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Are narrow-ranging species doomed to extinction? Projected dramatic decline in future climate suitability of two highly threatened species — Source link \square

Nicolas Dubos, Frédérique Montfort, Clovis Grinand, Marie Nourtier ...+9 more authors

Institutions: University of Paris, Residence Inn by Marriott, University of Kent, International Union for Conservation of Nature and Natural Resources ...+1 more institutions

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1 Are narrow-ranging species doomed to extinction? Probable dramatic and generalised decline in

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- 6 Nicolas Dubos^{1,2}, Frederique Montfort³, Clovis Grinand³, Marie Nourtier³, Gregory Deso⁴, Jean-Michel
- 7 Probst⁵, Julie Hanta Razafimanahaka⁶, Raphali Rodlis Andriantsimanarilafy⁶, Eddie Fanantenana
- 8 Rakotondrasoa⁶, Pierre Razafindraibe⁶, Richard Jenkins^{6,7} and Angelica Crottini⁸
- 9
- 10
- 11 1 Centre d'Ecologie et des Sciences de la Conservation (CESCO UMR 7204), Sorbonne Université, MNHN,
 12 55 rue Buffon, 75005 Paris, France.
- 13 2 INRAE (UMR TETIS), Maison de la télédétection, 500 rue Jean-François Breton 34093 Montpellier
 14 Cedex 5, France.
- 15 3 N'Lab, Nitidæ, Maison de la télédétection, 500 rue Jean-François Breton 34093 Montpellier Cedex 5
- 4 Association Herpétologique de Provence Alpes Méditerranée, Maison des Associations, 384 route de
 Caderousse, F-84100 Orange, France.
- 18 5 Association Nature and Patrimoine, 1 rue des amarantes résidence Valeriane 2, Bat C appartement 15.
 19 97490 Sainte Clotilde, Île de La Réunion, France.
- 20 6 Madagasikara Voakajy, B.P. 5181, Antananarivo (101), Madagascar.
- 21 7 Durrell Institute of Conservation and Ecology, School of Anthropology and Conservation, Marlowe
- 22 Building, University of Kent, Canterbury, CT2 7NR ; Current address: IUCN Global Species Programme,
- 23 219c Huntingdon Road, Cambridge, CB3 0DL, UK.
- 24 8 CIBIO, Research Centre in Biodiversity and Genetic Resources, InBIO, Universidade do Porto, Campus
- 25 Agrário de Vairão, Rua Padre Armando Quintas, No 7, 4485-661 Vairão, Portugal.

26 Abstract

- 27 Aim
- 28 Narrow-ranging species are usually omitted from Species distribution models (SDMs) due to statistical
- 29 constraints, while they are predicted to be particularly vulnerable to climate change. The recently available
- 30 high-resolution climate and land use data, along with recently developed methods enable to increase the
- 31 eligibility of narrow-ranging species for SDMs, provided their distribution is well known. We fill a gap of
- 32 knowledge on the effect of predicted climate change on narrow-ranging species.
- 33 Location
- 34 Madagascar and Reunion Island.
- 35 Time period
- 36 Response to recent climate (1995-2019) and prediction on future climate (2070).
- 37 Major taxa studied
- 38 The golden mantella frog Mantella aurantiaca and the Manapany day gecko Phelsuma inexpectata.
- 39 Methods
- 40 We modelled the distribution of two threatened narrow-ranging species for which the distribution of their
- 41 occurrence records is well documented. Our modelling scheme included a range of processes susceptible to
- 42 address statistical issues related to narrow-ranging species (model complexity, evaluation, background
- 43 extent, prevalence, extrapolation).
- 44 Results
- 45 We predict an alarming decline in climate suitability in the whole current distribution area of both species by
- 46 2070, potentially leading to a complete extinction even in the most optimistic scenarios. We identified the
- 47 areas with the best climate suitability in the future, but these remain largely suboptimal regarding species

48 climatic niche. The high level of habitat fragmentation suggests that both species likely need to be at least49 partly translocated.

