Title: Extreme lead tolerance in an urban lizard

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Abstract

Lead (Pb) is an extremely toxic heavy metal pollutant pervasive in many environments with

serious health consequences for humans and wildlife. Identifying organisms that can serve as

biomonitors of lead contamination and models to understand mechanisms of lead tolerance is

therefore an important goal. We used the Cuban brown anole lizard (*Anolis sagrei*) in New

Orleans, USA, to investigate the consequences of variation in environmental lead exposure at the

physiological and gene regulatory levels. We found that brown anole blood lead levels are

associated with soil lead contamination, and that their blood lead levels are extreme. At our most

contaminated site, the mean and highest individual blood lead levels were over two times higher

than the next highest values reported in free-living vertebrates. However, this lead exposure did

not decrease performance in whole-organism traits commonly affected by lead (balance, sprint

speed, endurance). A 60-day lead dosing experiment demonstrated that the brown anole blood

lead threshold for decreased whole-organism performance is actually an order of magnitude

higher than the highest mean field concentration. Transcriptomic analysis of brain and liver from

lizards in high and low lead sites revealed relatively minor effects of lead exposure. However,

several of the differentially expressed genes function in metal ion homeostasis and oxygen

carrying capacity, pointing to cellular mechanisms that may contribute to high lead tolerance.

The brown anole may be the most lead-tolerant vertebrate known to date and has the potential to

be a powerful model system to help us better understand how organisms cope with heavy metal

exposure.

Keywords: urban ecology, *Anolis*, heavy metals, locomotion, pollution, ecotoxicology, gene

expression

Introduction

Anthropogenic activity has resulted in extensive heavy metal contamination in water, soil, and

air, contributing to significant declines in human and wildlife health (Ali and Khan 2019). Lead

(Pb) specifically is a major concern due to its potent toxicity (Needleman 2004). Among many

effects, lead is neurotoxic and genotoxic, causes neuromuscular disfunction and reduced

cognition, diminished reproduction, anemia, and kidney, liver, and heart disease (Needleman

2004). Reflecting this toxicity, the California condor (*Gymnogyps californianus*) dwindled to

near extinction due to lead poisoning in the 1980s (Finkelstein et al. 2012), and the CDC

currently states that no level of blood lead is acceptable for children (CDC 2022). Lead use in

paints, gasoline, and industrial processes has been greatly reduced in recent decades (Levin et al.

2021). However, lead does not break down and therefore many cities globally continue to have

very high lead contamination due to historical deposition (Clune et al. 2011). As a result, finding

organisms that can serve as biomonitors of lead contamination, as well as models of lead

susceptibility and tolerance, remains a pressing issue (Tovar-Sánchez et al 2018).

We used the invasive Cuban brown anole lizard (*Anolis sagrei*) in New Orleans,

Louisiana, USA, as a model system to investigate effects of urban lead contamination,

integrating data on field lead exposure, experimental lead dosing, physiology, and functional

genomics. New Orleans provides an ideal location of study because of decades of research into

fine-scale geographic variation in soil lead contamination across the city (Figure 1A; Mielke et

al. 2016). Soil lead in New Orleans covaries with tissue lead in humans and birds and is

associated with detrimental outcomes (Mielke et al. 2016, McClelland et al. 2019, Hitt et al.

2023). Brown anoles are common in cities across the Southeastern United States, including New

Orleans, are amenable to both field and laboratory study, and already serve as a model system in

urban, physiological, and evolutionary ecology (Battles et al. 2019, Stroud et al. 2019, Deery et

et al. 2021). Lizards are underrepresented in studies of heavy metal pollution, especially with

respect to lead (Grillitsch and Schiesari 2016). Tissue-environment lead associations are known

in some species (Burger et al. 2004, Nasri et al. 2016), but physiological studies of lead exposure

in lizards are rare (Holem et al. 2006, Salice et al. 2009, Grillitsch and Schiesari 2016, Indest et

al. 2018). Therefore, tissue lead toxicity thresholds in lizards, and by extension their toxicity

thresholds relative to better-studied taxa such as birds and mammals, is poorly known.

