Understanding glycaemia and glycation levels in birds: diet and life history traits associations

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ABSTRACT

The pace of life hypothesis (POL) suggests that organisms' life history, physiological and behavioural traits should co-evolve. In this context, glycaemia (i.e., blood glucose levels) and its consequences to ageing through its reaction with proteins and other compounds (i.e. glycation) constitute physiological traits which remain under-investigated in an evolutionary comparative framework, notably in relation to life history traits and ecological factors. Although their blood glucose levels are twice as high as in mammals, birds live longer when accounting for their body mass. We thus performed a comparative analysis on glucose and albumin glycation rates on 87 bird species from 22 orders, in relation to life history traits (body mass, clutch mass, maximum lifespan and developmental time) and diet. Glucose levels correlated positively with albumin glycation rates in a non-linear fashion, suggesting resistance to glycation in species with higher glucose levels. Plasma glucose levels decreased with increasing body mass and increased with maximum lifespan before reaching a plateau. Finally, terrestrial carnivores showed higher albumin glycation compared to omnivores. Our results open an important new avenue of research to help us to better understand the striking condition of hyperglycaemic yet long-lived birds.

INTRODUCTION

The pace of life (POL) hypothesis [1][2][3] postulates that organisms behavioural/physiological characteristics and life histories have co-evolved in answer to specific ecological conditions along a continuum. According to these characteristics, the pace of life can be either slow (e.g. large body mass, late maturation, slow growth rate, small number of large offspring and high longevity) or fast (e.g. small body mass, early maturation, high growth rate, a large number of small offspring and a short lifespan (see [4][5][6][7]). Studies have often tested predictions of POLS by focusing on metabolic rate (e.g. [8][9][10][11]). However, key energy substrates related to metabolic performance, such as glucose concentrations in tissues, have been largely overlooked (but see [12][13]). Notably, glucose, a central energy source for many organisms, plays a pivotal role in metabolism and its circulating blood or plasma levels correlate positively with metabolic rate (e.g. [14] for interspecific comparison; [15] for healthy humans).

Research focusing on plasma glucose becomes highly relevant as high glycaemia can entail costs that accelerate ageing. Along with other reducing sugars, glucose can react non-enzymatically with free amino groups of proteins, lipids or nucleic acids, a reaction known as glycation [16][17].

This reaction, after several molecular reorganizations, can lead to the formation of a plethora of

molecules called Advanced Glycation End-products (AGEs) [18]. AGEs can form aggregates in

tissues that contribute to several diseases related to ageing like diabetes (e.g., reviews by

[19][20][21][22]). Also, AGEs are known to promote a proinflammatory state through their

action on specific receptors called RAGEs [23][24].

Remarkably, birds show circulating glucose levels much higher than other vertebrate groups, on

average twice as high as mammals [25]. Such relatively high glucose levels might support the

higher metabolic rates [26][10] and body temperatures observed in birds compared to mammals

[27][28]. In addition, elevated glycaemia is thought to be an adaptation to flight, providing birds

with rapid access to an easily oxidizable fuel during intense bursts of aerobic exercise (see

[29][30]), such as during take-off and short-term flapping flights [31][32]). However, the

presence of high glycaemia in birds is also paradoxical, given their remarkable longevity

compared to their mammalian counterparts — living up to three times longer than mammals of

equivalent body mass [33][10].

Several non-excluding hypotheses have been proposed to explain how birds apparently resist

the pernicious effects of a high glycaemic state. One possibility is that birds might resist protein

glycation through a lower exposure of lysine residues at the surface of their proteins [34].

Another hypothesis suggests that increased protein turnover may play a role [35][36].

Additionally, birds might benefit from more effective antioxidant defences [37][38][39]

or even the presence of "glycation scavengers" such as polyamines which circulate at high

concentration [32]. Moreover, it is generally considered that RAGEs are not present in avian

tissues, suggesting possible adaptive mechanisms that would allow birds to avoid the

inflammatory consequences associated with RAGE activation [40][41]. However, Wein et al. [42]

identified a putative candidate for RAGE and a case of AGE-induced inflammation, warranting

further investigation.