- 50 Main conclusions
- 51 Climate change may not only drive range contractions or distribution shifts in narrow-ranging species, but
- 52 may lead to the complete extirpation of suitable environments across their entire region. This study suggests
- 53 that the level of threats of data-poor and narrow-ranging species already identified as threatened may be
- 54 underestimated, especially in heterogeneous tropical areas. We stress the need to develop sampling
- 55 campaigns and implement proactive actions for narrow-ranging species in the tropics.
- 56
- 57 Keywords: Climate change, Ensemble of Small Models, Habitat restoration, Jackknife, Madagascar,
- 58 Mantella aurantiaca, Phelsuma inexpectata, Reunion Island, Species distribution models, Translocation

59

60 Introduction

61 Climate change is predicted to become the main driver of biodiversity loss it in the next decades (Bellard et 62 al., 2012). Species Distribution Models (SDMs) are probably the most common approach used to predict the 63 impact of future climate change on species. They are used to predict current and future environmental 64 suitability, and provide guidelines for the identification of priority areas for protection (Leroy et al., 2014), 65 habitat restoration and species (re)introduction/translocations (e.g., Bellis et al., 2020; Draper et al., 2019; 66 Westwood et al., 2020). Habitat restoration and translocations are two ecological engineering techniques 67 enabling the restoration of depleted populations. Translocation programs will become increasingly needed in 68 the face of climate change, especially for species with small distribution ranges (Thomas, 2011). In this 69 regard, SDMs help identifying suitable receptor sites that meet the species' habitat requirements while 70 accounting for climate suitability (Bellis et al., 2020). In highly degraded environments a combination of 71 habitat restoration and translocation may be needed to avoid species extinctions. This may be the case in 72 highly fragmented tropical systems, where the number of narrow-ranging species is higher and where climate 73 change effects are expected to be stronger (Tewksbury et al., 2008).

74 The impact of future climate change is largely understudied in endangered narrow-ranging species. This is 75 mainly due to the difficulty to model their current and future distribution because of low sample sizes and 76 subsequent little spatial replicates when fitted on climate data (Botts et al., 2013; Platts et al., 2014; Breiner 77 et al., 2015; Galante et al., 2018). Both these factors lead to statistical constraints that withdraw these species 78 from eligibility for SDMs. However, the omission of narrow-ranging species in SDMs may be problematic in 79 terms of conservation planning, because the area encompassing their distribution may be downplayed (Platts 80 et al., 2014). Although narrow-ranging species are known to be more vulnerable to climate change (Pearson 81 et al., 2014), few studies have provided quantitative assessment of climate change impacts on these (but see 82 Alamgir et al., 2015; Zhang et al., 2020).

83 The recent availability of high-resolution climatic data (e.g., Fick & Hijmans, 2017; Karger et al., 2017),

84 along with high-resolution land cover data (e.g., Vieilledent et al., 2018) is offering new opportunities for

85 modelling the distribution of these species (Lannuzel et al., 2021). However, despite a probable increase in

86 statistical power, there is still the possibility for SDMs to produce misleading results due to spatial sampling 87 bias (Phillips et al., 2009). The effect of sample bias may be particularly strong in rare (or poorly known) 88 species with small sample sizes, because models are more influenced by each occurrence data that is used 89 (Pearson et al., 2007). A number of techniques were recently developed to account for sampling bias, often 90 implying the filtering of occurrence or environmental data (Gábor et al., 2019) or non-random pseudo-91 absence selection (Phillips et al., 2009). However, data filtering can become problematic for species with 92 low sample sizes (Vollering et al., 2019), especially when species distribution is highly localised (Inman et 93 al., 2021) and is not recommended in absence of evidence of bias in occurrence data (Gábor et al., 2019). 94 Similarly, non-random pseudo-absence selection is not always effective (Dubos et al., 2021) and tends to 95 make predictions worse in narrow-niche species (Inman et al., 2021). On the other hand, the reliability of an 96 SDM is more driven by the quality of the data than the implementation of models (Araùjo et al., 2019). 97 Therefore, the best option for narrow-ranging SDMs may be to select species for which the distribution is 98 well known. 99 Given the predicted magnitude of climate change, along with the narrow thermal tolerance of tropical species 100 (Tewksbury et al., 2008), we may not only expect a reduction or a geographical shift in suitable conditions 101 for narrow-ranging species, but the extirpation of suitable conditions across the entirety of their distribution 102 range. Here we fill a gap of knowledge regarding the impact of future climate change on narrow-ranging 103 species using two species for which the distribution is particularly well documented. These were the 104 Manapany day gecko Phelsuma inexpectata, classified as Critically Endangered and the golden mantella frog 105 Mantella aurantiaca formerly classified as Critically Endangered, now classified as Endangered after the 106 inclusion of one locality record which increased its extent of occurrence. Both species are in continued 107 decline (Probst & Turpin, 1997; Crottini et al., 2019), live in highly fragmented areas (respectively in 108 Reunion Island and central Madagascar) and are in urgent need for conservation actions. Given the 109 important, long-term efforts invested to document their distribution, we assume that the geographic 110 information for these species is nearly comprehensive and unbiased. We tested whether climate change will 111 'only' drive range reductions/shifts, or lead to a total extirpation of their suitable areas. We eventually

