



Drying microclimates threaten persistence of natural and translocated populations of threatened frogs

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Abstract

Defining species habitat requirements is essential for effective conservation management through revealing agents of population decline and identifying critical habitat for conservation actions, such as translocations. Here we studied the habitat-associations of two threatened terrestrial-breeding frog species from southwestern Australia, *Geocrinia alba* and *Geocrinia vitellina*, to investigate if fine-scale habitat variables explain why populations occur in discrete patches, why *G. alba* is declining, and why translocation attempts have had mixed outcomes. We compared habitat variables at sites where the species are present, to variables at immediately adjacent sites where frogs are absent, and at sites where *G. alba* is locally extinct. Dry season soil moisture was the most important predictor of frog abundance for both species, and explained why *G. alba* had become extinct from some areas. Sites where *G. alba* were present were also positively associated with moss cover, and negatively with bare ground and soil conductivity. Modelling frog abundance based exclusively on dry season soil moisture predicted recent translocation successes with high accuracy. Hence, considering dry season soil moisture when selecting future translocation sites should increase the probability of population establishment. We propose that a regional drying trend is the most likely cause for *G. alba* declines and that both species are at risk of further habitat and range contraction due to further projected regional declines in rainfall and groundwater levels. More broadly, our study highlights that conservation areas in drying climates may not provide adequate protection and may require interventions to preserve critical habitat.

Keywords Amphibian conservation · Climate · Habitat · Population decline · Soil moisture · *Geocrinia*

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Introduction

Defining the environmental factors that determine species distributions is invaluable for effective species conservation and management. Population declines due to environmental change are prevalent worldwide (Pereira et al. 2012) and so understanding the environmental limits of species is critical for predicting future biodiversity. Species-habitat associations can be used to identify threatening processes and enable management actions that may halt or reverse declines, and ultimately prevent species extinctions (Caughley 1994). For example, defining species' habitat requirements can improve recovery actions, such as conservation translocations (Morris et al. 2015; McCulloch and Norris 2001). A key challenge for recovering threatened species is identification of suitable reintroduction sites, and low habitat suitability or a lack of consideration of micro-habitat are prime reasons for translocation failure (Germano and Bishop 2009; Bennett et al. 2013). Yet despite potential benefits to conservation, the fine-scale habitat associations that determine a species occurrence are often unknown.

The availability of detailed information on environmental requirements is particularly important for amphibians. Over 40% of amphibian species are threatened (IUCN Red List 2020), and the causes of declines and local extinctions are not universally known (Stuart et al. 2004). Being ectothermic vertebrates, the body temperature of amphibians is largely determined by their microenvironment, and their permeable skin and unshelled eggs make them especially sensitive to water loss and pollutants (Hillman et al. 2009). As a consequence, amphibians are highly sensitive to changes in habitat and are more vulnerable to changes in climate than other vertebrates (Blaustein et al. 2010; Walls et al. 2013; Rolland et al. 2018). Detailed knowledge of species-habitat associations can hence improve our understanding of how the amphibians may respond to future changes, such as hotter and/or drier climates, and can aid recovery actions such as translocations (Li et al. 2013).

White- and orange-bellied frogs (*Geocrinia alba* and *Geocrinia vitellina*; Wardell-Johnson and Roberts, 1989) are terrestrial-breeding species with highly restricted distributions in south-western Australia (Fig. 1). The species are allopatric with at least 6 km between their range edges and occur in naturally fragmented sub-populations clustered along drainage lines and headwater streams. *Geocrinia alba* is Critically Endangered, with over half the known populations going extinct in recent decades, and extant populations continuing to decline (TSSC 2019). Considerable habitat clearance and land-use change in the region has been attributed to earlier extinction events, but more recently populations are disappearing in areas within conservation reserves that are apparently undisturbed (Page et al. 2018). *Geocrinia vitellina* is known from a very small area (< 6 km²) and is listed as Vulnerable, however populations have remained relatively stable over time and are all within a contiguous national park (Department of Parks and Wildlife 2015). The causes of the patchy distributions of both species are unclear, as is the reason why some *G. alba* populations are declining in areas with apparently intact habitat.

Conservation translocations of head-started juvenile *Geocrinia* began in the 2000's and aimed to increase the number of sub-populations, but have had varying outcomes (Department of Parks and Wildlife 2015). One translocation site for *G. alba* has become a large self-sustaining population, whilst few or no calling males have been detected at other sites in the years following translocation. Juvenile frogs were all released into suitable looking riparian habitats near existing populations during spring and involved the release of a comparable number of individuals. Why some translocations have been more

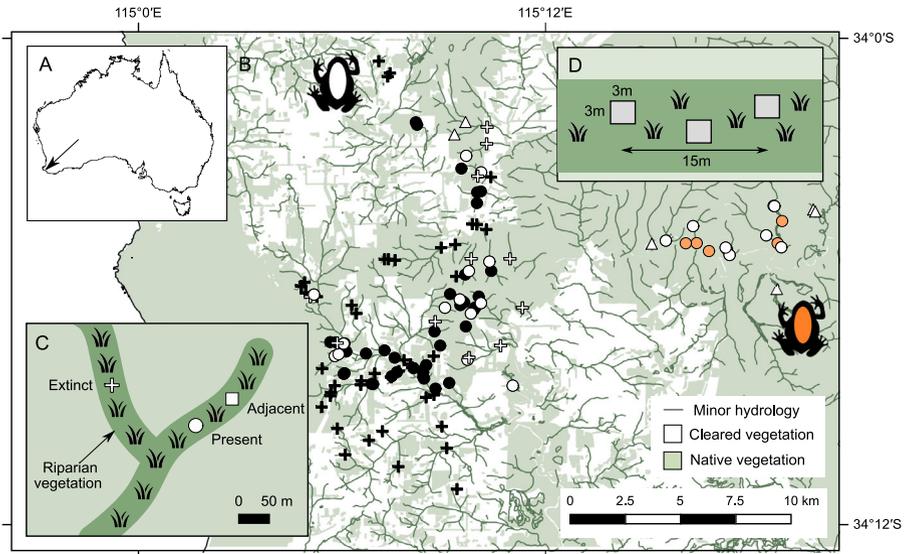


Fig. 1 Map of **a** the study area in southwest Western Australia **b** showing present (circles), extinct (crosses) and translocated (triangles) *Geocrinia alba* (black) and *Geocrinia vitellina* (orange) populations and sites sampled in this study (white symbols), **c** an example of the sampling design along drainage lines, showing sampled present site where frogs occur (circle), an immediately adjacent site (square) where frogs are absent, and a nearby extinct site (cross), and **d** an example of sampling quadrats at a site. Population locations and hydrology layers provided by Western Australian Department of Biodiversity Conservation and Attractions

successful than others has not been determined, and not knowing what makes a translocation site successful is currently constraining conservation management.