To date, several studies have measured protein glycation levels in birds, but these have included

a reduced number of species and are merely descriptive [43][44][45][46][47][48][49][50][51].

Additionally, many of these studies employ commercial kits that were not specifically designed

for avian species, making it challenging to interpret the results [50]. These factors limit the ability

to draw general conclusions from these studies. Here, we performed a comparative analysis on

87 species belonging to 22 orders and assessed whether and how bird glycaemia and glycation

rates are linked to ecological and life-history traits. While doing this, we checked whether

glycation levels are influenced by circulating plasma glucose concentrations. In addition, diet is

a key factor likely to influence glycaemia (see e.g. [52] but [13] found no significant effects), and

glycation. We hypothesized that carnivorous birds would exhibit higher glycaemia and glycation

levels than omnivorous or herbivorous species. This hypothesis is based on the evidence that a

low-carbohydrate, high-protein, high-fat diet is associated with hyperglycaemia and reduced

insulin sensitivity in some vertebrates (e.g.; [53][54][55]). This may be attributed to high levels

of constitutive gluconeogenesis, which has been confirmed in certain raptors [56][57].

Conversely, species with high sugar intake, such as frugivores or nectarivores, are expected to

exhibit high glycaemia, as observed in hummingbirds [46]. In line with POL theory, we also

hypothesized that species with a slower POL (hence slower metabolism and/or having specific

resistance mechanisms) would show lower glycaemia and higher glycation resistance (glycation

levels adjusted for glycaemia) compared to species with a faster POL. Hence, we included four

parameters in our analysis: body mass, maximum lifespan, clutch mass and developmental time.

Body mass is one of the main factors affecting life history, with higher body mass associated

with "slower" strategies and vice versa [58]. Previous studies have reported a negative

relationship between body mass and glycaemia [59][31][12][13], although [46] and [60] found

non-significant trends in birds). Therefore, we predicted that bird species that live longer,

develop more slowly and invest less per reproductive event should show lower plasma glucose

levels and albumin glycation rates.

MATERIALS AND METHODS

Species and sample collection

A total of 485 individuals from 87 species were included in this study. A detailed list with the

provenance of the samples is provided in ESM 3, with a textual description in ESM 6.

Blood samples were collected from the brachial or tarsal vein, or from the feet in the case of

swifts, using heparin-lithium as anticoagulant. Samples were centrifuged at 4°C, 3500 g for 10

min and plasma was aliquoted when a large volume was available. Subsequently, they were

transported on dry ice to the IPHC (Institut Pluridisciplinaire Hubert Curien) in Strasbourg and

stored at -80º C until analysis. Glycaemia was measured on plasma with a Contour Plus®

glucometer (Ascensia diabetes solutions, Basel, Switzerland) and expressed in mg/dl. Due to

methodological constraints, glycaemia values of 96 individuals from 40 species (coming from

different sources) were not measured as described. For these species, mean plasma glucose

values were obtained from the ZIMs from Species360 (Species360 Zoological Information

Management System (ZIMS) (2023), zims.Species360.org).

Glycation levels

Glycation levels for each individual were determined using liquid chromatography-mass

spectrometry (LC-MS). Briefly, 3 µl of plasma were diluted with 22 µl of distilled water containing

0.1% of formic acid, followed by injection of 5 µl into the system. Detailed information regarding

parameters and data processing can be found in [50]). The glycation values used in the analyses

represent the total percentage of glycated albumin, obtained by adding the percentages of

singly and doubly glycated albumin. In cases where albumin glycation values dropped below the

limit of detection, resulting in missing data, these individuals (10 individuals from 4 species) were

excluded from the statistical analyses, as outlined below.

Data on ecology and life-history

Diets of individual species were extracted from the AVONET dataset ([61], coming from[62], and

those adapted from [63]) and life history traits from the Amniote Database [64]. After realizing

that diet classifications from AVONET were not always suitable for our purposes, minor changes

were made after consulting the original Wilman et al. [63] database (see ESM 6).

