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3	Conservation translocations of fauna in Aotearoa New Zealand: a review
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22	Running head: Conservation translocations in Aotearoa New Zealand
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24	Abstract
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26	There have been numerous declines and extinctions of native fauna in Aotearoa New Zealand
27	since human settlement. Against this background of loss there have been remarkable
28	advances in conservation management, including the use of conservation translocations to
29	reduce extinction risk and restore depauperate ecosystems. Here, we review conservation
30	translocations in Aotearoa New Zealand. Our review assembles knowledge from Aotearoa
31	New Zealand's rich history of faunal translocations and describes six key considerations for
32	successfully establishing translocated populations: 1) What values will be met by a
33	translocation? 2) What is the natural and conservation history of the translocation candidate?

3) Does the release site habitat match that of the proposed source population, and if not, why

35	is the release site considered appropriate and can management ameliorate differences? 4) Will
36	dispersal be a problem? 5) Will genetic management be required and how realistic is it that
37	this management will be implemented? 6) What do future developments mean for the
38	management of translocated populations? We discourage a focus on any single element of
39	translocation planning but rather encourage all people involved in translocations, particularly
40	decision makers, to explicitly recognise that successful translocations typically have multiple,
41	values-based objectives. We also support recommendations that the principles of good
42	translocation decision-making are embedded in government policy.
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- 44 Keywords: Conservation translocation, reintroduction, restoration, decision making
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46 Introduction

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48 There have been numerous declines and extinctions of native fauna in Aotearoa New Zealand (hereafter, Aotearoa) following human settlement (Caughley 1989; Holdaway 1989). For 49 50 example c. 50% of all native bird and frog species have become extinct since the first humans arrived (Caughley 1989; Holdaway 1989), and remaining extant, native species show varying 51 52 levels of vulnerability to exotic pests (Innes et al. 2010). This history of extinction and drastic reduction in population size and range is recounted in Māori whakataukī (proverbs) such as 53 54 "Ko te huna i te moa - destroyed like the moa", (Wehi et al. 2018) and by Diamond (1984) who stated that "New Zealand doesn't have an avifauna, just the wreckage of one". 55

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Despite these losses there have been remarkable advances in conservation management, 57 including the use of conservation translocations, defined as the intentional movement of 58 animals from one place to another for a conservation benefit (referred to as "translocation" 59 60 hereafter). The increasing use of translocations in Aotearoa has been enabled by advances in large-scale pest eradication and control (here, pest primarily refers to exotic mammalian 61 62 predators and competitors, but also includes other unwanted harmful vertebrates, 63 invertebrates, plants and pathogens). Multi-species eradications have been completed on large and small islands (Towns & Broome 2003). Fenced sanctuaries provide islands of virtually 64 pest-free habitat on the main islands of Aotearoa, within which most significant pests are 65 absent most of the time (Innes et al. 2019). Such sanctuaries are also often isolated from 66 adjacent unmanaged habitat (Innes et al. 2019). The number of unfenced mainland sites under 67 varying forms of protection is also increasing every year (Innes et al. 2019) and the 68

Government's 2016 announcement of Predator Free 2050 will likely lead to an increase in
control of some pests, especially rats (*Rattus* spp.), stoats (*Mustela erminea*) and possums
(*Trichosurus vulpecula*). This will result in a gradient of pest density from areas with
complete eradication/zero density, to areas with lower densities than are currently achievable.
Surprisingly, there has been little detail about what a predator-free Aotearoa might look like,
but implicit is the goal of exchanging pest biomass for native and endemic biomass.

- 75 Translocations are an important tool for achieving this goal.
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77 Aotearoa conservationists are very good at doing translocations to pest free islands. However, 78 progress is also being made in the translocation of some forest birds to sites where community conservation initiatives have restored the available habitat through pest control, 79 planting and translocations. Many such projects have successfully re-established high-density 80 populations of native wildlife, particularly forest birds. A critical limitation is that most of 81 82 these sites are small (c.100–1000ha), and mice (*Mus musculus*) have rarely been eradicated, or even effectively controlled, which is problematic for the recovery of endemic lizards, 83 84 amphibians, invertebrates, bats and some threatened plants. In contrast, the bulk of our biodiversity is contained within vast areas (1000s of hectares) of back country conservation 85 86 estate which is both harder to protect and harder for the public to engage with. The Department of Conservation (DOC) "Tiakina Ngā Manu/Battle for our Birds" programme is 87 achieving impressive conservation gains by controlling pests over huge areas of habitat (c. 88 500 000 ha in 2022), in conjunction with species-focussed mainland recovery programmes 89 90 (e.g. kakī / black stilt (Himantopus novaezelandiae) and kākāriki karaka / orange-fronted parakeet (Cyanoramphus malherbi)). Nevertheless, biodiversity continues to decline in vast 91 92 tracts of land, especially non-forested habitats, that remain unprotected.

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94 The current situation on the main islands of Aotearoa is neatly captured by Caughley's (1994) small population and declining population paradigms. Our small, protected populations, 95 which by definition include all translocated populations, are subject to the risks of being 96 small, including pest incursions, dispersal, extreme weather events, novel pathogens, and loss 97 98 of genetic diversity. In contrast, many of our large mainland populations are declining because of the pervasive impacts of pests. The ongoing tension in conservation management 99 in Aotearoa lies in deciding how to allocate resources to maintain small populations, because 100 this seems generally easier and currently achievable, while also continuing to manage the 101 large areas of habitat on the main islands that contain the bulk of our biodiversity, a much 102

harder challenge largely dependent on the continued use of aerially applied toxins. Bothapproaches are necessary.

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Small, intensively protected populations provide insurance against further declines and can 106 serve as source populations for natural colonisation of, or translocation to, pest-free habitats 107 108 when these become available. Such sites also provide a glimpse of what a predator-free Aotearoa might look like and are critical tools for engaging the general public in conservation 109 management (Parker 2008). In contrast, ongoing pest control in large mainland areas is 110 111 essential for protecting biodiversity not readily protected on islands, or in small intensively protected areas. When these large mainland areas are released from the pervasive effects of 112 pests (primarily a question of social licence and technical advances) they will further buffer 113 threatened species against the challenges of small population size. 114

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116 In this paper we use our collective experience as practitioners and conservation scientists to focus on small population management in Aotearoa, specifically translocated populations that 117 118 have been established following local extinction. The DOC translocation proposal document captures the principles of sound translocation practice, including those described in the IUCN 119 120 "Guidelines for reintroductions and other conservation translocations" (IUCN 2013). However, these principles are not currently captured in DOC policy (Parliamentary 121 Commissioner for the Environment 2017), which sometimes compromises the ability of DOC 122 to assess and approve translocation proposals. This is important, especially as we move 123 beyond translocations to typical sites (islands and relatively small protected mainland areas), 124 towards release sites with much more uncertainty, e.g. very large areas (1000s of hectares) of 125 contiguous habitat, and also urban (van Heezik & Seddon 2018), and rural landscapes. 126

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128 We want more successful translocations and we think the best way to achieve this is to explicitly define a clear set of measurable, a priori, fundamental, values-based objectives at 129 130 the outset of each translocation (Box 1; Ewen et al. 2014, 2023). Common biological values 131 include fundamental objectives such as reducing extinction risk and restoring depauperate ecosystems. However, mana whenua and local community values can be equally critical and 132 central to translocation success. Therefore, achieving project objectives requires careful and 133 measurable evaluation of all factors that might contribute to translocation success, and an 134 understanding of the species and people specific time scales over which such factors might 135 act, rather than focussing on single factors and arbitrary timeframes. We also note the 136

137 increasing demand for translocations and that some might proceed with different fundamental

- 138 objectives to those posited above. However, a translocation cannot be considered successful
- 139 if a population fails to establish, even though uncertainty means that this sometimes happens.
- 140 Conservation translocations are not easy and many fail (Miskelly & Powlesland 2013) but
- 141 these failures are informative for future efforts to establish populations.
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We draw together knowledge that has been gained from the rich history of fauna
translocations in Aotearoa and outline six key considerations (Box 1) for translocation
decision-making: 1) What values will be met by doing a translocation? 2) What is the natural
and conservation history of the translocation candidate? 3) Does the release site habitat match
that of the proposed source population and, if not, why is it appropriate and can management
ameliorate differences? 4) Will dispersal be a problem? 5) Will genetic management be

required and how realistic is it that this management will be implemented? 6) What do futuredevelopments mean for the management of translocated populations?

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These key considerations can be applied to most fauna and we apply them to several species and species groups that have been translocated in Aotearoa (Table 1). The examples we use, and the perspectives we bring, mainly relate to terrestrial birds, simply because these have been translocated for conservation more frequently than other taxa (Miskelly & Powlesland 2013; Swan et al., 2016; Rayne et al., 2020). However, significant information gaps exist, even for bird translocations.

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1. What values will be met by doing a translocation?

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Translocations are most frequently conducted on public land administered by national or 161 local government, and they usually involve the use of at least some public money. 162 Accountability for the management of translocated species is vested in government, i.e. DOC, 163 and is bound by a commitment to give effect to Te Tiriti o Waitangi / The Treaty of 164 Waitangi. Therefore, at minimum, there is a legal requirement to consult with mana whenua 165 166 (iwi, hapū and whānau with customary authority over an identified area) about every translocation, including ongoing management of source populations, translocated 167 populations, and release sites. However, engagement may extend far beyond economic and 168 legal obligations, especially where translocations emerge from trusted relationships and 169 recognise the deep connections that mana whenua, and local communities, share with 170

populations and places (Bioethics Panel 2019). Translocations can therefore contribute to
realising multiple fundamental objectives, including those that are responsive to the needs
and aspirations of mana whenua and local communities (Parker 2008; McMurdo Hamilton et
al 2021; Fischer et al. 2023; Parker, Parlato et al. in press).

