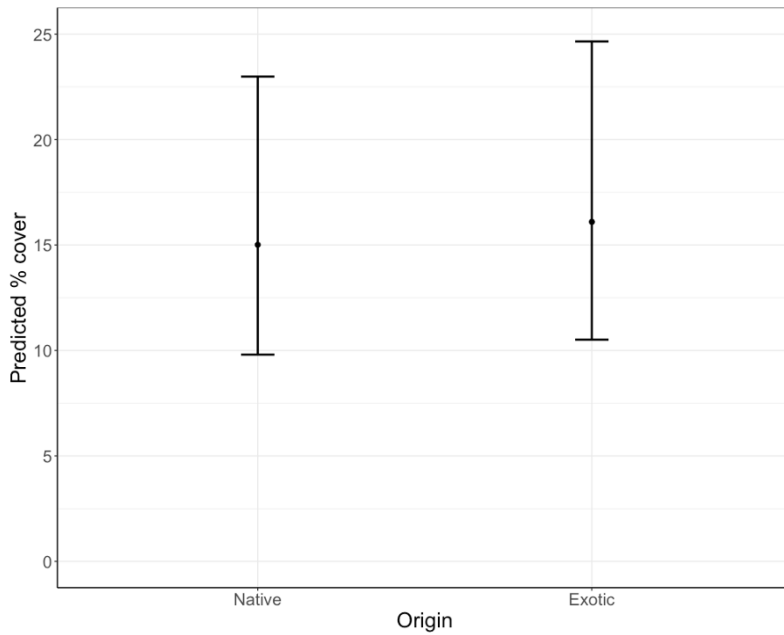


Figure F3. Predicted associations between vegetation cover and species origin from the analysis of vegetation cover as a function of flow regimes.

Solid points are mean estimates of cover for each functional group and error bars bound 95% confidence intervals. Note that the y-axis is truncated to a maximum value of 20%, which reflects the relatively low values of average cover given many zeros in the data set.

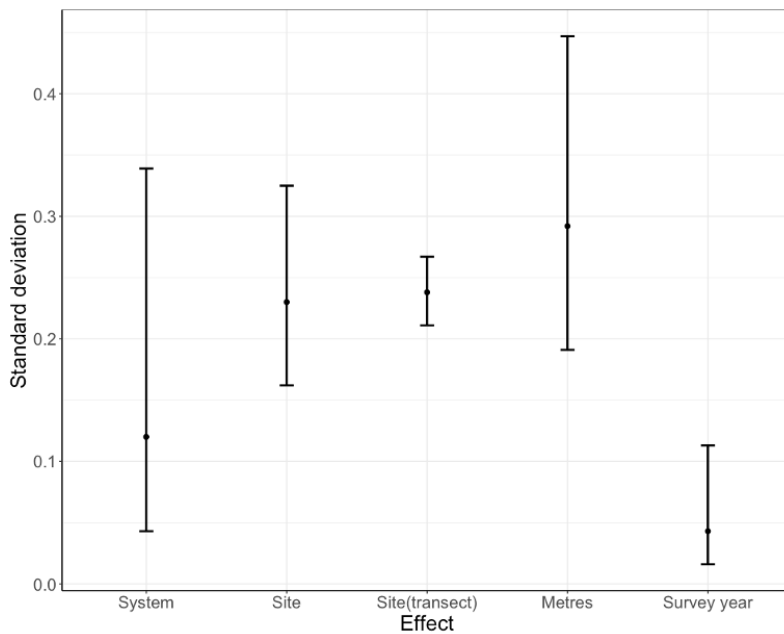


Figure F4. Estimated standard deviations of random intercepts for different factors included in the analysis of vegetation cover as a function of flow events (spring and summer freshes). Points are mean estimated standard deviations and bars bound 95% confidence intervals.

Standard deviations of random effects reflect the amount of variation among levels of each random factor, with high values highlighting the level/s at which variation is most pronounced.