Patterns of population occurrence and persistence for both natural and translocated populations of *G. alba* and *G. vitellina* may relate to fine-scale differences in habitat. Both species exhibit extreme site philopatry, with 90% of individuals moving less than 20 m within and between years (Driscoll 1997). This low dispersal is evident from genetic studies that showed almost no gene flow between populations (Driscoll 1998a). As the species are specialised terrestrial breeders with entirely endotrophic (non-feeding) development (Mitchell 2001; Anstis 2010), their lifecycle is completed in and immediately adjacent to breeding sites. Breeding sites are vulnerable to extremes of flooding and desiccation, as eggs develop in moist soils but cannot be submerged (Wardell-Johnson and Roberts 1993). Hence if a habitat becomes unsuitable, these species have very limited ability to disperse to suitable habitat elsewhere.

Habitat floristics do not distinguish between persisting and extinct populations (Pauli 1999), but habitat structure may play a key role in where frogs occur. Both species live and breed under moss, litter and other vegetation (Driscoll 1998a; Conroy 2001) and therefore the availability of these features is likely important in determining their presence. Likewise, microclimate characteristics of nest sites and soil parameters, such as pH and conductivity, are other factors known to influence amphibian occurrence (Wyman 1988; Mitchell 2002a; Dodd 2010) that could also play a role in determining the distribution of *G. alba* and *G. vitellina*.

In this study our aim was to identify if variation in fine-scale habitat attributes could explain why populations of *G. alba* and *G. vitellina* occur in discrete patches along

seemingly suitable riparian habitat, why *G. alba* is declining and has become absent from some sites whilst *G. vitellina* sites are more stable, and why translocation efforts for both species have had mixed outcomes. Based on current knowledge of the species, we explored four hypotheses to explain their persistence at sites. We hypothesised that frog occurrence, population declines, and translocation success are driven by differences in (1) site hydrology, (2) micro-climate, (3) habitat structure, and/or (4) soil characteristics. We predicted that sites where frogs persist would have higher year-round soil moisture, lower temperature extremes, more ground cover and soils with lower conductivity. We then use our findings on habitat associations based on occupied frog sites and successful translocation sites to provide recommendations for recognising critical habitat, and to identify key threats to the persistence of both species.

Methods

Study area

Geocrinia alba and *Geocrinia vitellina* occur along drainage areas and headwater streams in the Margaret River region in southwest Australia (Fig. 1). Drainage lines typically consist of dense riparian and rhizomatous vegetation with a shrub overstorey (e.g. *Astartea fascicularis*, *Taxandria linearifolia*, *Homalospermum firmum*), surrounded by jarrah (*Eucalyptus marginata*) and marri (*Corymbia calophylla*) forests, and are characterised by shallow water flows in winter (Wardell-Johnson and Roberts 1993). The region has a Mediterranean climate with a distinct wet period from May to October, and a dry period from November to April characterised by very little rainfall and high temperatures (Fig. 2). Cool wet winters and early spring coincide with frog breeding activity, and metamorphosis from terrestrial nest sites occurs from November to January.

Site selection

Geocrinia alba and *G. vitellina* habitats were sampled across four distinct site types, selected with reference to long-term acoustic monitoring records maintained by the Western Australian Department of Biodiversity, Conservation and Attractions (DBCA) (Fig. 1). The *G. alba* and *G. vitellina* monitoring program has been identified as an exemplary monitoring program for threatened frogs in Australia (Scheele and Gillespie 2018), with over 1760 surveys conducted across 150 sites since 1983. Frog populations are monitored annually or biannually, and monitoring includes point counts (where frog abundances are estimated from a single location) at all sites, as well as linear counts at selected sites, which involves counting all individuals and locating the first and last calling male in a breeding chorus along a drainage line. Translocation sites have census monitoring, where all calling individuals are located and recorded.

Present sites ($n = 21$) were locations where populations of *G. alba* or *G. vitellina* have consistently occurred, based on acoustic surveys conducted in spring. Present sites were selected at random from a list of all known present sites, where they were accessible (i.e. property access was permitted) and separated from another present site by at least 500 m. Adjacent sites ($n = 21$) were unoccupied sites along potential riparian habitat for frogs, either 50 m directly upstream or downstream of a present population site.

While typical adult movements between years are less than 20 m, the distance of 50 m is within the dispersal range of these species (Driscoll 1997), and so adjacent sites were

