Diving efficiency at depth and pre-breeding foraging effort increase

with haemoglobin levels in gentoo penguins

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Abstract

Individual differences in oxygen storage and carrying capacity have been associated with

fitness-related traits and, for air-breathing aquatic animals, to diving ability and foraging success.

In winter, many seabirds must replenish the energy reserves they have depleted during the

breeding period. Thus, winter foraging efficiency can influence their upcoming breeding

behaviour. Using gentoo penguins (*Pygoscelis papua*), we investigate (1) if inter-individual

variation in diving efficiency (proportion of time spent at the bottom) is associated with indices

of oxygen storage and carrying capacity (haemoglobin, haematocrit, body mass), and (2) if

measures of pre-breeding foraging effort (mean trip duration, total time at sea, and vertical

distance travelled) are associated with these oxygen indices and breeding status. Haemoglobin

was positively correlated with diving efficiency, particularly for deeper dives, and only penguins

with high haemoglobin levels frequently dove at depth ≥ 140 m. Such differences could affect

resource access. However, potentially because reaching deep offshore waters requires travelling

more than foraging nearshore, vertical distance travelled pre-breeding increased with

haemoglobin levels. The relationship with haematocrit was non-linear, suggesting that

commonly-used analyses may be inappropriate for this index. We found that early-laying

penguins spent less time at sea prior to nesting than non-breeding penguins, suggesting that more

efficient foragers lay earlier. Given that diving efficiency at depth is linked to aerobic capacity,

anthropogenic activities taking place in either nearshore or offshore waters (e.g., shallow water

fisheries, offshore oil rigs) may have differing impacts on individuals. Further understanding

these links could help the conservation of diving species.

Running page head: Diving efficiency and effort in penguins

Keywords: Diving efficiency, oxygen storage and carrying capacity, breeding participation,

gentoo penguin, haemoglobin, haematocrit, body mass, time-depth recorder

1. Introduction

Intra-specific variation in foraging and movement behaviour is ubiquitous, and can have

important ecological consequences at both the individual and population levels (Bolnick et al.

2003, Ceia & Ramos 2015, Phillips et al. 2017, Shaw 2020). Such variation can be caused by

complex relationships with the environment, intra- and inter-specific interactions (e.g.,

competition), and physiological differences (Shaw 2020). Differences in oxygen storage and

carrying capacity can be particularly important, as they have been associated with variation in

movement behaviour, habitat selection, and fitness-related traits such as annual breeding success

and arrival at breeding grounds (Crossin et al. 2013, 2015, Minias 2015). For diving animals,

oxygen storage and carrying capacity can limit diving ability, and thus access to resources may

be physiologically restricted (Roncon et al. 2018). For such animals, the time spent diving and

the rate of mass gain have been linked to indices of aerobic capacity (Crossin et al. 2015).

Diving animals have a range of adaptations to maximize their aerobic capacity and

extend their breath-holding capability (Roncon et al. 2018). Haemoglobin is an oxygen-carrying

protein present in red blood cells, and thus its level is a crucial determinant of the rate of oxygen

delivered to tissues (Minias 2015). Haemoglobin levels are often used as a proxy of aerobic

capacity and physiological conditions in birds (Minias 2015), and is elevated in bird species that

dive (Minias 2020). Prolonged breath-holding capacity is often linked to oxygen storage. Oxygen

stores typically increase with haemoglobin concentration, blood volume, myoglobin

concentration in skeletal muscles, muscle mass, and lung volume (Mirceta et al. 2013, Roncon et

al. 2018). As larger individuals possess more blood and muscle, body mass is often positively

associated with oxygen storage capacity (Ponganis & Kooyman 2000, Cook et al. 2013, Mirceta

et al. 2013, Polito et al. 2015, Camprasse et al. 2017). In addition, consistent with generally

observed allometric patterns of metabolism, larger body mass is often associated with lower

oxygen usage per unit mass (Glazier 2005, Hudson et al. 2013). As such, total blood

haemoglobin concentration (referred as Hb), haematocrit (percentage of red blood cells in blood,

referred as Hct), and body mass can be important determinants of a breath-holding diver's

physiological performance and can limit an animal's diving ability (Crossin et al. 2015,

Chimienti et al. 2017).

Seabirds display high levels of individual variation in foraging and movement (Ceia &

Ramos 2015, Phillips et al. 2017), and diving seabirds provide a unique opportunity to link

differences in oxygen storage and carrying capacity to variation in diving and movement

behaviour, and ultimately to differential breeding participation. Many seabirds severely deplete

their energy reserves during breeding and moulting periods and must replenish these reserves by

foraging intensively in winter (Sorensen et al. 2009, Crossin & Williams 2016). Consequently,

how efficient an individual is in acquiring food in winter can influence multiple aspects of its

breeding capacity and behaviour (Daunt et al. 2006, Shoji et al. 2015). For example, individuals

that spend less time foraging in winter, and thus are assumed to be more efficient foragers, breed

earlier (Daunt et al. 2006). Understanding how individuals vary in their foraging efficiency and

behaviour as they prepare for breeding can help assess the possible impacts natural and

anthropogenic ecosystem changes can have on populations (Durell 2000, Bearhop et al. 2006,

Cury et al. 2011, Phillips et al. 2017).

Here, using a population of gentoo penguins (*Pygoscelis papua*), we explore how

individual variation in winter diving behaviour relates to indices of oxygen storage and carrying

capacity, and how these relate to their breeding status. We use gentoo penguins because they

show considerable individual variation in winter movement behaviour (Baylis et al. 2021).

Gentoo penguins consume a wide variety of locally available prey found at a range of depths

down to approximately 200 m (Tanton et al. 2004, Thiebot et al. 2011, Camprasse et al. 2017).

While they are considered a generalist species, many individuals have a specialized diet and

foraging behaviour (Polito et al. 2015, Camprasse et al. 2017, Handley et al. 2017, Herman et al.