To test for an association between environmental lead contamination and lizard lead

exposure, we measured blood lead concentrations of brown anoles from sites that vary in soil

lead level across New Orleans. To determine if field lead exposure affects lizard whole-organism

physiology, we tested for associations between blood lead level and multiple metrics of

locomotor performance in field-caught animals. To experimentally determine blood lead toxicity

thresholds, we measured multiple aspects of brown anole physiology during a 60-day lead dosing

study in the laboratory. Finally, to investigate the effects of lead, and potential mechanisms of

lead tolerance, at the molecular level, we sequenced transcriptomes of brown anoles from high

and low lead contamination areas. We found that brown anoles in New Orleans have extreme

blood lead levels and lead tolerance, while gene expression patterns highlight molecular

mechanisms that may contribute to tolerance. Brown anoles may therefore be the most lead

tolerant vertebrate known to date and provide a compelling vertebrate model system to

investigate mechanisms of heavy metal tolerance.

Methods

Field Sampling and Animal Husbandry. Brown anoles were captured by hand in New Orleans,

Louisiana and transported to Tulane University. Animal were housed individually in plastic

cages (23 L x 15 W x 16 H cm) with a wooden dowel provided as a perch in fluctuating

temperature (35/24ºC, day/night) and humidity (60% daytime, 75% nighttime) controlled

Percival chambers on a 14:10 light:dark cycle, conditions that mimic an average summer day in

New Orleans (Deery et al. 2021). Lizards were watered twice daily and fed crickets dusted with

calcium powder three times per week.

Tissue Lead Analysis. We measured lead levels in the blood and bone of field animals. Blood

samples (~75 µl) were collected from the post-orbital sinus using heparinized capillary tubes

(Kimble Chase; Sharma et al. 2014). Each sample was immediately transferred to a trace-metal-

clean microcentrifuge tube and stored at -20ºC until processing. For bone lead analysis, animals

were euthanized via cervical dislocation and whole femur, tibia, and fibula were dissected out

and stored in a trace metal clean micro-centrifuge tubes at -20ºC until processing.

Tissue lead levels of lizards collected from 2019 – 2021 were measured at the UC Santa
Cruz Plasma Analytical Laboratory following established trace metal clean techniques and using
ultra-pure reagents as outlined in Hitt et al. (2023). Blood samples were weighed and then dried
overnight at 60ºC to obtain dry weight, then cold digested for 10 hours in 100 µL (blood volume
<50 µL) or 150 µL (blood volume >50 µL) of concentrated HNO_3 (optima, Fisher Scientific).
30% H_2O_2 (ultrex, JT Baker) and ultrapure water was added to each sample for a HNO_3 to H_2O_2
ratio of 2:1 and a final HNO_3 concentration of 6%. Samples were vortexed and left to sit
overnight before analysis. Lead concentrations were determined by inductively coupled plasma
mass spectrometry (ICP-MS), measuring masses of ^{208}Pb and ^{205}Tl isotopes (used as an internal
standard). For quality control, approximately 20 µL of NIST SRM 955c (lead in blood, level 2)
was digested using the methods described above with an average recovery of 94% ± 3.4% RSD
(N = 4). Frozen bone samples were thawed overnight then combined for each individual and
weighed. They were digested for 6 h with sub-boiling concentrated HNO_3 in closed Teflon vials
then evaporated to dryness and diluted with 5% HNO_3. Lead concentration was then analyzed as
described above.

Blood lead levels in lizards collected in 2022 and 2023 were analyzed at the Louisiana
Animal Disease Diagnostic Lab. 15 µL of blood were diluted in a solution of 0.1% Triton-X100
acidified to 0.2% with HNO_3, resulting in a 21-fold dilution. Seronorm-Trace Elements (lead in
blood, level 2, 303 ng/ml Pb) was used as a standard reference material and digested using the
methods above with an average recovery of 103% RSD. Clotted samples were first weighed and
digested with 600 µL of 35% Trace Metal HNO_3 overnight at 85ºC. Samples were then diluted to
a 1/60 solution with 0.2% HNO_3. Lead concentrations were determined using a Perkin Elmer
PinAAcle 900Z Zeeman graphite furnace spectrometer.