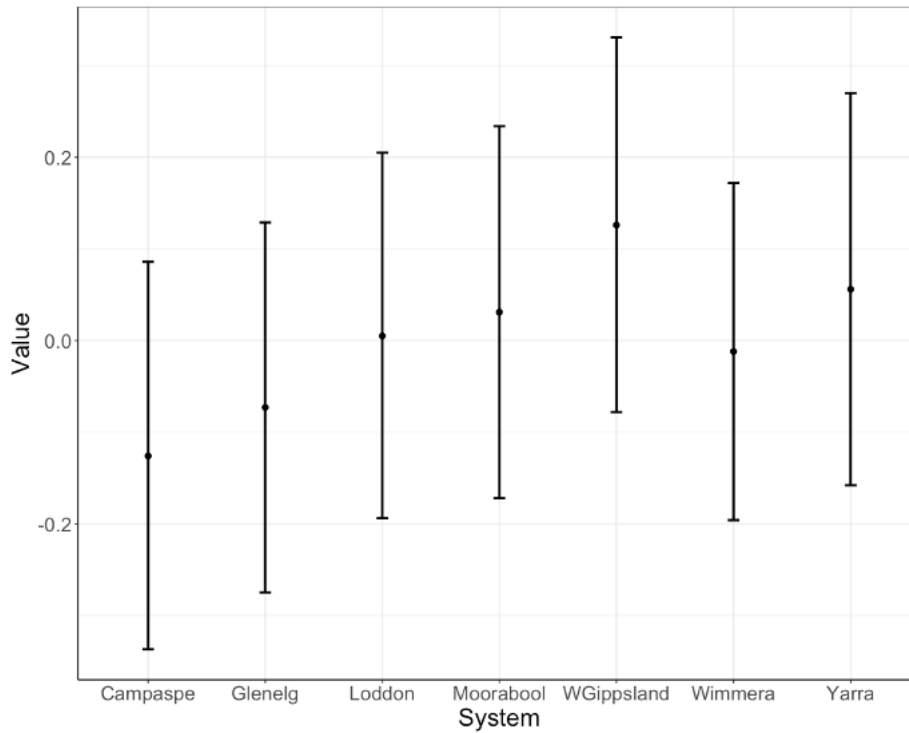


Figure F5. Estimated random intercepts for different systems included in the analysis of vegetation cover as a function of flow events (spring and summer freshes).

Points are mean estimated intercepts on the link scale and bars bound 95% confidence intervals. Random intercepts reflect the deviation of each level (here, different systems) from the average value over all levels. Higher values indicate generally higher-than-average cover and low values the opposite. Systems may include multiple waterways (see main text for sampling details).

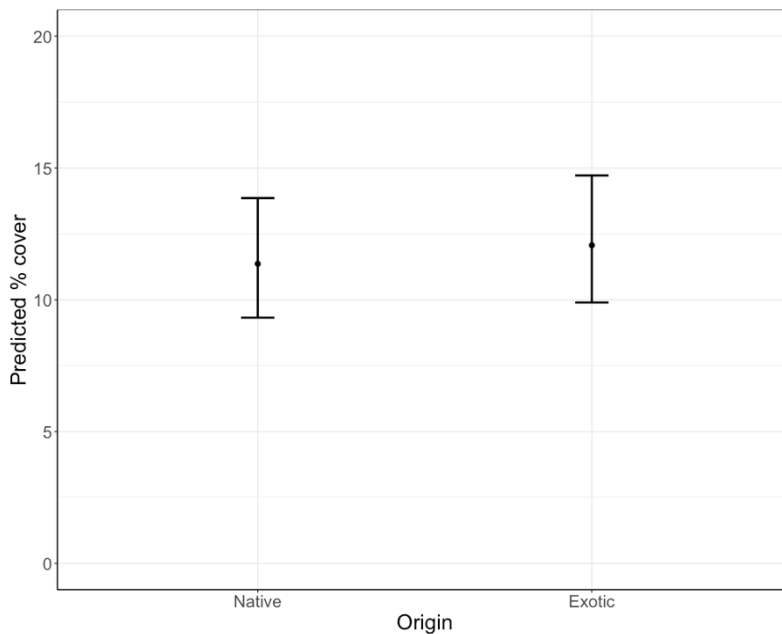


Figure F6. Predicted associations between vegetation cover and species origin from the analysis of vegetation cover as a function of flow events.