- 112 identify the most suitable candidate areas for habitat restoration and translocation across their respective
- 113 regions.
- 114

115 Methods

116 Occurrence data

117 *Phelsuma inexpectata* – The Manapany day gecko is endemic to the south of Reunion Island. We retrieved 118 31 occurrence data from literature (Bour et al., 1995). Since then, the surroundings of the known distribution 119 range of the species were regularly visited (an 11 km-long coastal band). Two localities corresponding to 120 introduced populations were identified west of the current range, which we added to the data (Deso, 2001; 121 Porcel et al., 2021). Recent sampling campaigns enabled to find additional occupied habitats (Dubos 2010) 122 but did not add any occurrence point after aggregation at the resolution of the environmental variables (30 123 arc seconds). Therefore, we assume that the sample occurrence of the species is nearly comprehensive. The 124 total number of 30 sec. occupied pixels resulted in 15 presence points. 125 Mantella aurantiaca - The golden mantella frog is distributed in central-Eastern Madagascar (region of 126 Moramanga; Fig. 1). We obtained 131 occurrence data from Piludu et al. (2015). Those included compiled 127 published data from surveys conducted between 2001 and 2007 (Bora et al., 2008; Randrianavelona et al., 128 2010) and new locations from additional surveys conducted between 2008 and 2013. More recent surveys 129 conducted between 2014 and 2019 enabled to add 39 occurrences. The region has been extensively surveyed 130 and it is very likely that most occupied habitats were identified. After aggregation to match the resolution of 131 environmental data, sample size resulted in 101 occurrence points.

132

133 *Climate data*

We used the 19 bioclimatic variables for 30 arc sec (approximately 900m) resolution of the current climate
data and of the 2070 projections from CHELSA (Karger et al., 2017; Fig. S1, S2). We decided to include all

136	the 19 variables because both temperature and precipitation are related to the species' biology, including
137	those related to indices of variability (e.g., cyclones drive mortality in Phelsuma and heavy rains drive
138	reproduction in Mantella; Vinson, 1975; Randrianavelona et al., 2010). We used three Global Circulation
139	Models (GCMs; i.e., BCC-CSM1-1, MIROC5 and HadGEM2-AO) and two greenhouse gas emission
140	scenarios (the most optimistic RCP26 and the most pessimistic RCP85; Fig. S3, S4). We also ran sets of
141	models using the WorldClim Global Climate Data (Fick & Hijmans, 2017), with the same GCMs and
142	scenarios to account for potential effect of the baseline data in our predictions.

143

144 Distribution modelling

145 We modelled and projected species distributions with the Biomod2 R package (Thuiller et al. 2009), using 10 146 modelling techniques: generalised linear model (GLM), generalised additive model (GAM), classification 147 tree analysis (CTA), artificial neural network (ANN), surface range envelop (SRE, as known as BIOCLIM), 148 flexible discriminant analysis (FDA) and random forest (RF), Multiple Adaptive Regression Splines 149 (MARS), Generalised Boosting Model (GBM) and Maximum Entropy (MaxEnt). We generated five 150 different sets of randomly-selected pseudo-absences (random generation is recommended for rare and 151 specialised species; Inman et al., 2021). We selected one variable per group of inter-correlated variables to 152 avoid collinearity (Pearson's r > 0.7) and assessed the relative importance of each variable kept with 10 153 permutations per model replicate (total = 500). The variables included in the final models were those with a 154 relative importance > 0.2 across at least 50% of model runs. We predicted species distributions with an 155 ensemble of small models approach (ESM; Breiner et al., 2015). We ran sets of bivariate models, i.e. 156 including all pairwise combinations of the selected variables, and produced an ensemble model with the 157 mean predictions across all models weighted by their respective AUC (see below). This method is advocated 158 for rare species and enables to reduce model complexity without reducing the explanatory power. We set 159 three runs of cross validation (except for the jackknife procedure; see below). We ran a first set of models 160 (1), hereafter referred to as 'Base' setting, with 1000 pseudo-absences and a background covering the entire 161 respective island of each species. We ran a second set (2) where pseudo-absences were down-weighted to

