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Blood analytes of hawksbill sea turtles (*Eretmochelys imbricata*) from Florida waters: reference intervals and size-relevant correlations

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Assessments of health variables in wild animal populations have evolved into important tools for characterizing spatiotemporal population trends and fitness, effects of stressors, diseases, and ecosystem health. Blood as a sample matrix can be obtained fairly non-invasively in the field, with preservation and sample processing techniques that allow for readily available routine and advanced diagnostic testing of blood. For wild-caught hawksbill sea turtles (*Eretmochelys imbricata*) foraging in southeastern Florida, USA, the objectives of this study were to (1) establish reference intervals for hematological and 24 plasma biochemical analytes, (2) determine length- and body condition-specific relationships with blood analytes, and (3) determine how water temperature influenced plasma biochemical analytes. Reference intervals were established for clinically normal juvenile ($n=26$) and subadult ($n=39$) hawksbills, with descriptive data reported for adult turtles ($n=3$). Although subadults (mainly captured at Palm Beach County) were heavier and larger with greater body depth, juveniles (mainly captured at Monroe County) had a higher body condition index. Positive length-specific correlations were identified for packed cell volume, eosinophils, aspartate aminotransferase, phosphorus, cholesterol, glutamate dehydrogenase, total protein, albumin, and globulins, with negative correlations including alkaline phosphatase, creatine kinase, calcium, calcium to phosphorus ratio, and glucose. Subadults had less frequent morphological features of red blood cell regeneration compared to juveniles. These findings provide insight into life-stage class differences regarding hematopoiesis, antigenic stimulation, somatic growth, dietary shifts, nutritional status, osmoregulation, metabolism, physical activity or stress levels, and possible habitat differences. Life-stage class is the likely driver for the observed blood analyte differences, in addition to influences from water temperature. The data herein offer baseline information for a snapshot in time for critically endangered hawksbills inhabiting the Florida reef system and for answering individual- and population-relevant questions of relevance to conservation and population management.

KEYWORDS

health assessment, hematology, marine turtle, plasma biochemistry, physiology, protein electrophoresis, somatic growth

1. Introduction

Six sea turtle species are known to occur in waters off the east and west coasts of Florida, USA (Foley et al., 2003; Eaton et al., 2008). After olive ridleys (*Lepidochelys olivacea*), the critically endangered (Mortimer and Donnelly, 2008) hawksbill sea turtle (*Eretmochelys imbricata*) is the rarest species to occur in Florida (Meylan and Redlow, 2006; Mortimer and Donnelly, 2008), with this aggregation being the second most northern of the Atlantic Ocean population (Meylan et al., 2011; Wood et al., 2013). The Atlantic hawksbill population appears to be genetically distinct from the Pacific population (Wood et al., 2013). Hawksbills are believed to occupy the Florida reef system due to warm water transport by the Florida current from the Gulf of Mexico and the Caribbean (Meylan and Redlow, 2006; Blumenthal et al., 2009). Although hawksbill nesting in Florida is uncommon (Meylan and Redlow, 2006), juvenile and subadult life-stage classes are known to inhabit Florida's waters year-round (Wood et al., 2013, 2017). The Florida Continental Reef Tract provides refuge for hawksbills and is home to numerous sponges and octocoral species on which these animals forage (Moyer et al., 2003; Banks et al., 2008; Wood et al., 2017). Hawksbill turtles in Florida waters originate from Mexico and the Caribbean via the Florida Current and/or Gulf Stream, recruiting from oceanic to neritic habitats predominantly in the Florida Keys (i.e., Monroe County) as small juveniles. It is apparent that some move along the Southeast Florida Continental Reef Tract to eventually reach their northern terminus in Palm Beach County, Florida, as they advance through their subadult life-stage (Wood et al., 2013). Then, upon reaching sexual maturity, the young adults undertake reproductive migrations out of Florida, and are only seen transiently thereafter.

Despite the known occurrence of these animals in Florida, no assessment of overall health using blood analytes has occurred for this aggregation to date. Health assessments have been conducted for wild-caught hawksbills from Oman (nesting adults) (Alkindi et al., 2002), Brazil (nesting adults) (Goldberg et al., 2013), the Persian Gulf (nesting adults) (Ehsanpour et al., 2015), Australia (immature foraging) (Whiting et al., 2014), the Galápagos Archipelago (immature and adult foraging) (Muñoz-Pérez et al., 2017), Mexico (nesting adults) (Salvarani et al., 2018), Belize (immature) (Crooks et al., 2023) and in rehabilitating hawksbills from the United Arab Emirates (juvenile) (Hampel et al., 2009; Caliendo et al., 2010). Reference intervals using American Society for Veterinary Clinical Pathology (ASVCP) guidelines (Friedrichs et al., 2012) have only been established for one wild-caught hawksbill population in Australia (Whiting et al., 2014); however, several other hawksbill studies report measures of central tendency and full range of values (Goldberg et al., 2013; Ehsanpour et al., 2015; Salvarani et al., 2018). Establishment of blood analyte reference intervals in sea turtles is important, as various stressors that sea turtles face continue to intensify; therefore, health assessments are becoming increasingly valuable to conservation efforts as they provide baseline data for future population health and fitness comparisons in the face of environmental changes, increasing stressors, and potential disease outbreaks (Deem et al., 2001; Aguirre and Lutz, 2004; Wikelski and Cooke, 2006; Cooke and O'Connor, 2010; Deem and Harris, 2017; Reséndiz and Lara-Uc, 2018; Mashkour et al., 2020; Page-Karjian et al., 2020; Perrault et al., 2020), in addition to improving clinical decision-making in individual animals undergoing veterinary care (Delgado et al., 2011).

Coral cover in the Florida Continental Reef Tract, where hawksbills are known to inhabit, is declining at alarming rates due to climate change, disease, and deteriorating water quality (Ruzicka et al., 2013; Voosen, 2019; Neely et al., 2021); therefore, establishing baselines of hawksbill health variables is necessary to understand physiological responses of animals and future population and ecosystem challenges that may result from habitat degradation or loss. For wild-caught juvenile and subadult hawksbills foraging in southeastern Florida, USA, the objectives of this study were to (1) establish reference intervals for hematological and 24 plasma biochemical analytes, (2) determine length- and body condition-specific relationships with blood analytes, and (3) determine how water temperature influenced plasma biochemical analytes.

2. Materials and methods

2.1. Ethical procedures

This study was reviewed and authorized by the National Marine Fisheries Service (NMFS) [Permit #22988], Florida Fish and Wildlife Conservation Commission (FWC) [Marine Turtle Permits #021 and #077], Florida Keys National Marine Sanctuary [Research Permit #175], and University of Florida's Institutional Animal Care and Use Committee (IACUC) [#201706823]. All handling and sampling procedures of sea turtles were performed according to NMFS and FWC regulations (NMFS SEFSC, 2008).

2.2. Capture technique, morphometrics, and sample collection, processing, and analysis

Hawksbill turtles were hand-captured during snorkel and/or scuba surveys between Jun 2017 and Oct 2020 along the 2–30 m deep nearshore reefs of the Southeast Florida Continental Reef Tract from Jupiter, Florida, USA (26.987°N, –80.033°W) to Key West, Florida, USA (23.673°N, –82.981°W) (Figure 1). The water temperature was recorded at the time and depth of each capture with a Sherwood Wisdom 2 (Sherwood Scuba®, Santa Ana, CA) dive computer while on scuba, and/or the vessel's onboard surface water temperature gauge when snorkeling in water less than 9 m deep. Primarily, two counties in Florida were utilized for this study including Palm Beach (towns of Jupiter and West Palm Beach) and Monroe (towns of Islamorada and Key West), which are known foraging areas of this species in Florida (Meylan and Redlow, 2006; Wood et al., 2013, 2017). One turtle was captured in Miami, Florida, USA (Miami-Dade County) (25.690°N, –80.085°W). This individual was included in analyses of the Monroe County turtles due to the close proximity of the two locations. Turtles captured at depths exceeding 10 m (>1 atm) were brought to the surface at a rate of ≤9 m/min. Turtles were transferred to a research vessel within 20 min of capture and examined for the presence of internal passive integrated transponder (PIT) tags.