Solid points are mean estimates of cover for each functional group and error bars bound 95% confidence intervals. Note that the y-axis is truncated to a maximum value of 20%, which reflects the relatively low values of average cover given many zeros in the data set.

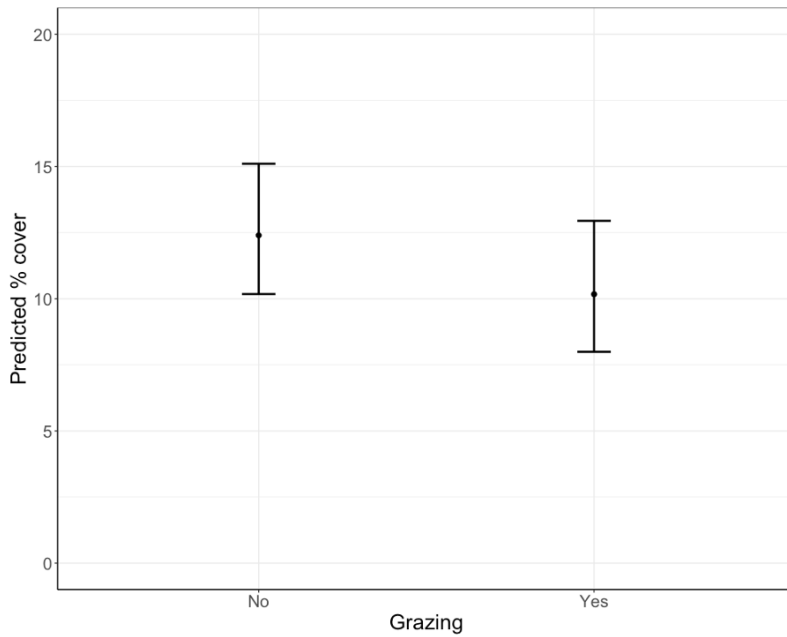


Figure F7. Predicted associations between vegetation cover and the occurrence of livestock grazing from the analysis of vegetation cover as a function of flow events (Yes: grazing occurs; No: no grazing occurs).

Solid points are mean estimates of cover for each functional group and error bars bound 95% confidence intervals. Note that the y-axis is truncated to a maximum value of 20%, which reflects the relatively low values of average cover given many zeros in the data set.

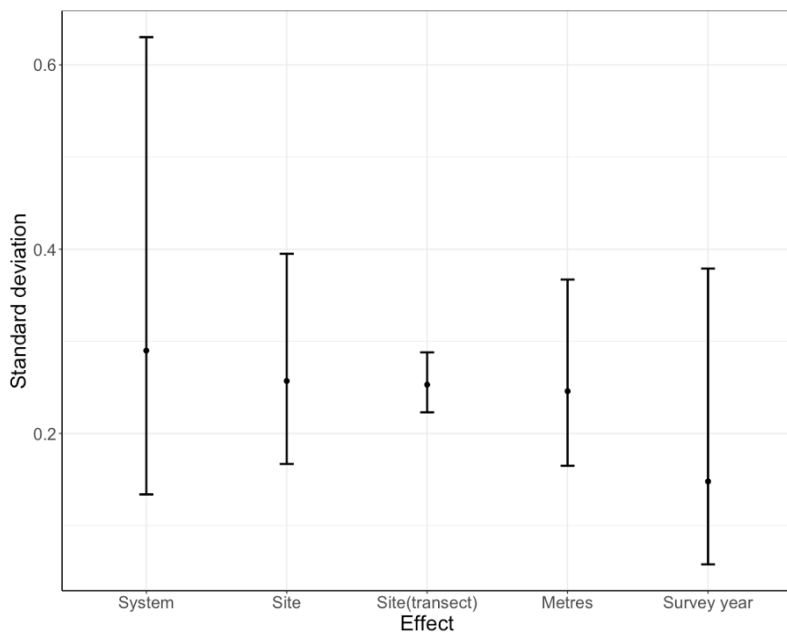


Figure F8. Estimated standard deviations of random intercepts for different factors included in the analysis of vegetation species richness.