162 equal presence data for the former two (setting prevalence to 0.5) referred to as 'equal total weight' (ETW). 163 We reperformed the first two model sets but with a background covering the southern part of Reunion Island 164 and the eastern forested part of Madagascar (see 'Land use data' section), without, and with pseudo-absence 165 down-weighting, respectively referred to as (3) 'Restricted background' and (4) 'ETW-Restricted 166 background'. Eventually, we ran a (5) Base model set with the Worldclim baseline climate instead of Chelsa, 167 referred to as 'Worldclim baseline'. We used a jackknife for *P. inexpectata*, a procedure that is appropriate 168 for small sample sizes (Pearson et al., 2007; Galante et al., 2018). The jackknife consists in running n 169 iterations corresponding to the number of occurrence data, removing one occurrence for each run of 170 calibration. Models are evaluated with the withheld occurrence. We did not used it for *M. aurantiaca* given 171 that the number of occurrence data was sufficient for a standard procedure, and used an 80% calibration 172 subset. For both species we present five ensemble models of current distribution (of varying background, 173 prevalence and baseline climate), and 30 ensemble models of predicted future distribution (of varying 174 background, prevalence, baseline climate, GCM and RCP).

175

176 Model evaluations – Ideally, performance evaluations are based on block-cross validation to limit spatial 177 autocorrelation at large scales. In our case, species distributions are highly localised and the use of spatial 178 splits would result in a small number of blocks and strong unbalances between blocks. We therefore used 179 random splits for M. aurantiaca (and a jackknife for P. inexpectata). We assessed model performance using 180 the Area Under the Operative Curve (AUC), the True Skill Statistics (TSS) and the Boyce index (the latter 181 was not computed for the jackknife procedure, because this index is based on Spearman's coefficients and 182 cannot be estimated on the basis of a single location). For ensemble models, we excluded models for which 183 the AUC was below 0.9 and for which the Boyce index was below 0.5 (Gillard et al., 2017). 184 We performed a multivariate environmental similarity surfaces analysis (MESS) to determine whether

185 models are well informed for predictions on novel (future) data. We eventually quantified the uncertainty

186 related to model input parameters (modelling technique, pseudo-absence distribution, number of pseudo-

- absences generated, pseudo-absence down-weighting, GCMs, RCPs and baseline climate data) by computing
 the standard deviation of the suitability scores between model predictions.
- 189
- 190 Land use data
- 191 We accounted for the habitat requirements of our model species by applying a filter to the projected climate
- 192 suitability based on land use and land cover data (e.g., Gillard et al., 2017). This enabled to minimise model
- 193 complexity while remaining biologically realistic and relevant for conservation applications.
- 194 Reunion Island We used very high-resolution land cover categories (Urban, agricultural, natural, water) at
- 195 1.5m resolution (resampled at 15m for computing purposes) derived from remote sensing (Dupuy and
- 196 Gaetano, 2019; Fig. S5). Since *P. inexpectata* can be found in both urbanised and natural areas and avoids
- 197 agricultural areas (Probst & Turpin, 1997), we excluded the latter land use type only. This is relevant also for
- 198 conservation applications because habitat restoration is more difficult to implement in agricultural areas.
- 199 Madagascar Mantella aurantiaca is exclusively found in swampy forested areas (Randrianavelona et al.,
- 200 2010). We filtered our suitability maps to include only cells corresponding to rainforests. We obtained forest
- 201 cover data from a study that combined global tree cover loss data with historical national forest (resolution
- 202 30m; Vieilledent et al., 2018; Fig. S6). We used forest cover of the year 1990 for current distribution models
- 203 because it corresponds approximately to the period of the oldest occurrence data (some areas were deforested
- since then). For future predictions, we used the latest forest cover data available, i.e. 2017.
- 205

206 Results

- 207 Variable selection and model performance
- 208 Three to four variables were selected in every cases (Table 1; Fig. S7, S8). The current distribution of both
- species was well predicted (*P. inexpectata*: mean AUC = 0.96, mean TSS = 0.91; *M. aurantiaca*: mean
- Boyce = 0.82, mean AUC = 0.97, mean TSS = 0.94; Fig. S9, S10). We excluded 5076 poorly performing

- 211 models out of 18750 for P. inexpectata, and 343 out of 4140 for M. aurantiaca. The median suitability score
- 212 at the presence points for *P. inexpectata* was 836 (911 when removing the two introduced populations) and
- 213 783 for *M. aurantiaca*. Models consistently identified the current distribution range between all runs for both
- species, as shown by the uncertainty map (SD intermodel suitability scores at the presence points of 168 for
- 215 *P. inexpectata* and 113 for *M. aurantiaca*; Fig. S11, S12).
- 216 Table 1. Selected variables for two narrow-ranging species in Reunion Island and Madagascar. Variables
- 217 were selected on the basis of 10 permutations per modelling technique and pseudo-absence set (total = 500).