Whole Organism Performance. With field-collected animals, we tested for associations
between individual blood lead level and performance in whole-organism traits known to be
affected by lead exposure in other vertebrates: balance, sprint speed, and endurance (Yu et al.
1996, Burger et al. 2004, Mansouri and Cauli 2009, Mansouri et al. 2013). To measure balance,
lizards were run on a 1.2 m long horizontal wooden dowel (Spezzano and Jayne 2004) covered
with fine mesh. We counted the number of times an individual slipped, defined as anytime the
hind legs lost traction such that the pelvis moved to the side of the dowel (Spezzano and Jayne

2004). Runs were filmed using an Apple iPhone 8 Plus. Lizards were run twice with 15 min

between runs. The dowel was 1.2 m off the ground and foam was placed underneath in case

individuals fell or jumped. The diameter of the dowel was 0.79 cm or 0.64 cm for lizards with a

knee-to-knee width of $\geq$ 27 mm and $\leq$ 26 mm, respectively, to account for inter-knee distance

(Spezzano and Jayne 2004). Trials were conducted at 32ºC, the body temperature of maximum

field activity for *A. sagrei* in New Orleans (Ryan and Gunderson 2021).

Sprint speed was measured by video analysis of lizards running up a 1.8 m wooden track

set at a 37º angle and covered with fine mesh for traction (Gunderson et al. 2018). Perpendicular

lines were drawn every 12.5 cm and the fastest interval was taken as an individual's maximum

speed. Each lizard was run twice at 32ºC and filmed at high speed (120 frames per second) with

an Apple iPhone 8 Plus. Runs were 15 min apart.

We measured endurance as time to exhaustion (s) while lizards ran at 32ºC on a

motorized treadmill (DogPacer MiniPacer) at 0.80 km^{-1} (Lailvaux et al. 2018). Lizards were

placed on the treadmill while tethered with thread around the waist and induced to run by gently

tapping their tails. Exhaustion was indicated by a loss of righting response. Endurance was

measured twice per lizard with 24h between trials and the mean endurance time was used in

further analysis.

Lead Dosing Study. In 2023, N = 50 males were captured at low lead contamination Site B

(Figure 1). After one week of lab acclimation, individuals were randomly assigned to one of five

treatment groups (N = 10/treatment). Treatments were administered daily by oral gavage (Holem

et al. 2006, Salice et al. 2009) for 60 days. The control group received distilled water, while the

treatment groups received 1, 10, 100, or 500 mg lead/kg bodyweight, respectively. Following

methods previously used in lizards (Holem et al. 2006, Salice et al. 2009, Indest et al. 2018), lead

solutions were prepared by diluting lead acetate trihydrate (ThermoFisher Scientific Chemicals)

in distilled water (ThermoFisher Scientific Chemicals). We measured seven traits/endpoints:

symptomatic lead poisoning, endurance, sprint speed, balance, kidney size, liver size, and testis

size (Yu et al. 1996, Burger et al. 2004, El-Said et al. 2008, Mansouri and Cauli 2009, Salice et

al. 2009, Gonick 2011, Mansouri et al. 2013, Hora and Wuestefeld 2023). Symptomatic lead

poisoning was defined as loss of appetite for one week and/or presence of oral ulcers (El-Said et

al. 2008). Animals were removed from dosing if they showed symptomatic lead poisoning.

Endurance was measured as described above three times: the day prior to dosing, after 30 days of

dosing, and after 60 days dosing (i.e., the end of the experiment). Individuals removed early due

to lead poisoning had their endurance measured on the day of removal. Sprint speed and balance

were measured as described above at 60 days on individuals in the control, 1 mg/kg, and 10

mg/kg treatment groups. Blood samples were collected from all individuals prior to the

beginning of dosing and after all performance measurements at the end of dosing. Blood lead

level did not differ among groups prior to dosing (F-value = 1.05, P = 0.394). Animals were

anaesthetized with isoflurane and euthanized via cervical dislocation, after which the liver,

kidney, and testes were removed and wet weight was measured with a digital balance.

With our dosing data, we calculated the blood lead toxicity threshold for a trait as the

highest blood lead level of an individual within the No Observed Effect Concentration (NOEC)

treatment for that trait. NOEC is defined as the highest lead dose that does not cause a significant

change from the control group (Buekers et al. 200).

Statistical Analyses of Lead and Performance Data. All statistical analyses were performed

using R Statistical Software (v4.1.2; R Core Team 2021). For the field collected animals, we

tested for differences in blood and bone lead level between lizards from high and low lead sites

using multiple linear regression with site soil lead level ('high' or 'low') as a predictor variable

and sex and snout-vent-length (SVL) as covariates. We tested for an association between blood

and bone lead level within individuals with a Pearson product-moment correlation test. We tested

for associations between individual blood lead level and endurance and sprint speed with

multiple linear regression with sex and SVL as covariates. For tests with balance, negative

binomial models were used with sex and relative leg length (residual knee-to-knee width relative

to SVL) as covariates.

