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4 Biological validation of fecal corticosterone metabolites as a non-invasive stress

5 assessment in translocated California valley quail (*Callipepla californica*)

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Lay summary

18 This study biologically validates the use of fecal corticosterone metabolites as a non-invasive

19 method for detecting stress in quail, demonstrates diurnal variation in quail stress levels,

20 confirms that translocations elevate stress which likely impacts success, and establishes a new

21 non-invasive tool for stress measurement in quail research, management, and conservation.

Abstract

U.S. quail species are vulnerable to population declines as a result climate change, habitat loss, and habitat fragmentation; all of which can induce physiological stress. Additionally, population restoration techniques (PRTs), like translocations, also induce stress. Traditional stress assessments include capturing and handling birds to extract blood, methods which are inherently stressful and can compound stress analyses. However, the stress hormone corticosterone is metabolized from the blood and excreted in feces as fecal corticosterone metabolites (FCMs). FCMs have been used as a non-invasive measurement of stress in a variety of species, but must be validated for each species. The objective of this study was to biologically validate the use of FCMs as a non-invasive measurement of stress-hormone levels in California valley quail (*Callipepla californica*). Reference and treatment quail were acclimated for 3 wks in an outdoor aviary. Subsequently, treatment quail were subjected to a simulated, 48-h translocation, a common and stressful PRT. Fecal samples were collected every 4 h and processed using an enzyme immunoassay. Mean FCM concentrations of treatment quail (41.50 ±16.13 ng/g) were higher than reference FCM concentrations (24.07 ±10.4 ng/g). These results biologically validate the use of FCMs as a non-invasive method to assess stress hormone levels in California valley quail, demonstrate diurnal variation in quail stress levels, and confirms that quail translocations are a stressful PRT. Ultimately, this research validates a new non-invasive tool for stress measurement to advance quail research, management, and conservation.

Keywords: biological validation, California valley quail, conservation, corticosterone, fecal corticosterone metabolites, physiology, population restoration, stress hormones, translocation

Introduction

Stress is a factor often overlooked in U.S. quail population declines and population restoration techniques. U.S. quail populations are vulnerable to decline as a result of climate change (Guthery *et al.*, 2000; Tanner *et al.*, 2017; Wilsey *et al.*, 2019), habitat loss, and habitat fragmentation (Brennan and Covington, 1994; Hernández *et al.*, 2013; Pope and Heekin, 2017); all of which are stressful to quail as indicated by increased levels of blood corticosterone, (CORT, 17-deoxycortisol), a steroid hormone produced in the cortex of the adrenal gland (Creel, 2001; Suorsa *et al.*, 2003; Tempel and Gutiérrez, 2004; Cockrem, 2007, 2013; Sheriff *et al.*, 2011; Wingfield *et al.*, 2017; Palme, 2019; van Vliet *et al.*, 2020). For example, elevated CORT levels have been observed in birds as a result of anthropogenic disturbances (Thiel *et al.*, 2011; Jankowski *et al.*, 2014; Arlettaz *et al.*, 2015) and extreme climatic events (Wingfield *et al.*, 2017), which are expected to increase in frequency and severity due to global climate change (Seneviratne *et al.*, 2012; IPCC, 2021). Stress from climate change can have synergistic effects, multiplying the deleterious impacts of other stressors (Felton *et al.*, 2009; Şekercioğlu *et al.*, 2012).

Elevated stress levels can directly influence bird populations by altering foraging, predator avoidance, fledging, and other behaviors (Cyr and Romero, 2007; Crespi *et al.*, 2013; Wingfield *et al.*, 2017). Elevated CORT levels in birds can also lead to increased predator encounters (Stephens *et al.*, 2004), which further increases stress levels (Clinchy *et al.*, 2013). Extreme stress can induce a type 1 allostatic overload, where birds decrease or eliminate behaviors (e.g., reproduction) that are not immediately necessary for survival (McEwen and Wingfield, 2003) by increasing gonadotropin-inhibitory hormone, reducing gonadotropin-

releasing hormone (GnRH), and displaying a reduced sensitivity to GnRH and luteinizing

hormone (Sapolsky *et al.*, 2000; Wingfield and Sapolsky, 2003; Dickens and Bentley, 2014);

though the mechanisms are not completely understood (Wingfield and Ramenofsky, 2011; Lattin

et al., 2016). For example, exposing zebra finches *(Taeniopygia guttata)* to stress stimuli

increased CORT levels and decreased testosterone, resulting in reduced reproductive behavior

(Lynn *et al.*, 2010). Capturing and holding wild European starlings (*Sturnus vulgaris*) blocked a

natural shift in the hypothalamic-pituitary-adrenal (HPA) axis from early breeding (gonadal

development) to active breeding activity, which inhibited breeding (Dickens and Bentley, 2014).