For missing data after this stage, we extracted values, in order of preference, from the AnAge

database [65] or the Birds of the World encyclopaedia [66], by calculating mean values if an

interval was given, and then averaging if male and female intervals were provided separately.

For European species, longevity records have always been checked against the most recent

Euring database [67]. Maximum lifespans from *Chionis minor* and *Eudyptes chrysolophus* (not

available from the previous sources) were extracted from the Centre d'Etudes Biologiques de

Chizé, Centre National de la Recherche Scientifique (CNRS-CEBC) database. Several species from

the zoo had maximum lifespan values available on the ZIMs from Species 360 (Species360

Zoological Information Management System (ZIMS) (2023), zims.Species360.org), so these were

also compared with the data we had from other sources.

In the case of variables for which two different sources provided different records, an average

value was calculated. For the maximum lifespan value, available sources were cross-checked

and the highest value was always used. In the absence of satisfactory support from another

source, values for maximum lifespan indicated as being anecdotal and of poor quality have been

excluded from the analyses. A table with the adequate citations for each value is provided as

part of the online available data (see Data accessibility section). When the source is AVONET

[61], this database is cited, and when it is Amniote Database [64], we cite the sources provided

by them, so the references can be checked in [64]. A list with all the additional references not

coming from any of these databases or not provided by their authors (and that are not already

in the main text) is given as ESM 4.

Statistical analyses

All analyses were performed in R v.4.3.2 [68]. Alpha less than 0.05 was considered as significant,

and between 0.05 and 0.1 as trends. All models were run with 6×10^6 iterations, with a thinning

interval of 50 and a burn-in of 1000. The models were simplified by eliminating quadratic terms

where they were not significant and selecting models with lower AIC (Akaike Information

Criterion) and BIC (Bayesian Information Criterion). For the phylogenetically controlled analyses,

consensus trees (each with the species included in the model) were obtained by using the

consensus.edges function (from the *phytools* package from R [69] from a total of 10,000 trees

downloaded from Birdtree.org (option "Hackett All species", [70]). The consensus trees were

included as *pedigree* in the models. We performed models with either glucose or glycation as

dependent variables and with the diet and life history traits as predicting variables. Models

testing the effects of age and sex on glucose and glycation levels and the number of exposed

lysine residues on glycation were also carried. A detailed description of all of them is given on

the ESM 6.

RESULTS

Plasma glucose and albumin glycation variability across the birds' tree

Plasma glucose and albumin glycation values varied considerably across species (**Figure 1.A and**

1.B, respectively). We found significant differences between orders for both glucose (F=30.54,

p-value<0.001) and albumin glycation (F=21.9, p-value<0.001) with Passeriformes (mean=354.6,

se=±5.024 n=90, min=165, max=488; all values in mg/dl) and Apodiformes showing the highest

average glucose values (mean=350.5, se=±7.63, n=40, min=287.5, max=414.5) and the

Rheiformes, Suliformes and Phoenicopteriformes the lowest average values (mean=157.2,

se=±19.46, n=6, min=105, max=212). For glycation, Apodiformes had the highest average values

(mean=24.87, n=40 se=±0.59, min=12.46 max=28.73) while Phoenicopteriformes had the lowest

average levels (mean=10.57, se=±0.80, n=22, min=8.57, max=12.41). Graphs and a table with

the difference between groups are shown on ESM2 (**Figure ESM2. 3 and Table ESM2. 1**). Raw

outcomes of all models are available on ESM1.

Figure 1.A Average plasma glucose values in mg/dl from all the species used in this study (some of them coming from ZIMs database; see methods) with the Orders they belong to. B Average albumin glycation rate for all the species measured in this study. Tree from a consensus on 10,000 trees obtained from Hackett All species on Birdtree.org, including 87 species from 22 orders (see methods).