For the dosing study, we first tested for differences in blood lead level among treatment

groups prior to dosing with a one-way ANOVA (there were none; P = 0.394). We tested for

differences in blood lead level within and among treatment groups due to dosing using a two-

way repeated measures ANOVA. One individual in the 500 mg/kg treatment group was found

deceased on day 10 of dosing and was excluded from the analyses since a second blood sample

could not be collected. We tested for effects of lead dosing on physiological performance at the

treatment and individual level. For treatment-level analyses, treatment group was a predictor

variable, and for individual-level analyses, individual blood lead level was a predictor variable.

For endurance, linear mixed effects models were used with endurance as the response variable,

day of measurement and SVL as covariates and individual as a random effect. Two individual-

level analyses were conducted for endurance, one with data from all treatments including

animals that were removed early due to lead poisoning, and another with only the animals that

made it through the entire study. We also tested for differences in sprint speed and balance at the

end of the study among the Control, 1 mg/kg, and 10 mg/kg animals at the treatment and

individual levels. We tested for differences in balance among treatment groups using a general

linear model with a Poisson distribution. We tested for differences in liver, kidney, and testis

mass among treatments using one-way ANOVAS. Organ masses were normalized to body mass

(organ wet mass/total body mass) before analysis.

Transcriptome Analysis. To investigate the cellular processes affected by lead exposure, and to

provide insight into how the lizards may tolerate high lead levels (see Results), we tested for

differential gene expression of lizards collected from a high (Site F) and a low (Site B) lead

contamination area in 2021 (Figure 1; N = 5 males and N = 5 females per site). Forebrain (males)

and liver (males and females) transcriptomes were sequenced. We chose brain due to the

neurotoxicity of lead and liver due to its role in detoxification (Needleman 2004, Hora and

Weustefeld 2023). Forebrain was only collected from males due to small female size. After one

week in the laboratory under our standard housing regime (see above), individuals were

anaesthetized as described above and the target tissues were immediately dissected out, placed in

RNAlater, flash frozen in liquid nitrogen, and stored at -80°C. Total RNA was extracted using

the Qiagen RNeasy Plus Universal Tissue Mini Kit. Transcriptome libraries were prepared and

sequenced at Novogene Corporation Inc (Sacramento, CA) using 150-base pair paired-end

sequencing on the Illumina NovaSeq 6000 platform. Sequencing resulted in an average of

$25{,}569{,}255 \pm 3{,}909{,}485$ reads per transcriptome.

We used Rcorrector to correct random sequencing errors in reads and removed kmers

with errors that were unfixable (Song and Florea 2015). Trimmomatic was used to filter the

resultant sequences based on quality, which was assessed for each read with a 4bp sliding

window (Bolger et al. 2014). Sequences were trimmed when Phred33 quality scores fell below

an average of 15 within the window. Adaptor sequences, if detected, were removed. Quality-

controlled sequence reads were mapped to the *Anolis sagrei* genome (AnoSag2.0; Geneva et al.
2021) using STAR (Dobin et al. 2013).

We used the DESeq2 package (Love et al. 2014) to identify differentially expressed
genes (DEG) between high and low lead groups based on sex and tissue type. We filtered out
low expression genes, and only conducted analyses on those with a count threshold of 10 in at
least three samples (Love et al. 2014). False discovery rate (FDR) correction for multiple
comparisons was applied to maintain a table-wide alpha of 0.05 (Storey and Tibshirani 2003).
Genes were considered differentially expressed when the absolute value of log2 fold expression
difference between high and low lead groups was $\geq$ 1 and FDR was $\leq$ 0.05 (Love et al. 2014).
Volcano plots were generated using EnhancedVolcano (version 1.16.0; Blighe et al. 2022). Gene
Ontology enrichment analyses were conducted on differentially expressed genes using Enrichr
(Kuleshov et al. 2016).

Results and Discussion

Field Lead Exposure. Lizard blood lead levels varied with soil lead contamination across New
Orleans (Figures 1A, B). In the two sites with the most data (low lead contamination Site B and
high lead contamination Site F; N = 40/site), blood lead in the high contamination site ($\bar{x} \pm$ SD;
955±877 µg/dL) was over 23 times higher than that of lizards from the low contamination site
(40.4 ± 50.6 µg/dL; P = < 0.001; Figure 1B; Table S1). Additionally, blood lead levels across the
city frequently exceeded the reported toxicity thresholds (Buekers et al. 2009) for whole-
organism physiological effects in birds (71 µg/dL) and mammals (18 µg/dL; Figure 1B). The
highest individual blood lead level was 3192 µg/dL, which is 2.3 X higher than the highest
individual blood lead levels that we could find reported for an individual free-ranging vertebrate,
a griffon vulture (*Gyps fulvus*) that died within 24 hours of capture (Carneiro et al. 2016; Figure
1B).