Species	Model set	Selected variables
Phelsuma inexpectata	Base	Bio1, Bio15, Bio18
_	Restricted	
	background	Bio1, Bio15, Bio18
	Worldclim baseline	Bio1, Bio3, Bio14, Bio16
Mantella aurantiaca	Base	Bio1, Bio3, Bio4, Bio5
	Restricted	
	background	Bio1, Bio18, Bio19
	Worldclim baseline	Bio10, Bio11, Bio14, Bio16

- 218
- 219 *Future climate suitability*
- 220 All scenarios indicated an important decrease in climate suitability in the entirety of the current range of both
- species by 2070 (Fig. 1-6). On average across scenarios and GCMs, suitability scores decreased by 59% for
- 222 P. inexpectata (Fig. 7) and 73% for M. aurantiaca at the presence points (Fig. 8). For M. aurantiaca, only
- 223 one scenario out of 30 showed high suitability values within the current range (i.e. the MIROC5 GCB,
- 224 RCP85, Chelsa baseline, restricted background).
- 225 In Reunion Island, predictions were the most variable between GCMs, but most ensemble models predicted a
- 226 low climate suitability across the island. The highest suitability was found for the MIROC5 GCM,
- 227 distributed along a band at higher altitudes (Fig. 1b). However, the MESS analyses indicated novel
- 228 conditions in the most suitable areas, suggesting model extrapolations in these areas for this GCM (Fig.
- 229 S13). According to the response curves (Fig. S14) and the predicted decrease in summer precipitations (Fig.
- 230 S3c-d), climate suitability may have been overestimated.

In Madagascar, predictions varied the most with the background extent. Apart from the restricted background sets, none of the ensemble models identified any suitable area in the future. Nevertheless, we consistently identified a thin forest band in central Madagascar, south the current distribution, as the most suitable area across GCMs and scenarios (Fig. 2b). Models identified clear climatic windows, with no issue related to model extrapolation (Fig. S15, S16).



Fig. 1 Current (a) and projected future (b) environmental suitability for *Phelsuma inexpectata* using Chelsa as baseline climate data, no prevalence setting and the whole island as a background. Black points represent the occurrence data. Future projections (2070) were estimated from three Global Circulation models (GCM) and two RCP scenarios. Left panels represent the most optimistic scenario (RCP26) and right panels represent the most pessimistic scenario (RCP85). Models produced similar results when setting prevalence to



Fig. 2 Current (a) and projected future (b) environmental suitability for *Phelsuma inexpectata* using Chelsa as baseline climate data, no prevalence setting and restricted background. Black points represent the

- 244 occurrence data. Future projections (2070) were estimated from three Global Circulation models (GCM) and
- 245 two RCP scenarios. Left panels represent the most optimistic scenario (RCP26) and right panels represent the
- 246 most pessimistic scenario (RCP85). Models produced similar results when setting prevalence to 0.5 (Equal
- total weights).

248



249 Fig. 3 Current (a) and projected future (b) environmental suitability for *Phelsuma inexpectata* using

250 Worldclim as baseline climate data. Black points represent the occurrence data. Future projections (2070)

251 were estimated from three Global Circulation models (GCM) and two RCP scenarios. Left panels represent

the most optimistic scenario (RCP26) and right panels represent the most pessimistic scenario (RCP85).



Fig. 4 Current (a) and projected future (b) environmental suitability for *Mantella aurantiaca* using the Chelsa baseline climate, no prevalence setting and the whole island as a background. Black points represent the occurrence data. Future projections (2070) were estimated from three Global Circulation models (GCM) and two RCP scenarios. Top panels represent the most optimistic scenario (RCP26) and bottom panels represent the most pessimistic scenario (RCP85). Models produced similar results when setting prevalence to 0.5 (Equal total weights).



Fig. 5 Current (a) and projected future (b) environmental suitability for *Mantella aurantiaca* using the Chelsa baseline climate, no prevalence setting and a restricted background. Black points represent the occurrence data. Future projections (2070) were estimated from three Global Circulation models (GCM) and two RCP scenarios. Top panels represent the most optimistic scenario (RCP26) and bottom panels represent the most pessimistic scenario (RCP85). Models produced similar results when setting prevalence to 0.5 (Equal total weights).