Bird glycemia appears affected by body mass and maximum lifespan

After controlling for intraspecific variation in our analyses, we found that variation in glucose levels were significantly explained by variations in body mass and the quadratic component of mass-adjusted maximum lifespan, but not by variations in diet, clutch mass and developmental time (referring to the model including life-history trait variables; **Table 1**). Heavier species had lower glucose levels (see **Figure 2.A**, drawn with the estimates from the model without life history traits, which includes more species). Glucose levels increase with increasing mass-adjusted lifespan until reaching a plateau (**Figure 2.B**). Models that did not consider interspecific

variability show no significant effect of any of the aforementioned variables on glucose levels
(see ESM1).

Table 1. Final glucose model with diet, body mass, and life history traits as explanatory variables, including
the significant quadratic effect of maximum lifespan. Posterior means, CI_{95} and pMCMC from a
phylogenetic MCMC GLMM model including n=326 individuals of 58 species. Both glucose and body mass
are log transformed and life history traits are residuals from a linear model of log body mass and log of
the trait in question. Body mass was also centred to better explain the intercept, as 0 body mass would
make no biological sense. The intercept corresponds to the Omnivore diet, being used as the reference
as it is considered the most diverse and "neutral" group for this purpose. Significant predictors are
indicated in bold.

	Estimates	Lower 95% CI	Upper 95% CI	Sampling effort	pMCMC
Intercept	**2.409**	**2.310**	**2.508**	**118632**	**< 0.001**
Diet: Carnivore terrestrial	-0.009	-0.077	0.059	119800	0.796
Diet: Aquatic predator	-0.004	-0.079	0.071	119800	0.932
Diet: Herbivore	-0.055	-0.175	0.061	119800	0.345
Diet: Frugivore/granivore	-0.032	-0.177	0.115	119800	0.667
Centered log(Body mass)	**-0.045**	**-0.084**	**-0.007**	**118572**	**0.025**
Maximum lifespan	0.108	-0.001	0.220	119800	0.056
Maximum lifespan^2	**-0.588**	**-0.998**	**-0.150**	**119800**	**0.009**
Clutch mass	-0.114	-0.260	0.034	119800	0.122
Developmental time	0.027	-0.133	0.193	122686	0.746

Figure 2. Plasma glucose levels (in mg/dl) variation as a function of **A** species mean centred body mass and **B** residual maximum lifespan. Both glucose and body mass are log transformed. Maximum lifespan (in years) is given as the residues of a linear model with body mass (in grams), both log_{10} transformed, so the effects of body mass on longevity are factored out. Different bird orders are indicated by symbols, as specified on the legends at the right side of the graphs. **Figure 2.A** uses the values and estimates from the glucose model without life history traits (n=389 individuals from 75 species), while **Figure 2.B** uses only the data points employed on the complete model (n=326 individuals of 58 species).

Bird albumin glycation is related to glycemia, diet and clutch mass

After controlling for intraspecific variability, we found that diet was affecting variation in
albumin glycation rates, with carnivorous species having higher glycation levels than omnivorous
species (**Table 2**; **Figure 4.B**). There were trends for species with greater investment into
reproduction (i.e. clutch mass) to have higher glycation levels (**Table 2**), and glycation levels to
increase with glucose levels (**Table 2**). The relation between glycation and glucose levels was
significant in the model that did not include life history traits (β=0.127, CI_{95}[0.0217, 0.232],
P_{MCMC}=0.0188; n=389 individuals from 75 species; **Figure 3**). Given that both plasma glucose and
albumin glycation rates are logarithm transformed and the estimated slope of their relationship
is lower than one, this implies that birds with higher glucose levels have relatively lower albumin
glycation rates. Models that did not consider intraspecific variability show no effects of the diet,
body mass, life history traits or glucose levels on albumin glycation rates (ESM1).

Additional analyses looking at the number of exposed lysines in the albumin sequence of a
species show no effect of this variable on albumin glycation rates (β=0.243, CI_{95}[-0.106, 0.578],
P_{MCMC}=0.154).