Figure 1. Soil and lizard lead levels in New Orleans. (A) Soil lead concentrations in New Orleans[8] with our study sites indicated. (B) Blood lead concentrations of brown anoles by study site. Dashed red lines: reported whole-organism physiological toxicity thresholds for birds (71 µg/dL) and mammals (18 µg/dL; Buekers et al. 2009). Blue dashed line: the highest individual blood lead level from a live free ranging vertebrate that we could find in the literature (1384 µg/dL; griffon vulture (*Gyps fulvus;* Carneiro et al. 2016). (C) Bone lead levels of lizards collected from Sites B and F. (D) Relationship between bone and blood lead levels for the subset of individuals from Sites B and F for which we have both measures.

Blood lead reflects shorter-term exposure because lead is absorbed into the bloodstream within hours of ingestion and has a half-life that ranges between 3 and 30 days in wildlife and humans (Elder et al. 2014). To determine if the lizards are exposed to lead over longer timescales we measured lead in bone, which is incorporated during development and can remain for years (Needleman 2004). Bone lead was significantly higher in high contamination Site F (39.9 ± 38.1

µg/g) than in low contamination Site B (4.67 ± 5.83 µg/g; P = 0.012; Figure 1C), indicative of

long-term exposure in high contamination areas. Furthermore, individual bone and blood lead

levels were significantly correlated ($r = 0.86$, $t_{(18)} = 7.16$, $P < 0.001$; Figure 1D), consistent with

lizards residing within their respective sites throughout their lives. Indeed, wild brown anoles

rarely move more than a few meters from their natal site (Calsbeek 2009) and adults have small

home ranges (mean = 17.6 m^2 for males; Schoener and Schoener 1982). Such high philopatry

and adult site fidelity is the norm for *Anolis* lizards generally (Schoener and Schoener 1982), and

therefore these lizards (and perhaps lizards more broadly; Talent et al. 2002) are strong

candidates to serve as biomonitors of environmental lead contamination.

To place the blood lead levels of brown anoles in New Orleans into a broader context, we

compiled all mean blood lead levels we could find reported for live, free-living adult vertebrates

(minimum sample size N = 10). If multiple values were found for the same species, only the

highest mean value was included. Brown anoles in New Orleans are clearly exceptional (Figure

2). The mean lead level in Site F is over three times higher than the next highest mean value

(Nile crocodiles, $\bar{x} = 250$ µg/dL; Humphries et al. 2022), and 10 times higher than the peak mean

level in California condors that experienced population declines due to lead exposure (130

µg/dL; Finkelstein et al. 2012; Figure 2). How these lizards attain lead is currently unknown.

High lead levels in crocodiles and scavenging birds like condors and vultures are linked to the

ingestion of large pieces of lead associated with fishing and hunting equipment (Pain and Mateo

2019, Humphries et al. 2022). That pathway of lead exposure is not possible for brown anoles as

small (adult mass = 1.5 – 8 g) insectivores that feed on live prey. Instead, they must ingest it

from a combination of diet, drinking water, and breathing or ingesting lead-contaminated dust.

Arthropods do accumulate lead in lead-contaminated areas (Khan 2019), although a significant

portion of that lead may be stored in the exoskeleton in a form not bioavailable to predators

(Inouye et al. 2007). Ingestion of lead in dust is a very common means of lead exposure for

humans and wildlife (Mielke et al. 2016), and brown anoles spend the majority of their time near

and on the ground in close proximity to contaminated soils.

Figure 2. Mean blood lead concentrations of free-living vertebrates. The brown anole value is from high lead contamination Site F (Figure 1). Data for other taxa were reported in the literature (minimum N = 10). If multiple values for a species were found we included only the highest. See Dataset S1 and References S1 for more information.