2017). There is a high likelihood of intraspecific competition for prey around colonies, as gentoo

penguins are not migratory and thus forage year-round relatively close to their breeding colonies

(Clausen et al. 2005, Bearhop et al. 2006, Kokubun et al. 2010). In addition, given their

magnitude, certain commercial fisheries in the Falkland Islands could reduce the availability of

important prey species in winter (Clausen & Pütz 2003). The number of breeding pairs in the

Falkland Islands fluctuates drastically, with changes of 50 % occurring in a single year (Pistorius

et al. 2010, Stanworth & Crofts 2019). Changes in population breeding success are thought to be

linked to prey availability (Pistorius et al. 2010) and broad-scale climatic variation (Baylis et al.

2012).

To understand the link between oxygen storage and carrying capacity, pre-breeding

diving and foraging behaviour, and breeding activity, we investigate (1) if diving efficiency

(proportion of the time within a dive spent at the bottom) is associated with indices of oxygen

storage and carrying capacity (Hb, Hct, body mass), and (2) if measures of pre-breeding foraging

effort (mean trip duration, total time at sea, and vertical distance travelled) differ between

individuals based on their breeding status and indices of oxygen storage and carrying capacity.

We use time-depth recorders (TDRs) to quantify diving efficiency and foraging effort. As

attaching such devices to animals can add drag and affect their performance, behaviour, and

reproductive success (Taylor et al. 2001, Beaulieu et al. 2010, van der Hoop et al. 2014, Wilson

et al. 2015), we also evaluate whether tagged individuals differ from control birds in terms of

breeding activity, Hb, Hct, and mass.

2.   Methods

2.1 Data collection

We captured 66 adult gentoo penguins post-moult (April 4 – 10, 2018) from two colonies

in the Falkland Islands: Rookery Sands, Race Point (51.4345°S, 59.1081°W, n=33) and Tamar

Point, Pebble Island (51.3241°S, 59.4523°W, n=33). We weighed each captured penguin using a

10 kg Pesola scale, took a blood sample (< 1.5 ml) from the left brachial vein using heparinized

syringes fitted with 25-gauge needles, and marked them with black hair dye (L'Oreal Preference)

to facilitate re-identification. We equipped each penguin with a Sirtrack K2G 173A SWS KIWISAT

202B Argos tag (34g, 55 x 27 x17 mm, Sirtrack, Havelock North, New Zealand) and a Lotek

LAT1800 TDR tag (13.6 g, 62 x 13 x 13 mm, Lotek Wireless Inc, St. John's, Canada). As in

Handley et al. (2018) and Baylis et al. (2021), we attached the two devices to midline back

feathers using overlapping layers of TESA$^{®}$ tape (Beiersdorf AG, GmbH, Hamburg, Germany)

and sealed the tape seams with cyanoacrylate glue (Supplement 1).

During the breeding season (September 27 - October 21, 2018), we recaptured 35 of the

previously tagged penguins and 35 unmarked penguins, the latter serving as controls in our

analysis of the potential effects of tagging (Table 1). As described above, we weighed each

penguin, took a blood sample, and marked the control bird with hair dye. We retrieved any

devices still attached to the penguins. Twenty-one penguins had TDR devices still attached (7

Race Point, 14 Pebble Island, Table 1), one of which still had an Argos tag attached (from Race

Point). We note that we were unable to collect sufficient blood samples from two individuals

with a TDR tag, resulting in only 19 of the individuals with TDR data also having blood data

from the breeding period (5 Race Point, 14 Pebble Island, Table 1).

The research was conducted under Falkland Islands Scientific Research Licence

(R12/2017) and conformed to guidelines from the Canadian Committee on Animal Care

(University of British Columbia Animal Care Permit A17-0243 and Dalhousie University

Animal Care Permit 17-100).

2.2 Indices of oxygen storage and carrying capacity

Using the blood samples from the breeding period, we used haematocrit (Hct) and

haemoglobin (Hb) as physiological indicators of blood oxygen storage and carrying capacity. As

Hct is based on blood cytology and Hb is based on blood biochemistry, each measure provides

different insights into the aerobic capacity of individuals (Fair et al. 2007, Kaliński et al. 2011,

Minias 2015). Hct (percent packed red blood cell volume) was determined from fresh whole

blood in heparinized capillary tubes centrifuged for 5 min (Centrifuge; Brinkmann Instruments,

Ontario, Canada) at 10000 g and measured using digital callipers (± 0.01 mm). Hb (total blood

haemoglobin concentration in g/dL) was determined using the cyanmethemoglobin method

(Drabkin & Austin 1932), where 5 μl of fresh whole blood was pipetted (Gilson Pipetman P2;

Gilson Medical Electronics, France) into 1.25 ml Drabkin's reagent (D541-6V, Brij 35 solution

B4184-100ml; Sigma-Aldrich, Missouri, US). We ran triplicate samples and read absorbance at

540 nm with a microplate spectrophotometer (BioTek PowerWave 340, Vermont, USA). The

inter-assay coefficient of variation was 1.70 % between plates.

We used the body mass measured during the breeding period as a proxy of oxygen

storage capacity. We did not scale body mass to structural body size to generate a morphometric

index because (1) oxygen stores (e.g., blood volume) and dive duration have been shown to

increase with body mass itself (e.g., Hassrick et al. 2010), and (2) mass alone has been shown to

be a more reliable measure of body condition in birds (Schamber et al. 2009, Labocha & Hayes

2012). We identified the sex of the individuals by sending blood samples to Viaguard[†], which

performs DNA testing. We note that sex results were unavailable for three of the 19 penguins for

which we had TDR and blood data (Table 1).

2.3 Dive data processing

The TDR tags recorded depth (0.25 m resolution), temperature (0.01°C resolution), and

wet/dry state at 1 s intervals. We wanted to focus on the time period that could be linked to the

blood samples. Since the maximum life span of red blood cells in birds is estimated to be 35 – 45

days (Rodnan et al. 1957), we limited the analyses to dives made within 40 days prior to

recapture. Given that the last Argos transmission was August 16, 2018, well before this

threshold, and very few tags transmitted locations past May, we do not consider the Argos data

further.