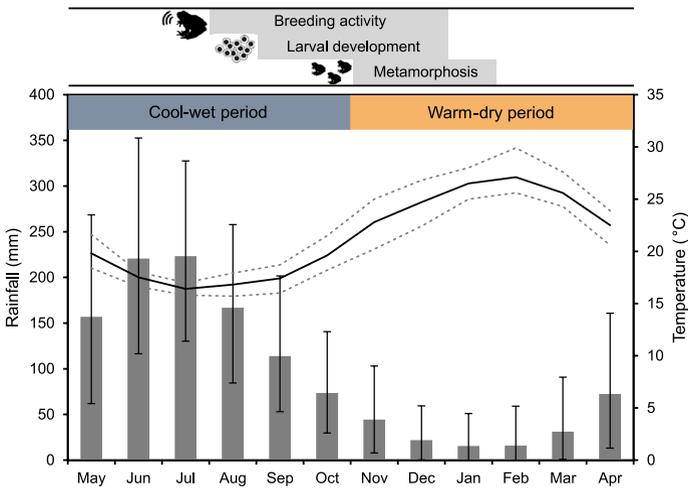


Fig. 2 Monthly averages (with 5% and 95% percentiles) of rainfall (vertical bars) and temperature (solid line) for the Margaret River region, showing distinct cool-wet winter and warm-dry summer periods. The relative timing of breeding activity (calling), larval development and metamorphosis of *G. alba* and *G. vitellina* (based on Conroy 2001; Anstis 2010) are shown by horizontal bars. Climate data are from weather stations at Witchcliffe (temperature from 1999 to 2018) and Forest Grove (rainfall from 1925 to 2018), located approximately 5 and 3 kms from the study area (Bureau of Meteorology 2019)

potentially available to frogs, but could be confirmed as unoccupied due to their proximity to present sites that are routinely surveyed. Upstream or downstream sites were chosen at random (by the flip of a coin), unless only one direction fitted our selection criteria (i.e. there was only unoccupied habitat in one direction). Extinct sites ($n = 11$) were nearby sites where the species had occurred in the past but was not recorded by the DBCA for at least three years prior. Translocation sites ($n = 6$) were sites where head-started juvenile frogs (ranging from 68 to 145 individuals) had been introduced between 2010 and 2017 in locations where they had not been recorded previously (Table S1).

Estimates of frog abundance

We estimated frog abundance from acoustic monitoring data collected by DBCA staff during September–November, when the majority (76–96%) of calling males in a population can be reliably detected (Driscoll 1998b). Frog call surveys were conducted after dusk by experienced DBCA staff, when the maximum number of males' call (Driscoll 1998b). The number of calling males was estimated over a 10-min period, and was considered to be a proxy for frog abundance that reflected relative differences in population size across sites. The number of males calling at a site was allocated to one of six categories (0, 1–4, 5–10, 11–20, 21–50, > 50), and the minimum value in each category was used as the estimate of male abundance for each site. This was the most conservative approach and was more informative than presence-absence data as it distinguished sites with only a few calling males from sites that supported much larger choruses. At translocation sites, frog abundance was estimated using a similar call survey technique but a censusing approach was employed, where the exact number of calling males were identified by marking their positions.

Habitat characteristics

Habitat surveys were conducted towards the end of the warm-dry period, from April 14 to May 2 2018, following pilot surveys to determine the number of quadrats and samples required to quantify characteristics of a site. Habitat variables were sampled in three 3×3 m quadrats at each site. The quadrats were established along a 15 m transect running along the edge of the riparian vegetation (7.5 m apart), and then positioned at a random distance from the edge (into the riparian habitat) (Fig. 1). At each quadrat, fifteen habitat characteristics were measured relating to our four hypotheses; site hydrology, micro-climate, habitat structure, and soil characteristics (Table 1).

Soil moisture (volumetric water content, $\text{m}^3 \text{m}^{-3}$) was measured in the top 6 cm of soil using a hand-held moisture meter (ICT International, MP306). Five measurements were taken, one in each corner and one in the centre of the quadrat, and then were averaged per quadrat. Additional soil moisture measurements were taken at the end of the wet period (December 18–20 2018), but only in the central quadrat. The soil moisture meter was calibrated using soil samples from five sites that included all soil texture categories (see below). Soils were air-dried then wetted to create a range of moisture levels and the meter value was recorded and compared to the actual water content by oven drying the sample at 105°C for 24 h. Bulk density was determined with the core method using the same samples as were used for calibration of the meter.

Table 1 Fine-scale habitat variables used to investigate habitat associations of *G. alba* and *G. vitellina* populations in south–west Western Australia

Category (Hypothesis)	Variable	Description
Hydrology	SoilMoist_Dry ^a	Volumetric soil water content ($\text{m}^3 \text{m}^{-3}$) measured at the end of the dry season, April 2018
	SoilMoist_Wet	Volumetric water content ($\text{m}^3 \text{m}^{-3}$) measured following the wet season, December 2018
Micro-climate	TempMax ^a	Average daily soil temperature maximum ($^\circ\text{C}$) in the hottest month, February 2019
	TempRange	Average daily soil temperature range ($^\circ\text{C}$) in the hottest month, February 2019
Habitat structure	CanopyCover ^a	Average canopy cover (%)
	BareGround ^a	Average cover of bare ground (%)
	Moss ^a	Average cover of moss (%)
	SedgeCover	Average cover of sedges (%)
	WoodyDebris	Average cover of coarse woody debris and logs (%)
	LitterCover ^a	Average cover of leaf litter (%)
Soil properties	LitterDepth	Average depth of the litter layer (cm)
	Texture	Soil texture classification based on five categories ranging from sand (1) to clay (3)
	OrganicCarbon	Soil organic carbon (%)
	Conductivity ^a	Soil electrical conductivity (dS m^{-1})
	pH ^a	Soil pH (CaCl_2)

^aVariables used in the final generalised linear mixed-effect modelling

One HOBO temperature logger (UA-001-08) was placed in each sites' central quadrat to obtain soil surface temperature data at hourly intervals. Loggers were placed in the most open patch of the quadrat, buried into the top layer of soil (0–2 cm) under any present surface covering (e.g. leaf litter) and secured with a survey pin. Canopy cover was estimated at each quadrat using a densiometer held approximately 1.2 m above ground level at each cardinal direction, and then averaged for each quadrat. Ground cover was estimated as the percent area of bare ground, moss, sedge, leaf litter, and coarse woody debris within the quadrat, and litter depth was measured at three random points using a ruler and then averaged for the quadrat. Soil samples were collected by scraping away any litter or moss covering and extracting three cores (of the top 10 cm) from within each quadrat. Soil samples were bulked per site and analysed for texture (scale 1–3; Sand-Clay), organic carbon (%), conductivity (dS m^{-1}), and pH (CaCl_2) by an accredited soil laboratory (CSBP Soil and Plant Analysis Laboratory).