The resultant reduction in reproduction in birds contributes to population declines.

Population restoration techniques (PRTs) can also increase CORT levels. U.S. quail

species are economically important gamebirds (Johnson *et al.*, 2012; Wszola *et al.*, 2020) and

part of a $3.7 billion per year upland gamebird hunting industry (Sportsmen's Alliance, 2021).

For over 150 years, a variety of PRTs like translocations have been used in attempts to bolster

populations (Hernández *et al.*, 2013; Gomez and Reyna, 2017; Whitt *et al.*, 2017). Quail

Translocations in general are rarely successful (Perez *et al.*, 2002; Scott *et al.*, 2013; Whitt *et al.*,

2017), especially when used for reintroduction (Martin *et al.*, 2017). Although these PRTs have

rarely resulted in long-term success, there is increasing interest in translocation of wild quail

(Stephenson *et al.*, 2011; Martin *et al.*, 2017; Sisson *et al.*, 2017). Numerous hypotheses for the

low success rate have been proposed (Sokos *et al.*, 2008; Martin *et al.*, 2017), but little is known

about how stress influences translocation success.

Translocation consists of capturing, handling, holding, transporting, and releasing quail to

a novel site, each of which can result in acute and chronic stress, independent of other processes

(Buchanan, 2000; Fazio and Ferlazzo, 2003; Jones *et al.*, 2005; Dickens *et al.*, 2009, 2010;

Wingfield and Romero, 2011; Batson *et al.*, 2017; Martin *et al.*, 2017). For example, short

translocations may induce acute stress; however, the duration of a quail translocation is typically

1-3 days, which increases the occurrence of chronic stress (Dickens *et al.*, 2010).

Chronic stress can disrupt the fight or flight response as a result in downregulation of

hormonal responses to acute and chronic stressors, (Romero and Butler, 2007; Dickens *et al.*,

2009; Martin *et al.*, 2017). For example, European starlings caught in the wild and placed into

captivity lost the ability to produce a fight or flight response when presented with a stress

stimulus (Dickens and Romero, 2009). In addition, when the fight or flight response was

chronically stimulated it negatively impacted cardiovascular health, leading to hypertension due

to surplus catecholamine exposure (Rupp, 1999). Both chronic and acute stress responses can

inhibit translocated quail survival (Armstrong and Seddon, 2008; Dickens *et al.*, 2010; Breed *et*

al., 2019), and lead to failure of the translocation by increasing the susceptibility to predation

(Dickens *et al.*, 2010; Martin *et al.*, 2017).

Weight loss has been observed when translocating wild birds to novel environments

(Rich and Romero, 2005; Dickens *et al.*, 2009; Fischer *et al.*, 2018), and is a symptom of chronic

stress. For example, European starlings lost 5–15% of their body weight after being exposed to a

14-d regime of multiple stressors during a translocation (Awerman and Romero, 2010).

California valley quail (*C. californica*) exhibited a mean weight loss of 14.3% when translocated

from Idaho to Texas (Reyna *et al.*, 2020). This is important because body mass can influence

survival, and the overall success of translocations. For example, Cirl buntings (*Emberiza cirlus*)

with a higher body weight at capture were more likely to survive in their new habitat than those

with lower body weights (Fountain *et al.*, 2017). Reducing the amount of stress during a

translocation could reduce weight loss and increase post-release survival (Warwick *et al.*, 2006).

The traditional method of measuring stress in birds requires capturing, handling, and

extracting blood samples; inherently stressful events (Arnold *et al.*, 2008), followed with

measuring blood CORT levels. It is assumed that if this process occurs within 3 min of initial

handling, CORT levels will be indicative of the physiological condition of the bird prior and will

not represent the elevated stress incurred during the procedure (Littin and Cockrem, 2001;

Cockrem and Silverin, 2002). However, CORT levels quickly rise during handling and peak

within 15–30 min. In addition, handling associated with blood extractions adds to the cumulative

stress experienced by the bird, and nearby birds, independent of the extraction.