To characterise the diving behaviour of foraging individuals, we extracted the dive

metrics from the TDR series using the software Divebomb (1.0.7, Nunes 2019) in Python

(3.4.10, Python Software Foundation 2019). We first corrected the dive data for drift using a zero

offset (Luque & Fried 2011), and we identified the following dive metrics: start of the dive, dive

duration, maximum depth, bottom time, and post-dive surface interval. To remove the shallow

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[†] https://www.accu-metrics.com/animal-services/p/avian-dna-bird-sexing-test

dives most likely associated with travelling, we discarded all dives shallower than 5 m (Kokubun

et al. 2010, Lee et al. 2015, Carpenter-Kling et al. 2017). Following Kokubun et al. (2010) and

Lee et al. (2015), we eliminated the inclusion of long surface periods unlikely to be associated

with active foraging by only considering dives with post-dive surface interval < 200 s. Through a

visual assessment of 100 random dives across all individuals, we quantified that Divebomb gave

precise measurements of decent, bottom, accent, and surface phases of dives 94 % of the time.

Divebomb produced minor errors distinguishing the bottom phase in cases where undulations in

depth occurred over the transition from decent to accent phases.

2.4 Behavioural observations

To assess the reproductive status of penguins, teams of two observers monitored the

breeding behaviour of marked birds. Through daily colony scans and the continuous monitoring

of arriving individuals, we identify which individuals were present at Race Point from September

27, 2018, to October 24, 2018, and Pebble Island from September 29, 2018, to October 24, 2018.

Between 13h00 – 19h00 daily, we conducted focal follows on each individual observed in the

colony. The focal follows were at least 5 min long and lasted until the presence or absence of

eggs in an occupied nest could be confirmed. The pair bonding behaviours we monitored

included nest building, copulation, pair calling, and pair bowing (Williams 1995). Gentoo

penguins in the Falkland Islands lay their first egg from mid-October to early November (Otley

et al. 2004, Black et al. 2018). As we left the field site before we could acquire the lay date of

many penguins, we classified penguins into three categories of reproductive status: non-

breeding, breeding, and early-laying individuals. Penguins classified as non-breeders were found

most often at different locations than nesting penguins, typically towards the edge of colonies,

and rarely displayed pair-bonding behaviours. Penguins categorized as breeders were observed

conducting pair-bonding behaviours at stable nest site locations. Early-laying penguins were

observed to be incubating at least one egg prior to October 24, 2018. We note that we had

individuals of both sexes in each reproductive category and that none of the penguins had laid an

egg at Race point at the time of departure of field observers.

2.5 Linking diving efficiency to oxygen storage and carrying capacity indices

We used a generalized additive mixed model (GAMM) to determine the relationship

between diving efficiency at different maximum depths and our three previously described

indices of oxygen storage and carrying capacity (Hb, Hct and body mass). We defined diving

efficiency as $\frac{bottom\ time}{dive\ duration\ +\ post-dive\ surface\ interval}$ (Simeone & Wilson 2003, Lescroël & Bost

2005, Lee et al. 2015). Using the R package mgcv (Wood 2011), we used a thin-plate regression

spline to model the non-linear relationship between diving efficiency and maximum dive depth.

To investigate the potential interactions between maximum depth and the indices of oxygen

storage and carrying capacity (Hb, Hct, and mass), we included a tensor product term (with cubic

regression spline basis) between maximum depth and each index. To be able to decompose the

interactions with maximum depth from the potential main effects of each index, we included a

separate tensor product and a thin-plate regression spline for the relationship with diving

efficiency and each index (i.e., used s() + ti() in mgcv). To account for potential biological or

behavioural differences in sex and colony, we first included both terms as fixed effects in our

GAMM and assessed their importance. As sex was not deemed an important covariate for this

analysis (see Supplement 2) and three individuals were missing sex information, our final

analysis did not include sex and included these three individuals. Thus, our final GAMM was

applied to the data of 19 individuals. We also included an individual ID random effect on the

intercept to account for the fact that we had multiple dives per penguin.

The diving data were highly temporally autocorrelated, and fitting GAMMs with

complex autoregressive-moving-average (ARMA) functions are computationally demanding. For

example, running our full GAMM with an ARMA function took multiple days to run on an iMac

3.6 GHz 10-Core Intel Core i9, and we ran into multiple convergence problems. Thus, we

assessed how much sub-sampling was required to remove the autocorrelation in the residuals of

the full model and found that we had to keep only one dive every 70 dives to completely remove

the autocorrelation in the residuals. We used this 70-dive sub-sampling scheme, which resulted

in having on average 85 dives per individual. To identify which of the covariates of the GAMM

should be selected, we used the shrinkage method with a double penalty (i.e., select = "TRUE" in

mgcv; Marra & Wood 2011), which penalizes both unnecessary wiggliness and linear effect of a

covariate. This approach has many advantages over stepwise variable selection and model subset

selection, such as not limiting the combinations of covariates explored, and has been previously

used to select covariates in diving models (Marra & Wood 2011, Photopoulou et al. 2014).

There was some correlation and concurvity between the covariates. As such, we explored

whether, and how, the results changed when creating GAMMs with only one of the important

oxygen indices at a time. See Supplement 3 for details. In addition, the residuals of the final

GAMM displayed evidence of heteroskedasticity. This variation in variance can be attributed to

the fact that our final GAMM modelled proportion data (as defined, diving efficiency ranges

between 0 and 1) with a normal error distribution. As such, we also explored whether, and how,

the results changed if we used a GAMM with a beta regression. Supplement 4 details how the

results remained largely unchanged.