Statistical analysis

All statistical analyses were carried out in R (R Core Team 2019). To visualise differences in habitat characteristics across site types we used non-metric multidimensional scaling (NMDS). Plots were produced identifying site type (present, adjacent or extinct) using the 'vegan' package (Oksanen et al. 2019). Analysis of similarity (ANOSIM) was used to test whether there was a significant difference between habitat characteristics and site types.

To investigate which habitat features best explained frog abundance we used generalised linear mixed-effect models (GLMMs) with Poisson error distributions. We conducted three sets of analyses—first comparing present versus adjacent sites for *G. alba* and *G. vitellina* to investigate why frog populations occur in discrete patches, and then comparing present versus extinct sites for *G. alba* to examine possible mechanisms of *G. alba* decline. The response variable was the minimum estimated abundance of frogs at a site, and there were 15 potential explanatory habitat variables thought to be important in determining frog abundance. To avoid over parametrisation, we reduced correlated pairs of variables ($r > 0.5$) and retained one with the strongest effect on the model as a proxy (Booth et al. 1994). The reduced predictor variables set included dry season soil moisture, maximum temperature, canopy cover, bare ground, moss cover, litter cover, soil conductivity and soil pH (Table 1). Predictors were scaled and centered to have a mean of 0 and standard deviation of 1 (Harrison et al. 2018), allowing for comparison of effect sizes (Grueber et al. 2011). The areas containing paired sites (e.g. a present and nearby adjacent site) within the same catchment were included as a random effect. Models were fitted using the 'glmer' function in the package 'lme4' (Bates et al. 2015).

All possible variable subsets were created and ranked based on Akaike information criterion adjusted for small sample size (AIC_C). Model averaging was then conducted on the subset of models with high support ($\Delta\text{AIC}_C \leq 6$ with the top ranked model, which includes the 95% confidence model set (Richards et al. 2011) using the 'modavg' function in the 'MuMIn' package (Bartoń 2019)). Rather than single best model methods, this allowed for model selection uncertainty, as inference was based on the entire candidate model set (Burnham and Anderson 2002). The relative importance of each variable across all models was determined by the sum of Akaike weights (W_i) (Burnham and Anderson 2002). Model diagnostics were investigated using the 'DHARMA' package (Hartig 2020) and indicated no overdispersion, outliers, or heteroscedasticity of the model residuals. We

checked for collinearity among fixed effects by calculating variable inflation factors in all top models using ‘vif’ function in the ‘car’ package (Fox and Weisberg 2019).

Once dry season soil moisture was identified as one of the best explanatory variables of frog abundance (consistently significant for both species), we tested the accuracy of model predictions for translocation outcomes by comparing them with the actual outcomes of translocations (note that data from translocation sites were not used in the models described above). As the number of years since frogs were first introduced varied across translocation sites, we initially used abundance estimates from two years following the initial release of frogs, as this could be applied to all translocation sites. However, as the 2-year results did not differ from those using frog abundances from the year of the habitat surveys (2017), we used the more recent estimates from 2017. Generalised linear models of frog abundance were made for each species with dry season soil moisture as the explanatory variable, using the data from naturally occurring present and absent sites described above. Predictions of frog abundance were then made for existing translocation sites based on their dry season soil moisture, and the accuracy of predictions was investigated using Spearman’s rank-order correlation. We also converted predicted and actual frog abundances into categories in order to construct a confusion matrix using the ‘carot’ package (Kuhn 2018). A cut-off value of ≥ 10 frogs was used to define ‘success’ at a translocation site based on the criteria defined in the *Geocrinia* Recovery Plan (Department of Parks and Wildlife 2015).

Results

Differences between *G. alba* and *G. vitellina* sites

Analysis of microhabitat characteristics using NMDS showed there was a significant difference in the habitat characteristics of present, adjacent and extinct sites (Fig. 3; ANOSIM global $r = 0.1802$, $p = 0.001$). Present sites of both species had considerable overlap and were more similar to each other than their paired adjacent sites (50 m upstream or downstream) and extinct *G. alba* sites. Present sites for *G. vitellina* had some separation along the first ordination axis indicating higher soil moisture, moss and soil electrical conductivity than at the present sites for *G. alba* (Table S3). Adjacent and extinct sites were also separated mostly along the first ordination axis, signifying these sites had more bare ground, warmer temperatures and lower soil moisture and moss cover than present sites (Fig. 3).

Habitat associations

Frog abundance was positively associated with dry season soil moisture for both *G. alba* and *G. vitellina*, and dry season soil moisture was present in all top models ($\Delta AIC_C \leq 6$) across all three model sets (Table 2; Fig. 4). Dry season soil moisture also had the highest relative importance and effect size for models comparing *G. alba* present vs extinct sites and *G. vitellina* present vs adjacent sites, and was equally important in *G. alba* present vs adjacent sites (Table 2).

Geocrinia alba abundance (comparing present sites with extinct sites) was positively associated with moss cover and pH but the effect size was smaller. The confidence interval for pH was almost zero, specifying less evidence of an effect. Compared to adjacent sites, *G. alba* abundance was negatively associated with the area of bare ground and soil