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Fig. 6 Current (a) and projected future (b) environmental suitability for *Mantella aurantiaca* using the Worldclim baseline climate. Black points represent the occurrence data. Future projections (2070) were estimated from three Global Circulation models (GCM) and two RCP scenarios. Top panels represent the most optimistic scenario (RCP26) and bottom panels represent the most pessimistic scenario (RCP85).



Fig. 7 Variation in climate suitability scores at the occurrence points of *Phelsuma inexpectata* under current
climate and 2070 climate. We show the variability in model predictions related to the climate scenario (RCP
2.6 versus RCP 8.5), the Global circulation model (3 GCMs), the background extent (wide versus restricted),
prevalence setting (no setting versus Equal Total Weights, ETW) and baseline climate (Chelsa versus
Worldclim).



Fig. 8 Variation in climate suitability scores at the occurrence points of *Mantella aurantiaca* under current
climate and 2070 climate. We show the variability in model predictions related to the climate scenario (RCP
2.6 versus RCP 8.5), the Global circulation model (3 GCMs), the background extent (wide versus restricted),
prevalence setting (no setting versus Equal Total Weights, ETW) and baseline climate (Chelsa versus
Worldclim).

283

284 Discussion

285 Dramatic and widespread decline in climate suitability

286 We predict a strong decline of climate suitability in the whole current distribution area of both species by

- 287 2070. The subsequent high extinction risk in these species is not surprising, since small occupied areas are
- 288 known to be good predictors of vulnerability to climate change (Pearson *et al.*, 2014), but forecasting models
- were lacking and we contribute to fill this gap. We also predict that few or no zone will be suitable in the
- 290 future across the entirety of their respective regions. If those species are not given the opportunity to adapt.

291 Depending on the Global Circulation model and the scenario, future predictions still identified areas with 292 suboptimal climatic conditions. In Reunion Island, the higher suitability found at ca. 20km west from the 293 current distribution was presumably driven by an increase in precipitation. *Phelsuma inexpectata* is mostly 294 present where precipitation is the lowest (< 500mm during the wet season, Fig., S10), probably because the 295 impact of cyclones—a driver of high mortality rates in *Phelsuma* species in the Mascarene (Vinson, 1975)— 296 is less severe. The identified area coincides with the region with the driest conditions in Reunion Island in 297 the future, according to these GCMs (Fig. S1). In all model sets, the MIROC5 GCM showed a higher-but 298 suboptimal—climate suitability around the southern coast of the island, with the highest suitability along a 299 band following the upper lands. This prediction is likely driven by the increase in overall temperature, along 300 with reduced precipitation. The species lives under the hottest and driest conditions of the island, which fits 301 the thermal requirements of the species for reproduction in captivity (28°C; McKeown, 1993). However, we 302 believe this GCM is questionable because (1) the reduction in precipitation does not reflect the possible 303 global increase in cyclone risk, and (2) the MESS analysis and response curves suggest an overestimation of 304 the suitability scores. Note that this scenario (i.e. RCP 8.5)—which was the least pessimistic for P. 305 *inexpectata*—is of interest to explore the widest range of possibilities in the future, but is considered unlikely 306 (Hausfather & Peters, 2020).

307 In Madagascar, the current distribution is mainly explained by both winter (i.e. the dry season) and summer 308 (i.e., the wet season) conditions with a short interval of suitability (Fig. S15). An important decline in climate 309 suitability may therefore be driven by small changes in either winter or summer conditions. During winter, 310 individuals migrate up the hills to shelter under dead woods and leaves and presumably enter a state of torpor 311 or hibernation (Randrianavelona et al., 2010; Edmonds et al., 2020). The narrow climatic window may 312 represent optimal conditions that minimise the risk of desiccation during this period of low activity. The 313 species is also dependent on summer conditions, with a narrow window of suitable temperature and 314 precipitations. This is highly consistent with a field study which recorded a surface temperature of 20-23°C 315 at the occupied sites (Edwards et al., 2019) and an experimental design which found a decrease in activity 316 when temperature deviates from 21.5°C (Edwards, 2019) with warm and rainy conditions (Fig. S12).

317 Summer corresponds to the period of reproduction, where females lay eggs under dead leaves on slopes, 318 which are likely washed down into temporary pond during heavy rain episodes. This may explain the 319 dependence with summer high precipitation and temperatures. This combination of climatic conditions is 320 unlikely to be met in the future, except for the MIROC5 GCM under the RCP85 scenario (but this scenario is 321 unlikely, Hausfather & Peters, 2020). However, we consistently identified potential suboptimal areas in the 322 central-south of the eastern forest corridor and in one small area in the north east (corresponding to the top of 323 mountains). For both species P. inexpectata and M. aurantiaca, the high level of habitat fragmentation 324 associated to agricultural areas may limit potential distribution shifts, which calls the need for human 325 intervention.