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176 The challenge is that the values and objectives underlying translocations usually seem obvious to the project instigators, managers and decision makers. However, they might 177 overlook key fundamental objectives of mana whenua and local communities. For example, a 178 179 manager trained in modern science might see a translocation as an opportunity to reduce 180 extinction risk or restore a depauperate ecosystem. In contrast, mana whenua might see it as an expression of rangatiratanga (sovereignty, authority, self-determination), kaitiakitanga 181 (guardianship), and the restoration of mauri (not easily defined but sometimes translated as 182 "life essence") (McMurdo Hamilton et al 2021, Fischer et al. 2023; Parker, Parlato et al. in 183 184 press). A community conservation group or private landowner might simply want a particular species living in their area. These objectives might seem similar, but this should not be 185 186 assumed, nor will they necessarily be measured in the same way. This is critical because a review by Ewen et al. (2014) found that the setting, reporting and, measurement of objectives 187 188 is highly variable among reintroduction programmes. Furthermore, most are rooted in 189 Western science with little mention of other values. Ewen et al. (2014) also noted that 190 fundamental objectives (the things we want, e.g. reduce extinction risk) were often mixed with means objectives (how we get what we want, e.g. do a translocation), and are not 191 192 measured in an appropriate way, nor even explicitly stated. In the case of Predator-Free NZ, the name states a means objective (and has led some to believe the project to be short-193 sighted), but the fundamental objective is clear: a landscape dominated by indigenous 194 biodiversity (Department of Conservation 2020). 195

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197 Given this complex decision environment Ewen et al. (2014, 2023) characterise a

198 conservation translocation as a sequence of decisions, and argue that poor planning,

implementation, and monitoring is a consequence of not approaching the decision-making

200 process in a deliberate and rational manner. They, along with several other authors, advocate

a more structured approach to decision making (Maguire 1986; McCarthy et al. 2012;

202 Converse et al. 2014). Structured decision making is an iterative process whereby uncertainty

is addressed by 1) defining clear objectives and how they will be measured; 2) identifying a

range of possible management alternatives; 3) predicting the outcomes of the chosen

management alternatives relative to the stated objectives; 4) evaluating trade-offs and
uncertainty; 5) implementing the optimal management alternative and monitoring its results
(Fig. 1) (Gregory et al. 2012; Ewen et al. 2014, 2023). This approach to decision making has
been characterised as "*a formalisation of common sense for decision problems which are too complex for informal use of common sense*" (Keeny 1982).

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Structured decision making is useful only if all people with connections to, or who might be 211 impacted by, a translocation are directly involved in setting fundamental and means 212 213 objectives for the project, and then deciding between management alternatives as to how these might be achieved. For example, translocation planning for pekapeka / short tailed bats 214 (Mystacina tuberculata) was initiated at Te Kiri marae alongside Ngāti Manuhiri who led the 215 216 kõrero on a mātauranga Māori (Māori knowledge, wisdom) fundamental objective for assessing translocation options (McMurdo Hamilton et al 2021). Similarly, recovery planning 217 218 for the kuaka / Whenua Hou diving petrel (Pelecanoides whenuahouensis) was initiated on Takutai o Te Tītī marae with Kai Tahu seeing kuaka translocation as one means to express 219 220 rangatiratanga and exercise kaitiakitanga (Fischer et al. 2023). On Rēkohu / Wharekauri / Chatham Islands the translocation of karure / kakaruia / black robins (Petroica traversi) to 221 222 reduce extinction risk is viewed by Moriori as consistent with their principles and values. 223 Black robin translocation also recognises Ngāti Mutunga o Wharekauri as Treaty Partners and provides a means for the broader Chatham's community to reconnect to black robins, a 224 vital source of local identity (Parker, Parlato et al. in press). 225

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Ultimately, meaningful engagement and decision sharing with mana whenua, and local 227 communities, provides a means to deepen support, interest, and engagement in conservation. 228 However, resourcing is often limited for genuine relationship-building, given the substantive 229 230 costs, time and energy needed (e.g., for hui (meetings) and site visits). Where translocations are initiated by DOC they might cover this cost (Fischer et al. 2023). But translocations 231 232 initiated outside of DOC can result in poorly resourced community conservation groups asking poorly resourced mana whenua for time and energy. It is difficult to know how to 233 resolve this, other than increasing funding bids to cover all translocation costs, although it 234 could also be argued that these initiatives are contributing to national conservation objectives 235 and might therefore deserve government assistance. 236

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2. What is the natural and conservation history of the translocation candidate?

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One obvious starting point for setting biological objectives and informative performance 240 measures is understanding the natural and conservation history of the candidate species 241 (Ewen et al. 2023). For example, North Island (NI) toutouwai / NI robins (Petroica longipes) 242 have persisted at sites on the main islands of Aotearoa with no predator management whereas 243 244 NI tieke / NI saddlebacks (Philesturnus rufusater) have been extinct on the mainland for >120 years (Heather & Robertson 2015). These two species clearly differ in their ability to 245 tolerate pests and therefore require different performance measures for pest control (a means 246 247 objective), even though the fundamental biological objectives for translocating these species, 248 typically to reduce extinction risk or restore a depauperate ecosystem, are often the same 249 (Table 1).

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However, it can be extremely difficult to determine why a translocation failed. One way is to 251 252 model vital rates from another species to estimate the vulnerability of the focal species to pests. For example Parlato and Armstrong (2018) used NI toutouwai data to predict rat-253 254 tracking indices that might correlate with NI treke translocation success. Alternatively, factors other than pests might lead to translocation failure. For instance, of nine korimako / 255 256 bellbird (Anthornis melanura) translocations (Miskelly & Powlesland 2013) only one appears to have been successful. While several factors might have contributed to these failures it is 257 unequivocal that dispersal from the release site has been a critical factor, even at sites where 258 some breeding occurs. Given such low success it is questionable whether any further 259 260 translocations of korimako are justified unless there is a significant change in methods or understanding, especially given their ability to naturally recolonise protected sites (Brunton et 261 al. 2008). Clearly, if a species has rarely or never been translocated then the outcomes of 262 previous translocations are not useful indicators of future outcomes. In these cases, the 263 264 translocation of other species, along with the ecology and conservation history of the candidate species, will have to be assessed against extirpation history, vulnerability to pests, 265 266 dispersal abilities and other habitat requirements. However, there will naturally be a higher degree of uncertainty regarding establishment and persistence of the translocated population. 267 268

269 270 3. Does the release site habitat match that of the proposed source population? If not, why is the release site considered appropriate and can management ameliorate differences?

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Conservation translocations are typically, but not always, carried out within the former range
of a species, i.e. are reintroductions (IUCN 2013), following local extirpation, and where
natural recolonisation is unlikely on a time scale acceptable to site managers, mana whenua
and local communities. Clearly, the conditions that we understand/predict a species needs to
persist must be present in the release area. However, these conditions might also be provided
by management, for example the provision of supplementary food to translocated hihi
(*Notiomystis cincta*) (Ewen et al. 2013).

280

281 Unfortunately, the concept of habitat is often misused and poorly defined in translocation planning (Stadtmann & Seddon 2018). Here, we use the definition of Hall et al. (1997), in 282 describing habitat "...as the resources and conditions in an area that produce occupancy -283 including survival and reproduction – by a given organism." This includes all physical (e.g. 284 climate, aspect) and biological (e.g. predators, vegetation associations, landscape 285 286 connectivity) aspects of an area where a species lives. Habitat quality refers to "...the ability of the environment to provide conditions appropriate for individual and population 287 persistence" (Hall et al. 1997), specifically survival, reproduction and population growth. 288 Habitat quality is a continuous variable, ranging from low quality to high quality, and can be 289 290 very difficult to define explicitly, although there are useful proxies (Hall et al. 1997). The finite rate of increase (λ) is the most direct measure of habitat quality, assuming density 291 292 dependence and genetic quality are accounted for. The most essential pre-requisite for translocation success is that λ is > 1 at low density, as the population will otherwise decline to 293 294 extinction. High quality habitat is typically perceived as places where animals formerly 295 occurred. However, habitat conditions need not replicate past states if they are predicted to 296 allow λ to be > 1 (Table 1).

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298 Pests are nearly always considered in translocation planning but are rarely explicitly defined as a habitat variable in Aotearoa, where discussions of habitat quality have focussed on 299 vegetation associations that animals are either known or assumed to rely on for survival, 300 while recognising that remnant populations do not necessarily survive in high quality habitat 301 302 (Griffith et al. 1989). However, any assessment of habitat quality in Aotearoa must consider the presence and density of pests because they have such a critical impact on the survival of 303 so many native and endemic species (Table 1; Innes et al. 2010; Richardson et al. 2014). 304 While other biological and physical habitat variables, especially vegetation associations, are 305

306 clearly essential, effective pest control is almost always a prerequisite for translocated307 populations to establish and persist.

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In Aotearoa, current management of mammalian pests includes three major regimes of 309 control: 1) total eradication on offshore islands; 2) Maintenance of pests at "zero density" 310 311 within fenced mainland sites, i.e. key pests are absent most of the time but when present they are quickly detected and removed; 3) Suppression of pest densities in unfenced mainland 312 areas relative to unmanaged sites (Byrom et al. 2016). These are not mutually exclusive and 313 314 there is often overlap between them. For example, peninsula fences, such as at Tāwharanui Open Sanctuary, are leaky but have extensive areas of pest control outside the fences. It's 315 hoped that this reduces incursions while also providing some protection for animals that 316 317 disperse outside the fence.