Within 5–10 min of returning to the vessel, up to 5 mL of blood were collected from the dorsal cervical sinus using 20-gauge, 1.5" needles and 7 mL lithium heparin vacutainer tubes (Becton Dickinson, Franklin Lakes, New Jersey, USA), following standardized safe blood volume withdrawal guidelines for sea turtles (NMFS SEFSC, 2008).

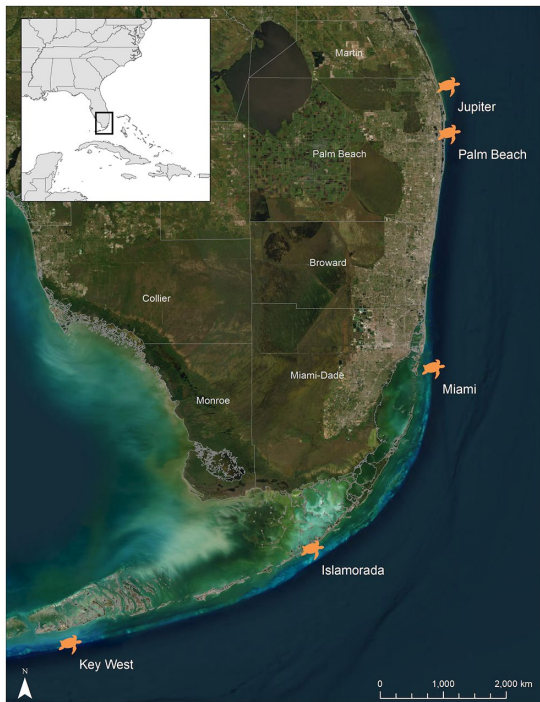


FIGURE 1
Locations of the five capture sites of hawksbill sea turtles (*Eretmochelys imbricata*) across Florida's, USA reef tract. Jupiter and Palm Beach are located in Palm Beach County, while Islamorada and Key West are located in Monroe County. One individual was captured off of a reef in Miami (Miami-Dade County). This individual was included in the Monroe County group for statistical analyses.



FIGURE 2
Photograph of a subadult hawksbill sea turtle (*Eretmochelys imbricata*) from Palm Beach County representative of the study turtles with associated normal epibiota coverage.

The sampling site was disinfected with alternating applications of povidone iodine and 70% isopropyl alcohol both before and after venipuncture. The vacutainers were then transferred to an insulated sleeve (i.e., bubble wrap) and placed on wet ice for up to 8 h until processing.

Following blood collection, standard (i.e., notch to tip) straight carapace length (SCL) and the maximum body depth (BD) of each subject were measured with aluminum calipers. Mass was estimated

to the nearest kg using a 300 kg digital scale. Body condition index (BCI) was calculated (Bjornndal et al., 2000), applying a correction factor of 10,000. Physical examination included assessment of the presence and location of conspicuous epibiota (e.g., macroalgae, coral, barnacles, etc.; Figure 2) and/or physical abnormalities, which were diagrammed and photographed. Prior to release, Inconel tags (National Band and Tag Co., Newport, Kentucky, USA) were placed on the trailing edge of the second distal scale of each front flipper, and/or one PIT tag (Biomark, Inc., Boise, Idaho, USA) was injected into the right front shoulder in all turtles of sufficient size (>30 cm SCL). All turtles were released ~20 min after sample collection, at or near the capture site.

Upon return to the laboratory, whole blood was well-mixed and packed cell volume (PCV) was determined using the average of two microhematocrit tubes after centrifugation for 5 min at 7,500g (12,000 rpm) using a ZipCombo microhematocrit centrifuge (LW Scientific, Lawrenceville, Georgia, USA). Blood films were prepared from well-mixed whole blood, air-dried, and stained with Wright-Giemsa (Harleco®, EMD Millipore, Billerica, Massachusetts, USA). Blood film evaluation included white blood cell (WBC) estimate (Weiss, 1984) using an eyepiece of 18 mm diameter, a 200 white blood cell differential including heterophils, lymphocytes, monocytes, eosinophils, and basophils, and morphological evaluation of red blood cells (RBC), WBCs, and thrombocytes. The heterophil:lymphocyte ratio was calculated.

The remaining whole blood was spun at 1350g (3,300 rpm) in a Champion E33 centrifuge (Ample Scientific, LLC, Norcross, Georgia, USA) for 6–8 min. Plasma was separated from the RBC and WBC, placed in cryovials, and stored in a standard freezer (−20°C) until shipment on dry ice after 48 h to the University of Florida for storage in an ultralow freezer. After 8–434 d (mean ± SD = 139 ± 102 d), samples were then shipped to the University of Miami Avian and Wildlife Laboratory for biochemical analyses using an Ortho 250XR (Ortho Clinical Diagnostics, Rochester, New York, USA) dry slide chemistry analyzer. Biochemical analytes of interest included alkaline phosphatase (ALP), amylase, aspartate aminotransferase (AST), bile acids, blood urea nitrogen (BUN), calcium, phosphorus, chloride, cholesterol, creatine phosphokinase (CK), gamma-glutamyl transferase, glucose, glutamate dehydrogenase (GLDH), lipase, magnesium, potassium, sodium, triglycerides, and uric acid. The calcium:phosphorus ratio was calculated.

Plasma protein electrophoresis was conducted using the SPIFE 3000 system (Helena Laboratories, Inc., Beaumont, Texas, USA). Fractions of interest included albumin and total globulins. The albumin:globulin ratio was calculated.

3. Statistical analyses

Statistical analyses were performed using MedCalc® statistical software (version 19.6, Ostend, Belgium) and SPSS for Windows (version 27; Chicago, Illinois, USA). Measures of central tendency and range are reported for all blood analytes in Standard International (SI) units. Reference intervals (95% with associated 90% confidence intervals) were determined for sea turtles of juvenile and subadult life-stage classes that fit the inclusion criteria for the study (Moore et al., 2020), using parametric methods based on recommendations by Friedrichs et al. (2012) for sample sizes ≥20, but <40. Normality

was assessed using the Shapiro-Wilk test, while outliers were detected using the Dixon-Reed test. Outliers were subsequently removed and when necessary, logarithmic or Box-Cox transformations were employed to generate accurate reference intervals. Several blood analytes could not be normalized to fit a Gaussian distribution; therefore, reference intervals for those analytes were calculated using the robust method.

Spearman correlations were used to determine relationships between individual hematological and plasma biochemical analytes. Differences in SCL by capture location (Palm Beach v. Monroe) and BCI in juveniles and subadults were assessed using independent samples t-tests. Linear regression analysis was used to assess the relationship of SCL to mass (using log-transformed data). Stepwise backward multiple regression was used to determine the impacts of SCL, BCI, water temperature (independent variables) upon capture on blood analytes (dependent variables). Outliers were determined using Tukey's method and subsequently removed from analyses. Transformations were employed as necessary. To determine if polychromasia, anisocytosis, basophilic stippling, and number of immature RBCs/100 mature RBCs were influenced by life-stage class, Kruskal-Wallis with Dunn's post-hoc comparisons were used.