326

327 Sources of uncertainty

328 The largest source of uncertainty was related to the GCM for *P. inexpectata*, and to the background extend 329 for *M. aurantiaca*. We followed a protocol that attempted to mitigate most sources of uncertainty, 330 corresponding to the acceptable standards defined in Araújo et al. (2019). The long term, extensive and 331 repeated efforts dedicated to species sampling enabled to define the current distributions of these two species 332 with high precision. Uncerainty map showed a high level of agreement between model replicates for current 333 projections ()We included the 19 bioclimatic variables for the selection process, which is not recommended 334 in most cases. However, we selected the most biologically meaningful variables among the groups of 335 intercorrelated variables. We used the finest resolution available for climate data (i.e. 30 arc sec) which 336 seems sufficient to discriminate suitable to unsuitable areas at the scale of Reunion Island and Madagascar. 337 The inclusion of high-resolution land use variables enabled to improve the realism of both distributions and 338 provide specific guidelines for conservation applications. In addition, we found a causal interpretation for the 339 selected variables, which, along with the high model performance, brings confidence in credibility of the 340 models. An important limitation is that we did not use scenarios of future land use. This may lead to an 341 overestimation of the available habitat in Madagascar, since the country is under important rates of 342 deforestation (Veilledent et al., 2018). In Reunion Island, most of the natural habitats is incorporated in

343 private properties or public gardens and in steep zones with limited access for agricultural practices. We 344 believe that the apparent stability in agricultural areas would maintain the applicability of our results in the 345 future. The inclusion of a range of different greenhouse-gas emission scenarios and GCMs showed an 346 important uncertainty in the future suitability for P. inexpectata. However, this uncertainty is mostly related 347 to one GCM (i.e. MIROC5), which we assume to be doubtful. The remaining ones consistently identified the 348 most suitable area by 2070. Uncertainty related to model design was mitigated by the limitation of model 349 complexity (with the ESM approach and the post-filtering technique), the removal of collinearity and the 350 testing of a range of input parameters (number of pseudo-absences sets and coss-validation subsets). Model 351 performance was assessed with random partitions for *M. aurantiaca*, while spatial partitions are 352 recommended. However, we argued that this methods is not appropriate for highly localised species due to 353 strong imbalance between spatial blocks. We used multiple evaluation metrics, including discrimination 354 (AUC and TSS) and reliability (i.e. calibration; Boyce index) metrics, all showing a high performance 355 overall. We did not account for species dispersal ability, because the purpose was not to predict potential 356 shifts, but to assess how suitable will be the climate in the future in order to identify candidate sites for 357 restoration and translocation. Both species have low dispersal ability and their habitat is highly fragmented. 358 Therefore, the potential for distributional shifts may be strongly limited and future projections must not be 359 interpreted as the future distributions of our study species. An important limitation may be the absence of 360 empirical knowledge on species thermal tolerances or other features of their climatic niche, adaptability and 361 plasticity, which prevents us from determining whether our future predictions underestimated the 362 environmental suitability.

363

364 A glimmer of hope

365 Despite the predicted low suitability in climate conditions, it is possible that species persist under changing 366 conditions through adaptation or plasticity (Chevin *et al.*, 2010; Hoffmann & Sgró, 2011). This hypothesis is 367 supported by the persistence of two introduced populations of *P. inexpectata* away from their current range in 368 suboptimal environments (Fig. 1). Their persistence may result either from physiological or behavioural

369 adaptation while benefiting from a combination of urban island effect and access to microclimate refuges in 370 anthropogenic structures, as it is the case for *Phelsuma grandis* in Florida (Fieldsend et al., under review). It 371 is also frequent that the current distribution of a given species represents only a fragment of its climatic niche 372 (e.g., Guisan et al., 2014). For instance, experimental design on M. aurantiaca showed no expression of a 373 thermal stress during periods of extreme heat (Edmonds et al., 2015). Therefore, predictions of climate 374 suitability may underestimate the bounds of our model species niches. Further studies are needed to better 375 characterise their climatic niche, and explore potential adaptive and plastic responses to changing conditions 376 in these species and, more generally, in other threatened narrow-ranging species. Meanwhile, we encourage 377 practitioners to implement conservation measures to grant those species a chance to adapt and persist. 378 Nevertheless, the rate of climate change is generally faster than that of animal adaptive responses (Radchuk 379 et al., 2019), which stresses the need for urgent actions. In the tropics, extinction risks may be greatly 380 reduced with the development of land conservation programs provided climate change is mitigated by inter-381 governmental actions (Hannah et al., 2020).