Points are mean estimated standard deviations and bars bound 95% confidence intervals. Standard deviations of random effects reflect the amount of variation among levels of each random factor, with high values highlighting the level/s at which variation is most pronounced.

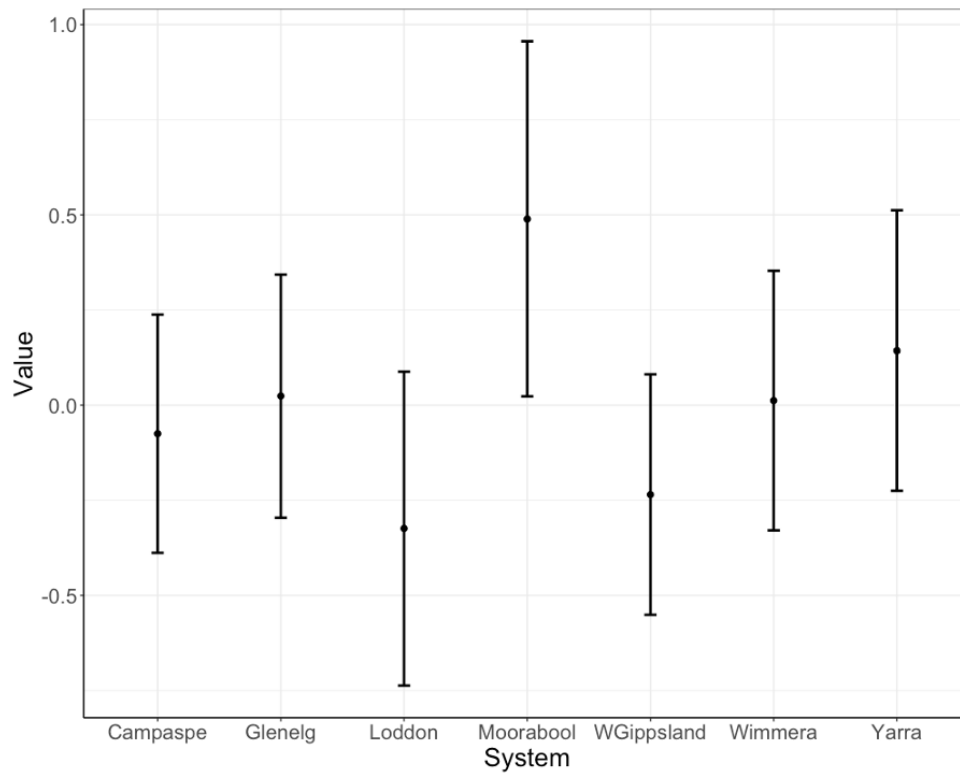


Figure F9. Estimated random intercepts for different systems included in the analysis of vegetation species richness.

Points are mean estimated intercepts on the link scale and bars bound 95% confidence intervals. Random intercepts reflect the deviation of each level (here, different systems) from the average value over all levels. Higher values indicate generally higher-than-average species richness and low values the opposite. Systems may include multiple waterways (see main text for sampling details).

Table F1. Parameter estimates for vegetation richness model. SE: standard error, SD: standard deviation, std: standardised, wpfg: water plant functional group.