4. Results

4.1. Physical examination and morphometrics

A total of 68 hawksbills were captured from 5 Jun 2017–30 Oct 2020. Water temperatures were similar for both sites from May–October, when turtles were typically captured: 23.3–29.4°C (mean 27.4°C) in Palm Beach County and 26.7–29.4°C (mean 28.1°C) in Monroe County, respectively. Inclusion criteria for study animals included the absence of overt acute or debilitating external injuries, normal behavior, visibly adequate body condition, and absence of difficulties during blood withdrawal (Figure 2). Morphometric results by sampling location are reported in Table 1. Turtles of <50 cm SCL were considered juveniles (n=26), those of 50–78 cm SCL were considered subadults (n=39), and turtles >78 cm were considered adults (n=3) (Boulon, 1994; Wood et al., 2017). Hawksbills from Palm Beach County were significantly heavier, had a significantly longer SCL (Figure 3), and a greater body depth than turtles from Monroe

County (p<0.001 in all cases; Table 1); turtles from Monroe County had significantly larger BCI (p<0.001; Table 1). Log-transformed mass and SCL strongly correlated (r²=0.989; p<0.001; Figure 4). Using a t-test (t(60)=-2.378; p=0.021), we found that juveniles had a significantly higher BCI (mean±SD: 1.16±0.10; range: 0.97–1.39) than subadults (mean±SD: 1.10±0.10; range: 0.94–1.32), and overall that BCI tended to decrease with SCL (y=-0.002x+1.24; r²=0.079; p=0.027; n=62).

Small patches of red and/or brown algae (Figure 2), particularly on the posterior carapace, were ubiquitous, while small patches of calcareous algae were less common. Fire coral colonies (*Millepora* sp.) were occasionally found on larger individuals. Fourteen turtles (20%) had dented or chipped marginal scutes; seven (10%) had at least one scalloped hind limb; one (1%) had a partially amputated (~50%) hind limb; and two (3%) had minor front flipper damage. All described injuries/abnormalities were completely healed and were not expected to impact blood analytes. All turtles appeared to be in robust condition as observed by the thickset nature of the soft tissues surrounding the neck and flippers and BCI scores. No fibropapilloma tumors or acute or healed injuries specific to shark predation were observed.

All turtles were considered to be clinically normal and fit the inclusion criteria for this study as they were all active and alert, had robust subjective body condition scores (Tristan and Norton, 2017), minimal epibiota, and minor injuries or shell abnormalities that most likely did not impact blood analytes.

4.2. Reference intervals

Mild (1+) hemolysis was present in seven samples, while mild (1+) lipemia was present in one sample. Neither degree of interferences is thought to affect dry chemistry analyses (Andreasen et al., 1997; Stacy and Innis, 2017; Stacy et al., 2019). Measures of central tendency and range of the measured blood analytes are reported in SI units for all three life-stage classes (juvenile, subadult, and adult; Tables 2–4, respectively; conventional units are reported in Supplementary Tables S1–S3), while reference intervals, due to the sufficient number of data points, are described for juveniles and subadults only (Tables 2, 3). Results of morphological evaluation of RBC, WBC, and thrombocytes are shown in Table 5.

Spearman correlations of hematological and biochemical analytes from all study turtles identified several statistically significant correlations of biological relevance: bile acids and AST (r_s=0.363; p=0.032; n=35), GGT and AST (r_s=0.310; p=0.010; n=68), GGT

TABLE 1 Morphometric data of in-water assessed hawksbill sea turtles (*Eretmochelys imbricata*) from Florida, USA.

Measurement	Palm Beach (5 Jun 2017–30 Oct 2020)				Monroe (21 Jul 2017–20 Sep 2020)				t	df	p
	Mean±SD	Median	Range	N	Mean±SD	Median	Range	N			
Mass (kg)	27 ± 11	28	11–48	35	9 ± 6	7	3–22	27	-8.378 ^a	54.2	<0.001
SCL (cm)	63.3 ± 9.4	63.9	46.9–84.4	40	40.8 ± 8.5	39.8	27.4–56.2	28	-10.073	66	<0.001
BCI	1.09 ± 0.09	1.09	0.94–1.32	35	1.18 ± 0.09	1.15	1.03–1.39	27	3.901	60	<0.001
Body depth (cm)	23.3 ± 3.8	23.4	16.4–31.1	36	15.1 ± 3.2	14.4	9.8–21.2	28	-9.206	62	<0.001

^aWelch's t-test was used as variances were not homogenous.

Individual metrics for each county are included. Sampling dates are included parenthetically next to the capture location. Statistical differences between sites, determined using independent samples t-tests, are also indicated. BCI, body condition index; df, degrees of freedom; SCL, standard straight carapace length; SD, standard deviation.

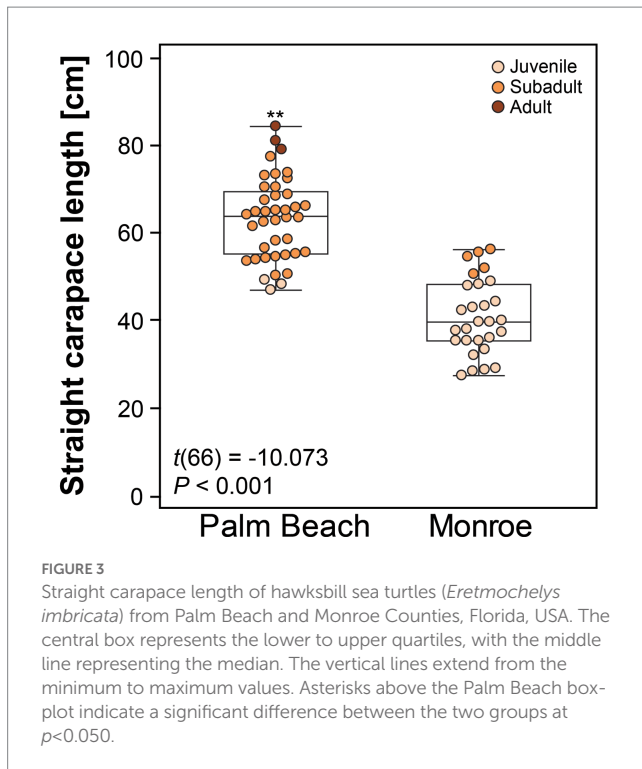


FIGURE 3
Straight carapace length of hawksbill sea turtles (*Eretmochelys imbricata*) from Palm Beach and Monroe Counties, Florida, USA. The central box represents the lower to upper quartiles, with the middle line representing the median. The vertical lines extend from the minimum to maximum values. Asterisks above the Palm Beach box-plot indicate a significant difference between the two groups at $p < 0.050$.

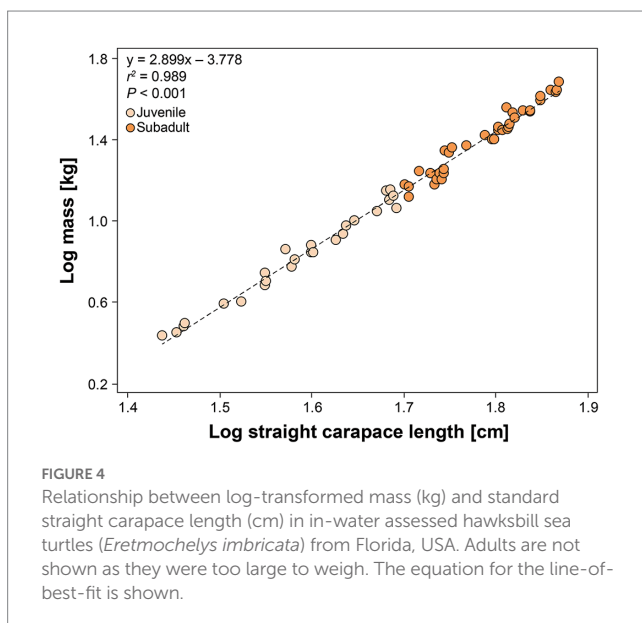


FIGURE 4
Relationship between log-transformed mass (kg) and standard straight carapace length (cm) in in-water assessed hawksbill sea turtles (*Eretmochelys imbricata*) from Florida, USA. Adults are not shown as they were too large to weigh. The equation for the line-of-best-fit is shown.

and cholesterol ($r_s = 0.288$; $p = 0.017$; $n = 68$), GLDH and AST ($r_s = 0.709$; $p < 0.001$; $n = 35$), magnesium and phosphorus ($r_s = 0.369$; $p = 0.002$; $n = 68$), uric acid and phosphorus ($r_s = 0.369$; $p = 0.002$; $n = 68$), uric acid and magnesium ($r_s = 0.325$; $p = 0.007$; $n = 68$), uric acid and sodium ($r_s = 0.328$; $p = 0.007$; $n = 68$), triglycerides and GGT ($r_s = 0.318$; $p = 0.008$; $n = 68$), and triglycerides and glucose ($r_s = 0.359$; $p = 0.003$; $n = 68$). There were no correlations between CK and AST ($r_s = 0.117$; $p = 0.340$; $n = 68$) or CK and phosphorus ($r_s = 0.158$; $p = 0.197$; $n = 68$).