As post-hoc analyses, we further explored how the deep diving behaviour of individuals

changed in relation to the oxygen storage and carrying capacity indices deemed important in the

GAMM above. As some individuals with low oxygen storage and carrying capacity may reduce,

or altogether forgo, doing deep dives (here defined as ≥ 140 m), we used simple generalized

additive models (GAMs) to assess whether the proportion of deep dives each individual

performed changed in relation to each of the selected oxygen indices. As high diving efficiency

can be achieved by either extending the bottom time or reducing surface recovery between dives,

we used new GAMMs to explore how bottom time and post-dive surface intervals for deep dives

changed in relation to each of the selected oxygen indices. For these new GAMMS, we looked at

the oxygen indices separately, as the smaller sample size did not allow to have both in the

models at once. We included an individual ID as a random effect to account for the fact that we

have more than one dive per penguin.

2.6 Associating measures of foraging effort with breeding status and oxygen storage and

carrying capacity indices

To determine if pre-breeding foraging behaviour changes between individuals based on

their breeding status and oxygen storage and carrying capacity, we investigated three measures

of foraging effort: mean trip duration, time at sea, and vertical distance travelled. To estimate

mean trip duration and time at sea, we first identified trips at sea using the binary wet-dry data

collected with the saltwater sensor of the TDR tags. To account for occasional jumps at sea,

which results in brief recordings of dry conditions, we averaged the wet-dry data using a 10-

minute rolling window. We classified any window value with an average > 0 as wet. We

terminated a trip just before a dry period lasting at least one hour, and the subsequent trip began

once a new wet value was recorded. The mean trip duration was the average duration of trips for

the 40 days prior to recapture, while the time spent at sea was the sum of all trip durations within

that period. Our last measure of foraging effort was the total amount of vertical distance travelled

for the time spent at sea ($\sum maximum\ depth\ (km) \times 2$; similar to Lescroël & Bost 2005,

Booth et al. 2018).

Using multiple linear regressions, we investigated whether each of these measures of

foraging effort (mean trip duration, time at sea, and vertical distance travelled) was associated

with breeding status, sex, and the three oxygen storage and carrying capacity indices (Hb, Hct,

mass). We only included individuals from Pebble Island since none of the Race Point individuals

laid eggs before the end of the field season. To identify the covariates that best explained the

data, we fitted models with all possible combinations of three of these covariates. Given the

small sample size (n=13), we compared the models using the Akaike Information Criterion with

correction for small sample size (AICc). We considered models that outperformed the null model

(model with no covariates) and had ΔAICc < 4 from the best model (Burnham and Anderson

2002). Next, we used Wilcoxon rank sum tests to assess whether there were differences in

oxygen indices across sex and breeding status. We used a Bonferroni correction to account for

the multiple comparisons.

2.7 Assessing the potential effects of tagging

To limit the potential influence of our tags on the behaviour, energy expenditure, or

condition of the animals (e.g., Wilson et al. 1986, Vandenabeele et al. 2015), we minimized drag

by using tags with a streamlined design and attaching them in an appropriate caudal position

(Bannasch et al. 1994). Furthermore, the overall weight of devices was < 1 % of the average

body mass recorded, and similar devices have not affected the diving behaviour of gentoo

penguins (Kokubun et al. 2010). However, especially because we deployed tags for an extended

period, we wanted to assess whether the data for the tagged penguin were representative of the

untagged animals and look for signs of tagging effects. To do so, we compared the physiology at

breeding and breeding status of tagged to untagged penguins. Specifically, we used Welch two-

sample t-tests to compare Hb, Hct, and mass at breeding. For the reproductive status, we used χ^2

goodness-of-fit tests, and because of the small sample size (37 individuals for 6 categories), we

estimated the p-value using a Monte Carlo method (Hope 1968). We note that we were unable to

sample sufficient blood for three of the recaptured penguins, as well as unable to measure Hct for

one untagged penguin and Hb for another untagged individual. Thus, while the mass comparison

included 35 tagged, and 35 untagged individuals, the comparisons for Hct and Hb included only

32 tagged and 34 untagged individuals. In addition, for the comparison of reproductive status, we

focussed only on penguins from Pebble Island since none of the penguins at Race Point laid an

egg before the end of our field season. Finally, we also quantify the proportion of tagged

penguins that increased in mass between capture and recapture and the average weight gain

during that period.

All data analyses were performed using R (4.2.1, R Core Team 2022). Values presented

in the results are mean ± SD unless otherwise stated, and statistical tests were considered

significant at a 0.05 level. Supplement 8 provides the code and data needed to reproduce the

analyses.

3. Results

3.1 Relationships between diving efficiency and indices of oxygen storage and carrying

capacity

Our GAMM indicated that diving efficiency had statistically significant relationships

with colony, as well as with the maximum depth of a dive, and its interaction with Hb and Hct

(Table 2, Fig. 1). Diving efficiency was highest at intermediate depths followed by a decline at

deeper depth (Fig. 1A). The interaction between maximum depth and Hb showed that while

penguins with higher Hb did not appear to have higher diving efficiency for shallow dives,

diving efficiency increased with Hb for deeper dives (Fig. 1B). In the post-hoc analyses, the

relationship between bottom time for deep dives and Hb was positive and statistically significant,

while the relationship between Hb and post-dive surface interval of deep dives was not

statistically significant (Table 3). The results of these post-hoc analyses indicate that this increase

in diving efficiency with Hb at deeper depth is primarily explained by an increase in bottom time

rather than a decrease in the post-dive surface interval (Fig. 2B-C). In addition, we found that the

proportion of deep dives (≥ 140 m) made by an individual increased with Hb (Fig. 2A), a

relationship that was statistically significant (Table 3).

The non-linear interaction between maximum depth and Hct was more complex, with

multiple and less easily interpretable peaks in diving efficiency across Hct values (Fig. 1C). The

post-hoc analyses revealed that neither bottom time, post-dive surface interval, nor the

proportion of deep dives showed a statistically significant relationship with Hct (Table 3).

Individuals displayed variation in diving and foraging beyond those described for diving

efficiency. For example, the maximum depth reached by individuals during the 40 days ranged

from 74.8 to 217.8 m, and trips at sea ranged from less than one day to 24 days. We provide a

general description of dive characteristics, foraging effort measures, and oxygen storage and

carrying capacity indices in Supplement 5.