Parameter	Coefficient	SE	95% CI	z	p
Fixed Effects					
(Intercept)	-3.88	0.26	(-4.39, -3.37)	-14.87	< .001
days above baseflow std	0.02	0.08	(-0.13, 0.17)	0.24	0.810
wpfg (Emergent)	1.20	0.23	(0.74, 1.65)	5.15	< .001
wpfg (Riparian)	2.94	0.20	(2.54, 3.33)	14.57	< .001
wpfg (Terrestrial)	3.87	0.20	(3.48, 4.26)	19.41	< .001
days above springfresh std	0.37	0.11	(0.15, 0.58)	3.33	< .001
origin (native)	0.28	0.01	(0.25, 0.31)	18.94	< .001
origin (unknown)	-3.94	0.10	(-4.13, -3.75)	-40.41	< .001
days above baseflow std^2	-0.01	0.02	(-0.05, 0.03)	-0.70	0.485
days above springfresh std^2	-0.07	0.03	(-0.13, -0.02)	-2.61	0.009
zone (baseflow_to_springfresh)	0.26	0.24	(-0.22, 0.73)	1.05	0.295
zone (below_baseflow)	2.35	0.24	(1.89, 2.82)	9.86	< .001
period (after_summer)	0.06	0.27	(-0.47, 0.60)	0.24	0.813
period (before_spring)	-0.10	0.29	(-0.68, 0.47)	-0.36	0.721
grazing (Y)	-0.22	0.10	(-0.42, -0.03)	-2.24	0.025
days above baseflow std × wpfg (Emergent)	-0.11	0.07	(-0.24, 0.02)	-1.60	0.109
days above baseflow std × wpfg (Riparian)	-0.22	0.06	(-0.33, -0.10)	-3.77	< .001
days above baseflow std × wpfg (Terrestrial)	-0.13	0.06	(-0.24, -0.02)	-2.23	0.026
wpfg (Emergent) × days above springfresh std	0.09	0.06	(-0.04, 0.21)	1.38	0.167
wpfg (Riparian) × days above springfresh std	0.03	0.06	(-0.08, 0.13)	0.47	0.635
wpfg (Terrestrial) × days above springfresh std	-0.08	0.06	(-0.19, 0.02)	-1.51	0.130
wpfg (Emergent) × zone (baseflow_to_springfresh)	0.43	0.28	(-0.11, 0.98)	1.55	0.121
wpfg (Riparian) × zone (baseflow_to_springfresh)	0.49	0.25	(7.74e-03, 0.98)	1.99	0.046
wpfg (Terrestrial) × zone (baseflow_to_springfresh)	-0.72	0.25	(-1.20, -0.23)	-2.91	0.004
wpfg (Emergent) × zone (below_baseflow)	-0.62	0.29	(-1.18, -0.06)	-2.18	0.030
wpfg (Riparian) × zone (below_baseflow)	-1.67	0.25	(-2.16, -1.19)	-6.76	< .001
wpfg (Terrestrial) × zone (below_baseflow)	-3.61	0.26	(-4.12, -3.10)	-13.93	< .001
zone (baseflow_to_springfresh) × period (after_summer)	0.08	0.33	(-0.57, 0.74)	0.25	0.805
zone (below_baseflow) × period (after_summer)	-0.13	0.33	(-0.78, 0.52)	-0.38	0.703
zone (baseflow_to_springfresh) × period (before_spring)	0.55	0.35	(-0.13, 1.24)	1.59	0.113
zone (below_baseflow) × period (before_spring)	0.18	0.35	(-0.50, 0.86)	0.52	0.606
wpfg (Emergent) × period (after_summer)	-0.01	0.32	(-0.64, 0.62)	-0.04	0.972
wpfg (Riparian) × period (after_summer)	-0.09	0.28	(-0.64, 0.46)	-0.31	0.755