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319 Pest densities at the release site must be within the tolerance of the translocated species (Table 1). For example, NI toutouwai can persist with moderate levels of ship rats (Rattus 320 321 *rattus*) but will have higher survival and reproduction rates if rats are reduced to low levels (\leq 5% tracking tunnel indices) before each breeding season, with mustelid control also likely to 322 323 be beneficial. NI toutouwai persist at some sites with ship rat tracking indices of > 25%, but female survival, reproductive output and ultimately population growth are reduced (Parlato & 324 325 Armstrong 2012, 2013). As well as reducing the likelihood of population persistence, slow population growth and loss of founders will increase the loss of genetic diversity. In stark 326 327 contrast, the current distribution of species such as tieke, hihi, and red-crowned kākāriki (Cyanoramphus novaezelandiae) indicates they are much more vulnerable to pests, as they 328 currently persist only in sites where pests have either been eradicated or reduced to zero 329 density. A further challenge when making translocation decisions is that the impact of 330 varying densities of pests is well understood for only a few bird species, poorly predicted for 331 many others, and virtually unknown for most invertebrates, lizards, amphibians and bats 332 333 (Table 1). For example, pest thresholds on the mainland, and population growth in response to pest control, have only been demonstrated for Otago skinks (Oligosoma otagense) and 334 grand skinks (Oligosoma grande) (Reardon et al. 2012), just two of 106 endemic lizard 335 species. 336

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Further habitat variables, including climate, altitude, aspect, and soil type will be associatedwith vegetation and might shift habitat quality from high to low, i.e. decrease the probability

of establishment and persistence, depending on the needs of the translocated species and their 340 ability to adapt to variable conditions. This might be especially difficult at sites that 341 experience climatic extremes relative to those with more benign conditions. Climate change 342 might also cause high-quality habitat to become low quality in the future. Furthermore, the 343 impact of these variables is not consistent across species. For example, NI toutouwai and NI 344 mātātā / fernbird (*Poodytes punctatus vealeae*) are evidently flexible in their habitat 345 requirements as they occupy a broad range of habitats and have been successfully 346 translocated between very different habitats. Productivity and population growth has varied 347 348 between sites, suggesting that some are better than others (Parlato & Armstrong 2012, 2013; KAP unpublished data), but they clearly tolerate a range of habitats during establishment and 349 persistence. In contrast, species such as hihi need protection from mammalian pests but the 350 vital rates of translocated hihi populations, and the fact that most require supplementary food 351 to establish and persist, indicate that there are also other currently unknown habitat 352 353 requirements (Ewen et al. 2013).

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355 Translocating animals to a habitat similar to their current habitat is likely to have a greater chance of success than translocating them to a different habitat. For instance Parlato and 356 357 Armstrong (2012, 2013) showed that translocation of NI toutouwai between habitats with 358 similar pest assemblages and vegetation associations had a small advantage over translocations between contrasting habitats. The similarity of the source and release site, the 359 objectives of the translocation, and the risk profile or level of uncertainty associated with the 360 translocation will also influence decisions about health screening. For example, translocations 361 between two mainland sites and/or relatively close inshore islands will have a relatively low 362 disease risk because their pathogen communities are probably similar (Sainsbury & Carraro 363 2023). In contrast, translocations between distant sites with different habitats could be 364 relatively high risk, especially if the recipient site has species that could be vulnerable to 365 novel pathogens. Ideally, there should also be an understanding of potential pathogen impacts 366 367 on the translocated species, and on conspecifics and heterospecifics at the release site, and/or 368 a documented history of health screening to inform decisions about health management (Parker et al. 2006; Ewen et al. 2007; Ortiz-Catedral et al. 2011; Ewen et al. 2012; Massaro et 369 al. 2012; Sainsbury & Carraro 2023). Unfortunately, this information is usually lacking or of 370 poor quality. 371

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4. Is dispersal a likely impediment to establishment and persistence?

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Individuals translocated to a managed site must stay there rather than dispersing into adjacent 375 unmanaged habitat where their likelihood of persistence will be much lower, or, in many 376 cases be zero. Habitat connectivity, and the ability for species to disperse between habitat 377 patches, is typically seen as a positive landscape feature and a desirable management 378 379 objective. However, dispersal from managed release sites into adjacent unmanaged areas appears to be an important cause of failure for many translocations (Richardson et al. 2014). 380 Dispersal generally affects population growth at two levels. First, post-release dispersal 381 382 following the initial release can cause the loss of individuals from the founding population, thereby reducing the probability of establishment and persistence. For example, an analysis of 383 14 reintroduced NI toutouwai populations showed that habitat connectivity was a key factor 384 determining individual establishment following translocation, with individuals released at 385 highly connected sites having a lower establishment probability than those at less connected 386 387 sites, such as islands or isolated forest patches (Parlato & Armstrong 2013). Second, natal dispersal, i.e. the loss of juveniles raised at the release site, can also reduce establishment and 388 389 persistence if juveniles move from managed to unmanaged sites (Richardson et al. 2014). Critically, the interaction of post-release dispersal and natal dispersal can limit population 390 391 growth, erode genetic diversity, and reduce the likelihood of the long-term persistence of a 392 translocated population.

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The dispersal of translocated species from release sites is highly variable and sometimes 394 395 difficult to predict (Table 1; Richardson et al. 2014; Innes et al. 2022). For instance, some birds are very strong dispersers regardless of habitat connectivity. These include korimako, 396 miromiro / tomtit (Petroica macrocephala), and red-crowned kākāriki (Table 1; Parker et al. 397 2004; Brunton et al. 2008; Ortiz-Catedral 2010) whereas others, such as NI toutouwai and NI 398 399 tieke, are less likely to disperse from sites with low connectivity (Table 1; Newman 1980; Richard & Armstrong 2010). The connectivity of the release site to surrounding unprotected 400 401 habitats therefore varies according to the dispersal ability of the species in question, making connectivity difficult to measure. The shape of the relationship between dispersal ability and 402 403 connectivity is also unknown for all species. However, we hypothesise that it will show a similar shaped curve as seen for other sources of mortality or loss to a managed population, 404 e.g. increasing predator density (Fig. 2). Many species, including some with relatively strong 405 dispersal abilities, rarely leave isolated sites such as islands or forest patches surrounded by 406 407 pasture (Table 1). In contrast, species with poor dispersal abilities can move out of protected

areas if connected to habitat that the species will willingly move through (Table 1; Richard &
Armstrong 2010), although this is likely to be a greater problem for birds and bats than
reptiles, amphibians and invertebrates.

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The best way to manage dispersal in contiguous landscapes is to manage as large an area as possible, including potential dispersal routes, through an integrated landscape management approach (Richardson et al. 2014). However, it is not currently known how big a site needs to be to accommodate post-release and natal dispersal in most species, and it will often be difficult, too expensive, or simply not feasible to protect very large sites. This currently limits the ability to translocate some species to large sites.

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A variety of alternative approaches have been used to try to reduce dispersal, albeit with 419 variable results. Holding animals in captivity at the release site (delayed release) has been 420 421 tried with many taxa, and many sites, but the results have been extremely variable. They have been generally ineffective for wild to wild releases, but sometimes useful when releasing 422 423 captive-reared animals (Parker, Dickens et al. 2012; Smuts-Kennedy & Parker 2013; Richardson et al. 2014, 2015; Parker et al. 2015). Supplementary feeding has also been used 424 425 with success for some species at some release sites (e.g. kākā (Nestor meridionalis), pāteke / brown teal (Anas chlorotis) Rickett et al. 2013) but has been less useful for others (e.g. hihi, 426 427 Richardson et al. 2014). Acoustic anchoring (playback of pre-recorded calls) was attempted with NI kokako (Callaeas wilsoni), NI toutouwai, and popokotea / whitehead (Mohoua 428 429 albicilla) in Aotearoa but was not effective (Leuschner 2007; Molles et al. 2008; Bradley et 430 al. 2011).

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Another option for mitigating the impact of dispersal in the establishment phase is the release 432 of large numbers of individuals, either in one big release or in a series of smaller releases 433 over several years. This is intuitively appealing but is rarely effective and there are many 434 examples where large numbers of animals have been released but the translocation has failed 435 (Miskelly & Powlesland 2013). For instance, single popokotea translocations of 40-100 birds 436 437 to isolated managed sites of up to 3300 ha have typically been successful. However, the translocation of 653 birds over 12 years into a 2450 ha protected block within the 17 000 ha 438 Waitākere Ranges appears to have been unsuccessful (KAP unpublished data). The 439 relationship between release group size and establishment is also unclear. This is because 440 high-quality sites where translocations are successful following the release of large numbers 441

442 of animals could have been equally successful if fewer animals were released. In contrast,

- 443 managers typically release fewer animals when they have less confidence in a site, creating a
- reporting bias towards success with larger releases (Armstrong & Seddon 2008; Armstrong &
- 445 Wittmer 2011). Translocating large numbers of animals in the knowledge that many will die
- also raises significant welfare and ethical issues, and may strain relationships, especially
- 447 where translocation is not essential for the management of the species in question.
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Ultimately, the best way to reduce dispersal is to release animals at isolated or relatively 449 450 isolated sites. However, the great challenge with managing dispersal is that we want translocated species to establish populations within large contiguous sites, and we want 451 individuals to be able to disperse freely between sites. This will protect against the problems 452 of populations being small and will largely remove the need for reinforcement translocations 453 for genetic management, i.e. natural dispersal via safe dispersal corridors will essentially act 454 455 as passive meta-population management. It will also provide new opportunities for populations in smaller sites. In the current environment safe corridors generally mean 456 457 protection from pests but as pest control improves other habitat variables will become more important. For example, what size, shape, and structure do corridors need to be to cater for as 458 459 wide a range of native species as possible? We recommend that the ability of animals to 460 safely disperse from intensively managed areas should be a performance measure for initiatives such as Predator Free 2050. Furthermore, dispersal pathways should be 461 incorporated into decisions about which landscapes to protect first. 462

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5. Will genetic management be required and how realistic is it that this will be implemented?