Blood film review data (Table 5) showed that polychromasia and anisocytosis significantly differed by life-stage class ($H(2) = 6.347$;

$p = 0.042$ for both categories), with subadults having a lower occurrence of both in comparison to juveniles ($p = 0.035$ for both categories). Basophilic stippling also significantly differed by life-stage class ($H(2) = 7.406$; $p = 0.025$), with subadults having a lower occurrence in comparison to juveniles ($p = 0.020$). Lastly, number of immature RBCs/100 mature RBCs significantly differed by life-stage class ($H(2) = 7.209$; $p = 0.027$), with subadults having a lower occurrence of immature RBCs in comparison to juveniles ($p = 0.033$).

4.3. Influence of straight carapace length, body condition, and water temperature upon capture on blood analytes

SCL showed significant positive relationships with PCV, eosinophils, AST, phosphorus, cholesterol, GLDH, total protein, albumin, and globulins and significant negative relationships with ALP, CK, calcium, calcium:phosphorus ratio, and glucose. BCI showed a significant negative relationship with PCV. Water temperature showed significant positive relationships with AST, cholesterol, GGT, glucose, lipase, triglycerides, and albumin. Complete statistical results are shown in Table 6.

5. Discussion

This study reports hematology and plasma biochemical blood analyte data for wild-caught hawksbills during four years of field work in the Atlantic Ocean, and evaluated influences of intrinsic (length, body depth, BCI) and extrinsic factors (water temperature), with life-stage class being identified as the most relevant factor for the observed blood analyte differences. The data herein provide an important baseline framework and insight into the physiology of the critically endangered hawksbill sea turtle from the reef system of southeastern Florida during a defined time frame. This work will provide a springboard for future assessments on fitness and stressor effects on this population and allows for comparisons to other geographical populations.

5.1. Physical examination and morphometrics

A thorough external physical examination is an essential component of sea turtle health assessment studies as it provides important information on body condition, activity level, mentation, physical abnormalities, and evidence of trauma or disease (Deem and Harris, 2017; Tristan and Norton, 2017; Page-Karjian and Perrault, 2021). These findings are endpoints needed for defining inclusion criteria in wild animal studies. In fact, some health assessment studies are solely based on physical examination and morphometric measurements (Maulida et al., 2017). All hawksbills captured during the study time frame fit the inclusion criteria for the study. The lack of overt injuries in hawksbills leading to exclusion of some of the study animals contrasts with other sea turtle species. For example, two of 36 immature Kemp's ridleys (*Lepidochelys kempii*) from Georgia, USA were excluded due to abnormal shell formation or monofilament entanglement (Perrault et al., 2020).

TABLE 2 Measures of central tendency, range, and reference intervals (with 90% confidence intervals for upper and lower limits) for hematological and plasma biochemical data (including protein electrophoresis) in Standard International units for in-water, juvenile (25–49.9cm standard straight carapace length) hawksbill sea turtles (*Eretmochelys imbricata*) from Florida, USA.

Analyte	Mean±SD	Median	Range	n	RI	LRL 90% CI	URL 90% CI
Hematology							
Packed cell volume [L/L]	0.30 ± 0.05	0.30	0.22–0.38	11	–	–	–
Immature RBC/100 mature RBC	6 ± 2	7	3–9	8	–	–	–
White blood cells [x 10 ⁹ /L]	8.35 ± 2.60	8.10	4.80–12.10	8	–	–	–
Heterophils [x 10 ⁹ /L]	4.33 ± 1.82	4.30	1.60–6.80	8	–	–	–
Immature heterophils [x 10 ⁹ /L]	0	0	0	8	–	–	–
Lymphocytes [x 10 ⁹ /L]	3.36 ± 0.78	3.00	2.40–4.40	8	–	–	–
Heterophil:lymphocyte ratio	1.26 ± 0.37	1.34	0.57–1.61	8	–	–	–
Monocytes [x 10 ⁹ /L]	0.53 ± 0.24	0.55	0.19–0.82	8	–	–	–
Eosinophils [x 10 ⁹ /L]	0.14 ± 0.13	0.17	0–0.31	8	–	–	–
Basophils [x 10 ⁹ /L]	0.04 ± 0.05	0.03	0–0.12	8	–	–	–
Biochemistry							
Alkaline phosphatase [µkat/L]	0.85 ± 0.28	0.85	0.32–1.39	26	0.38–1.34	0.23–0.55	1.17–1.49
Amylase [µkat/L]	11.59 ± 4.41	11.91	4.24–18.45	26	4.33–18.85	1.84–6.83	16.35–21.34
Aspartate aminotransferase [µkat/L]	2.37 ± 1.09	2.10	1.24–6.05	26	1.17–4.11 ^c	0.94–1.45 ^c	3.31–5.09 ^c
Bile acids [µmol/L]	–	2.1	<0.5–5.0 ^a	10	–	–	–
Blood urea nitrogen [mmol/L]	31.0 ± 4.9	30.0	21.8–41.4	26	23.0–39.0	20.2–25.7	36.2–41.7
Calcium [mmol/L]	2.4 ± 0.3	2.4	1.7–3.2	26	1.9–2.9	1.7–2.0	2.7–3.1
Phosphorus [mmol/L]	1.9 ± 0.3	2.0	1.4–2.5	26	1.5–2.4	1.3–1.6	2.2–2.6
Calcium:phosphorus ratio	1.26 ± 0.30	1.20	0.85–1.98	26	0.77–1.76	0.60–0.94	1.59–1.92
Chloride [mmol/L]	123 ± 4	123	115–134	26	116–130	114–118	128–132
Cholesterol [mmol/L]	–	1.63	<1.17–3.81	26	<1.17–3.08	<1.17	2.64–3.52
Creatine phosphokinase [µkat/L]	23.53 ± 15.25	22.34	5.93–88.84 ^b	26	8.45–33.38	4.07–12.83	29.01–37.76
Gamma-glutamyl transferase [µkat/L]	–	<0.08	<0.08–0.35	26	– ^d	– ^d	– ^d
Glucose [mmol/L]	6.0 ± 0.9	6.0	3.9–7.7	26	4.6–7.4	4.1–5.1	6.9–7.9
Glutamate dehydrogenase [µkat/L]	1.05 ± 0.57	0.83	0.47–2.13	10	–	–	–
Lipase [µkat/L]	–	0.35	<0.02–1.22	26	<0.02–1.20 ^f	<0.02–0.06 ^f	0.82–1.69 ^f
Magnesium [mmol/L]	3.6 ± 0.5	3.5	3.0–4.8	26	2.9–4.3 ^e	2.8–3.2 ^e	4.1–4.7 ^e
Potassium [mmol/L]	4.6 ± 0.5	4.6	3.8–5.5	26	3.7–5.5	3.4–4.0	5.2–5.8
Sodium [mmol/L]	158 ± 5	159	148–167	26	151–165	148–153	163–168
Triglycerides [mmol/L]	1.22 ± 0.43	1.17	0.49–2.15	26	0.51–1.94	0.27–0.76	1.69–2.18
Uric acid [mmol/L]	–	0.04	<0.01–0.05	26	0.01–0.05 ^e	<0.01–0.02 ^e	0.05–0.06 ^e
Protein electrophoresis							
Total protein [g/L]	33 ± 6	32	22–43	26	23–43	20–26	39–46
Albumin [g/L]	12.9 ± 2.0	13.1	7.0 ^c –17.3	26	10.0–16.3	9.1–10.9	15.4–17.3
Total globulins [g/L]	19.4 ± 4.9	18.9	11.4–27.2	26	9.8–28.9	7.1–12.6	26.2–31.7
Albumin:globulin ratio	0.73 ± 0.22	0.66	0.45–1.27	26	0.41–1.20 ^e	0.35–0.48	1.03–1.40

^a5.0 µmol/L was an outlier. The next highest value was 3.4 µmol/L.