wpfg (Terrestrial) × period (after_summer)	-0.33	0.28	(-0.87, 0.22)	-1.18	0.240
wpfg (Emergent) × period (before_spring)	0.04	0.34	(-0.64, 0.71)	0.10	0.917
wpfg (Riparian) × period (before_spring)	-0.11	0.30	(-0.69, 0.48)	-0.35	0.725
wpfg (Terrestrial) × period (before_spring)	0.21	0.30	(-0.37, 0.79)	0.71	0.477
(wpfg (Emergent) × zone (baseflow_to_springfresh)) × period (after_summer)	-0.12	0.39	(-0.88, 0.63)	-0.32	0.747
(wpfg (Riparian) × zone (baseflow_to_springfresh)) × period (after_summer)	-0.08	0.34	(-0.74, 0.59)	-0.23	0.819
(wpfg (Terrestrial) × zone (baseflow_to_springfresh)) × period (after_summer)	0.07	0.34	(-0.59, 0.74)	0.22	0.830
(wpfg (Emergent) × zone (below_baseflow)) × period (after_summer)	-1.46e-03	0.40	(-0.79, 0.79)	- 3.63e- 03	0.997
(wpfg (Riparian) × zone (below_baseflow)) × period (after_summer)	0.21	0.35	(-0.47, 0.89)	0.60	0.546
(wpfg (Terrestrial) × zone (below_baseflow)) × period (after_summer)	0.47	0.36	(-0.24, 1.18)	1.29	0.196
(wpfg (Emergent) × zone (baseflow_to_springfresh)) × period (before_spring)	-0.78	0.41	(-1.58, 0.01)	-1.93	0.054
(wpfg (Riparian) × zone (baseflow_to_springfresh)) × period (before_spring)	-0.56	0.36	(-1.26, 0.13)	-1.59	0.113
(wpfg (Terrestrial) × zone (baseflow_to_springfresh)) × period (before_spring)	-0.56	0.35	(-1.25, 0.13)	-1.60	0.110
(wpfg (Emergent) × zone (below_baseflow)) × period (before_spring)	-0.34	0.43	(-1.18, 0.49)	-0.81	0.420
(wpfg (Riparian) × zone (below_baseflow)) × period (before_spring)	-0.11	0.37	(-0.83, 0.61)	-0.30	0.768
(wpfg (Terrestrial) × zone (below_baseflow)) × period (before_spring)	-0.55	0.39	(-1.31, 0.21)	-1.41	0.158
Random Effects					
SD (Intercept: transect:site)	0.25				
SD (Intercept: site)	0.26				
SD (Intercept: metres)	0.25				
SD (Intercept: survey_year)	0.15				
SD (Intercept: system)	0.29				

Table F2. Parameter estimates for vegetation cover flow-regime model. SE: standard error, SD: standard deviation, std: standardised, wpfg: water plant functional group.

Parameter	Coefficient	SE	95% CI	z	p
Fixed Effects (Count Model)					
(Intercept)	1.30	0.22	(0.86, 1.73)	5.88	< .001
log hits tm1	0.49	2.75e-03	(0.49, 0.50)	178.72	< .001
days above baseflow std	-0.06	0.03	(-0.12, 2.59e-03)	-1.88	0.060
wpfg (Emergent)	0.48	0.04	(0.40, 0.56)	12.01	< .001
wpfg (Riparian)	0.30	0.04	(0.23, 0.38)	7.90	< .001
wpfg (Terrestrial)	0.39	0.04	(0.32, 0.47)	10.10	< .001
days above springfresh std	0.24	0.07	(0.10, 0.39)	3.25	0.001
days above baseflow std ²	0.05	0.02	(0.02, 0.08)	3.35	< .001
days above springfresh std ²	-0.16	0.02	(-0.20, -0.12)	-7.61	< .001
origin (native)	-0.07	6.65e-03	(-0.08, -0.06)	-10.48	< .001
origin (unknown)	-4.49	0.07	(-4.63, -4.36)	-67.13	< .001
days above baseflow std × wpfg (Emergent)	-0.11	0.03	(-0.16, -0.06)	-3.97	< .001
days above baseflow std × wpfg (Riparian)	-0.22	0.02	(-0.26, -0.17)	-9.04	< .001
days above baseflow std × wpfg (Terrestrial)	-0.23	0.02	(-0.27, -0.18)	-9.47	< .001
wpfg (Emergent) × days above springfresh std	0.49	0.06	(0.37, 0.62)	7.80	< .001
wpfg (Riparian) × days above springfresh std	0.52	0.06	(0.40, 0.63)	9.05	< .001
wpfg (Terrestrial) × days above springfresh std	0.51	0.06	(0.40, 0.62)	9.02	< .001
wpfg (Emergent) × days above baseflow std ²	-0.03	0.02	(-0.06, 4.15e-04)	-1.93	0.053
wpfg (Riparian) × days above baseflow std ²	9.48e-04	0.01	(-0.03, 0.03)	0.07	0.945
wpfg (Terrestrial) × days above baseflow std ²	-3.20e-03	0.01	(-0.03, 0.02)	-0.23	0.817
wpfg (Emergent) × days above springfresh std ²	-0.13	0.02	(-0.16, -0.09)	-7.32	< .001
wpfg (Riparian) × days above springfresh std ²	-0.09	0.02	(-0.12, -0.06)	-5.70	< .001
wpfg (Terrestrial) × days above springfresh std ²	-0.11	0.02	(-0.14, -0.07)	-6.62	< .001
Fixed Effects (Zero-Inflation Component)					
(Intercept)	3.09	0.06	(2.96, 3.21)	48.92	< .001
wpfg (Emergent)	-1.35	0.07	(-1.50, -1.21)	-18.48	< .001