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Genetic diversity maintains evolutionary potential by providing populations with long-term 467 capacity to adapt to changing conditions (Frankham et al. 2017; Forsdick et al. 2022). All 468 populations lose genetic diversity over time because of chance events, i.e. genetic drift. 469 470 However, small populations are especially vulnerable because they accumulate the mutations 471 required to replace lost alleles so slowly (Frankham et al. 2017; Forsdick et al. 2022). Inbreeding (mating between relatives) in small populations can also reduce survival and 472 reproductive success through inbreeding depression which, in turn, threatens population 473 persistence (Frankham et al. 2017; Forsdick et al. 2022). Translocated populations are 474 475 particularly susceptible to genetic drift and inbreeding depression. They also often impose a

genetic bottleneck on new populations because the founders only represent a portion of the
source population's genetic diversity. This effect is further compounded because the number
of founders that recruit and contribute to the new population is usually smaller than the
number released. In addition, translocated populations at small sites will always be small.

481 Therefore, careful consideration of genetic objectives is needed to minimise the loss of genetic diversity, to select a source population or populations, to define ongoing genetic 482 483 management, and to predict the genetic diversity of the translocated population (Weeks et al. 484 2015). It is also essential to clarify whether genetic objectives are fundamental or means based. For example, we are rarely interested in maintaining genetic diversity for its own sake, 485 i.e. as a fundamental objective. Rather our interest in genetic diversity is usually as a means 486 objective that contributes to the long-term persistence of the translocated population by 487 maintaining evolutionary potential and reducing extinction risk. If this is the case, then a 488 489 means objective might be to release enough animals to maximise genetic diversity in the 490 founders and therefore the evolutionary potential of the new population.

491

Alternatively, there are many reasons why small (≤ 100 individuals) translocated populations 492 493 are created, including because only small numbers of animals exist, ease of management, for advocacy, or simply that only small sites are available for release. In these cases, genetic 494 495 means objectives might include informed reinforcement translocations to maintain genetic diversity across a metapopulation. All management involves trade-offs. For example, the best 496 497 source populations are typically large, genetically diverse, and have no history of tight (<40 -100 individuals) and/or long-term bottlenecks, although bottlenecks are sometimes acceptable 498 499 if they were of short duration (Boessenkool et al. 2007). However, while obtaining animals 500 for translocation from a small, inshore island or fenced sanctuary might be relatively easy and 501 cheap, populations from such small sites are likely to have lower genetic diversity than those obtained from a large source population. Furthermore, the ongoing maintenance of a large 502 release site, and the translocation of a large diverse founder population, could be significantly 503 more expensive than managing a much smaller site with ongoing reinforcement 504 505 translocations, at least in the short to medium term.

506

One creative option is to combine populations that have low, but different, genetic diversity.
This approach was used by Heber et al. (2013) who mixed SI toutouwai (*Petroica australis*)
from two low diversity translocated populations, to increase diversity in a new translocated

population. Similarly, all translocated populations of NI tieke are descended from the last 510 surviving population on Taranga / Hen Island from which birds were translocated to 511 Whatupuke, Whakau / Red Mercury or Repanga / Cuvier Island (Parker, Anderson et al. 512 2012). However, the Repanga lineage is overrepresented with 15 descendent populations 513 followed by Whatupuke (7 descendent populations) and Whakau (2 populations). Therefore, 514 515 recent translocations have used multiple source populations including, where possible, underrepresented lineages, to maximise the genetic diversity in both new populations and the 516 517 metapopulation (Parker, Lovegrove et al. in press). Alternatively, if it is uncertain if animals 518 will establish and persist, lower value individuals (e.g. males or juveniles) can be released to 519 test a new site. If these survive, additional animals can be released to maximise genetic diversity in the medium to long-term. This approach has been used for hihi translocations 520 where the primary founders for new sites are juveniles from Tiritiri Matangi whereas birds 521 522 from the remaining wild population on Te Hauturu-o-Toi / Little Barrier Island, which are viewed as higher value by some, are reserved for reinforcement translocations to established 523 sites. Another option could be to increase the size of the release area through improved pest 524 525 control thereby enabling a larger population to establish and removing or reducing the need for reinforcement translocations. 526

527

Additional considerations include the genetic profile and history of the source population(s). 528 For instance, will the source population(s) provide genetically diverse individuals for 529 translocation? How many individuals are needed to capture that diversity? These questions 530 531 are not easily answered because they require high resolution genetic and demographic data for source populations and species. These data are usually lacking, especially for widely 532 distributed taxa that show significant geographic variation in genetics and demography, such 533 as lizards and invertebrates. However, even in the absence of such data, combining 534 knowledge of natural history, individual population history and theory allows reasonable 535 assumptions to be made (Weiser et al. 2013; Frankham et al. 2017). 536

537

Post-release monitoring is essential to determine how many animals a release site can
support. If the population remains small reinforcement translocations might be recommended,
but how easy will they be to achieve? The feasibility of reinforcement translocations is often
presented in a simplistic manner with little recognition of the cost and difficulties in getting
additional animals to recruit into an established population. Often, very large numbers of
individuals must be added to ensure that at least a few will be able to recruit and breed

(Weiser et al. 2013). This is because density dependence (Armstrong et al. 2005) or
behavioural barriers (Parker, Hauber et al. 2010; Parker, Anderson et al. 2012) often reduce
the recruitment of immigrants. As noted, releasing large numbers of animals while knowing

that only a few will survive raises welfare and ethical questions and can strain relationships.

549 Regardless of the method chosen to maintain genetic diversity, it is important to recognise that not every translocated population needs to have maximum or ideal genetic diversity. 550 551 Overall genetic diversity can also be represented and conserved within a metapopulation 552 connected via natural dispersal and/or management. This likely represents a more "natural" scenario (i.e. genetic diversity will not be equal across all natural populations, especially 553 when moving from the core of a species range to the edges), whilst also increasing options 554 for establishing and maintaining translocated populations that cater to a wide range of values 555 and objectives. 556

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6. What do future developments mean for the management of translocated populations?

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561 Translocations will continue to play an important role in conservation in Aotearoa. Experience and research will increase our understanding of the values driving translocations 562 including, but not limited to, cultural and societal desires, cost, animal welfare, genetic and 563 564 pathogen management, translocation techniques and dispersal. We also need to fill the significant knowledge gaps that exist for many species, especially invertebrates, lizards, 565 566 amphibians and bats (Table 1). In Aotearoa the biggest opportunities will come about through improved control of pests over large, unfenced areas of the mainland, including forests, 567 568 wetlands, dryland and braided river systems, and alpine zones (Table 1). This will provide 569 additional habitat for species that are currently in higher threat categories, along with further 570 options for the management and translocation of all species, especially habitat specialists such as whio / blue duck (Hymenolaimus malacorhynchos), kakī / black stilt, and pīwauwau / 571 rock wren (Xenicus gilviventris), and neglected fauna, such as lizards, amphibians, bats and 572 invertebrates. While opinions vary on the feasibility of effective pest control over vast 573 swathes of Aotearoa (Urlich 2015) it will clearly be a game changer if it can be achieved. 574 However, in the short term (c. 20 years) large (≥3000 ha) fenced sanctuaries will likely 575

protect the greatest diversity of biodiversity on the main islands, especially if mice can beeffectively controlled within them.

578

We also expect to see an increasing shift away from translocations for single-species 579 recovery toward those where the fundamental objective is ecosystem restoration (Parker 580 2013). Pathogens and predators, such as weka (Gallirallus australis), small rails (Rallus spp. 581 and Porzana spp.) and karearea / NZ falcons (Falco novaeseelandiae) are components of 582 583 Aotearoa ecosystems that are currently either not included in restoration plans or relegated to 584 some point in the distant future once their potential prey or host species are well established (Carpenter et al. 2021). It seems logical to plan ecosystem restoration sequences in stages so 585 that prey species are established before predators, but it is important to distinguish between a 586 pest, against which native species have few defences, and a native predator that they have co-587 evolved with over 1000s of years. For example, although pests have caused the extinction of 588 589 many large wētā populations elsewhere, translocated Mercury Island tusked wētā (Motuweta isolata) and the giant weta / wetapunga (Deinacrida heteracantha) have established in the 590 591 presence of very high densities of a natural predator, the NI tieke. Translocations of native predators require acceptance that there will be ongoing predation and possibly a reduction in 592 593 the population size, alongside changes in the behaviour of prey species. This will be difficult for some people to accept and could be problematic for very small prey populations, but it is 594 a logical objective for true ecosystem restoration. It might also require a change in thinking 595 about the management of native predator species, and pathogens, especially where there is a 596 597 perception that natural predators and pathogens must be controlled.