^b88.84 µkat/L was an outlier and was removed from calculation of reference intervals. The next highest value was 31.26 µkat/L.

^c7.0 g/L was an outlier and was removed from calculation of reference intervals. The next lowest value was 10.5 g/L.

^dReference intervals for gamma glutamyl transferase could not be calculated as the majority of values (54%) fell below detection limits.

^eReference intervals were calculated using logarithmic transformations, as data were non-normal.

^fReference intervals were calculated using Box-Cox transformations, as data were non-normal.

For analytes with n < 20, values are reported descriptively, as reference intervals cannot be calculated for sample sizes < 20. For analytes with values below the limits of detection, only median and range are reported; values of half of the detection limit were then assigned to generate reference intervals. Parametric methods for sample sizes ≥ 20, but < 40 were used to calculate reference intervals (Friedrichs et al., 2012), unless otherwise indicated in the footnotes. Normality was assessed using the Shapiro-Wilk test, while outliers were detected using the Dixon-Reed test. Three plasma samples had mild hemolysis (1+), while one sample had mild lipemia (1+), which are not considered to cause interference using dry chemistry analysis (Andreasen et al., 1997; Stacy and Innis, 2017; Stacy et al., 2019). CI, confidence interval; LRL, lower reference limit; RI, reference interval; SD, standard deviation; URL, upper reference limit.

BCI of all hawksbills in this study ranged from 0.94–1.39 (mean ± SD: 1.13 ± 0.10), which is similar to two aggregations of hawksbills from Puerto Rico (means of 1.16 and 1.18) (Diez and van Dam, 2002) and the Bahamas (mean ± SD: 1.17 ± 0.08; range: 1.05–1.41) (Bjorndal and Bolten, 2010), but higher than hawksbills from

Indonesia (mean ± SD: 1.06 ± 0.07; range: 0.92–1.14) (Maulida et al., 2017), and lower than hawksbills from the Monito cliff wall in Puerto Rico (mean: 1.24) (Diez and van Dam, 2002) and the Cayman Islands (Little Cayman mean ± SD: 1.25 ± 0.17; Grand Cayman mean ± SD: 1.24 ± 0.18) (Blumenthal et al., 2009). Norton

TABLE 3 Measures of central tendency, range, and reference intervals (with 90% confidence intervals for upper and lower limits) for hematological and plasma biochemical data (including protein electrophoresis) in Standard International units for in-water, subadult (50–78cm standard straight carapace length) hawksbill sea turtles (*Eretmochelys imbricata*) from Florida, USA.

Analyte	Mean ± SD	Median	Range	n	RI	LRL 90% CI	URL 90% CI
Hematology							
Packed cell volume [L/L]	0.38 ± 0.05	0.38	0.27–0.48	27	0.30–0.46	0.27–0.33	0.43–0.49
Immature RBC/100 mature RBC	3 ± 3	2	0–11	16	–	–	–
White blood cells [×10 ⁹ /L]	8.42 ± 1.94	8.15	5.30–12.30	16	–	–	–
Heterophils [×10 ⁹ /L]	3.91 ± 0.98	3.80	2.50–6.40 ^a	16	–	–	–
Immature heterophils [×10 ⁹ /L]	0	0	0	16	–	–	–
Lymphocytes [×10 ⁹ /L]	3.63 ± 1.19	3.55	1.50–6.00	16	–	–	–
Heterophil:lymphocyte ratio	1.18 ± 0.49	1.04	0.58–2.73 ^b	16	–	–	–
Monocytes [×10 ⁹ /L]	0.50 ± 0.27	0.51	0.12–1.20 ^c	16	–	–	–
Eosinophils [×10 ⁹ /L]	0.34 ± 0.14	0.34	0.05–0.62	16	–	–	–
Basophils [×10 ⁹ /L]	0.08 ± 0.13	0	0–0.44	16	–	–	–
Biochemistry							
Alkaline phosphatase [μkat/L]	0.70 ± 0.22	0.68	0.25–1.29	39	0.35–1.05	0.25–0.45	0.95–1.15
Amylase [μkat/L]	13.73 ± 4.17	12.94	8.73–34.05 ^d	39	9.03–17.35	7.85–10.20	16.17–18.52
Aspartate aminotransferase [μkat/L]	2.32 ± 0.92	1.97	1.05–5.43	39	1.10–3.59 ^{e,i}	0.97–1.30 ^{e,i}	2.99–4.29 ^{e,i}
Bile acids [μmol/L]	–	2.3	<0.5–9.2 ^f	21	<0.5–4.9 ^j	<0.5–0.9 ^j	3.8–6.2 ^j
Blood urea nitrogen [mmol/L]	30.3 ± 6.6	31.4	10.7–40.7	39	19.1–38.5 ⁱ	11.4–23.6 ⁱ	36.7–40.2 ^j
Calcium [mmol/L]	2.2 ± 0.3	2.2	1.8–3.1	39	1.9–2.7 ⁱ	1.8–2.0 ⁱ	2.5–2.8 ⁱ
Phosphorus [mmol/L]	2.2 ± 0.3	2.1	1.5–3.0	39	1.6–2.7	1.5–1.8	2.5–2.8
Calcium:phosphorus ratio	1.05 ± 0.19	1.01	0.71–1.53	39	0.73–1.37	0.64–0.82	1.28–1.46
Chloride [mmol/L]	122 ± 5	122	112–134	39	113–131	111–116	128–133
Cholesterol [mmol/L]	–	1.99	<1.17–5.59	39	<1.17–3.53 ^e	<1.17 ^e	2.95–4.02 ^e
Creatine phosphokinase [μkat/L]	19.12 ± 11.94	18.02	3.62–80.74 ^g	39	6.95–28.06	3.96–9.94	25.07–31.05
Gamma glutamyl transferase [μkat/L]	–	<0.07	<0.08–0.50 ^h	39	<0.08–0.18 ^e	<0.08 ^e	0.15–0.23 ^e
Glucose [mmol/L]	5.2 ± 0.6	5.2	4.0–7.1	39	4.4–6.2 ⁱ	4.2–4.6 ⁱ	5.9–6.5 ⁱ
Glutamate dehydrogenase [μkat/L]	1.98 ± 1.55	1.16	0.45–5.69	23	0.48–4.94 ⁱ	0.31–0.74 ⁱ	3.22–7.57 ⁱ
Lipase [μkat/L]	–	0.17	<0.02–0.80	39	0.03–0.53 ^j	0.02–0.05 ^j	0.40–0.68 ^j
Magnesium [mmol/L]	3.7 ± 0.4	3.7	2.5–4.6	39	3.0–4.4	2.8–3.2	4.2–4.6
Potassium [mmol/L]	4.4 ± 0.5	4.5	3.7–5.4	39	3.7–5.2	3.5–3.9	5.0–5.4
Sodium [mmol/L]	157 ± 5	156	148–178	39	147–165 ^e	144–151 ^e	162–168 ^e
Triglycerides [mmol/L]	0.97 ± 0.40	0.89	0.36–2.21	39	0.32–1.62	0.14–0.51	1.44–1.81
Uric acid [mmol/L]	–	0.04	<0.01–0.08	39	0.01–0.07 ^e	<0.01–0.02 ^e	0.06–0.07 ^e
Protein electrophoresis							
Total protein [g/L]	36 ± 4	36	27–43	39	29–43	27–31	41–45
Albumin [g/L]	14.7 ± 1.9	14.0	11.8–20.3	39	11.0–18.4	10.1–11.9	17.6–19.3
Total globulins [g/L]	20.8 ± 3.4	20.1	13.8–28.5	39	14.2–27.4	12.6–15.7	25.9–29.0
Albumin:globulin ratio	0.75 ± 0.15	0.74	0.47–1.25	39	0.52–1.06 ^h	0.47–0.56	0.98–1.16

(Continued)

TABLE 3 (Continued)

* 6.4×10^9 cells/L was an outlier. The next highest value was 5.1×10^9 cells/L.