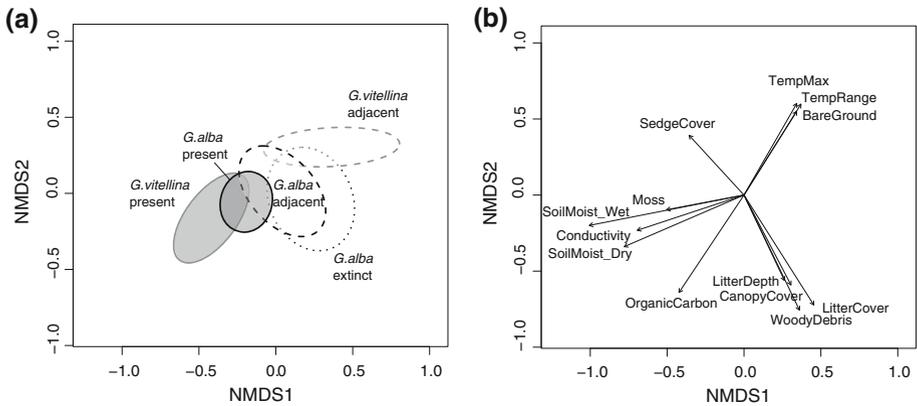


Fig. 3 Non-metric MDS ordination of site microhabitat characteristics showing **a** *G. alba* and *G. vitellina* present (shaded, solid line), adjacent (unshaded, dashed line) and extinct site types (unshaded, dotted line), and **b** vectors of environmental variables significantly related to the ordination ($p < 0.01$). Ellipses in a show 95% confidence intervals. Stress = 0.17

conductivity, which had the biggest effect sizes (Table 2; Fig. 5a, b). *Geocrinia alba* abundance was also positively associated with moss cover, similarly to present vs extinct sites (Fig. 5c). All four variables had equal relative importance (Table 2). For *G. vitellina*, several other variables were contained in the top models ($\Delta\text{AIC}_C \leq 6$), but confidence intervals of the estimates included zero, indicating weak evidence for an effect (Table 2).

Dry season soil moisture, which was present in all top models for both species, was strongly negatively correlated with maximum soil temperature ($t_{57} = -6.45$, $p < 0.001$, $r = -0.65$) and soil temperature range ($t_{57} = -8.23$, $p < 0.001$, $r = -0.74$) and was influenced by soil texture, with soils with higher clay content generally having higher soil moisture than sandier soils (Fig. S1).

Predicting translocation outcomes

Generalised linear models using dry season soil moisture as the sole predictor variable were good predictors of *G. alba* and *G. vitellina* abundances at translocation sites (Fig. 6). Predicted abundance was significantly correlated with actual abundances at translocation sites ($r_s = 0.89$, $p = 0.015$). Using a cut-off value of ≥ 10 calling males as a ‘successful’ translocation site, models of frog abundance exclusively based on dry season soil moisture correctly predicted translocation success or failure at with an overall accuracy of 0.83 (Table S5). Agreement between observed and predicted was substantial (Kappa = 0.67).

Discussion

Studies of species-habitat relationships often produce a myriad of complex associations, but here we were able to identify a single predictor that explained the majority of variation in frog abundance for both *G. alba* and *G. vitellina* and was validated using empirical data. Fine-scale habitat variables—particularly soil moisture during the warmer-drier months—explained not only the patchy distribution of each species, but also explained local extinctions of *G. alba* populations and translocation outcomes of *G. alba* and *G. vitellina*.

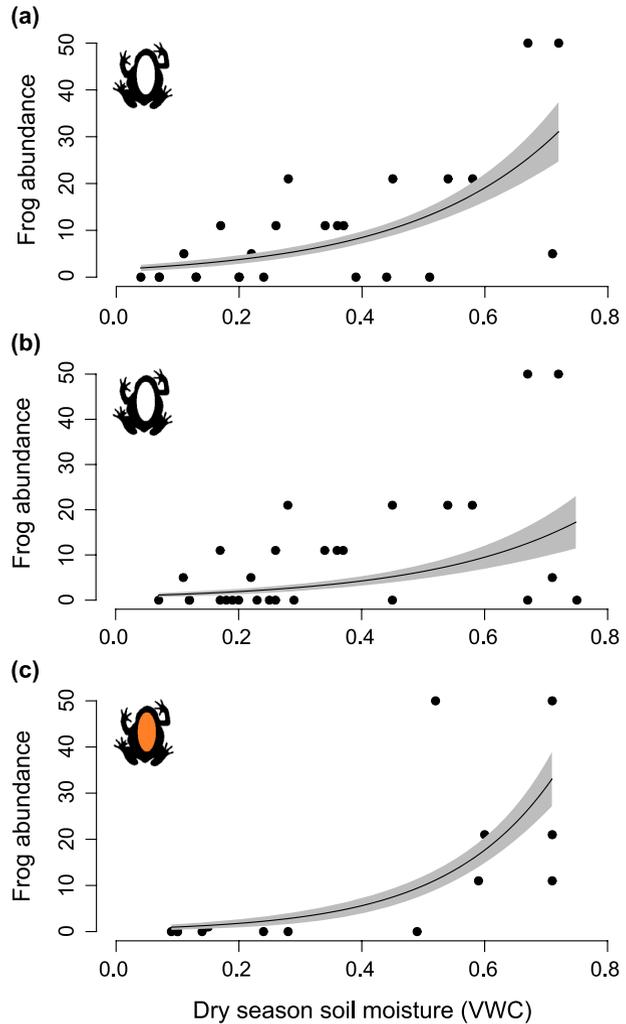
Table 2 Averaged model coefficients from the top model set predicting abundance of *G. alba* (present vs extinct sites and present vs adjacent sites) and *G. vitellina* (present vs adjacent sites)

Variable	Estimate	95% confidence interval		Relative importance	n models
		Lower	Upper		
<i>Geocrinia alba</i> —present vs extinct sites					
(Intercept)	0.8475	– 0.42	2.12		
SoilMoist_Dry	2.2326	1.03	3.43	1.00	26
Moss	1.1782	0.17	2.19	0.93	22
pH	1.0139	0.02	2.01	0.76	18
Conductivity	– 0.6349	– 1.31	0.04	0.48	13
CanopyCover	0.5146	– 0.14	1.17	0.39	12
TempMax	– 0.1974	– 1.60	1.20	0.13	5
BareGround	– 0.2788	– 1.04	0.48	0.11	5
LeafLitter	0.1326	– 0.75	1.02	0.09	5
<i>Geocrinia alba</i> —present vs adjacent sites					
(Intercept)	0.68	– 0.01	1.36		
Conductivity	– 1.78	– 2.44	– 1.12	1.00	9
BareGround	– 1.40	– 2.05	– 0.75	1.00	9
SoilMoist_Dry	1.02	0.45	1.60	1.00	9
Moss	0.61	0.14	1.08	1.00	9
CanopyCover	0.42	– 0.30	1.14	0.29	4
LeafLitter	0.33	– 0.33	1.00	0.19	3
pH	– 0.14	– 0.70	0.42	0.12	3
TempMax	– 0.01	– 0.53	0.51	0.12	2
<i>Geocrinia vitellina</i> —present vs adjacent sites					
(Intercept)	– 0.39	– 2.26	1.49		
SoilMoist_Dry	3.38	1.58	5.18	1.00	15
CanopyCover	1.19	– 0.52	2.90	0.31	6
BareGround	– 1.17	– 3.30	0.96	0.20	5
Moss	0.67	– 0.62	1.97	0.13	3
pH	0.47	– 0.47	1.42	0.13	3
LeafLitter	– 0.10	– 1.43	1.23	0.06	2
TempMax	0.41	– 1.14	1.97	0.09	3