Table 2. Final glycation model with diet, body mass, glucose and life history traits as explanatory variables.
Posterior means, CI_{95} and pMCMC from a phylogenetic MCMC GLMM model including n=326 individuals
of 58 species. Both glucose and body mass are log transformed and life history traits are residuals from a
linear model of log body mass and log of the trait in question. Body mass was also centred to better explain
the intercept. The intercept corresponds to the Omnivore diet, being used as the reference as it is
considered the most diverse and "neutral" group for this purpose. Significant predictors are indicated in
bold.

	Estimates	Lower 95% CI	Upper 95% CI	Sampling effort	pMCMC
Intercept	1.027	0.712	1.334	121,142	< 0.001
Diet: Carnivore terrestrial	0.101	0.014	0.187	119,800	0.023
Diet: Aquatic predator	0.036	-0.063	0.134	119,800	0.470
Diet: Herbivore	-0.009	-0.167	0.154	118,625	0.906
Diet: Frugivore/granivore	0.101	-0.103	0.302	119,800	0.317
Centered log(Body mass)	0.005	-0.045	0.057	119,800	0.833
log(Glucose)	0.101	-0.014	0.217	119,800	0.088
Maximum lifespan	0.084	-0.058	0.227	120,728	0.239
Clutch mass	0.173	-0.008	0.358	119,800	0.060
Developmental time	0.069	-0.149	0.282	118,454	0.519

Figure 3. Albumin glycation rates (as a percentage of total albumin) variation as a function of individual

plasma glucose values (mg/dl). Different bird orders are indicated by symbols, as specified on the legends

at the right side of the graphs. The values and estimates used are from the glycation model without life

history traits (n=389 individuals from 75 species).

Regarding diet, the only significant effect observed was a higher level of albumin glycation in

terrestrial carnivores compared with omnivores (see **Table 2 and Figure 4.B**).

Figure 4. Effects of diet on (**A**) plasma glucose levels and (**B**) albumin glycation in birds. Glucose levels are

given in mg/dl, while glycation levels are a percentage of total plasma albumin, which is found to be

glycated. Terrestrial carnivores showed significantly higher glycation levels than omnivores

(Estimate=0.101, p_{MCMC}=0.0231). Raw data shown.

We observed no significant effect of age relative to maximum lifespan nor sex on either
glycaemia or glycation (see **Figure ESM 2.1**).

DISCUSSION

Plasma glucose level and albumin glycation rate (co)variation suggests resistance mechanisms

Our findings show that glucose levels vary widely among different bird groups. Interspecific
differences are partly explained by the allometric relationship of glycaemia with body mass
(**Figure 2.A**), which has already been reported in previous studies [59][31][12][13]. Indeed,
Passeriformes, Apodiformes and some Columbiformes (e.g. *Nesoenas mayeri* holds the highest
value in our dataset) are found at the higher end of the glucose level continuum, in accordance
with their relatively small body mass and powered flight, while groups of larger birds such as
Phoenicopteriformes, Anseriformes, Rheiformes and Suliformes tend to show low glycaemia
levels. This pattern is similar for glycation, with some groups of large birds (such as
Phoenicopteriformes, Anseriformes and Suliformes) showing the lowest levels of glycated
albumin, while small birds (such as Apodiformes) had the highest values. Nevertheless, glycation
remains high in some birds in relation to their glucose levels, as in Rheiformes, or low as in
Psittaciformes or Passeriformes. The case of Procellariiformes, which are typically long-lived
birds, is particularly striking, with some species exhibiting some of the highest glycation levels
(*Calonectris diomedea* and *Macronectes giganteus*) and others some of the lowest (*Procellaria*
aequinoctialis). This suggests that, if birds are protected against the deleterious effects of high
glycaemia, certain species may have evolved mechanisms to efficiently prevent proteins to be
glycated at high rates while others may have evolved mechanisms to resist the consequences of
protein glycation. We should also bear in mind that some taxonomic groups may be
underrepresented in our study, or biases in species selection due to availability contingencies
(e.g. species common in zoos or present in European countries) may exist, so further studies
should target this underrepresented groups in order to confirm our predictions.
As expected for a non-enzymatic reaction, just by the law of mass action, bird albumin glycation
rates increase with plasma glucose concentration. However, the logarithmic nature of the
relationship, together with a scaling factor lower than 1, suggests that species with higher