598

599 There has also been considerable debate about the ongoing impacts of global climate change and how translocations can be used to conserve species whose habitat will deteriorate under 600 601 current climate change predictions (Hoegh-Guldberg et al. 2008; Seddon et al. 2009; Seddon 2010). In Aotearoa this would likely mean moving animals across latitudinal gradients, e.g., 602 603 between the North and South Islands. For instance, climate modelling suggests that the northern South Island, where hihi have never existed, might provide higher quality habitat in 604 605 the future than the North Island, to which they are currently restricted (Chauvenet et al. 2013). Any decision to undertake a translocation beyond a species' natural range will also 606 clearly raise challenges in setting appropriate objectives, especially if it would bring closely 607 related species into contact, although we note that this already happened for some species. 608

609

Another interesting proposition is using close relatives of extinct species as ecological 610 replacements in ecosystem restoration (Atkinson 1988). For example, the tutukiwi / Snares 611 Island snipe (Coenocorypha huegeli) has been translocated to replace the extinct tutukiwi / SI 612 snipe (Coenocorypha iredalei) and the NI kokako as a replacement for the presumed extinct 613 SI kōkako (*Callaeas cinerea*). SI takahē (*Porphyrio hochstetteri*) are also frequently 614 translocated to the North Island (Jamieson & Ryan 2001; Parker, Seabrook-Davison et al. 615 2010; Miskelly et al. 2012), although takahē translocations are motivated by species recovery 616 goals rather than as a replacement for the extinct moho / NI takahē (Porphyrio mantelli). It 617 618 has also been suggested that the introduced Australian brown quail (Synoicus ypsilophorus) could be a suitable ecological replacement for the extinct New Zealand quail (Coturnix 619 novaezelandiae) (Parker, Seabrook-Davison et al. 2010). These species, and others, might be 620 useful for restoring ecosystem functions, known or otherwise, along with restoring generally 621 depauperate ecosystems. In addition, genetic techniques are advancing to the point where de-622 623 extinction, the resurrection of functional proxies of extinct species, might become feasible (Seddon et al. 2014; Seddon 2017). This is a contentious issue and the objectives of any such 624 625 proposal would have to be very carefully considered, including the conservation benefit of diverting funds from extant species to de-extinction proposals (Bennett et al. 2017). 626

627

Emerging genomic tools will further enhance translocation decision making (Luikart et al. 628 629 2018; Funk et al. 2019; Forsdick et al. 2022; Moehrenschlager et al. 2023). Advanced highthroughput sequencing technologies, combined with rapidly decreasing costs, increased 630 capability and capacity in the conservation genetics community, can provide ready access to 631 10s to 10 000s of markers from across the entire genome, even for non-model species 632 (Harrisson et al. 2014; Galla et al. 2019). These genome-wide markers can increase resolution 633 for translocation questions previously answered using just a handful of neutral genetic 634 markers. For example, genomic markers can provide more robust estimates of relatedness for 635 pairing decisions in conservation breeding programmes that include translocations (e.g. Galla 636 et al. 2020). Similarly, genomic markers are increasingly used to identify suitable source 637 populations for translocations to enhance adaptive potential (e.g., McLennan et al. 2020; 638 639 Rayne et al. 2022). Indeed, the promise of characterising adaptive variation has also reignited debate over how we should source, or mix, populations to enhance adaptive potential (Kardos 640 et al. 2021). However, translating theory into practise remains difficult (Flanagan et al. 2017) 641 despite a surge of theoretical and simulation-based papers focussed on characterising 642

- adaptive variation (Funk et al. 2019; Hoelzel et al. 2019). For many threatened species, it
 may prove challenging to characterise adaptive variation at all (Forsdick et al. 2022).
- 645

Recent years have also seen the rise of a new era of conservation genomics that reintegratesthe structure and function of DNA (Deakin et al. 2019). For example, emerging chromosomic

648 approaches combine genomic data with cytogenetics (chromosome architecture),

epigenomics (histone modifications), and cell biology to reveal the mechanisms underpinning

behavioural and phenotypic traits under selection (Mérot et al. 2020). Although each of these

approaches come with caveats (Wold et al. 2023), genomic and chromosomic approaches are

a valuable addition to the translocation toolbox, particularly in the face of novel challenges

such as climate change (Hoffmann et al. 2021; Wold et al. 2021).

654

655 Conclusions

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The common perception that translocations are relatively easy and that success is assured is 657 658 not supported by the evidence either in Aotearoa (Miskelly & Powlesland 2013), or internationally (Griffith et al. 1989; Wolf et al. 1996; Fischer & Lindenmayer 2000). The 659 660 frequency of translocations is also increasing in Aotearoa (Cromarty & Alderson 2013), including calls for urban translocations (van Heezik & Seddon 2018). Furthermore, the 661 quality of translocation proposals presented to DOC is highly variable, with some being 662 poorly written, poorly thought out, or just a bad idea for the candidate species. The DOC 663 approval process itself also produces variable outcomes. The authors want to see more 664 successful translocations in Aotearoa, and we think that the six key considerations we present 665 here (Box 1) will help in achieving that success. 666

667

Ultimately, our goal is to encourage careful thinking in the formulation of translocation
objectives so that these capture the diverse values motivating translocations, together with
appropriate performance measures for determining success. We discourage a focus on any
single value. Instead, we encourage all people involved in translocations, particularly
decision makers, to explicitly recognise the multiple values-based objectives associated with
translocations (Box 1).

674

Haphazard conservation translocations can cause problems at the release site, for futuretranslocations, and in maintaining equitable relationships with mana whenua, local

communities, relevant agencies, and the public. All the translocations we have been involved 677 with have been guided by clear principles. However, the principles of good translocation 678 practice are not currently captured in DOC policy (Parliamentary Commissioner for the 679 Environment 2017). Furthermore, the fundamental objectives of many translocations have 680 rarely been stated explicitly or are dominated by singular means objectives. Therefore, a clear 681 and widely consulted translocation policy framework would enable DOC decision makers to 682 make better decisions about all translocations. This policy should 1) specifically acknowledge 683 that translocations are values based; 2) should be driven by an understanding of the problem 684 685 at hand; 3) require informed decisions between management alternatives (including rejecting translocation as a management tool for some species/programmes); 4) should be measured by 686 explicitly stated objectives with appropriate performance indicators. Ultimately, being clear 687 about what all relevant parties really want will set everyone on the right path towards the 688 landscape of Aotearoa being one that is once again dominated by indigenous biodiversity. 689

690

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692

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701

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703

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- Caughley G 1989. New Zealand and plant-herbivore systems: past and present. New Zealand
 Journal of Ecology 12: 3–10.
- Caughley G 1994. Directions in conservation biology. Journal of Animal Ecology 63: 214–
 244.
- Chauvenet ALM, Ewen JG, Armstrong D, Pettorelli N 2013. Saving the hihi under climate
 change: a case for assisted colonization. Journal of Applied Ecology 50: 1330–1340.
- Converse S, Moore C, Folk M, Runge M 2014. Optimal release strategies for cost-effective
 reintroductions. Journal of Wildlife Management 77: 1145–1156.
- Cromarty PL, Alderson SL 2013. Translocation statistics (2002-2010), and the revised
 Department of Conservation translocation process. Notornis 60: 55–62.
- Deakin JE, Potter S, O'Neill R, Ruiz-Herrera A, Cioffi MB, Eldridge MDB, Fukui K, Graves
 JAM, Griffin D, Grutzner F, Kratochvil L, Miura I, Rovatsos M, Srikulnath K,
- Wapstra E, Ezaz T 2019. Chromosomics: bridging the gap between genomes and
 chromosomes. Genes 108: 627.
- Department of Conservation 2020. Towards a predator free New Zealand: Predator free 2050
 strategy. Wellington, Department of Conservation. 44p.
- Diamond JM 1984. Distributions of New Zealand birds on real and virtual islands. New
 Zealand Journal of Ecology 7: 37–55.
- Ewen JE, Soorae PS, Canessa S 2014. Reintroduction objectives, decisions and outcomes:
 global perspectives from the herpetofauna. Animal Conservation 17: 74–81.
- Ewen JG, Thorogood R, Nicol C, Armstrong DP, Alley M 2007. Salmonella typhimurium in
 hihi, New Zealand. Emerging Infectious Diseases 13: 788–790.
- Ewen JG, Armstrong DP, Empson R, Jack S, Makan T, McInnes K, Parker KA, Richardson
 K, Alley M 2012. Parasite management in translocations: lessons from an endangered
 New Zealand bird. Oryx 46: 446–456.
- Ewen JG, Renwick R, Adams L, Armstrong DP, Parker KA 2013. 1980-2012: 32 years of reintroduction efforts of the hihi (stitchbird) in New Zealand. In: Soorae PS ed. Global
- re-introduction perspectives: 2013. Abu Dhabi, IUCN/SSC Re-introduction Specialist
- 771 Group & Environment Agency-ABU DHABI. 282 p.
- Ewen JG, Canessa S, Converse SJ, Parker KA. 2023. Decision-making in animal
- conservation translocations: biological considerations and beyond. In: Gaywood MJ,
- Ewen JF, Hollingsworth PM, Moehrenschlager A. eds. Conservation Translocations.
- Cambridge, Cambridge University Press. pp 175–186.