Physiological Performance of Field Exposed Lizards. Given their high lead levels, we predicted that field-exposed lizards would experience physiological detriments. Contrary to our expectation, blood lead was not associated with decreased performance (Figure 3). Blood lead was unrelated to balance (P = 0.276; Figure 3A; Table S2), positively related to sprint speed (P = 0.01; Figure 3B; Table S3), and trended towards a positive relationship with endurance (β = 0.04; P = 0.086 Figure 3C; Table S4). Brown anoles in New Orleans therefore appear to have greater lead tolerance than birds and mammals (Buekers et al. 2009). However, these data do not allow us to estimate their blood lead thresholds because no performance detriments were observed. To estimate brown anole lead toxicity thresholds, we turn to our experimental lead dosing experiment.

Figure 3. Physiological performance of individual field-caught brown anoles relative to blood lead level. (A) Balance. (B) Sprint speed. (C) Endurance.

Effects of Experimental Lead Dosing. Blood lead level increased in all groups over time consistent with treatment level (Figure 4A, Tables S5, S6). The final mean blood lead level of lizards in all of the lead-dosed treatments was higher than the mean blood level at our most contaminated field site (Figure 4A, Table S5). All lizards in the 100 and 500 mg/kg treatments exhibited symptomatic lead poisoning before the end of the experiment and were removed from dosing by day 40 and day 10, respectively. In contrast, no lizards exhibited symptomatic lead poisoning in the Control, 1 mg/kg, or 10 mg/kg treatments. The NOEC for symptomatic lead poisoning is therefore 10 mg/kg, and the blood lead toxicity threshold is 10,600 µg/dL (Table 1).

Figure 4. Effects of lead dosing on brown anole lizards. (A) Blood lead concentrations before

and at the conclusion of dosing. Post-dosing samples were collected at 60 days except for the

individuals removed early due to symptomatic lead poisoning in the 100 mg/kg and 500 mg/kg

treatment groups. Points and bars indicate means and standard errors, respectively. The dashed

line indicates the mean blood lead level of field-caught brown anoles from Site F, which had the

highest lead level of all field sites. (B) Locomotor endurance by treatment group over time. (C)

Sprint speed and (D) balance of lizards in the Control, 1 mg/kg, and 10 mg/kg treatment groups

at the conclusion of dosing. (E) Organ wet weight as percent of total body mass by treatment.

Table 1. Brown anole lead toxicity thresholds from the dosing experiment. NOEC = No Observed Effect Concentration. Blood lead toxicity threshold = highest individual blood lead concentration in the NOEC treatment. We do not have a direct estimate of NOEC for kidney and liver size because both organs were enlarged in the lowest-dose treatment, and we report the highest individual blood lead concentration in the Control treatment as the toxicity thresholds for both.

Endpoint	NOEC (mg/kg)	Blood lead toxicity threshold (µg/dL)
Symptomatic lead poisoning	10	10,600
Endurance	10	10,600
Sprint speed	10	10,600
Balance	10	10,600
Kidney size	< 1	130
Liver size	< 1	130
Testis size	10	10,600

Endurance differed by treatment (F = 4.81, P < 0.001), decreasing over time in the 500 mg/kg and 100 mg/kg treatments, but not in the others (Figure 4B). The NOEC for endurance is 10 mg/kg, and the estimated blood lead toxicity threshold is 10,600 µg/dL (Table 1). We also tested for an association between final individual blood lead level and final endurance. With data from all treatments, blood lead is negatively associated with endurance (blood lead range = 53 – 80,000 µg/dL; R^2 = 0.47, β = -0.007, P < 0.001; Figure S1A, Table S7). However, when animals from the treatments above NOEC were excluded, endurance and blood lead were unrelated (blood lead range = 53 – 10,600 µg/dL; R^2 = 0.05, β = 0.001, P = 0.432; Figure S1B).

Sprint speed and balance were unrelated to lead for the only animals that remained at the end of the experiment (the Control, 1 mg/kg, and 10 mg/kg groups) at both the treatment (both P > 0.325; Figures 4C, D; Tables S8, S9) and individual blood lead levels (both P > 0.781; Figures S1C, D; Tables S8, S9). We therefore cannot directly calculate NOEC or blood lead toxicity thresholds for these traits, though we can say that the NOEC and blood lead threshold are at least as high as for endurance (Table 1).

Given the dosing results, it is unsurprising that we found no whole-organism

physiological detriments of what at first appeared to be very high levels of field lead exposure

(Figure 3). The whole organism performance of these animals is not affected until blood lead

concentration is much higher than field levels. In our site with the highest lead levels, Site F, the

mean blood lead level was 955 µg/dL and the highest documented individual blood lead level

was 3192 µg/dL (Figure 1). However, we only began to see whole organism performance effects

for individuals in the 100 mg/kg treatment, which had a mean blood lead level 12x higher than

the field mean (Figure 2A, Table S5).