plasma glucose levels exhibit relatively greater resistance to glycation. This finding aligns with
previous research indicating that *in vitro* glycation levels of chicken albumin increase at a slower
rate than those of human albumin when glucose concentration is elevated [71]. Moreover, these
levels are consistently lower than those of bovine serum albumin regardless of glucose
concentration and exposition time [34]. As discussed in previous studies comparing chicken to
bovine serum albumin [34], or zebra finch to human haemoglobin [50], the lower glycation rates
observed in bird proteins may result from a lower number of glycatable amino acids (e.g. lysines)
in their sequence and/or their lesser exposure at the protein surface.

Our analyses indicate that average glycation levels increase with the number of glycatable
lysines in the albumin sequence, but not significantly. This may be attributed to the limited
number of species employed or the weak variation in the number of glycatable lysine residues,
which are mostly ranging from 33 to 39. An interesting exception are the 18 glycatable lysine
residues of flamingos (*Phoenicopterus ruber*), which also shows very low glycation levels
(mean=10.1%). However, the exceptions of 44 in zebra finches and 20 in godwits (*Limosa
lapponica*, used in place of *Limosa limosa*) are nevertheless associated with very similar average
glycation levels of 20.2% and 20.7% respectively.

Plasma glucose relates with longevity and may be influenced by reproductive
strategies

Our results were only partially in agreement with predictions from POLS theory. The only
previous comparative study of glycaemia and life history traits to our knowledge [13] shows no
relationship with mass-adjusted maximum lifespan in passerines. However, our study over 87
bird species on 22 orders revealed an increase in glucose with mass-adjusted longevity up to a
plateau (see **Figure 2.B**). Thus, the relationship between glucose and maximum lifespan may
depend on differences between bird orders or be tied to specific species particularities not
explored in our study. Such species particularities might involve additional undetermined
ecological factors that modify the relationship of glycaemia with longevity. Further exploration
of glucose metabolism in relation with lifestyle will bring further light on species-specific life-
history adaptations concerning glucose.

Regarding reproductive investment (i.e. clutch mass), our results show an apparent positive
linear relationship with glycation (see **Table 2**), while previous studies reported positive
relationships with glycaemia in passerines [13][72]. Interestingly, most of the species with high
clutch mass included in our study belong to the Anseriformes (**Figure ESM 2.2**). While these

species exhibit very low glycaemia and albumin glycation rates, they are also characterized by a particular reproductive strategy compared to passerines, for whom clutch mass does not imply the same in terms of parental investment. For instance, Anseriformes, unlike Passeriformes, are precocial and their offspring are not very dependent on parental care. Furthermore, they are capital breeders, accumulating the energetic resources required for reproduction in advance rather than collecting them during reproduction. These species typically have large clutch sizes and incubation is usually carried out by females, who store internal fat resources to support this main part of the parental investment. In addition, ducklings are born highly developed, reducing the amount of care required post-hatching (see e.g. [73][74]). Consequently, their dependence on glucose as a rapid energy source for reproduction may be lower, with investment in this activity likely more closely linked to lipid accumulation. This could explain why we did not detect previously reported effects of clutch mass on glucose levels, even if they may be present for glycation.

Terrestrial carnivores show a paradoxically increased albumin glycation rate without increased plasma glucose levels

Contrary to our expectation of finding differences across dietary groups, plasma glucose did not significantly vary with species diet. Nevertheless, this aligns with results previously reported by [13] for Passeriformes or by [52] for 160 species of vertebrates among which 48 were birds. Additionally, intraspecific data indicates that changes in diet composition hardly affects glycaemia in birds [75]. Together, these studies suggest that glycaemia is tightly regulated independently of dietary composition.