Variables in bold indicate those with 95% confidence intervals that did not span zero. The relative importance of each variable (sum of Akaike weights) and the number of models in the top set containing the variable are also displayed

Geocrinia alba and *G. vitellina* appear to share the same environmental niche. The similarity in their habitat associations is not surprising as they are the most closely related species within the *Geocrinia rosea* complex, which includes four direct-developing frog species endemic to southwestern Australia (Read et al. 2001). Despite no overlap in their distributions, NMDS analysis revealed that present sites of each species had significant overlap in their habitat attributes and were more similar to each other than to riparian sites only ~ 50 m from the edge of areas where frogs occur. This was particularly stark for *G.*

Fig. 4 Predicted abundance of *G. alba* **a, b** and *G. vitellina* **c** as a function of dry season soil moisture (volumetric water content, $\text{m}^3 \text{m}^{-3}$) based on **a** present versus extinct sites or **b** and **c** present versus adjacent sites. Circles are actual observations, solid lines are the model predictions, and grey areas denote 95% confidence intervals. Predicted abundance and confidence intervals were generated with a GLM without random effects, holding all other significant variables constant on their means



vitellina, where there was no overlap between present and adjacent sites. The presence of both species was positively related to soil moisture at the ends of the wet (December) and dry (April) seasons, and to moss cover, and was negatively associated with bare ground, temperature maximum and temperature range during summer (February). This demonstrates that both species require relatively cooler, wetter, and mossier sites, with more ground cover.

Dry season soil moisture measured in April was the principal habitat factor determining *G. alba* and *G. vitellina* occurrence and abundance at sites. This time is towards the end of a drier-hotter period (Fig. 2) and when the drainage areas where frogs occur are likely at their driest. Soil moisture varied considerably during this time, even across small distances, with sites where frogs were present having significantly higher soil moisture than adjacent patches in the same riparian habitat 50 m away. As the surrounding open woodland habitat is drier (E. Hoffmann, unpublished data), both species appear to be utilising patches along

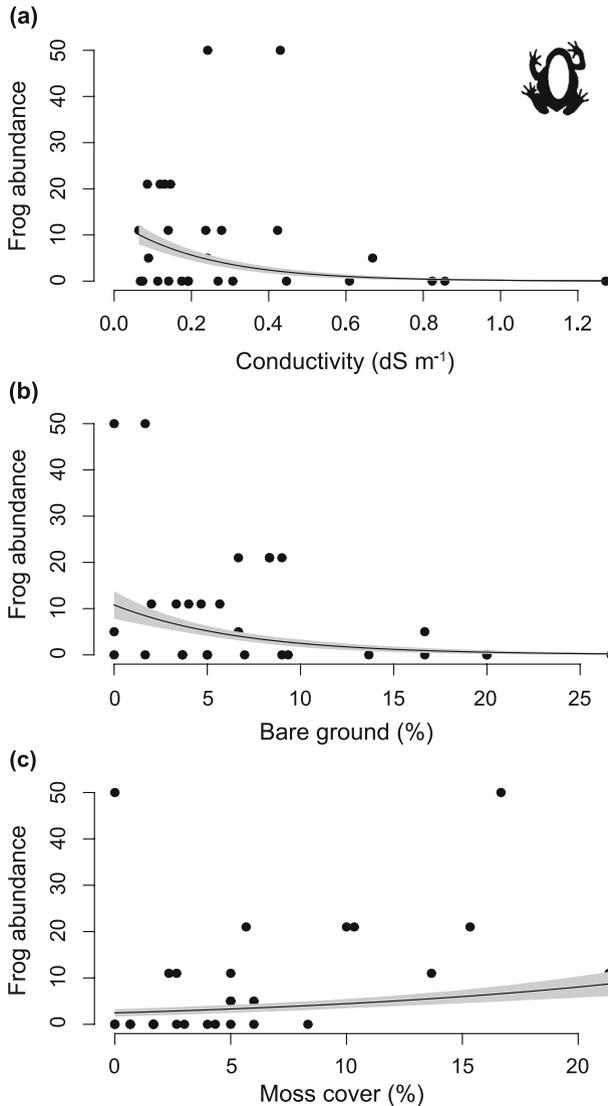
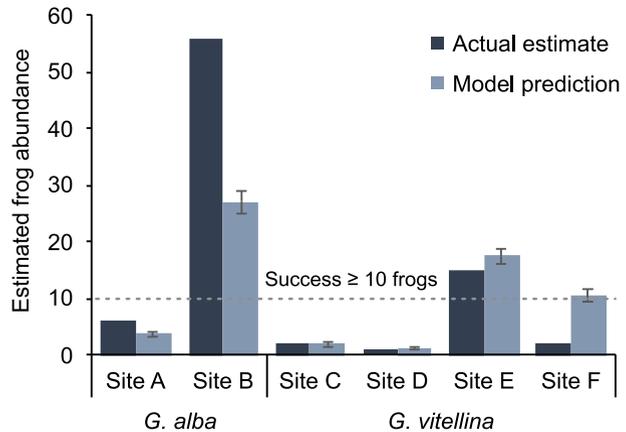