Parameter	Coefficient	SE	95% CI	z	p
wpfg (Riparian)	-2.73	0.07	(-2.86, -2.60)	-41.30	< .001
wpfg (Terrestrial)	-3.12	0.07	(-3.24, -2.99)	-47.16	< .001
Random Effects Variances					
SD (Intercept: transect:site)	0.25				
SD (Intercept: site)	0.67				
SD (Intercept: metres)	0.34				
SD (Intercept: survey_year)	0.20				
SD (Intercept: system)	0.23				

Table F3. Parameter estimates for vegetation cover flow-event model. SE: standard error, SD: standard deviation, std: standardised, wpfg: water plant functional group.

Parameter	Coefficient	SE	95% CI	z	p
Fixed Effects (Count Model)					
(Intercept)	1.28	0.13	(1.03, 1.54)	9.80	< .001
log hits tm1	0.52	2.86e-03	(0.51, 0.52)	180.40	< .001
zone (baseflow_to_springfresh)	-0.60	0.11	(-0.83, -0.38)	-5.26	< .001
zone (below_baseflow)	-7.81e-03	0.10	(-0.20, 0.18)	-0.08	0.935
wpfg (Emergent)	0.13	0.09	(-0.05, 0.31)	1.37	0.172
wpfg (Riparian)	0.17	0.09	(5.80e-03, 0.34)	2.03	0.043
wpfg (Terrestrial)	0.12	0.09	(-0.05, 0.28)	1.36	0.174
period (after_summer)	-0.26	0.12	(-0.49, -0.03)	-2.24	0.025
period (before_spring)	2.50e-03	0.87	(-1.70, 1.71)	2.87e-03	0.998
origin (native)	-0.06	6.69e-03	(-0.07, -0.05)	-8.97	< .001
grazing (Y)	-0.20	0.08	(-0.36, -0.03)	-2.35	0.019
zone (baseflow_to_springfresh) × wpfg (Emergent)	0.86	0.12	(0.62, 1.09)	7.06	< .001
zone (below_baseflow) × wpfg (Emergent)	0.16	0.11	(-0.05, 0.37)	1.53	0.127
zone (baseflow_to_springfresh) × wpfg (Riparian)	0.64	0.11	(0.42, 0.87)	5.61	< .001
zone (below_baseflow) × wpfg (Riparian)	-0.04	0.10	(-0.23, 0.16)	-0.37	0.708
zone (baseflow_to_springfresh) × wpfg (Terrestrial)	0.64	0.11	(0.41, 0.86)	5.56	< .001
zone (below_baseflow) × wpfg (Terrestrial)	-0.09	0.11	(-0.29, 0.12)	-0.82	0.411
zone (baseflow_to_springfresh) × period (after_summer)	1.13	0.15	(0.84, 1.42)	7.66	< .001