- Fischer J, Lindenmayer DB 2000. An assessment of the published results of animal
 relocations. Biological Conservation 96: 1–11.
- Fischer JH, Parker KA, Kenup CF, Davis T, Cole RA, Taylor GA, Debski I, Ewen JG. 2023.
- 779 Decision analysis for seabird recovery: navigating complexity across ecosystems,
- balancing competing values, and bridging the research implementation gap. Journal ofApplied Ecology 00: 1–14.
- Flanagan SP, Forester BR, Latch EK, Aitken SN, Hoban S 2017. Guidelines for planning
 genomic assessment and monitoring of locally adaptive variation to inform species
 conservation. Evolutionary Applications 117: 1027–1193.
- Forsdick NJ, Adams CIM, Alexander A, Clark AC, Collier-Robinson L, Cubrinovska I, Croll
 Dowgray M, Dowle EJ, Duntsch L, Galla SJ, Howell L, Magid M, Rayne A, Verry
- 787 AJF, Wold JR, Steeves TE. 2022. Current applications and future promise of
- 788 genetic/genomic data for conservation in an Aotearoa New Zealand context.
- 789 Wellington, Department of Conservation, 61p.
- Frankham R, Ballou JD, Ralls K, Eldridge MDB, Dudash MR, Fenster CB, Lacy RC,
 Sunnucks P 2017. Genetic management of fragmented animal and plant populations.
 Oxford, Oxford University Press. 401 p.
- Funk WC, Forester BR, Converse S, Darst C, Morey S 2019. Improving conservation policy
 with genomics: a guide to integrating adaptive potential into U.S. Endangered Species
 Act decisions for conservation practitioners and geneticists. Conservation Genetics
 20: 115–134.
- Galla SJ, Forsdick NJ, Brown L, Hoeppner MP, Knapp M, Maloney RF, Moraga R, Santure
 AW, Steeves TE 2019. Reference genomes from distantly related species can be used
 for discovery of single nucleotide polymorphisms to inform conservation
 management. Genes 10: 1–19.
- Galla SJ, Moraga R, Brown L, Cleland S, Hoeppner MP, Maloney RF, Richardson A, Slater
 L, Santure AW, Steeves TE 2020. A comparison of pedigree, genetic and genomic
 estimates of relatedness for informing pairing decisions in two critically endangered
 birds: implications for conservation breeding programmes worldwide. Evolutionary
 Applications 13: 991–1008.
- Gregory R, Failing L, Harstone M, Long G, McDaniels T, Ohlson D 2012. Structured
 decision making: a practical guide to environmental management choices. Oxford,
 Wiley-Blackwell. 299 p.

- Griffith B, Scott JM, Carpenter JW, Reed C 1989. Translocation as a species conservation
 tool: status and strategy. Science 245: 477–480.
- Hall LS, Krausman PR, Morrison ML 1997. The habitat concept and a plea for standard
 terminology. Wildlife Society Bulletin 25: 173–182.
- Harrisson KA, Pavlova A, Telonis-Scott M, Sunnucks P 2014. Using genomics to
 characterize evolutionary potential for conservation of wild populations. Evolutionary
 Applications 7: 1008–1025.
- Heather B, Robertson H 2015. The field guide to the birds of New Zealand. New Zealand,
 Auckland, Penguin Books. 464p.
- Heber S, Varsani A, Kuhn S, Girg A, Kempenaers B, Briskie J 2013. The genetic rescue of
 two bottlenecked South Island robin populations using translocations of inbred
 donors. Proceedings of the Royal Society B 280: 1–8.
- Hoegh-Guldberg O, Huhes L, McIntyre S, Lindenmayer DB, Parmesan C, Possingham HP,
 Thomas CD 2008. Assisted colonization and rapid climate change. Science 321: 345–
 346.
- Hoelzel AR, Bruford MW, Fleischer RC 2019. Conservation of adaptive potential and
 functional diversity. Conservation Genetics 20: 1–5.
- Hoffmann AA, Weeks AR, Sgrò CM 2021. Opportunities and challenges in assessing climate
 change vulnerability through genomics. Cell 184(6): 1420–1425.
- Holdaway RN 1989. New Zealand's pre-human avifauna and its vulnerability. New Zealand
 Journal of Ecology 12: 11–25.
- Innes J, Kelly D, Overton J. McC., Gillies C 2010. Predation and other factors currently
 limiting New Zealand forest birds. New Zealand Journal of Ecology 34: 86–114.
- Innes J, Fitzgerald N, Binny R, Byrom A, Pech R, Watts C, Gillies C, Maitland M, CampbellHunt C, Burns B 2019. New Zealand ecosanctuaries: types, attributes and outcomes.
 Journal of the Royal Society of New Zealand 49: 370–393.
- Innes J, Miskelly CM, Armstrong DP, Fitzgerald N, Parker KA, Stone ZL 2022. Movements
 and habitat connectivity of New Zealand forest birds: a review of available data. New
 Zealand Journal of Ecology 46: 1–21.
- 838 IUCN 2013. Guidelines for reintroductions and other conservation translocations. Version
- 839 1.0. Gland, Switzerland, IUCN Species Survival Commission. 57 p.
- Jamieson IG, Ryan CJ 2001. Island takahe: closure of the debate over the merits of
- 841 introducing Fiordland takahe to predator-free islands. In: Lee WG, Jamieson IG ed.

- 842 The takahe: Fifty years of conservation management and research. Dunedin,
- 843 University of Otago Press. Pp. 96–113.
- 844 Kardos M, Armstrong EE, Fitzpatrick SW, Hauser S, Hedrick PW, Miller JM, Tallmon DA,
- Funk WC 2021. The crucial role of genome-wide genetic variation in conservation.

Proceedings of the National Academy of Sciences 118: e2104642118.

Keeny R 1982. Decision analysis: an overview. Operations Research 30: 803-838

- 848 Leuschner N 2007. Ecology and behaviour of the whitehead (*Mohoua albicilla*) in its
- translocated ranges in New Zealand. Unpublished thesis, University of Auckland,Auckland. 109 p.
- Luikart GKM, Hand BK, Rajora OP, Aitken SN, Hohenlohe PA 2018. Population genomics:
 advancing understanding of nature. In: Rajora OP ed. Population Genomics: concepts,
 approaches, applications. New York, Springer. Pp. 3–79.
- Maguire L 1986. Using decision analysis to manage endangered species populations. Journal
 of Environmental Management 22: 345–360.
- Massaro M, Ortiz-Catedral L, Julian L, Galbraith JA, Kurenback B, Kearvell J, Kemp J, van
 Hal J, Elkington S, Taylor G, Greene T, van de Wetering J, van de Wetering M, Pryde
 M, Dilks P, Heber S, Steeves T, Walters M, Shaw S, Potter J, Farrant M, Brunton DH,
 Hauber M, Jackson B, Bell P, Moorhouse R, McInnes K, Varsani A. 2012. Molecular
 characterisation of beak and feather disease virus (BFDV) in New Zealand and its
 implications for managing an infectious disease. Archives of Virology 157: 1651–
- 862 1663.
- McCarthy MA, Armstrong DP, Runge MC 2012. Adaptive management of reintroduction. In:
 Ewen JG, Armstrong DP, Parker KA, Seddon PJ ed. Reintroduction biology:
 integrating science and management. Chichester, Wiley-Blackwell. Pp. 256–289.
- McLennan EA, Grueber CE, Wise P, Belov K, Hogg CJ 2020. Mixing genetically
- differentiated populations successfully boosts diversity of an endangered carnivore.
 Animal Conservation 23: 700–712.
- McMurdo Hamilton T, Canessa S, Clark K, Gleeson P, Mackenzie F, Makan T, Moses-Te
 Kani G, Oliver S, Parker KA, Ewen JG 2021. Applying a values-based decision
 process to facilitate comanagement of threatened species in Aotearoa New Zealand.
 Conservation Biology 35: 1162–1173.
- Mérot C, Oomen RA, Tigano A, Wellenreuther M 2020. A roadmap for understanding the
 evolutionary significance of structural genomic variation. Trends in Ecology &
 Evolution 35: 561–572.

- Miskelly CM, Powlesland RG 2013. Conservation translocations of New Zealand birds,
 1863-2012 Notornis 60: 3–28.
- Miskelly CM, Charteris MR, Fraser JR 2012. Successful translocation of Snares Island snipe
 (*Coenocorypha huegeli*) to replace the extinct South Island snipe (*C. iredalei*).
 Notornis 59: 32–38.
- Molles LE, Calcott A, Peters D, Delamare G, Hudson JD, Innes J, Flux I, Waas JR 2008.
 "Acoustic anchoring" and the successful translocation of North Island kokako
- (*Callaeas cinerea wilsoni*) to a New Zealand mainland site within continuous forest.
 Notornis 55: 57–68.
- Moehrenschlager A, Soorae P, Steeves TE. 2023. From genes to ecosystems and beyond:
 addressing eleven contentious issues to advance the future of conservation
- translocations. In: Gaywood MJ, Ewen JF, Hollingsworth PM, Moehrenschlager A.
- eds. Conservation Translocations. Cambridge University Press, Cambridge. pp 381–
 412.
- Newman DG 1980. Colonisation of Coppermine Island by the North Island saddleback.
 Notornis 27: 146–147.
- 892 Ortiz-Catedral L 2010. Homing of a red-crowned parakeet (*Cyanoramphus novaezelandiae*)
 893 from Motuihe Island to Little Barrier Island, New Zealand. Notornis 57: 48–49.
- 894 Ortiz-Catedral L, Prada D, Gleeson D, Brunton DH 2011. Avian malaria in a remnant
- population of red-fronted parakeets on Little Barrier Island, New Zealand. New
 Zealand Journal of Zoology 38: 261–268.
- Parker KA 2008. Translocations: Providing outcomes for wildlife, resource managers,
 scientists, and the human community. Restoration Ecology 16: 204–209.
- Parker KA 2013. Avian translocations to and from Tiritiri Matangi 1974-2013. New Zealand
 Journal of Ecology 37: 282–287.
- Parker KA, Hughes B, Thorogood R, Griffiths R 2004. Homing over 56 km by a North Island
 tomtit (*Petroica macrocephala toitoi*). Notornis 51: 238–239.
- Parker KA, Brunton DH, Jakob-Hoff R 2006. Avian translocations and disease; implications
 for New Zealand conservation. Pacific Conservation Biology 12: 155–162
- Parker KA, Hauber ME, Brunton DH 2010. Contemporary cultural evolution of a conspecific
 recognition signal following serial translocations. Evolution 64: 2431–2441.
- Parker KA, Seabrook-Davison M, Ewen JG 2010. Opportunities for non-native ecological
 replacements in ecosystem restoration. Restoration Ecology 18: 269–273.