Fig. 5 Predicted abundance of *Geocrinia alba* as a function of **a** soil conductivity, **b** bare ground and **c** moss cover, based on present versus adjacent sites. Solid lines are the model predictions, grey areas denote 95% confidence intervals, and circles are actual observations. Predicted abundance and confidence intervals were generated with a GLM without random effects, holding all other significant variables constant on their means

drainage lines that retain the highest moisture in the dry season. Soil water availability in summer–autumn may therefore be the defining factor that determines the patchy and restricted distributions for both species. Soil temperature (both maximum and range) was significantly correlated with soil moisture, and therefore drier areas were also warmer. The summer dry season may therefore be a critical period for *G. alba* and *G. vitellina* and potentially more important in determining population persistence than winter–spring

Fig. 6 Estimates of the number of calling males at *Geocrinia* translocation sites and predicted abundance (\pm standard error) based on models using dry season soil moisture as the predictor variable



breeding conditions. Currently, nearly all research conducted has focused on the *G. alba* and *G. vitellina* breeding period in spring (Driscoll 1996; Pauli 1999; Conroy 2001; Mitchell 2001), and little is known about their behaviour and habitat in the drier and warmer summer–autumn period.

Lower frog abundance or absence at drier sites could be due to desiccating conditions experienced by frogs and their egg masses. The physiological constraints of moisture on amphibians is well known (Shoemaker et al. 1992), and drier conditions at the end of the breeding period could lead to reduced or failed recruitment. Both species breed over a prolonged period following winter rains, laying eggs in moist depressions in the soil adjacent to winter streams. Egg laying and development can continue into December and January, several months into the drier period (Driscoll 1996; Conroy 2001) (Fig. 2). Incubation in drier conditions causes terrestrial frog eggs to lose water from their jelly capsules and embryos have higher rates of deformations and mortality (Bradford and Seymour 1988; Mitchell 2002b; Andrewartha et al. 2008). If soils remain saturated throughout embryonic and larval development, newly metamorphosed frogs emerge just prior to the warmest and driest time of year (around November–January, Fig. 2) and are vulnerable to desiccation due to their very small size (~ 0.03 g at metamorphosis) and large surface area to volume ratio. Juvenile amphibians consequently lose more moisture and experience lower survival than adults in hotter and drier landscapes (Cayuela et al. 2016). The juvenile stage is not only the most vulnerable, but also prolonged, lasting at least two years (Driscoll 1999). Both *Geocrinia* species rely on high juvenile survival due to low fecundity (with an average of 11 eggs per clutch) and because most adults only breed once (Driscoll 1999; Conroy 2001). Population viability modelling of *G. alba* and *G. vitellina* has indicated that a reduction in juvenile survival would have the biggest impact on population trends compared with other life stages (Conroy and Brook 2003). Therefore, we infer that frogs are at risk of desiccation during the dry phase, and juveniles are likely the most vulnerable life stage.

Local extinctions throughout the range of *G. alba* also appear to be driven by habitat differences, with extinct sites being significantly drier in summer and less mossy than present and adjacent sites. Hydrological change had been suggested as one of the main threats to *G. alba* (e.g. Driscoll 1996; Pauli 1999; Conroy 2001), but until this study, there were no quantitative data to evaluate the importance of site hydrology, specifically soil moisture, as a factor in *G. alba* declines. Considerable land clearance and land-use change

has occurred within *G. alba*'s range. It is estimated that 70% of potentially suitable habitat has been removed and most of the clearance has occurred relatively recently, between 1960 and 1980 (Pauli 1999; Page et al. 2018). Clearing of native vegetation may increase stream flows and groundwater levels, at least in the short term (Bari et al. 1996), but these increases may have been offset by extensive land-use change following vegetation clearance, such as expansions of orchards, forestry plantations, barriers to flow (e.g. roads) and installation of dams, and could be reducing flows to *G. alba* catchments (Department of Parks and Wildlife 2015). Despite the clear attribution of local extinctions of some populations to land clearance and adjacent land use change, most of the extinct *G. alba* sites sampled in this study were within stands of native vegetation and conservation estate (with the upstream catchment entirely within reserve or native vegetation block) and so should be less impacted by land clearance and hydrological alterations, such as dams. Consequently, the pattern of extinctions at drier sites may also reflect hydro-climatological changes that are occurring in the region on a broader scale.

There is mounting evidence of substantial climatic and hydrological change occurring in the region. South-west Western Australia has experienced a 15–20% decline in rainfall and a consequent 35–50% reduction in streamflow since the 1970s, with the biggest rainfall changes occurring at the start of the wetter winter period in May and June (Petroni et al. 2010; McFarlane et al. 2020). Reduced rainfall in winter is increasing the length of the 'no-flow' period in rivers of the region, with surface flows in ephemeral creek lines commencing later and ending earlier (e.g. Smettem et al. 2013). Furthermore, riparian habitats may also be impacted by decreasing groundwater levels in the region (McFarlane et al. 2020). Streamflow is dominated by winter rainfall, but some catchments in *Geocrinia* habitats overlie sedimentary aquifers (e.g. Upper Chapman Brook) and may receive groundwater inputs during summer (Department of Water 2015). If sites rely on groundwater seepage to retain moisture over summer and autumn, lower groundwater levels could be resulting in drier summer conditions. As we showed that drier soils had higher maximum temperatures and greater temperature ranges, drier sites are also likely to be more impacted by the warmer air temperatures that have been observed and predicted to increase in the region (Charles et al. 2010). Therefore, regional changes in climate and hydrology are likely to be resulting in drainage habitats receiving less surface water and groundwater inputs, and consequently, experiencing more extreme temperatures and drying. The changes being experienced in headwater systems may be amplified as the relationship between rainfall decline and streamflow can be highly non-linear (e.g. 1% rainfall decline \geq 3% runoff decline; McFarlane et al. 2020) and through a disconnection of groundwater-surface water connectivity due to declining groundwater levels (Petroni et al. 2010).