3.2 Foraging effort relationships with oxygen storage and carrying capacity indices, sex, and reproductive status

Two pre-breeding foraging effort measures were associated with oxygen storage and carrying capacity indices, sex, and reproductive status (Fig. 3, Table 4). The best model for vertical distance travelled indicated that it was increasing with Hb, and the best model for time at sea suggested that early-laying penguins spent less time at sea prior to capture than breeding and non-breeding penguins (Fig 4, Table 4, Supplement 6). However, Table 4 shows that other models had ΔAICc < 4, including the null model for vertical distance travelled. The presence of many competitive models is likely explained by the small sample size (13 individuals) and some of the relationships we observed between covariates. For example, given that the 8 males in our samples had a significantly lower average Hb level than the 5 females (W = 34, p-value = 0.045, n = 13; Fig. 4A), it is not surprising that the second-best model for vertical distance travelled showed that males travelled less than females. Similarly, early-laying penguins had, on average, a lower Hb level than non-breeding penguins (Fig. 4B). Although this difference was not significant (W = 15, p-value = 0.057, n = 8), it may contribute to the fact that the second-best model for time at sea did not include breeding status but included a positive relationship with Hb. Average trip duration had the null model as best model (Table 4), and thus was not considered further. We note that a small sample increases the chance of making a type II error (failing to reject the null hypothesis when it is false) and that AICc strongly penalizes model complexity, thus with a larger sample size more complex relationship may have been detected.

3.3 Potential effects of tagging

Almost all (32/35) tagged penguins recaptured gained weight during the winter, with an

average gain of 1.1 kg ± 0.7. However, on average the tagged penguins were significantly lighter

at recapture than the untagged penguins (tagged penguins: 6.9 kg ± 0.6, 5.7 - 8.2kg; untagged

penguins: 7.3 kg ± 0.5, 6.2 - 8.8 kg; $t_{66.498}$ = 2.812, p-value = 0.006, n = 70). We did not find any

other significant differences between tagged and untagged penguins. Their Hb and Hct values

were similar (Hb: $t_{62.863}$ = -0.524, p-value = 0.602, n = 66; Hct: $t_{63.633}$ = 1.480, p-value = 0.144, n

= 66), and their breeding status did not differ significantly (χ^2 = 1.167, simulated p-value =

0.604, n = 39). However, given the small sample size for the breeding status analysis (tagged

=18, untagged = 19), we note that, compared to the untagged penguins, there was a higher

percentage of tagged penguins that did not participate in breeding and a lower percentage of

tagged penguins that laid early (Supplement 7). Overall, our results suggest that it is unlikely the

relationships we found with blood oxygen storage and carrying capacity indices (i.e., Hb and

Hct) are affected by tagging, but that tagging is associated with reduced mass and that the effects

of tagging on breeding behaviour and reproductive success should be further explored. See

Supplement 7 for more details.

4. Discussion

Our results show that the diving efficiency of gentoo penguin was related to Hb, with the

efficiency of deep dives (≥ 140 m) increasing with Hb level. This increase in efficiency can be

explained by the fact that Hb limits the total amount of oxygen able to be carried by the blood

(Minias 2015, Roncon et al. 2018). Of the factors that could have affected diving efficiency,

higher Hb was primarily associated with an increase in the bottom time of the dive rather than

reduced recovery time (i.e., no decrease in post-dive surface interval with increased Hb levels

was observed). Gains in diving efficiency, and associated increased time at the bottom, are likely

important to gentoo penguins, as they increase the time available to search for and capture prey

during foraging dives. As prey are almost always encountered during the bottom phase of dives

(Simeone & Wilson 2003, Takahashi et al. 2004, 2008, Kokubun et al. 2010), increased bottom

time translates into time spent where prey is most likely to be encountered. For dives $\geq$ 140 m,

penguins with high Hb (20-31 g/dL) increased their mean bottom phase by 22 % compared with

penguins with low Hb (10-20 g/dL). Such an increase could impact foraging success.

The proportion of deep dives performed by penguins increased with Hb levels, with only

individuals with high Hb levels regularly diving 140 m or deeper. Vertical distance travelled pre-

breeding also increased with Hb levels, suggesting that individuals with high Hb levels may use

a deep diving strategy that consequently requires more diving both because they tend to dive

deeper and because they have to travel longer in order to reach deep waters further from shore.

Such individuals are accessing resources only available at deep depths, giving them access to a

wider variety of prey items and potentially decreasing intraspecific competition (e.g., Cimino et

al. 2016). Similar findings have been documented in other species. For example, macaroni

penguins (*Eudyptes chrysolophus*) forage in habitats within their aerobic capacity (Crossin et al.

2015), and Hb is correlated with higher-quality diets (reviewed in Minias 2015).

Diving efficiency displayed a complex non-linear relationship with Hct levels. The

complexity of this non-linear relationship could be associated with the counteracting effects of

high Hct levels. While a high Hct level is associated with a higher number of oxygen-carrying

red blood cells, it is also linked to increased blood viscosity, and thus increased cardiac workload

or decreased blood flow (Birchard 1997, Schuler et al. 2010, Reinhart 2016). In addition,

optimum Hct levels can be affected by factors such as age, sex, breeding status, and disease

(Reinhart 2016, Johnstone et al. 2017, Brown et al. 2021). In research employing Hct as an index

of body condition, higher values are often assumed to indicate better physiological performance

(reviewed in Fair et al. 2007, Minias 2015). In contrast, our results suggest that Hct has a

complex non-linear relationship with diving efficiency and that higher Hct may not always result

in higher diving efficiency. The use of Hct as a physiological indicator has been criticized due to

inconsistent relationships with performance and fitness-related traits in many species (reviewed

in Fair et al. 2007, Minias 2015, Johnstone et al. 2017). For example, intermediate, rather than

high Hct levels were linked to maximum longevity and recruitment rate in house wrens

(*Troglodytes aedon*, Bowers et al. 2014). Our results suggest that, while Hct relates to diving

efficiency, linear relationships are likely too simplistic. Instead, non-linear models should be

used to capture the complex relationship between these variables, though such models require a

sufficiently large sample size.