- Parker KA, Anderson MJ, Jenkins PF, Brunton DH 2012. The effects of translocationinduced isolation and fragmentation on the cultural evolution of bird song. Ecology
 Letters 15: 778–785.
- Parker KA, Dickens MJ, Clarke RH, Lovegrove TG 2012b. The theory and practise of
 catching, holding, moving and releasing animals. In: Ewen JG, Armstrong DP, Parker
 KA, Seddon PJ ed. Reintroduction biology: integrating science and management.
- 915 Chichester, U.K., Wiley-Blackwell. Pp. 105–137.
- Parker KA, Adams L, Baling M, Kemp L, Kuchling G, Lloyd B, Parsons S, Ruffell J,
- 917 Stringer I, Watts C, Dickens MJ 2015. Practical guidelines for planning and
- 918 implementing fauna translocations. In: Armstrong DP, Hayward MW, Moro D,
- 919 Seddon PJ ed. Advances in reintroduction biology of Australian and New Zealand

920 fauna. Clayton South, CSIRO Publishing. Pp. 255–272.

- Parker KA, Parlato EH, Fischer JH. *In press*. A structured decision-making approach for the
 recovery of karure / kakaruia / Chatham Island black robins (*Petroica traversi*).
 Wellington, Department of Conservation.
- Parker KA, Lovegrove TG, McClellend P. *In press*. Best practice techniques for the
 translocation of tīeke / saddlebacks (*Philesturnus* spp.). Wellington, Department of
 Conservation.
- Parlato EH, Armstrong DP 2012. An integrated approach for predicting fates of
 reintroductions with demographic data from multiple populations. Conservation
 Biology 26: 97–106.
- Parlato EH, Armstrong DP 2013. Predicting post-release establishment using data from
 multiple introductions. Biological Conservation 160: 97–104.
- Parlato EH, Armstrong DP 2018. Predicting reintroduction outcomes for highly vulnerable
 species that do not currently co-exist with their key threats. Conservation Biology 32:
 1346–1355.
- Parliamentary Commissioner for the Environment 2017. Taonga of an island nation: Saving
 New Zealand's birds. Wellington, Parliamentary Commissioner for the Environment.
 139 p.
- 838 Rayne A, Byrnes G, Collier-Robinson L, Hlows J, McIntosh A, Huika MRK, Rupene M,
- Tamti-Elliffe, Thoms C, Steeves TE 2020. Centring indigenous knowledge systems to
 re-imagine conservation translocations. People and Nature 2: 512–526.
- Rayne A, Blair S, Dale M, Flack B, Hollows J, Moraga R, Parata RN, Rupene M, TamatiElliffe P, Wehi PM, Wylie MJ, Steeves TE 2022. Weaving place-based knowledge for

- 943 culturally significant species in the age of genomics: Looking to the past to navigate
 944 the future. Evolutionary Applications 15: 751–772.
- Reardon JT, Whitmore N, Holmes KM, Judd LM, Hutcheon AD, Norbury G, Mackenzie D
 2012. Predator control allows critically endangered lizards to recover on mainland
 New Zealand. New Zealand Journal of Ecology 36: 141–150.
- Richard Y, Armstrong DP 2010. Cost distance modelling of landscape connectivity and gapcrossing ability using radio-tracking data. Journal of Applied Ecology 47: 603–610.
- 950 Richardson K, Doerr V, Ebrahimi M, Parker KA 2014. Considering dispersal in
- 951 reintroduction and restoration planning. In: Armstrong DP, Hayward MW, Moro D,
 952 Seddon PJ ed. Advances in Reintroduction Biology of Australian and New Zealand
 953 Fauna. Australia, CSIRO Publishing. Pp. 59–72.
- Richardson K, Castro IC, Brunton DH, Armstrong DP 2015. Not so soft? Delayed release
 reduces long-term survival in a passerine reintroduction. Oryx 49: 535–541.
- 956 Rickett J, Dey CJ, Stothart J, O'Connor CM, Quinn JS, Ji W 2013. The influence of
- 957 supplemental feeding on survival, dispersal and competition in translocated brown
 958 teal, or pateke (*Anas chlorotis*). Emu 113: 62–68.
- Sainsbury AW, Carraro C 2023. Animal disease and conservation translocations. In:
 Gaywood MJ, Ewen JF, Hollingsworth PM, Moehrenschlager A. eds. Conservation
- 961 Translocations. Cambridge University Press, Cambridge. pp 149–179.
- Seddon PJ 2010. From reintroduction to assisted colonization: moving along the conservation
 translocation spectrum. Restoration Ecology 18: 796–802.
- 964 Seddon PJ 2017. The ecology of de-extinction. Functional Ecology 31: 992–995.
- Seddon PJ, Armstrong DP, Soorae P, Launay F, Walker S, Ruiz-Miranda CR, Molur S,
 Koldewey H, Kleiman DG 2009. The risks of assisted colonization. Conservation
 Biology 23: 788–789.
- Seddon PJ, Moehrenschlager A, Ewen JG 2014. Reintroducing resurrected species: selecting
 DeExtinction candidates. Trends in Ecology & Evolution 29: 140–147.
- 970 Smuts-Kennedy C, Parker KA 2013. Reconstructing avian biodiversity on Maungatautari.
 971 Notornis 60: 93–106.
- 972 Stadtmann S, Seddon PJ 2018. Release site selection: reintroductions and the habitat concept.
 973 Oryx 54: 687–695.
- 974 Swan KD, McPherson JM, Seddon PJ, Moehrenschlager A 2016. Managing marine
 975 biodiversity: The rising diversity and prevalence of marine conservation
- translocations. Conservation Letters 9: 239–251.

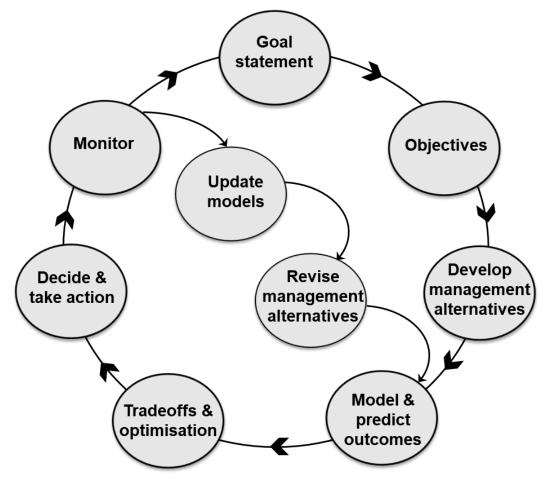
- 977 Towns DR, Broome K 2003. From small Maria to massive Campbell: Forty years of rat
 978 eradications from New Zealand islands. New Zealand Journal of Zoology 30: 377–
 979 398.
- 980 Urlich SC 2015. What's the end-game for biodiversity: is it time for conservation evolution?
 981 New Zealand Journal of Ecology 39: 133–142.
- van Heezik Y, Seddon PJ 2018. Animal reintroductions in peopled landscapes: moving
 towards urban-based species restorations in New Zealand. Pacific Conservation
 Biology 24: 349–359.
- Weeks AR, Moro D, Thavornkanlapachai R, Taylor HR, White NE, Weiser EL, Heinze D
 2015. Conserving and enhancing genetic diversity in translocation programs. In:
 Armstrong DP, Hayward MW, Moro D, Seddon PJ ed. Advances in reintroduction
 biology of Australian and New Zealand fauna. Australia, CSIRO Publishing. Pp. 127–
 140.
- Wehi PM, Cox MP, Roa T, Whaanga H 2018. Human perceptions of megafaunal extinction
 events revealed by linguistic analysis of indigenous oral traditions. Human Ecology
 46: 461–470.
- Weiser EL, Grueber CE, Jamieson IG 2013. Simulating retention of rare alleles in small
 populations to assess management options for species with different life histories.
 Conservation Biology 27: 335–344.
- Wold J, Galla S, Eccles D, Hogg CJ, Koepfli K-P, Lec ML, Guhlin J, Price K, Roberts J,
 Steeves T 2021. Expanding the conservation genomics toolbox: incorporating
 structural variants to enhance functional studies for species of conservation concern.
 Molecular Ecology 30: 5949–5965.
- Wold JR, Guhlin JG, Dearden PK, Santure AW, Steeves TE (2023). The promise and
 challenges of characterising genome-wide structural variants: A case study in a
 critically endangered parrot. Molecular Ecology Resources 00:1–18
- Wolf CM, Griffith B, Reed C, Temple SA 1996. Avian and mammalian translocations:
 update and reanalysis of 1987 survey data. Conservation Biology 10: 1142–1154.
- 1005

Box 1. Six key considerations for conservation translocations in Aotearoa New Zealand. Of
these, the first is the most critical because, if done correctly, all other considerations, both
listed and unlisted, derive from this.