Steroid hormones, like CORT, can also be measured from urine, feces, hair, and feather

samples (Bortolotti *et al.*, 2008; Sheriff *et al.*, 2011). One effective, non-invasive, and

increasingly popular method to evaluate stress is by measuring fecal corticosterone metabolites

(FCMs) (Romero and Remage-Healey, 2000; Tempel and Gutiérrez, 2004; Möstl *et al.*, 2005;

Dickens *et al.*, 2009; Sheriff *et al.*, 2010). Measuring FCMs as an assessment of stress has been

successful in a wide range of avian studies (Washburn *et al.*, 2003; Fletcher *et al.*, 2018; Sokół

and Koziatek-Sadłowska, 2020). Excretion of FCMs varies between species, as does the

suitability of different immunoassays to accurately detect CORT, therefore methods must be

validated for each species (Palme *et al.*, 1996; Wasser *et al.*, 2000; Palme, 2019). Physiological

validation of measuring FCM concentrations requires injecting a known quantity of CORT into

an animal (Touma and Palme, 2005), an invasive and stressful procedure (Palme, 2019;

Mohlman *et al.*, 2020). Biological validation of measuring FCM concentrations is a non-invasive

procedure where feces is collected before and after an event that increases HPA activity for an

extended time, resulting in increased CORT (Touma and Palme, 2005; Palme, 2019).

The goal of this study was to biologically validate the use of FCMs as a non-invasive

measurement of stress in California valley quail using a simulated translocation as the stressor.

This is the first biological validation of FCMs as a tool to evaluate stress in a new world quail

species. Mohlman et al. (2020) performed a physiological validation of FCMs with northern

bobwhite (*Colinus virginianus*), a close relative, but were unsuccessful at biological validation.

Materials and methods

Flight-ready captive-reared valley quail ($N = 5$ females, $N = 8$ males), 16–24 wks in age, were

acquired from a breeder (ZKD Game Birds, West Point, Texas, USA) in October, 2020. Quail

were housed and acclimated in an outdoor aviary (Quail Hotel, Fannin Fabrication, Bonham, TX,

USA) for 3 wks. Birds were provided with gamebird feed (Gamebird Starter and Grower, M-G

Inc., Weimar, TX, USA) and water *ad libitum,* and enrichments including social and

environmental stimuli (Hawkins et al. 2001). This mimicked the time and conditions in which

quail were held prior to the first shipment of trapped birds during a 2019 translocation (Reyna *et*

al., 2020). Acclimation was assumed when birds exhibited typical behavior (e.g., eating,

drinking, roosting, grooming; Reyna and Newman, 2018).

To biologically validate FCMs in valley quail, reference (low stress) fecal samples were

obtained from acclimated valley quail. The floor of the aviary was lined with a plastic sheet and

fecal samples were collected daily. Once collected, individual fecal samples were placed into a

plastic storage bag (Ziploc freezer quart, S. C. Johnson & Son, Inc., Racine, WI) and labeled

with the date, time, and sex of the bird. Temperature, humidity, and presence of direct sunlight

were recorded on the plastic storage bag, since these factors can degrade hormone detection

probability (Shipley *et al.*, 2019). Immediately after collection and labeling, samples were stored

in a freezer at -20 °C until FCM analysis (Wasser *et al.*, 1988; Messmann *et al.*, 1999; Khan *et*

al., 2002).

For biological validation, quail were transported to the animal care facility at Texas

A&M University-Commerce to undergo a simulated 48-h translocation, a stressful event (Martin

et al. 2017). Each quail was given a unique leg band, and had their age, sex, and weight recorded.

Quail were held individually in research-approved breeding pens located in a temperature and

light-controlled room with a 12L:12D photoperiod. To ensure accuracy and reduce

contamination, fecal samples were collected from the underlying waste pan trays 4–5 times daily

(Millspaugh and Washburn, 2004; Mohlman *et al.*, 2020). The frequency of feces collection

ensured a representative sample, since FCMs may fluctuate diurnally (Breuner *et al.*, 1999).

Individual fecal samples were placed in a plastic storage bag and labeled with the bird ID

number, sex, time, and date of collection. Steroid hormone concentrations can be affected by

storage duration (Millspaugh and Washburn, 2004) and bacteria (Lexen *et al.*, 2008). Therefore,

high stress fecal samples were immediately stored in a freezer (-20 °C) until FCM analysis

(Wasser *et al.*, 1988; Messmann *et al.*, 1999).