382

383 Conservation application

384 In Reunion Island, we identified two potential areas suitable for habitat restoration of Phelsuma inexpectata 385 around the south-western coast and along a mid-altitude band. Despite climate conditions are predicted to be 386 suboptimal, we believe these might represent the best options to ensure the long-term persistence of the 387 species. Habitat restoration should be focussed on natural habitats invaded by non-native plants and in 388 urbanised areas. Habitat restoration will consist in promoting the spread of native species (Pandanus utilis, 389 Latania lontaroides, Scaevola taccada and Psiadia retusa) for which the species depends on (Bour, 1995). 390 Finally, given the high level of habitat fragmentation, it is likely that the design of translocation programs 391 will also represent a relevant aspect to consider for management.

In Madagascar, the area with the highest future climate suitability is located at the south of the currentdistribution, along the eastern rainforest corridor between the current distribution area and the north-west of

394 the Vatovavy-Fitovinany region. The forest cover within the distribution range of *M. aurantiaca* has 395 experienced a continued decline, regardless of the conservation status of the inhabited area (Piludu et al., 396 2015; Vieilledent *et al.*, 2018). We recommend to reinforce the level of protection and to improve 397 governance and management of the Mangabe (Moramanga region) and the Marolambo (Vatovavy-398 Fitovinany region) reserves. This can be achieved by promoting the development of alternative economic 399 solutions through the development of valuable and sustainable activities that mitigate the rate of conversion 400 of natural areas, the long-term management of soil fertility, and by considering the development of 401 ecotourism. This species was included into a program that enabled to develop amphibian husbandry 402 capacities in Madagascar, and that succeeded to establish captive bred colonies for this species 403 (Rakotonanahary et al. 2017). Efforts of captive husbandry should be maintained (and possibly expanded to 404 other species) and we encourage the design of translocation programs accounting for both local habitat 405 characteristics and future climate suitability. Further study of biotic interaction between M. aurantiaca and 406 other amphibians is needed to assess the consequences of its introduction.

407

408 Are all endangered narrow-ranging species doomed to extinction?

409 Narrow-ranging species usually live under very specific environmental conditions and are the most 410 vulnerable to climate change (Botts et al., 2013). Not only they are more prone to face distribution shifts and 411 range contractions, but they might also disappear due to the strong alteration of their climatic envelope 412 throughout their entire region. This may be the case for most narrow-ranging species with a specialised niche 413 in regions with heterogeneous climates and high levels of endemism such as Madagascar but also Central 414 America, South East Asia, tropical islands and more generally in tropical rainforests and mountains (Kier et 415 al., 2009). In such regions, small shifts in climatic conditions may induce important changes in local 416 environmental suitability for endemic—often specialised—species (e.g., Raxworthy et al., 2008). The risk is 417 greater in tropical regions where species live closer to the upper bound of their thermal tolerance (Tewksbury 418 et al., 2008; Sekercioglu et al., 2012; Dubos et al., 2019). Species that are already under threats are at the 419 greatest risk because of a large array of synergistic effects (Sekercioglu et al., 2012). This might be the case

420	for our two model species, which are threatened by habitat destruction and invasive species (Dubos, 2013;
421	Piludu et al., 2015). Among the few Critically Endangered species for which the impact of climate has been
422	investigated, most predict severe reductions in the climate suitability (e.g., Alamgir et al., 2015; Zhang et
423	al., 2020), with the suitable range of the giant salamander Andrias davidianus predicted to decrease by more
424	than two thirds by 2050 (Zhang et al., 2020). This species has faced a number of threats and climate change
425	may be leading to imminent extinction despite a larger distribution range. Despite recent efforts developed
426	for the monitoring of rare species in the tropics (e.g., Dubos et al., 2020), there is still an important lack of
427	species occurrence data in these regions (Feeley & Silman, 2011). With the newly available high-resolution
428	climate and land use data, the spectre of eligible species for SDMs has enlarged. We urge filling this data
429	void by starting to assess the effect of climate change for narrow-ranging species at broader taxonomic
430	scales, promote field investigations, assess species thermal requirements, and develop proactive conservation
431	actions.
432	
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436	
437	Conflict of Interest Statement
438	The authors declare no conflict of interests.
439	
440	Data availability statement
441	The dataset and the scripts used in this analysis are available as R objects in the supporting information.
442	

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