Although terrestrial carnivore species did not have significantly higher glycaemia levels, they nonetheless demonstrated significantly higher albumin glycation rates, suggesting a potential susceptibility to protein glycation. Besides unique structural features, this phenomenon could be due to lower albumin turnover rates in terrestrial carnivores. Studies in humans have linked higher oxidative damage to albumin with lower turnover rates (reviewed in [76]), which may extend to other species and post-translational modifications of albumin, such as bird's albumin glycation. However, the contrary outcome would be anticipated if protein intake were high as seen in carnivorous species [77][76].

Interestingly, the contrast between terrestrial and aquatic predators, which did not show such high glycation rates, suggests the beneficial influence of a substantial difference in food composition or the pre-eminence of another ecological factor, yet to be determined, which may

explain lower glycation levels in aquatic predators. In terms of food composition, the proportion
of essential n-3 PUFA (polyunsaturated fatty acids), particularly long-chain ones such as DHA and
EPA compared with n-6 PUFA and short-chain n-3 PUFA, is different in aquatic and terrestrial
environments [78] and therefore in the diet of predators in each environment (e.g. [79]). For
instance, low n-6 PUFA/n3- PUFA ratio have been shown to decrease insulin levels and improve
insulin resistance in humans [80]. On the other hand, while some studies report that an increase
in dietary n3-PUFA reduces the levels of glycated haemoglobin (HbA1c) in humans, this effect
was not detected in many other studies and therefore remains inconclusive (reviewed in [81]).
The detailed study of the fat composition of the diets of terrestrial and aquatic predators, in
particular the ratio between different types of PUFA, merits more attention. In addition,
micronutrient content, such as circulating antioxidant defences, should also be taken into
consideration. Indeed, a positive relationship between oxidative stress and glycated levels of
bovine serum albumin has been reported [82]. One hypothesis is that terrestrial predators have
higher systemic oxidative stress levels compared to other species, which may be explained by
defects in their antioxidant defences. Uric acid is one of the main non-enzymatic antioxidants in
birds [83][84][85], and uric acid levels are especially high in carnivore species that have a rich
protein diet [86][84][87][88][89]. We should therefore tale into account other important
antioxidants such as vitamin E, vitamin A and carotenoids, which may be less abundant in the
diets of terrestrial carnivores (e.g. [90], but see [49]). The question of whether the diet of aquatic
carnivores provides a better intake of antioxidants would therefore requires a more detailed
description of dietary habits. The herbivorous diet, meanwhile, despite the expected possibility
contributing to lower glycaemia and glycation levels due to higher levels of PUFA compared to
SFA (saturated fatty acids) (effects reviewed in [81] for humans) and higher fibre content (see
[91]), did not lead to significantly lower levels of either of these two parameters. However, the
relatively low sample size in that group or the presence of outliers as *Nesoenas mayeri*, with the
highest glucose levels of this dataset, makes the interpretation of the obtained results somehow
limited. Therefore, further research should be carried out on these species to determine if the
expected pattern would emerge with a better sampling.
Finally, differences between captive and wild populations could be considered as a source of
variation in glucose or glycation levels, due to a more sedentary lifestyle in captivity, with lower
activity levels and a higher food intake likely to lead to increased circulating glucose levels in
captive individuals. Additionally, differences in nutrient intake, such as antioxidants, specific
fatty acids or amino acids, between captive and wild populations could contribute to variations

in glycation levels, as we saw above. Nevertheless, we believe that this factor may not

significantly affect our findings, as our study encompasses species from both captive and wild

populations across various diet groups, particularly those exhibiting significant differences (e.g.,

omnivores versus terrestrial carnivores).