While values of both Hb and Hct aligned with previously reported values for gentoo

penguins (20.8 ± 5.4 g/dL and 49.2 ± 4.7 % in our study, n = 66, ; 20.7 ± 1.6 g/dL and 50 ± 4 %

in Ibañez et al. 2015), they were not closely correlated to one another (see Supplements 2-4).

Because mature red blood cells have similar amounts of Hb, we expected a correlation between

Hct and Hb. However, ecological stressors can affect these indices differently (Bańbura et al.

2007, Wagner et al. 2008, Johnstone et al. 2017), and weak correlations have been documented

in other seabird species (e.g., Kaliński et al. 2011). While some anaemic birds with low Hb

concentrations may be unable to improve their condition, some will actively regenerate red blood

cells (Jaensch & Clark 2004, Fair et al. 2007, Campbell et al. 2010). In such cases, whether Hb

and Hct will remain correlated will be affected by erythropoiesis. Erythropoiesis increases the

number of immature red blood cells, which are larger and initially have lower Hb content,

disproportionately increasing Hct but not Hb (Wagner et al. 2008, Campbell et al. 2010,

Johnstone et al. 2017). We identified one individual that may have demonstrated evidence of

compromised health. This non-breeding penguin had the lowest Hb and yet the fourth highest

Hct. Anaemia can be associated with additional health conditions, such as heavy parasite loads or

severe injury (Vleck et al. 2000, Jaensch & Clark 2004, Fair et al. 2007, Clark 2015). Despite

showing no apparent signs of injury at the time of re-capture, this anaemic individual was one of

the only individuals that lost mass between April and October. Such an example supports

experimental designs that measure both Hb and Hct when assessing individual health (Johnstone

et al. 2017).

According to optimal foraging theory, penguins maximize their bottom time and

minimize energy spent travelling down to where prey is located (Stephens & Krebs 1986, Wilson

et al. 2010, Zimmer et al. 2010). As such, shallow dives should be preferred, requiring less travel

time and energy to swim down to depth (Shepard et al. 2009, Zimmer et al. 2010). Deeper dives

are only profitable when prey capture rates, size, or caloric value outweigh the energetic costs of

diving (Sala et al. 2014b). In our data, peak diving efficiency occurred in shallow foraging

habitat at approximately 60 m deep (Fig. 1A), which is deeper than previously reported values

(25-30 m; Lee et al. 2015). In line with theory, this putative optimal depth (60 m) was repeatedly

visited by penguins (see Supplement 5). However, potentially to reach larger and/or energy-rich

prey, many penguins foraged far deeper than this putative optimal depth (maximum depth

reached by individuals: 166 m $\pm$ 29).

Diet studies suggest that deeper dives may provide greater access to larger prey that may

be more valuable to gentoo penguins than the smaller prey in shallower habitat. Clausen et al.

(2005) compared the diet of gentoo penguins in colonies around the Falkland Islands with

surveys of prey abundance during the 2000 pre-breeding season. They highlight preferential

foraging of squid and fish over krill, which has also been highlighted in the historical diet of

gentoo penguins (McMahon et al. 2019). Prey likely to be essential for Pebble Island and Race

Point penguins includes bentho-demersal lobster krill (*Munida gregaria*), found at depths 26-116

m, and Patagonian squid (*Doryteuthis gahi*)*,* found at depths 60-96 m, both of which were in

high abundance close to the colonies in 2000 (Clausen et al. 2005). A deeper squid species

(*Moroteuthopsis ingens*), found at depths 100-166 m, and various rock cod species

(*Patagontothen spp.*), found at depths 40-205 m, are abundant further offshore and were found in

their diet in 2000 (Clausen et al. 2005). These prey species vary in size and energy density. For

example, the Patagonian squid found at shallower depth is smaller (max length = 28.0 cm,

Palomares & Pauly 2023) but has a slightly higher energy density (5.0 kJ/g; Ciancio et al. 2007)

than rock cod species such as *Patagonotothen ramsayi* (max length = 44.4 cm, Froese & Pauly

2023; 4.7 kJ/g, Ciancio et al. 2007). In a more recent assessment of prey, rock cods were found

to be a principal prey item for gentoo penguins in the Falkland Islands (Handley et al. 2016). In

addition, underwater footage demonstrated that gentoo penguins do not deviate from their path

for small prey such as lobster krill, but actively chase larger prey items such as large squid and

fish species (Handley et al. 2018). Generally, for marine species, prey size increases with depth

(Masello et al. 2010, Miller et al. 2010).

Using diving efficiency as a measure of diving abilities implicitly assumes that animals

typically dive to their aerobic dive limit and does not account for the numerous factors

influencing dive time and efficiency. For example, predators adapt foraging strategies based on

prey conditions (e.g., Magellanic Penguins, *Spheniscus magellanicus*; Sala et al. 2014a) and may

abandon a dive early in poor prey conditions (e.g., Thums et al. 2013, Viviant et al. 2016). Such

behavioural factors likely explain the low diving efficiency at shallow depths. However, deep

diving is more energetically costly to undertake (Shepard et al. 2009, Zimmer et al. 2010). As

such, oxygen storage and carrying capacity are more likely to influence the behaviour of animals

foraging at deeper depths (Thompson & Fedak 2001, Viviant et al. 2016) and may explain why

the relationship with Hb is more apparent at depth.

Gentoo penguins that laid early spent significantly less time at sea than non-breeding

penguins, a difference we attribute to higher foraging efficiency rather than higher breeding

obligations. Participating in breeding compels individuals to defend nest locations and partake in

pair-bonding behaviours, requiring many species of penguins to fast during this time (Williams

1995). While gentoo penguins do not fast during breeding, they increase their time on land to

maintain a nest site for approximately two weeks before egg-laying (Black 2016). All monitored

gentoo penguins, except one non-breeder, were captured at least 11 days (14.5 ± 2.2 days,

September 30 to October 07) before eggs were first seen in the colony (October 18). Thus, while

early-laying penguins may have been maintaining nest sites for a few days at the end of the TDR

observation period, most of the 40-day data would be prior to nesting and may be associated with

early-laying penguins requiring less time to meet energetic demands.