^b2.73 was an outlier. The next highest value was 1.50.

^c 1.20×10^9 cells/L was an outlier. The next highest value was 0.78×10^9 cells/L.

^d34.05 μ kat/L was an outlier and was removed from calculation of reference intervals. The next highest value was 19.79 μ kat/L.

^eData for aspartate aminotransferase, cholesterol, gamma glutamyl transferase, sodium, and uric acid could not be normalized; therefore, the robust method was used to generate reference intervals.

^f9.2 μ mol/L was an outlier and was removed from calculation of reference intervals. The next highest value was 5.2 μ mol/L.

^g80.74 μ kat/L was an outlier and was removed from calculation of reference intervals. The next highest value was 31.76 μ kat/L.

^h0.50 μ kat/L was an outlier and was removed from calculation of reference intervals. The next highest value was 0.32 μ kat/L.

ⁱReference intervals were calculated using logarithmic transformations, as data were non-normal.

^jReference intervals were calculated using Box-Cox transformations, as data were non-normal.

For analytes with $n < 20$, values are reported descriptively, as reference intervals cannot be calculated for sample sizes < 20 . For analytes with values below the limits of detection, only median and range are reported; values of half of the detection limit were then assigned to generate reference intervals. Parametric methods for sample sizes ≥ 20 but < 40 were used to calculate reference intervals (Friedrichs et al., 2012), unless otherwise indicated in the footnotes. Normality was assessed using the Shapiro-Wilk test, while outliers were detected using the Dixon-Reed test. Three plasma samples had mild hemolysis (1+), while one sample had mild lipemia (1+), which are not considered to cause interference using dry chemistry analysis (Andreassen et al., 1997; Stacy and Innis, 2017; Stacy et al., 2019). CI, confidence interval; LRL, lower reference limit; RI, reference interval; SD, standard deviation; URL, upper reference limit.

and Wyneken (2015) suggest that BCI scores of < 1.00 are indicative of emaciation in sea turtles, and since all turtles in the present study were considered robust based on visual evaluation, BCI scores should be specifically developed for individual sea turtle species, geographical locations/populations, and life-stage classes (Nishizawa and Joseph, 2022).

There is evidence that hawksbill turtles migrate during their development and growth from small juveniles in the Florida Keys along the Southeast Florida Continental Reef Tract to subadults in their northern range in Palm Beach County, Florida (Wood et al., 2013). The observed statistically significant difference between capture sites based on SCL and thus differentiating juveniles mainly captured in Monroe County vs. subadult and adult turtles mainly captured at Palm Beach County supports this further. The very small overlap of SCL (Figure 3) between juveniles and subadults at both capture sites also substantiates that secondary recruitment of juveniles from the Keys to the north occurs in this population.

BCI was significantly higher in hawksbills from Monroe County compared to those from Palm Beach County, despite turtles from Palm Beach County having higher SCL, mass, and body depth. These trends are driven by life-stage class distribution between the two sites, as juveniles had a significantly higher BCI than subadults, and more juveniles (23 of 28 total turtle captures; 82%) were caught in Monroe compared to Palm Beach County (3 of 40 total turtle captures; 8%). These morphometrical data trends in hawksbills suggest fast somatic growth rates in juvenile hawksbills since smaller juveniles had higher BCI potentially indicating greater food intake. A comparison of forage items of the two Florida hawksbill aggregations from this study has not been reported, therefore it is unknown if dietary or habitat-related differences between the Palm Beach County and Monroe County sites have influenced the results. Differences in BCI by capture locations and by carapace length have previously been reported in green turtles (*Chelonia mydas*) (Diez and van Dam, 2002; Peig and Green, 2010; Lamont and Johnson, 2021). It is likely that dissimilarities in BCI of sea turtles are mainly driven by differences in environmental conditions, population density, disease prevalence, and/or forage availability (Bjorndal et al., 2000; Diez and van Dam, 2002; Labrada-Martagón et al., 2010; Rossi et al., 2019; Lamont and Johnson, 2021).

5.2. Hematology

PCV is considered an indicator of overall fitness (Stamper et al., 2005). Based on data from captive animals, Frair (1977) suggested that hawksbills have an overall lower PCV than other sea turtle species (range: 0.17–0.42 L/L) and attributed this to dissimilarities in size and growth. In foraging hawksbills from Australia and nesting hawksbills from Brazil, PCV ranged from 0.12–0.41 L/L and 0.34–0.40, respectively (Goldberg et al., 2013; Whiting et al., 2014). The PCV of all hawksbills in the current study ranged from 0.22–0.48 L/L. Mild anemia in sea turtles is defined as PCV ranging from 0.19–0.25 L/L, with severe anemia occurring in patients with PCV ≤ 0.12 L/L (Stacy and Innis, 2017); therefore, it seems that wild-caught juvenile hawksbills may have naturally occurring lower PCV (Frair, 1977; Whiting et al., 2014), under due consideration that lymph contamination during blood sampling cannot always be reliably excluded in sea turtles (Stacy and Innis, 2017). Of note, the methodology of obtaining PCV data should be considered, as some studies report discrepancies between hematocrit (Hct) data (using automated analyzers) and the manually obtained PCV (*via* spun capillary tubes), with Hct data being considered inaccurate (i.e., lower) *via* automated methods in sea turtles (Muñoz-Pérez et al., 2017; Stacy and Innis, 2017).

While the ranges of PCV established in this and other studies overlap, it also appears that there may be life-stage class differences in this species. For example, we observed a strong positive relationship between PCV and SCL in Florida hawksbills, indicating that RBCs presumably increase in length, width, and volume as turtles mature and oxygen demands increase with longer dive times as described in other studies (Frair, 1977; Stamper et al., 2005; Perrault et al., 2016; Stacy et al., 2018). This trend of increasing PCV with turtle size has been documented in six of the seven sea turtle species (Frair, 1977; Wood and Ebanks, 1984; Casal et al., 2009; Rousselet et al., 2013; Perrault et al., 2016; Stacy et al., 2018).

In comparison to subadults, juvenile hawksbills had higher incidences of RBC polychromasia (i.e., increased basophilic color), anisocytosis (i.e., variation in RBC size), and basophilic stippling (i.e., punctate basophilic cytoplasmic inclusions), and a greater proportion of immature RBC in relation to total mature RBC. These morphological features of RBC are often associated with active erythroid production (Stacy et al., 2011), an observation also described in younger turtles of other sea turtle species and reptiles in general (Fleming et al., 2020; Perrault et al., 2020; Stacy and Harr, 2020; Perrault et al., 2022).

TABLE 4 Measures of central tendency and range for hematological and plasma biochemical data (including protein electrophoresis) in Standard International units for in-water, adult (>78cm straight carapace length) hawksbill sea turtles (*Eretmochelys imbricata*) from Florida, USA.