FCM concentrations of reference and stressed quail fecal samples were measured using

an Enzyme Immunoassay (EIA) Kit (Arbor Assays, Ann Arbor, MI, USA). The kit was validated

for use on CORT extracted from dry fecal extracts on a multi-species design (Jackson, 2015;

Fletcher *et al.*, 2018), and included detailed instructions for extracting CORT from fecal samples

and analyzing FCM concentrations. To determine the assays suitability for valley quail, 10 fecal

extracts were pooled from the valley quail simulated translocation and a serial dilution ($N = 6$

dilutions) was performed. FCM concentrations from the pooled fecal extracts were parallel to the

slope of the standard curve, verifying suitability for valley quail. The sensitivity of the EIA was

determined to be 18.19 pg/ml. Optical density was calculated using a Synergy LX Multi-mode

microplate reader (Bio Tek Instruments, USA), and entered into MyAssay web

(https://www.myassay.com), to calculate FCM concentration in pg/ml. FCM concentrations

reported in pg/ml were converted to ng/g for comparison to other studies.

All animals were handled in accordance with procedures outlined in the Guide for the

Care and Use of Laboratory Animals (National Research Council, 2010), and Texas A&M

University-Commerce Animal Use Protocol P20-013.

Statistical Analysis

All statistical analyses were conducted in R (version 4.0.2, R Foundation, Austria). Shapiro-

Wilk's statistical test was used to check for normality (Zar, 1996). Welch's 2-sample t-test was

used to calculate the p-value, mean, and 95% confidence interval between 2 variables with

normal distributions. Mann-Whitney U-test was used to compare 2 variables with non-normal

distributions. Kruskal-Wallis test was used to compare multiple variables with non-normal

distributions, followed by Dunn's test with Bonferroni correction for pairwise comparisons.

Correlations were tested using Pearson's product-moment correlation with Bonferroni correction

(Zar, 1996). All weight data are presented as mean ± SD. Results were significant at alpha <0.05.

RESULTS

Mean ±SD FCM concentrations of valley quail during the simulated translocation were

36.21 ±18.56 ng/g, and higher than reference FCM concentrations (24.07 ±10.4 ng/g; $N = 13$

quail; Mann-Whitney U, $Z = 4.20$, $P < 0.001$; Fig. 1). Mean 48-hour weight loss was 17.8 ±6.3 g,

or 12 ±4.4% of initial body mass.

Fig 1. Biological validation of using fecal corticosterone metabolites in translocated

California valley quail. Fecal corticosterone metabolites (FCM) concentrations (mean ±SD)

were extracted from scat collected from acclimated quail (*N*=13) prior to (Reference) and during

a 48-hr simulated translocation (Stressed). FCM concentrations from stressed birds were higher

than reference FCM concentrations (Mann-Whitney *U*, *Z* = 4.20, *P* < 0.001). Asterisk indicates

statistical difference.

Valley quail with higher initial body mass prior to the simulated translocation

experienced lower mean FCM concentrations (r^2 = 0.14, *P* < 0.001), and lower percentage of

initial mass lost (r^2 = 0.13, *P* < 0.001) during the simulated translocation. FCM concentrations

positively correlated with percent of initial body mass loss (r^2 = 0.27, *P* < 0.001). No difference

in FCM concentrations (Mann-Whitney U, $Z = 0.15$, $P = 0.44$), total mass lost (Welch's t-test, P

$= 0.58$), or percentage of initial mass lost (Welch's t-test, $P = 0.29$) was observed based on sex.

Valley quail FCMs varied diurnally during the simulated translocation (Fig. 2), with

higher concentrations from 10:00–18:00 and lower concentrations from 20:00–06:00 (Kruskal-

Wallis, H = 62.3, $P < 0.001$). Mean FCM concentrations were not different between the first 24 h

and last 24 h (Mann-Whitney U, $Z = 1.86$, $P = 0.32$).

Fig 2. Diurnal variation in fecal corticosterone metabolite (FCM) concentrations

from California valley quail during a simulated 48-h translocation. FCM concentrations

correlated with photoperiod (12L:12D) and quail circadian rhythm. Letters indicate statistical

groupings.

Discussion

This study was the first biological validation of fecal corticosterone metabolites as a non-

invasive assessment of stress for a new world quail species. When exposed to a simulated

translocation, valley quail experienced a 72.4 % increase in FCMs compared to reference

concentrations, indicating that this protocol successfully detected an increase in FCMs during a

stressful event. These results are comparable to the 73% increase in FCMs recorded in northern

bobwhites during a physiological validation (Mohlman *et al.*, 2020). However, the increase in

FCM levels is lower than that recorded in many other avian species. This isn't surprising because

excretions of metabolites can vary between species (Palme *et al.*, 1996; Wasser *et al.*, 2000;

Palme, 2019). For example. experimentally stressed European starlings showed a ~100%

increase in FCMs (Cyr and Romero, 2008). Following capture and holding for a veterinary

examination, African penguins (*Spheniscus demersus*) showed a 155–349% increase in FCMs.