Acquisition of food resources during the pre-breeding period is essential, as breeding

birds must expend additional energy to defend their nest and produce eggs (Williams 2012).

Successful acquisition of high quality resources relates to earlier laying, which can influence

reproductive success (Sorensen et al. 2009). For example, early breeding European shag

(*Phalacrocorax aristotelis*) spend less time foraging in winter than other individuals (Daunt et al.

2006), successfully breeding Adélie penguins (*Pygoscelis adeliae*) are more efficient foragers

(Lescroël et al. 2010), and breeding Manx shearwater (*Puffinus puffinus*) spend less time in

winter foraging and flying than those that defer breeding (Shoji et al. 2015). In species such as

the Adélie penguin, non-breeding individuals were 90 % of the mass of breeders (Vleck & Vleck

2002). We found no significant differences in the average mass between non-breeding, breeding,

and early-laying penguins. However, in line with what we would expect if early-laying penguins

were more efficient foragers, the non-breeding penguins in our sample were 93 % the weight of

early-laying penguins. As such, the decrease in time at sea for early-laying gentoo penguins

could indicate that these individuals may achieve food requirements in less time during the pre-

breeding period.

While none of the indices of oxygen storage and carrying capacity were in the best model

for time spent at sea (the best model only included breeding status), Hb, Hct, and mass were

included in models with substantial support (ΔAICc < 2), and there are known relationships

between these indices and reproduction. In many bird species, Hb levels in females decline

during the egg production and laying period (reviewed in Minias 2015). Given the observed

increase in diving efficiency with Hb levels, such reduction in Hb could create important trade-

offs between foraging capacity and reproduction. Other penguin species, such as macaroni

penguins, have evolved extreme size dimorphism due to similar trade-offs between aerobic

condition and egg production (Jubinville et al. 2020). As the physiology of each sex is affected

differently by breeding (Williams 2012, Desprez et al. 2018), future studies with a larger sample

size could explore sex-specific associations between Hb levels, foraging capacity, and breeding,

and further disentangle these potential trade-offs.

Foraging is an essential activity for survival, and understanding how an individual's

physiology and health affect dive performance, selection of foraging habitat, and breeding

participation is essential to understanding the effects natural and anthropogenic ecosystem

changes can have on populations. In the Falkland Islands, the continental shelf is a popular

fishing ground, and gentoo penguins are occasionally caught as bycatch and potentially compete

for prey with fishing industries (Clausen & Pütz 2003, Trathan et al. 2015). Their extended

winter range could overlap with exploration for hydrocarbons, and there is a consideration of

further developing fisheries inshore (Augé et al. 2015). Given the associations between an

individual's Hb and Hct levels and their diving and foraging patterns, these human activities

could affect portions of the population differently. For example, reducing prey availability

nearshore could be especially detrimental for penguins with low oxygen storage and carrying

capacity.

According to the island-wide census of 14 colonies, nesting pair numbers and breeding

success in 2018 were above the long-term annual average recorded since 2003 (Stanworth and

Crofts 2019), suggesting that the penguins in our study experienced relatively prosperous

environmental conditions. However, in a year of low environmental conditions, the costs of an

inferior strategy could be more prevalent (Fronstin et al. 2016, Storey et al. 2017). Therefore,

further understanding the relationship between body conditions and diving efficiency, and its

impact on foraging success and breeding participation across a range of environmental

conditions, may be crucial in understanding breeding participation and population dynamics in

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Table 1. The number of individuals tagged, recaptured, and sampled as control. We state how

many individuals were captured and recaptured at each colony, the number of individuals for

which we had blood samples, the number that could be sexed, and the female-to-male sex ratio

(F:M). For the recaptured penguins, we present in brackets the number of individuals for which

the TDR tags were still attached at recapture.

	Tagged	Recaptured				Control		
		Recaptured (with tag)	Blood (with tag)	Sexed (with tag)	F:M (with tag)	Blood	Sexed	F:M
Race Point	33	16 (7)	13 (5)	11 (4[*])	2:9 (0:4)	15[**]	14	6:8
Pebble Island	33	19 (14)	19 (14)	18 (13)	10:8 (5:8)	20	19	7:12
Total	66	35 (21)	32 (19)	29 (17)	12:17 (5:12)	35[**]	33	13:20

[*] One individual that was sexed did not have blood at recapture; it was sexed based on blood at

capture. As we use blood at recapture for the indices of oxygen storage and carrying capacity,

this individual could not be included in the final GAMM modelling the relationship between

diving efficiency and indices of oxygen storage and carrying capacity.

[**] We were unable to measure the Hb of one penguin and the Hct of another penguin.

Table 2. Results from the final GAMM for diving efficiency. We present the estimate, standard

error (SE) of the parametric coefficient (intercept and fixed effect), as well as their associated

test statistic, and p-value. We also present the estimated degrees of freedom (EDF), test statistic,

and p-value associated with the smooth terms of the GAMM. For model selection, we used the

shrinkage method with a double penalty, which allows non-important terms to be shrunk to the

null. A total of 19 penguins were included in this analysis (5 from Race Point and 14 from

Pebble Island).

Term	Parametric coefficient				Significance of smooth term		
	Estimate	SE	t	p-value	EDF	F	p-value
Intercept	0.295	0.006	47.606	< 0.001			
Colony (Race point)	0.034	0.014	2.474	0.014			
s(Depth)					7.856	33.369	< 0.001
s(Mass)					1.000	0.005	0.943
s(Hct)					1.000	2.302	0.129
s(Hb)					1.000	0.140	0.708
ti(Depth, Mass)					1.000	2.613	0.106
ti(Depth, Hct)					10.675	3.985	< 0.001
ti(Depth, Hb)					9.683	5.110	< 0.001

Table 3. Parameter estimates and test statistic values for the post-hoc GAMs relating the

proportion of deep dives to each Hb and Hct, as well as for the post-hoc GAMMs relating either

post-dive surface interval of deep dives or bottom time of deep dives to each Hb and Hct. For the

proportion of deep dive, 19 penguins were included in the GAMs (5 from Race Point and 14

from Pebble Island). For the post-dive surface interval and bottom time, only 16 were included in

the GAMMs because three individuals did not make dives $\geq$ 140 m.