Analyte	Mean \pm SD	Median	Range	n
Hematology				
Packed cell volume [L/L]	0.36 \pm 0.02	0.36	0.34–0.38	3
Immature RBC/100 mature RBC	2 \pm 2	3	0–4	3
White blood cells [$\times 10^9$ /L]	6.67 \pm 2.15	5.90	5.00–9.10	3
Heterophils [$\times 10^9$ /L]	3.40 \pm 0.75	3.30	2.70–4.20	3
Immature heterophils [$\times 10^9$ /L]	0	0	0	3
Lymphocytes [$\times 10^9$ /L]	2.60 \pm 0.95	2.10	2.00–3.70	3
Heterophil:lymphocyte ratio	1.35 \pm 0.22	1.35	1.14–1.57	3
Monocytes [$\times 10^9$ /L]	0.54 \pm 0.41	0.41	0.20–1.00	3
Eosinophils [$\times 10^9$ /L]	0.20 \pm 0.08	0.20	0.12–0.27	3
Basophils [$\times 10^9$ /L]	0	0	0	3
Biochemistry				
Alkaline phosphatase [μ kat/L]	0.53 \pm 0.20	0.65	0.30–0.65	3
Amylase [μ kat/L]	18.12 \pm 2.69	17.54	15.78–21.06	3
Aspartate aminotransferase [μ kat/L]	1.89 \pm 0.42	2.10	1.40–2.17	3
Bile acids [μ mol/L]	–	<0.5	<0.5–2.4	2
Blood urea nitrogen [mmol/L]	27.7 \pm 8.4	27.1	19.6–36.4	3
Calcium [mmol/L]	2.0 \pm 0.3	2.1	1.7–2.3	3
Phosphorus [mmol/L]	2.0 \pm 0.03	2.0	1.9–2.0	3
Calcium:phosphorus ratio	1.03 \pm 0.17	1.07	0.85–1.19	3
Chloride [mmol/L]	122 \pm 4	120	119–127	3
Cholesterol [mmol/L]	2.50 \pm 0.68	2.62	1.76–3.11	3
Creatine phosphokinase [μ kat/L]	10.62 \pm 2.72	9.55	8.60–13.71	3
Gamma-glutamyl transferase [μ kat/L]	–	<0.08	<0.08–0.45	3
Glucose [mmol/L]	5.55 \pm 0.60	5.38	5.05–6.22	3
Glutamate dehydrogenase [μ kat/L]	0.52 \pm 0.03	0.52	0.50–0.54	2
Lipase [μ kat/L]	–	0.05	<0.02–0.10	3
Magnesium [mmol/L]	3.2 \pm 0.7	3.4	2.4–3.7	3
Potassium [mmol/L]	4.3 \pm 0.3	4.2	4.1–4.7	3
Sodium [mmol/L]	154 \pm 3	153	152–158	3
Triglycerides [mmol/L]	1.62 \pm 0.86	1.23	1.03–2.61	3
Uric acid [mmol/L]	0.02 \pm 0.01	0.03	0.01–0.03	3
Protein electrophoresis				
Total protein [g/L]	39 \pm 3	39	36–41	3
Albumin [g/L]	16.4 \pm 0.8	16.1	15.8–17.3	3
Total globulins [g/L]	21.4 \pm 2.3	20.6	19.6–23.9	3
Albumin:globulin ratio	0.82 \pm 0.09	0.84	0.72–0.89	3

For analytes with values below the limits of detection, only median and range are reported. One plasma sample had mild hemolysis (1+), which is not considered to cause interference using dry chemistry analysis (Andreasen et al., 1997; Stacy and Innis, 2017; Stacy et al., 2019). SD, standard deviation.

Eosinophils in this study made up a small proportion (range: 0–8%) of the overall leukogram ($<0.70 \times 10^9$ cells/L), which is much lower than in similarly sized wild-caught hawksbills from the Galápagos (Muñoz-Pérez et al., 2017). We also observed a significant positive relationship between SCL and absolute eosinophils. While the exact

function of reptilian eosinophils has not been established, they are thought to aid in immune stimulation and phagocytosis of parasites (Stacy et al., 2011; Rousselet et al., 2013; Stacy and Innis, 2017; Stacy and Harr, 2020). Therefore, the observed correlation may be due to antigenic stimulation or influences from environmental factors (Stacy

et al., 2011), with increasing antigenic stimulation as turtles grow and age (e.g., increased incidence of spirorchid infection in larger turtles) (Deem et al., 2006; Casal et al., 2009; Innis et al., 2010).

5.3. Electrolytes and minerals

Although plasma electrolyte and mineral concentrations in hawksbill turtles from this study fell within normal ranges reported for other sea turtle species (Stacy and Innis, 2017), several physiologically relevant relationships with SCL were observed. Similar to juvenile loggerheads (*Caretta caretta*) in North Carolina (Kelly et al., 2015), phosphorus concentrations positively correlated with SCL in hawksbills, suggesting bone and/or somatic growth, given that there were no positive size-relevant correlations with CK, the main enzyme of muscle tissues (Stacy and Innis, 2017). The observed negative correlations of calcium and calcium to phosphorus ratio with SCL suggest differences in somatic growth rates as turtles grow and age (Bolten and Bjørndal, 1992), bone metabolism, or possibly osmoregulation (i.e., associated with changes in salt gland regulation with somatic growth). Hawksbill turtles have an oceanic-neritic development pattern and recruit back to nearshore environments at a small size (20–35 cm carapace length) (Bolten, 2003). Additionally, regional differences are a known major driver for somatic growth in the species (Bjørndal et al., 2016). In contrast to hawksbill SCL, phosphorus in oceanic-juvenile loggerheads correlated negatively (Stacy et al., 2018) and calcium to phosphorus ratio in immature Kemp's ridleys (Perrault et al., 2020) correlated positively with carapace length. Possible considerations for these differences include species, life stage-class, and dietary or habitat variations. The observed positive correlations of magnesium with phosphorus and uric acid with magnesium, phosphorus, and sodium suggest potential clinical utility of these analytes in the assessment of renal function in sea turtles, particularly in the presence of normal CK activities (Stacy and Innis, 2017).

5.4. Tissue enzyme activities

Tissue enzyme activities showed several metabolically important correlations with SCL and water temperature in this study. The positive correlations of AST and GLDH with SCL, with the absence of CK associations that would indicate release from muscle tissue, suggest that larger turtles with higher hepatic volume likely have greater tissue enzyme release into plasma, similar to hawksbills in the Persian Gulf and Brazil (Goldberg et al., 2013; Ehsanpour et al., 2015). The negative correlation of SCL with CK further supports the conclusions on mineral correlations, in that the observed correlations with SCL and calcium and phosphorus suggest bone growth differences in smaller turtles rather than growth of muscle tissue.

The negative correlation of ALP with SCL was unexpected, since ALP activities are known to be higher in younger, growing mammals (Allison, 2022); however, this tissue enzyme is widely distributed in sea turtle tissues and isoenzymatic activities of ALP in bone tissue have not been assessed to date in reptiles (Anderson et al., 2013; Petrosky et al., 2015; Adamovicz et al., 2019). The tissue enzymes AST, GGT, and lipase correlated positively with water temperature, which could be driven by higher physical activity or somatic growth rate in smaller turtles.

TABLE 5 Morphological evaluation of red blood cells (RBC), white blood cells, and thrombocytes for in-water, hawksbill sea turtles (*Eretmochelys imbricata*) of three life-stage classes from Florida, USA.

Morphological finding	Juveniles	Subadults	Adults
Polychromasia*	Mild: 100% (8/8)	Absent: 25% (4/16) Minimal: 31% (5/16) Mild: 44% (7/16)	Absent: 33% (1/3) Mild: 66% (2/3)
Anisocytosis*	Mild: 100% (8/8)	Absent: 25% (4/16) Minimal: 31% (5/16) Mild: 44% (7/16)	Absent: 33% (1/3) Mild: 66% (2/3)
Basophilic stippling*	Mild: 8/8 (100%)	Absent: 19% (3/16) Rare: 6% (1/16) Minimal: 38% (6/16) Mild: 38% (6/16)	Absent: 33% (1/3) Mild: 66% (2/3)
Immature RBC/100 mature RBC*	6 ± 2 (3–9)	3 ± 3 (0–11)	2 ± 2 (0–4)
Erythrocyte morphology	NSCF	NSCF	NSCF
Leukocyte morphology	NSCF	NSCF	NSCF
Thrombocytes	Adequate: 100% (8/8)	Adequate: 100% (16/16)	Adequate: 100% (3/3)

For immature RBC/100 mature RBC, mean ± standard deviation is reported, with the range parenthetically. NSCF, no significant cytological findings. Asterisks denote statistically significant differences between juveniles and subadults at $p < 0.050$.