Wild Dickcissels (*Spiza americana*) experienced a 1,700% increase in FCM concentrations

within 24 h of having leg harness transmitters attached (Suedkamp Wells *et al.*, 2003). Valley

quail and northern bobwhite quail are both new world quail and their similar stress response

during validation, and differences from other taxa were expected.

The correlation of weight loss with FCM levels further supports the association of

chronic stress with weight loss in birds (Cyr and Romero, 2007; Dickens *et al.*, 2009; Awerman

and Romero, 2010). Quail in our study lost ~12% of body weight which is consistent with weight

loss in translocated starlings (5-15%) and wild valley quail (14.3 %) in an actual translocation of

similar duration [60], indicating that translocation weight loss may primarily be a result of stress.

Similar to our study, Reyna et al. (2020) reported that larger birds lost less weight during

translocation. Larger birds may be healthier overall and future studies may benefit from

considering a minimum weight for translocation.

By collecting fecal samples 4–5 times daily during the simulated translocation, we were

able to capture diurnal variation in FCM concentrations. While not previously observed in new

world quail (Mohlman *et al.*, 2020), diurnal variation in corticosterone metabolites has been

recorded in a variety of species (Florant and Weitzman, 1980; de Jong *et al.*, 2001; Touma *et al.*,

2004; Bosson *et al.*, 2009; Sheriff *et al.*, 2009), generally corresponding to the circadian rhythm

of the animal. The diurnal variation in valley quail FCM concentrations was similar to variations

observed in broiler chickens (de Jong *et al.*, 2001), with lower concentrations at night and

elevated concentrations during the day. This trend was expected since valley quail are active in

the day and inactive during the night (Leopold, 1977).

257   This study verifies that translocations are stressful events for quail that result in

significant weight loss; results that suggest stress influences translocation success. Chronic stress

is associated with increased mortality (Cyr and Romero, 2007; Dickens *et al.*, 2010), and animal

weight at release is a predictor of survival (Fountain *et al.*, 2017). Further, stress is a major driver

of reproduction, which is one major indicator of translocation success (Griffith *et al.*, 1989;

Dickens *et al.*, 2010). Future studies could benefit by focusing efforts on stress mitigation to

increase long-term translocation success.

264   The use of FCM as a non-invasive measurement in new world quail could be extended

beyond translocations to benefit quail conservation. There is considerable environmental and

economic interest in conserving and restoring populations of quail and other gamebird species.

Increased plasma CORT can be indicative of environmental disturbances and habitat-related

metabolic challenges (Homyack, 2010; Shipley *et al.*, 2022). For example, FCMs have been used

as an indicator of declining habitat quality in greater sage-grouse (*Centrocercus urophasianus*;

Rabon *et al.*, 2021). FCMs have also been used to detect elevated CORT levels associated with

extreme weather events, noise related to fossil fuel extraction (Cinto Mejia *et al.*, 2019), and

other anthropogenic disturbances (Thiel *et al.*, 2008, 2011; Blickley *et al.*, 2012; Arlettaz *et al.*,

2015; Formenti *et al.*, 2015). Through the use of innovative field collection methods (Shipley *et*

al., 2019), FCMs may be used to assess chronic stress levels in wild quail populations, evaluate

responses to multiple stressors (e.g., extreme drought), and map stress levels across a landscape

to identify focus areas for conservation (Rabon *et al.*, 2021). Ultimately, the use of FCMs could

alert quail biologists to the presence of environmental stressors before population demographics

are impacted (Ellis *et al.*, 2012). Clearly, FCMs have enormous potential as a non-invasive

indicator of stress in quail and as a tool for improving quail research, restoration, management,

and conservation.

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Data Availability

The data underlying this article will be shared on reasonable requests to the corresponding author.

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Fig 1. Biological validation of using fecal corticosterone metabolites in translocated California valley quail. Fecal corticosterone metabolites (FCM) concentrations (mean ±SD) were extracted from scat collected from acclimated quail (N=13) prior to (Reference) and during a 48-hr simulated translocation (Stressed). FCM concentrations from stressed birds were higher than reference FCM concentrations (Mann-Whitney U, $Z = 4.20$, $P < 0.001$). Asterisk indicates statistical difference.

Fig 2. Diurnal variation in fecal corticosterone metabolite (FCM) concentrations from California valley quail during a simulated 48-h translocation. FCM concentrations correlated with photoperiod (12L:12D) and quail circadian rhythm. Letters indicate statistical groupings.

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