Parameter	Coefficient	SE	95% CI	z	p
zone (below_baseflow) × period (after_summer)	0.25	0.13	(-0.01, 0.51)	1.87	0.062
zone (baseflow_to_springfresh) × period (before_spring)	-1.64	0.93	(-3.46, 0.18)	-1.77	0.077
zone (below_baseflow) × period (before_spring)	-0.18	0.87	(-1.90, 1.53)	-0.21	0.835
wpfg (Emergent) × period (after_summer)	0.07	0.13	(-0.19, 0.32)	0.52	0.604
wpfg (Riparian) × period (after_summer)	0.01	0.12	(-0.22, 0.25)	0.11	0.912
wpfg (Terrestrial) × period (after_summer)	0.12	0.12	(-0.12, 0.35)	0.97	0.330
wpfg (Emergent) × period (before_spring)	-2.49	0.93	(-4.31, -0.66)	-2.67	0.008
wpfg (Riparian) × period (before_spring)	-0.31	0.87	(-2.02, 1.39)	-0.36	0.719
wpfg (Terrestrial) × period (before_spring)	0.65	0.87	(-1.05, 2.35)	0.75	0.455
(zone (baseflow_to_springfresh) × wpfg (Emergent)) × period (after_summer)	-1.08	0.16	(-1.39, -0.77)	-6.82	< .001
(zone (below_baseflow) × wpfg (Emergent)) × period (after_summer)	-0.13	0.15	(-0.42, 0.16)	-0.87	0.387
(zone (baseflow_to_springfresh) × wpfg (Riparian)) × period (after_summer)	-0.99	0.15	(-1.28, -0.69)	-6.61	< .001
(zone (below_baseflow) × wpfg (Riparian)) × period (after_summer)	0.07	0.14	(-0.20, 0.34)	0.49	0.622
(zone (baseflow_to_springfresh) × wpfg (Terrestrial)) × period (after_summer)	-1.02	0.15	(-1.31, -0.73)	-6.86	< .001
(zone (below_baseflow) × wpfg (Terrestrial)) × period (after_summer)	-0.28	0.15	(-0.57, 3.19e-03)	-1.94	0.053
(zone (baseflow_to_springfresh) × wpfg (Emergent)) × period (before_spring)	3.80	0.99	(1.86, 5.73)	3.85	< .001
(zone (below_baseflow) × wpfg (Emergent)) × period (before_spring)	2.32	0.94	(0.48, 4.15)	2.47	0.014
(zone (baseflow_to_springfresh) × wpfg (Riparian)) × period (before_spring)	1.79	0.93	(-0.02, 3.61)	1.93	0.053
(zone (below_baseflow) × wpfg (Riparian)) × period (before_spring)	0.45	0.88	(-1.27, 2.16)	0.51	0.610
(zone (baseflow_to_springfresh) × wpfg (Terrestrial)) × period (before_spring)	1.17	0.93	(-0.65, 2.99)	1.26	0.208
(zone (below_baseflow) × wpfg (Terrestrial)) × period (before_spring)	-1.68	0.89	(-3.42, 0.06)	-1.89	0.058
Fixed Effects (Zero-Inflation Component)					
(Intercept)	2.97	0.07	(2.84, 3.10)	45.68	< .001
wpfg (Emergent)	-1.31	0.08	(-1.46, -1.17)	-17.47	< .001

Parameter	Coefficient	SE	95% CI	z	p
wpfg (Riparian)	-2.66	0.07	(-2.79, -2.52)	-38.99	< .001
wpfg (Terrestrial)	-3.07	0.07	(-3.20, -2.94)	-45.04	< .001
Random Effects Variances					
SD (Intercept: transect:site)	0.24				
SD (Intercept: site)	0.23				
SD (Intercept: metres)	0.29				
SD (Intercept: survey_year)	0.04				
SD (Intercept: system)	0.12				