- What values will be met by doing a translocation? <u>All translocations are values</u> <u>based so these values should be explicitly stated</u>. For example, a translocation might reinstate rangatiratanga, kaitiakitanga or mauri, reduce extinction risk, restore a depauperate ecosystem or reconnect a local community with the target species.
- 2. What is the natural and conservation history of the species being translocated?
- 3. Does the release site habitat (e.g. pests, vegetation associations, pathogens) match that of the proposed source population? If not, why is the release site considered appropriate? Can management ameliorate differences?
- 4. Is dispersal likely to be an impediment to establishment and persistence?
- 5. Will genetic management be required and how realistic is it that this will be implemented (e.g. increase the number of founders, conduct reinforcement translocations or increase the size of the management area)?
- 6. What do future developments (e.g. improved pest control or emerging genomic tools) mean for the management of translocated populations?

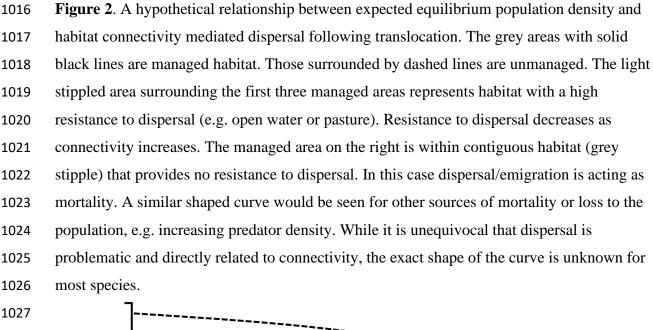
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- 1010 Figure 1. Steps in the structured decision-making process for conservation translocations
- 1011 (adapted from Gregory et al. 2012). Note the double loop learning whereby monitoring might
- 1012 lead to a revision of management alternatives.



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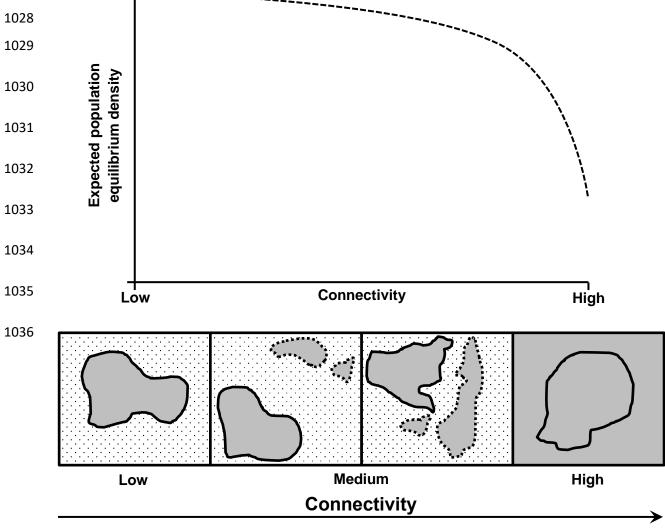


Table 1. Five of the six key considerations (see Box 1) for some terrestrial species/species groups that have been translocated in Aotearoa New Zealand. We have not specified key consideration number one because values are project specific. However, for the species listed they usually include objectives such as minimise extinction risk, restore depauperate ecosystems, restore mauri and reconnect local communities with the translocated species. Knowledge is patchy, even for many bird species, and there is a lot of uncertainty to resolve, especially for herpetofauna and invertebrates. In particular, other habitat variables, such as ideal vegetation associations, are often difficult to resolve until suitable pest control is in place. NI – North Island; SI – South Island.

Translocated species or species group	2. Pest thresholds based on extirpation and management history	3. Habitat match required?	4. Ability to disperse when connectivity is:			5. Will genetic management be required?	6. What future developments will assist translocation of this species?
			High	Medium	Low		speces.
Kiwi spp.	Key pest species controlled to low density, typically mustelids	Not necessarily - occupy a diverse range of habitats	High	High	?	Possibly, depending on the size of the recipient site, the number and source of founders and vital rates post release	Improved pest control Safe dispersal corridors
Weka spp., particularly NI and buff weka		Not necessarily - occupy a diverse range of habitats	High	High	?	Possibly, depending on the size of the recipient site, the number and source of founders and vital rates post release	Improved pest control Safe dispersal corridors Acceptance that weka are a natural, endemic predator that co-evolved with all other indigenous species and ecosystems.
Whio / blue duck		Unknown. Archaeological evidence suggests whio might have once used a more diverse range of habitats	High	High	?	Unknown?	Improved pest control Safe dispersal corridors Understanding dispersal behaviour Trial translocations to test habitat plasticity

Toutouwai / robin spp.	Multi-species pest control	No, but vital rates vary	High	?	Low	Possibly, depending on	Improved pest control
	to low density, typically including ship rats, mustelids, possums and cats, sometimes including ungulates and pigs. Mice usually present,	among sites, suggesting some, especially damp lowland forest with thick leaf litter, are better than others				the size of the recipient site, the number and source of founders and vital rates post release	Safe dispersal corridors
Yellow crowned kākāriki	sometimes at high density Control is sometimes delivered seasonally (e.g. over the bird breeding season)	Not necessarily - occupy a diverse range of habitats	High	High	?	Possibly, depending on the size of the recipient site, the number and source of founders, vital rates post release and dispersal distance to other populations	Improved pest control Safe dispersal corridors Understanding dispersa behaviour
Popokōtea / whitehead		Not necessarily - occupy a diverse range of habitats	High	Moderate	Low	Unlikely, except at very small sites (<50ha)	Improved pest control Safe dispersal corridor
Mohua / yellowhead		Not necessarily - occupy a diverse range of habitats	High?	?	Low	Possibly, depending on the size of the recipient site, the number and source of founders, vital rates post release and dispersal distance to other populations	Improved pest control Safe dispersal corridors
Titipounamu / rifleman		Not necessarily - occupy a diverse range of habitats	?	?	Low	Unlikely, except at very small sites (<50ha)	Improved pest control Safe dispersal corridor
Kākā		Kākā are mobile and use a wide range of habitats but their core requirements are unclear	High ¹	High ¹	High	Possibly, depending on the size of the recipient site, the number and source of founders, vital rates post release, dispersal distance to other populations and	Improved pest control Safe dispersal corridors

					propensity to mix with other populations	
North Island kōkako	Kōkako persist in a wide range of habitats but large (≥2000ha) diverse forested habitats are likely optimal habitats	High	?	Low	Possibly, depending on the size of the recipient site, the number and source of founders, vital rates post release and dispersal distance to other populations	Improved pest control Safe dispersal corridors
Pekapeka / Short-tailed bats	Short-tailed bats use a variety of habitats but the full extent of their habitat plasticity is unknown	High?	?	?	Unknown – no translocated populations have persisted	Development of successful translocation techniques
Mainland herpetofauna, e.g. Northern spotted skinks and the infrapunctatum complex, jewelled and forest geckos, Hochstetter's frog	Unknown	?	?	?	Unknown – will depend on source populations, founder size and vital rates at new sites	Improved pest control, especially of mice Improved understanding of basic biology and ecology, including habitat plasticity Improved profile and funding by government and the broader community
Mainland invertebrates	Unknown	?	?	?	Unknown – will depend on source populations, founder size and vital rates at new sites	Improved pest control, especially of mice Improved understanding of basic biology and ecology, including habitat plasticity Improved profile and funding by government and the broader community

Tīeke / saddleback spp.	Multi-species pest control to eradication or zero density of all mammalian pests with the probable exception of mice (as is typical of all mainland fenced sanctuaries).	Not necessarily - occupy a diverse range of habitats	High	Low	Low	Unlikely with careful selection of founder populations but very small populations might benefit from periodic reinforcement translocations	Improved pest control Safe dispersal corridors
Hihi		Unknown but large, intact and diverse forested habitats are likely optimal	High	Moderate ?	Low	Possibly, depending on the size of the recipient site, the number and source of founders and vital rates post release	Improved pest control Improved understanding of habitat requirements Safe dispersal corridors
Kākāpō		Historically occupied a wide variety of habitats	High	?	?	Possibly, depending on the size of the recipient site, the number and source of founders and vital rates post release	Improved pest control, especially over very large landscapes
Highly threatened herpetofauna, e.g. McGregor's, robust, and Whitaker's skink, Duvaucel's gecko, tuatara	Multi-species pest control to eradication or zero density of all mammalian pests, including mice.	Unknown	?	?	?	Unknown – will depend on source populations, founder size and vital rates at new sites	Improved pest control, especially of mice Improved understanding of basic biology and ecology, including habitat plasticity Improved profile and funding by government and the broader community
NZ snipe		Unknown, but could likely persist in a wide range of habitats when key pests are absent	?	?	?	Possibly, depending on the size of the recipient site, the number and source of founders and vital rates post release	
Large native and endemic threatened invertebrates,		Unknown	?	?	?	Unknown – will depend on source populations,	Improved pest control, especially of mice

e.g. giant wētā, weevils and beetles

founder size and vital rates at new sites

Improved understanding of basic biology and ecology, including habitat plasticity Improved profile and funding by government and the broader community

¹Dispersal of translocated kākā has been moderated through the provision of supplemental food