CONCLUSION AND PERSPECTIVES

In conclusion, the avian plasma glucose levels measured here are generally higher or in the high

range of values recorded for mammalian species [25]. Our study also concludes that there is

considerable variation in plasma glucose levels and albumin glycation rates between bird

species, with those with the highest glucose levels showing greater resistance to glycation. The

correlation between plasma glucose and life history traits proves complex, primarily influenced

by its inverse association with body mass, along with non-linear, mass-independent effects of

longevity and potentially clutch mass. Finally, although diet does not explain plasma glucose

levels, terrestrial carnivores have higher albumin glycation rates. Whether these intriguing

results are explained by specific effects of certain dietary components, such as PUFAs and/or

antioxidants, remains to be determined. Differences in plasma glucose levels and albumin

glycation rates across bird species are only partially consistent with predictions according to the

POL hypothesis. Further investigation is needed to elucidate the correlation between these traits

and specific life conditions, such as reproductive strategy, migration patterns, flight mode or

detailed diet composition. In addition, more in-depth exploration of high glycaemic groups such

as the Passeriformes and other small birds, which make up a significant proportion of avian

species, could provide valuable new insights. Similarly, investigating groups such as the

Phoenicopteriformes or Anseriformes, which are at the other end of the glycaemic-glycation

spectrum, could shed light on the origin of differences between avian orders. Furthermore,

notable variations were observed between species within orders, as in the Procellariiformes,

which given their high mass-adjusted longevity, seems to suggest that some species may not be

intrinsically resistant to glycation, but rather to its adverse consequences for health. Future

studies should focus on specific species to unravel the physiological mechanisms mediating the

effects of blood glucose and protein glycation on life-history trade-offs, in particular mechanisms

that may vary between taxa and contribute to characteristic adaptations in birds to mitigate the

potentially negative effects of relative hyperglycaemia.

CONFLICTS OF INTEREST

None declared.

DATA ACCESSIBILITY

All data used, including references for the values taken from bibliography or databases, will be available as Electronic Supplementary Material (ESM5).

ETHICS STATEMENT

This study followed all the legal considerations, with the ethic authorisations from the French Ministry of Secondary Education and Research, n°32475 for the zebra finches sampling, the Swiss Veterinary Office (FSVO) nº 34497 for the Alpine swifts, the Ethics Committee of the University of Extremadura (licenses112//2020 and 202//2020) and the Government of Extremadura (licenses CN0012/22/ACA and CN0063/21/ACA) for the godwits and terns from Spain, and Sampling in Terres Australes et Antarctic Françaises was approved by a Regional Animal Experimentation Ethical Committee (French Ministry of Secondary Education and Research permit APAFIS #31773-2019112519421390 v4 and APAFIS#16465–2018080111195526 v4) and by the Comité de l'Environnement Polaire and CNPN (A-2021-55). The samples from the Mulhouse zoo were taken by its licensed veterinary with capacity number: 2020-247-SPAE-162.

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AUTHORS' CONTRIBUTIONS

F. Criscuolo and F. Bertile conceived the idea, directed most of the sample collection and logistics and contributed significantly to the writing. A. Moreno-Borrallo contributed to the development of the questions, gathered the diet and life-history data, performed the statistical analyses, and led the writing. S. Jaramillo Ortiz and C. Schaeffer performed the mass spectrometry analyses for protein glycation measurements, and S. Jaramillo Ortiz contributed to the glucose measures and commented on the manuscript. T. Boulinier. and V. A. Viblanc coordinated the ECONERGY and ECOPATH polar programs, organised the collection of samples on subantarctic seabirds, and commented on the manuscript. B. Rey contributed with the collection of samples from Parc des Oiseaux (Villars-les-Dombes, France) and commented on the manuscript. P. Bize leads the monitoring of Alpine swifts' populations from which the samples were obtained, he helped collecting them and commented on the manuscript. J. S Gutiérrez and J. A. Masero contributed with samples from Spain and part of the statistic scripts and commented on the manuscript. B. Quintard leads the health monitoring of Mulhouse zoo bird collection, organized and realized

most of Mulhouse zoo samplings and commented on the manuscript. All authors gave their
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