All lead-dosed treatments had larger kidneys (F-value = 56.47, P < 0.001) and livers (F-

value = 12.79, P < 0.001) than Controls, including the 1 mg/kg and 10 mg/kg groups that

exhibited no whole-organism evidence of lead effects (Figure 4E). In contrast, testes were only

enlarged in the 100 mg/kg and 500 mg/kg treatments (F-value = 29.50, P < 0.001; Figure 4E).

The NOEC for effects on liver and kidney size are therefore lower than for whole-organism traits

and cannot be directly estimated, but we know the NOEC must be less the 1 mg/kg (Table 1).

We report the blood lead toxicity threshold as 130 µg/dL, the highest individual concentration in

the Control group (Table 1). Whether or not the liver and kidney responses reflect an adaptive

response to lead or pathology are unclear and require further investigation. The kidney and liver

remove and sequester heavy metals (Gonick 2011, Hora and Wuestefeld 2023) and increased

organ mass may improve detoxification (Hora and Wuestefeld 2023). However, enlarged organs

can also reflect heavy metal-induced tissue damage (Habeebu et al. 2000, Gonick 2011). The

response of testis mass to dosing was similar to that of whole-organism traits, and had the same

thresholds (NOEC = 10 mg/kg; blood lead = 10,600 µg/dL; Table 1). Testes do not generally

function in heavy metal detoxification or adaptive sequestration, so changes in testis mass likely

reflects harm from lead exposure (Salice et al. 2009, Heidaeri et al. 2021).

Differential Gene Expression of Field Exposed Lizards. The high lead tolerance of brown

anoles raises the questions of how this tolerance is achieved. Lead exposure in vertebrates

typically causes the differential expression of hundreds if not thousands of genes within a given

tissue (Schneider et al. 2012, Shi et al. 2020). Genes commonly affected by heavy metal

exposure are those related to the cellular stress response (e.g., antioxidants and molecular

chaperones; Needleman 2004, Gupta et al. 2010), transport of key ions that heavy metals can

replace within biological processes (Needleman 2004), and genes that function in the binding

and sequestration of metals (Gonick 2011). Indeed, tolerance for lead in vertebrates has been

linked to metallothioneins (MT), a family of metal-binding proteins that can function in

sequestering lead in a nontoxic form (Gonick 2011).

We found relatively few genes differentially expressed (DE) in high versus low lead

brown anoles. In male forebrain, there were only four DE genes in high lead lizards, two

upregulated and two downregulated (Figure 5A). In male liver, 51 genes were DE, with 41

upregulated and 10 downregulated (Figure 5B). In female liver, 23 genes were DE, 14

upregulated and 9 downregulated (Figure 5C).

Figure 5. Differential gene expression of lizards from high compared to low lead

contamination sites. Lizards were from high contamination Site F and low lead contamination

Site B. Highlighted genes function in ion binding and transport, the oxidative stress response, or

oxygen binding and transport. (A-C) Volcano plots of differential expression in the (A) male

forebrain, (B) male liver, and (C) female liver. Cutoff for significance (red dots) is False

Discovery Rate < 0.05 and an absolute log2 fold change $\geq$ 1. (D) Venn diagram of unique and

shared differentially expressed genes across sex and tissue type.

There were no significantly enriched Gene Ontology (GO) categories across all DE genes
after correction for multiple comparisons (Table S10). However, six Biological Processes neared
significance (0.10 > P > 0.05), including Intracellular Iron Ion Homeostasis (GO:0006879) and
Oxygen Transport (GO:0015671). Many of the DE genes are involved in the binding, transport,
and homeostasis of metals and other ions (*alas2, frrs1, ypel3, plscr1, tmem161b, cacnb3, slc7a9,*
slc4a1, slc38a4, slc22a13, slc11a1), and are also DE under heavy metal exposure in other taxa
(He et al. 2009, Plusquin et al. 2012, Schneider et al. 2012, Truong et al. 2018, Smith et al.
2023). The response of several solute carrier genes (*slc*) is intriguing. SLCs are transmembrane
proteins that function in ion transport into and out of cells and have been implicated in heavy
metal susceptibility and tolerance (He et al. 2009, Schneider et al. 2012). We found no DE of
metal-binding metallothionein genes.