If drying of some *G. alba* habitats is due to regional changes in climate or hydrology, why is *G. vitellina* not declining? *Geocrinia vitellina* occurs in sites that are less sandy than *G. alba* sites and thereby retain higher soil moisture. *Geocrinia vitellina* also occurs on drainage lines that may receive groundwater inputs from the deeper Yarragadee and Leederville aquifers, buffering these areas from drying over summer (CSIRO 2009). Therefore, *G. vitellina* habitats may be less vulnerable to hydrological changes. However, *G. vitellina* also appears to be highly restricted to wetter patches within riparian drainage habitats in the dry period, as adjacent areas where frogs didn't occur were significantly drier. They also share the same specialised breeding requirements, limited dispersal ability, and very small area of occurrence as *G. alba*. Therefore, whilst only *G. alba* has shown extreme declines, both species are threatened by habitat and range contraction associated with climatic and hydrological changes occurring in the region. For example, groundwater

tables in the Blackwood plateau underlying *G. vitellina* habitat are currently declining at ~ 1.2 m/year and are predicted to drop 10 m in some areas by 2030 (CSIRO 2009). More recent population monitoring has also indicated that some *G. vitellina* populations are showing signs of decline (K. Williams, unpublished observation).

In addition to soil moisture levels, habitat structure and other soil parameters were also important in determining *G. alba* abundance. Most notably, compared with sites where *G. alba* were present, extinct sites had less moss cover. Adjacent areas were also associated with lower moss cover, as well as more bare ground and higher soil conductivity. More moss and ground cover at sites where *G. alba* are present would provide frogs with more habitat for nest construction, as well as greater protection from desiccation (Glime and Boelema 2017). The electrical conductivity levels in the drainage lines were generally low (< 1 dS m⁻¹) but indicated that *G. alba* prefer areas with lower soil conductivity. Conductivity is an indicator of salinity, which can cause osmotic stress (Smith et al. 2007) and can decrease growth and survival of amphibians at higher concentrations (Chinathamby et al. 2006; Kearney et al. 2012).

Management implications

Here we have demonstrated that gaining a detailed understanding of species habitat-associations can provide vital information for conservation management. Studies of habitat requirements typically compare occupied and unoccupied habitats, but we additionally evaluated the outcomes of translocations in the context of habitat variability. This approach highlighted that fine-scale habitat attributes are likely to be a key driver of translocation success, as we were able to hindcast translocation outcomes with high accuracy using only dry season soil moisture as the predictor variable. Therefore, using dry season soil moisture to select translocation sites should increase the likelihood of success of new translocations. We suggest that sites should have soil moisture contents of at least 40% during the drier period to sustain a population of > 10 calling males. Very few frogs were recorded at sites where soil moisture content was below 20%, indicating that sites below this moisture threshold are unlikely to support *G. alba* or *G. vitellina* populations long-term. These recommendations are tentative in recognition that our predictions are based on one year of data, and the variability and extremes the sites experience throughout the year is unknown. Further, soil moisture per se has limited biological relevance to amphibians, as soil water potential drives water availability and varies substantially with soil type (Shoemaker et al. 1992). However, as soil moisture content can be measured more simply than water potential, it is a practical method to rapidly detect relatively wetter and drier areas.

More broadly, we highlight that conservation areas may not provide a buffer against wider regional threats, such as drying brought about by climate change. This corroborates our related study showing that physiological tolerance limits of *G. alba* and *G. vitellina* are being breached in atypically warm periods (Hoffmann et al. in review), and other studies that have recorded impacts of drought on amphibians within conservation estates in other regions of Australia (Scheele et al. 2012) and the world (McMenamin et al. 2008; Cayuela et al. 2016). Mitigating broad scale impacts of changing hydrology and climate (e.g. those that are occurring across entire species ranges) including in protected areas, is an imposing management challenge. Some suggested strategies include providing microclimate refugia through watering actions, such as sprinkler systems to wet soils or artificial filling of wetlands (see Shoo et al. 2011 for a summary), which may help alleviate pressures at key sites in the short term. Groundwater levels are heavily linked to rainfall but also vegetation

(CSIRO 2009). Reversing groundwater declines, as well as increases in streamflow, could potentially be achieved by a reduction in upland vegetation via thinning (Jones et al. 2018) and has been trialled in Jarrah forests in Western Australia (e.g. Stoneman, 1993), but requires consideration of other biodiversity values. Another conservation approach when habitats become marginal due to climate change is assisted colonisation, where species are moved to areas outside of their indigenous range that will become suitable in the near future (Mitchell et al. 2013; Gallagher et al. 2015). Our findings suggest that the conditions that led to the loss of *G. alba* populations have not alleviated, and that sites where the species became extinct are not suitable for recolonisation, making assisted colonisation worthy of consideration. A priority should be to detect areas that currently stay moist or retain shallow surface water year-round and to evaluate their resilience to future hydrological change, as these areas may provide important refugia for the species into the future.

Our findings also have implications for other taxa and locations. Local extinctions of *G. alba* are potentially among the first signs of impacts of changes to the hydrology of drainage areas and headwaters on regional biodiversity, and the impacts on other water-dependent fauna and flora that occur in these systems is unknown (e.g. threatened *Engaewa* spp.—endemic burrowing crayfish). We also detected critical habitat differences on an extremely fine-scale, just tens of meters along drainage lines. These distances are much finer in scale than the focus of much climate and species distribution modelling, emphasising the need to consider modelling habitat variables at the scale most relevant to that species. Many mid-latitude areas of the world are experiencing a drying and warming trend, and our study highlights the vulnerability of sedentary and specialised water-sensitive species in an era of rapid environmental change.

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Data availability Due to the sensitive nature of the species locations the datasets from the current study are not publicly available but are available from the corresponding author on reasonable request.

Code availability Not applicable.

Compliance with ethical standards

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

Ethical approval This research was completed under a DBCA Regulation 4 Permit and University of Western Australia Animal Ethics Approval RA/3/100/1554.

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