Term	Parametric coefficient				Significance of smooth terms		
	Estimate	SE	t	p-value	EDF	F	p-value
Proportion of deep dives (Hb)							
Intercept	0.064	0.015	4.122	0.001			
Colony (Race point)	0.019	0.035	0.544	0.595			
s(Hb)					2.999	3.862	< 0.001
Post-dive surface interval (Hb)							
Intercept	117.404	8.353	14.055	< 0.001			
Colony (Race point)	-7.190	19.744	-0.364	0.717			
s(Hb)					1	0.670	0.415
Bottom time (Hb)							
Intercept	91.018	2.808	32.41	< 0.001			
Colony (Race point)	2.964	7.808	0.38	0.705			
s(Hb)					1	6.498	0.012
Proportion of dives (Hct)							
Intercept	0.074	0.024	3.086	0.007			
Colony (Race point)	-0.019	0.046	-0.400	0.694			
s(Hct)					0.653	0.189	0.127
Post-dive surfaces interval (Hct)							
Intercept	119.317	7.942	15.023	<0.001			
Colony (Race point)	6.773	16.444	0.412	0.681			
s(Hct)					1	1.39	0.241
Bottom time (Hct)							
Intercept	90.738	3.543	25.613	< 0.001			
Colony	-9.817	7.771	-1.263	0.209			

(Race point)				
s(Hct)		1	0.401	0.528

Table 4. Model selection results for the multiple linear regressions relating three measures of

foraging effort to oxygen store indices, sex, and breeding status. We present the number of

parameters estimated (k), the Akaike Information Criterion with correction for small sample size

(AICc), and the difference in AICc from the best model (ΔAICc). While we explore all possible

combinations of three covariates for each measure, we only show in the table the null model (no

covariates) and models that are either with ΔAICc < 4 of the best model (model with lowest

AICc value) or all the models that outperformed the null model if the null model had a ΔAICc ≥

4. Note that no models with breeding status are included in the table for vertical distance

travelled and trip duration, as none of them had a ΔAICc small enough to be part of the subset

presented. A total of 13 penguins (all from Pebble Island) were included in these analyses.

Response	Covariates	k	AICc	ΔAICc
Vertical distance travelled	Hb	3	171.0	--
	Sex	3	171.4	0.34
	Hct	3	172.6	1.56
	(Null)	2	172.7	1.69
	Hb + Hct	4	173.8	2.76
	Hb + Sex	4	173.8	2.78
	Hct + Sex	4	173.9	2.92
Time at sea	Breeding status	4	66.0	--
	Hb + Hct + Mass	5	67.1	1.17

	Hb + Mass	4	67.9	1.91
	Hb	3	68.3	2.32
	Breeding status + Mass	5	69.8	3.82
	Breeding status + Sex	5	70.3	4.36
	Breeding status + Hb	5	70.4	4.47
	(Null)	2	70.5	4.51
Trip duration	(Null)	2	59.0	--
	Hct	3	59.4	0.40
	Hb + Hct	4	60.4	1.48
	Sex	3	61.8	2.82
	Hb	3	62.0	3.00
	Mass	3	62.3	3.39
	Hct + Mass	4	62.8	3.83

Figure 1. Slices of predictions from the final GAMM. Panel A displays how the predicted diving

efficiency changes in response to varying maximum depth values of a dive. The blue band

represents the 95% confidence intervals. The points show the observed values. Panel B shows

how diving efficiency is predicted to change with both depth and Hb, while panel C shows how

diving efficiency is predicted to change with both depth and Hct. For both these panels, the

colour represents the diving efficiency value (yellow representing highest efficiency) and the

grey bands represent the standard error (more opaque grey representing higher standard error).

The areas with larger standard error values (i.e., more opaquely shaded) should be interpreted

more cautiously. A total of 19 penguins were included in this analysis (5 from Race Point and 14

from Pebble Island).

Figure 2. Association between Hb and the proportion of dives an individual made at deeper

depth (≥ 140 m, panel A), as well as changes in deep dives' post-dive surface intervals (panel B)

and bottom times (panel C). A total of 19 penguins were included in the analysis from panel A (5

from Race Point and 14 from Pebble Island), but only 16 were included in panels B and C

because three individuals did not make dives ≥ 140 m. The blue lines represent the predicted

value, and the blue bands represent the 95% confidence intervals. The points represent the

observed values.

Figure 3. Relationship between Hb and reproductive status and two of the pre-breeding foraging

effort measures for the penguins from Pebble Island. Panel A shows the relationship between the

sum of the vertical distance travelled while at sea and Hb. The line shows the linear regression

line, the band is the 95% confidence interval. Panel B shows the relationship between the time

spent at sea and the reproductive status. In both panels, the points are the observations. A total of

13 penguins (4 non-breeding, 5 breeding, and 4 early-laying penguins from Pebble Island) were

included in these analyses.

Figure 4. Variation in the three oxygen storage and carrying capacity indices across sex (panels

A, C, E) and breeding status (panels B, D, F). Panels A and B show the boxplots (with black

points identifying outliers) for Hb, panels C and D for Hct, and panels E and F for mass. In all

panels, the grey points are the observations. To be able to see overlapping observation, we added

919 horizontal jitter to the points. The values presented are the p-values associated with the Wilcoxon

920 rank sum tests associated with the pairwise comparisons. A total of 13 penguins (4 non-breeding,

921 5 breeding, and 4 early-laying penguins from Pebble Island) were included in these analyses.