One of the key consequences of lead exposure is anemia (Fuller and Wiley 2018) because
lead interferes with heme production and decreases hematocrit, hemoglobin, and erythrocyte
levels in vertebrates (Kwong et al. 2004), including western fence lizards (Salice et al. 2009).
Key genes related to oxygen carrying capacity were upregulated in high lead anoles: *alas2* in
female liver, and *hba1* and *hbe1* in both male and female liver (two of only four genes
differentially expressed in both sexes). ALAS2 is a rate-limiting enzyme in heme production
within red blood cells (Tanimura et al. 2016). *Hba1* produces alpha globin 1, one of two globins
that form the mature hemoglobins that carry oxygen within vertebrate red blood cells. *Hbe1* is
not part of primary hemoglobin function, but *hbe1* upregulation in mice ameliorates sickle cell
disease and pathological underproduction of hemoglobin (thalassemia; Wang et al. 2016).
Furthermore, there is evidence for parallel natural selection on the *hbe1* gene in populations of
humans, dogs, yaks, and horses adapted to high-altitude hypoxia (Liu et al. 2019).

None of the top GO terms were associated with stress, and few stress-response genes
were upregulated in high lead lizards. There were some key exceptions. *fkbp5*, *map3k15*, and
gtf2h2 are stress-related genes upregulated in male liver. *Fkbp5* produces an HSP90 cochaperone
integral to the stress response (Gupta et al. 2010) and is also upregulated in carp cells in response
to cadmium (Luan et al. 2022). *map3k15* is involved in stress-induced apoptosis, while *gtfh2* is
involved in DNA repair (Tulay et al. 2015, Liu et al. 2022). *pfkm*, upregulated in female liver,
can be induced by oxidative stress (Tang et al. 2012). None of the antioxidant genes frequently

associated with heavy metal exposure, such as sodium oxide dismutase, catalase, or glutathione
peroxidase, were DE.

Overall, we found minimal gene regulatory response to high levels of environmental lead
exposure in brown anoles. Given the lack of higher-level physiological effects (Figure 3) this is
perhaps not surprising, and further reinforces that these animals are very lead tolerant. Among
the few genes that were differentially expressed, there are intriguing clues as to how the lizards
tolerate lead. In particular, plasticity in the expression of solute transport genes and genes
involved in oxygen carrying capacity are clearly candidates for further study.

Conclusion

We found that brown anoles are reliable lead biomonitors (Burger et al. 2004) and provide
evidence that they have extreme lead tolerance. Indeed, they may be the most lead tolerant
vertebrate known. Many questions remain. Because of the scant attention paid to lizard heavy
metal tolerance, we do not know if our findngs reflect only brown anoles in New Orleans, brown
anoles in general, the genus *Anolis*, or squamates more broadly (Moore et al. 2025). The high
lead tolerance we observed could result from evolution in response to urban lead exposure.
Heavy metal tolerance can evolve rapidly (Merritt and Bewick 2017), and *Anolis* lizards are
known to quickly physiologically evolve in response to environmental change (Gunderson and
Leal 2012, Campbell-Staton et al. 2017). Regardless, brown anoles represent a compelling, novel
vertebrate model for developing a mechanistic understanding heavy metal tolerance with
extensive resources available (Rasys et al. 2019, Geneva et al. 2021). Our findings also point to a
need for greater consideration of heavy metals in the rapidly growing field of urban ecology and
evolution (Rivkin et al. 2019). Cities across the globe have high levels of heavy metal pollution
(Clune et al. 2011), but far more effort is devoted to factors such as urban heat islands, light and
noise pollution, structural habitat modification, and invasive species (Rivkin et al. 2019). Taking
steps to better understand the effects of heavy metals in urban areas, and how they interact with
other well-known urban stressors, outside of a traditional ecotoxicological context will likely
yield important insights (Schneeweiss et al. 2023, Sigmund et al. 2023).

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Competing Interests

The authors declare no competing interests.

Author Contributions

A.B., A.G. and J.K. conceived the project. A.B. and A.J.F. conducted physiological experiments.
A.B. and K.S. conducted bioinformatic analyses. A.B. and A.G. wrote the paper with input from
other authors.

Data Availability

Supplementary information is available for this paper. Data presented here are available in a
GitHub repository (https://github.com/ablanchette23/Brown-anole-data.git)

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