Activities of GLDH have rarely been reported in sea turtles (March et al., 2018) and the clinical utility of this enzyme is currently unknown (Stacy and Innis, 2017). Green turtles undergoing rehabilitation in Australia showed strong correlations between plasma GLDH and AST activities, suggesting the potential use as indicators for hepatocellular injury (March et al., 2018). This assumption is further supported by the observed correlations between GLDH and AST, but not CK in hawksbill turtles from this study. Additionally, in eastern box turtles (*Terrapene carolina carolina*), GLDH activities were highest in liver, followed by kidney and gall bladder, with activities in skeletal muscle being among the lowest of the ten tissues examined (Adamovicz et al., 2019). The observed positive correlations of GGT with AST and cholesterol, bile acids with AST, and triglycerides with GGT and glucose highlight the potential for the clinical utility of these analytes for the diagnosis of liver disorders in hawksbill turtles. As such, concurrently increased GLDH, GGT, and/or AST activities, increased triglycerides, cholesterol, and/or bile acids, along with normal CK could support the clinical diagnosis of liver abnormalities, as supported by general assumptions in clinical chemistry interpretations in sea turtles (Stacy and Innis, 2017).

5.5. Lipids

Plasma lipid concentrations in sea turtles often differ by life-stage class. Similar to green turtles and loggerheads (Hasbún et al., 1998; Labrada-Martagón et al., 2010; Delgado et al., 2011; Prieto-Torres

TABLE 6 Relationship of blood analytes to straight carapace length (SCL), body condition index (BCI), and water temperature upon capture of in-water, hawksbill sea turtles (*Eretmochelys imbricata*) from Florida, USA.

Analyte	Adj. r^2	N	p	Significant predictor(s)	r	p
Hematology						
Packed cell volume	0.63	36	<0.001	SCL	0.69	<0.001
				BCI	-0.56	<0.001
Eosinophils	0.27	22	0.007	SCL	0.55	0.007
Biochemistry						
Alkaline phosphatase	0.15	62	0.001	SCL	-0.41	0.001
Aspartate aminotransferase ^a	0.06	59	0.035	Water temperature	0.28	0.035
Creatine phosphokinase	0.12	61	0.004	SCL	-0.37	0.004
Calcium	0.15	62	0.001	SCL	-0.40	0.001
Phosphorus	0.07	62	0.022	SCL	0.29	0.022
Calcium:phosphorus ratio ^a	0.18	62	<0.001	SCL	-0.44	<0.001
Cholesterol	0.25	62	<0.001	SCL	0.37	0.003
				Water temperature	0.44	<0.001
Gamma-glutamyl transferase ^a	0.10	60	0.010	Water temperature	0.33	0.010
Glucose	0.39	62	<0.001	SCL	-0.56	<0.001
				Water temperature	0.41	0.001
Glutamate dehydrogenase ^a	0.16	32	0.013	SCL	0.44	0.013
Lipase ^a	0.20	62	0.001	Water temperature	0.43	0.001
Triglycerides	0.23	62	<0.001	Water temperature	0.46	<0.001
Protein electrophoresis						
Total protein	0.20	62	<0.001	SCL	0.46	<0.001
Albumin	0.27	62	<0.001	SCL	0.51	<0.001
				Water temperature	0.28	0.027
Globulins	0.09	62	0.011	SCL	0.32	0.011

^aLog-transformed.

Stepwise backward multiple regressions were employed. Results of significant predictors are shown. adj, adjusted.

et al., 2013; Rousset et al., 2013), positive correlations of plasma cholesterol and SCL were also observed in hawksbill turtles from this study. These observations suggest BCI differences due to life-stage class as observed in leatherbacks (*Dermochelys coriacea*), whereby foraging individuals had higher plasma lipids compared to nesting turtles (Harris et al., 2011), likely due to little or lack of foraging during the nesting season (Perrault et al., 2014).

5.6. Glucose

Glucose concentrations in hawksbills from Florida were similar in range to juveniles after successful rehabilitation and in adult nesting hawksbills (Caliendo et al., 2010; Goldberg et al., 2013), and were negatively correlated with SCL, a trend that has also been observed in loggerheads in captive care (Kakizoe et al., 2007; Rousset et al., 2013). With the additional positive correlation with water temperature, considerations for higher plasma glucose in smaller turtles include higher activity (i.e., in warmer waters), comparatively increased stress level during capture and handling (i.e., smaller turtles often require a short duration chase before capture that is not typically required in larger

turtles), or, given similar trends in captive loggerheads, differences in metabolic rates between life-stage classes (Goldberg et al., 2013; Rousset et al., 2013). Specifically for water temperature, possible higher carbohydrate consumption during periods of higher temperatures is presumed to result in higher metabolic rate, as described in green turtles from Mexico and loggerheads from North Carolina, USA (Stamper et al., 2005; Labrada-Martagón et al., 2010; Kelly et al., 2015).

5.7. Proteins

Similar to other sea turtles, we found that SCL positively correlated with total protein, albumin, and globulins (Hasbún et al., 1998; Osborne et al., 2010; Delgado et al., 2011; Whiting et al., 2014; Kelly et al., 2015; Stacy et al., 2018; Perrault et al., 2020), trends that can be explained by nutritional, seasonal, and/or environmental differences (Frair and Shah, 1982), or as a result of immune system maturation due to antigenic stimulation as turtles grow and age (Innis et al., 2010; Osborne et al., 2010; Perrault et al., 2014). Larger hawksbills foraging in Honduras preferred poriferan prey (e.g., *Meloplhus ruber*), with smaller individuals foraging mainly on algae

(Berube et al., 2012). Forage items of the two hawksbill aggregations in Florida from this study have not been reported and could also influence size-relevant correlations with plasma protein concentrations. Lastly, we found that temperature and albumin were positively correlated in hawksbills from this study possibly due to higher activity levels and increased food consumption during warming periods, a trend that has also been observed in Blanding's turtles (*Emydoidea blandingii*) from Illinois (Andersson et al., 2021).

6. Conclusion

This study reports blood analyte data from the critically endangered hawksbill sea turtle encompassing three life-stage classes inhabiting the Florida reef system and gives insight into basic physiological metrics for this population during the study period of 2017–2020. With the defined inclusion criteria used herein, the population was assessed as “clinically normal,” thus offering to identify various aspects of sea turtle biology and physiology that are influenced by extrinsic and intrinsic factors. These data will support answering population-driven questions (e.g., future spatio-temporal comparisons, conservation management) in addition to their utility for individual animals (e.g., stranding, rehabilitation).

Data availability statement

The original contributions presented in the study are included in the article/[Supplementary materials](#). Further inquiries can be directed to the corresponding author.

Ethics statement

This study was reviewed and authorized by the National Marine Fisheries Service (NMFS) [Permit #22988], Florida Fish and Wildlife Conservation Commission (FWC) [Marine Turtle Permits #021 and #077], Florida Keys National Marine Sanctuary [Research Permit #175], and University of Florida's Institutional Animal Care and Use Committee (IACUC) [#201706823]. All handling and sampling procedures of sea turtles were performed according to NMFS and FWC regulations.

Author contributions

NIS, JRP, and LDW conceptualized the study and wrote the original manuscript. LDW led all fieldwork and sample collection.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Supplementary material

The Supplementary material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fevo.2023.1199688/full#supplementary-material>

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