



Wind Drifting vs. Pulsating Swimming Jellyfish: Respiratory Metabolism and Composition Differences in *Physalis physalis*, *Velella velella*, *Aurelia aurita*, and *Pelagia noctiluca*

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Physalia physalis and *Velella velella*, are among the few marine organisms that harness the wind for their locomotion, whereas other cnidarian jellyfish make use of their pulsating bell-shaped bodies to propel themselves through the seas. We investigate their composition and metabolism compared with two species of pulsating scyphozoan jellyfish, *Aurelia aurita* and *Pelagia noctiluca*. Protein (P), lipid (L), carbohydrate (K), and derived energy content (Ec), provided information on the biochemical composition of these species and their relevance as prey. Physiological respiration (R) from oxygen consumption. As well as potential respiration (Φ) from the electron transport system (ETS) activity and the derived respiratory carbon demand (RCD) and heterotrophic energy transformation (HET), allow the comparison of the impact of these two types of propulsion on the metabolism, along with the impact of these organisms as predators. In this study it was found that these hydrozoans depicted a different biochemical composition relative to other gelatinous zooplankton. Lower water content at around 90% was observed, while WM-specific P, L, K, and Ec were higher, showcasing new aspects of these species as prey. The lower R/P in *P. physalis* and *V. velella* (1.8 ± 0.7 and $2.9 \pm 1.1 \mu\text{L O}_2 \text{ h}^{-1} \text{ mg Prot}^{-1}$, respectively) and the low R/ Φ , around 0.1, indicate lower respiration in wind-driven propulsion compared to pulsation-driven propulsion. Additionally, these results encourage the use and research on enzymatic techniques that are particularly useful for gelatinous research, and the calculation of RCD and HET helps in understanding the physiology and role played by the organisms as predators from carbon and energy perspectives.

Keywords: hydrozoa, scyphozoa, neuston, oxygen consumption, ETS, respiration, jellyfish locomotion

Abbreviations: AE, Assimilation efficiency; ATP, Adenosine triphosphate; C, Carbon; DM, Dry mass; Ec, Energy content; ETS, Electron transport system; F, C:CO₂ conversion; HET, Heterotrophic energy transformation; INT, 2-p-iodophenyl-3-p-nitrophenyl-5-phenyltetrazolium chloride; K, Carbohydrate; L, Lipid; NADH, Nicotinamide adenine dinucleotide reduced; NADPH, Nicotinamide adenine dinucleotide phosphate reduced; Oe, Number of electron pairs needed to reduce one mole of O₂; P, Protein; p-HET, Potential HET; PO, The modified P/O ratio, ATP produced per O; R, Aerobic respiration; RCD, Respiratory carbon demand; RQ, unitless ratio of CO₂ to O₂ during R; t, Number of hours in a day; WM, Wet mass; ΔG_{ATP} , Gibbs free energy associated with ATP; Φ , Potential respiration.

INTRODUCTION

There is a growing interest in gelatinous zooplankton in ocean ecosystems (Chiaverano et al., 2018; Ruzicka et al., 2020; Wright et al., 2021). It is known that their role as predator and prey extends from the deep ocean (Choy et al., 2017) to the neuston (Bieri, 1966). The neuston, also known as pleuston, is the ecosystem located on the ocean surface, comprising of a unique community of floating organisms, such as *Sargassum*-type seaweeds, wind-propelled cnidarians, and other floating invertebrates. This highly dynamic surface layer fosters the exploration of various lifestyles and locomotion strategies and connects disparate atmospheric and oceanic habitats (Helm, 2021). However, it is also, heavily impacted by human waste (Egger et al., 2021).

Research conducted in various ecosystems has demonstrated jellyfish consumption of fish eggs, fish larvae, other fish life-stages, multiple zooplankton species, and other members of pelagic environments (Bieri, 1966; Purcell, 1984, 1997, 2003; Purcell and Arai, 2001; Hansson et al., 2005; Boero, 2013; Purcell et al., 2015; Choy et al., 2017; Zeman et al., 2018). Jellyfish have been described as the main competitor of zooplanktivorous fish (Pauly et al., 2009), thereby known as predators as well as competitors of several commercial fish species. Substantial efforts have been made to quantify and assess the impact of this consumption (Larson, 1987; Morand et al., 1987; Schneider, 1989; Purcell, 1997; Uye and Shimauchi, 2005; Ishii and Tanaka, 2006; Han et al., 2012; Iguchi et al., 2017; Nagata and Morandini, 2018). Furthermore, several studies have investigated the formation of dense aggregations by some jellyfish and their impact on prey populations (Malej, 1989a; Mills, 1995; Schneider and Behrends, 1998; Hansson et al., 2005; Ishii and Tanaka, 2006; Condon et al., 2011; Boero, 2013; Iguchi et al., 2017; Schiariti et al., 2018). Quantifying consumption rates of these organisms will aid in the assessment of their effect on the recovery of populations in areas where jellyfish are proliferating (Purcell et al., 2007; Richardson et al., 2009; Boero, 2013). Carbon demand, associated with the respiratory activity can be calculated from the respiration measurements, which can help in the estimation of the impact on prey populations (Purcell et al., 2010; Lilley et al., 2014; Iguchi et al., 2017).

However, the trophic importance of gelatinous zooplankton as prey is presently experiencing a paradigm shift (Hays et al., 2018). In some cases, the energy content in jellyfish is insufficient to explain the predation observed. It is hypothesized that this may occur due to their high water content (Thiebot and McInnes, 2020). Yet, multiple organisms predate on them, such as turtles, nudibranchs, cirripeds, crustaceans, birds, cephalopods, sharks and fish (especially sunfish), which have been known to associate with cnidarian jellyfish by either living symbiotically, feeding upon them, or both (Bieri, 1966; Jenkins, 1983; Arai, 1988, 2005; Frick et al., 2009; Phillips et al., 2017; Griffin et al., 2019; Thiebot and McInnes, 2020). This begs further research on capture practices, opportunism, and self-medication using jellyfish. Hays et al. (2018) and Thiebot and McInnes (2020) agree that further studies on the gelatinous prey's dietary "value" are needed.

Jellyfish, or planktonic cnidarians, are usually envisioned as umbrella-shaped (bell), gelatinous organisms, such as *Aurelia aurita* or *Pelagia noctiluca* that efficiently propel themselves across the ocean by contracting their bells (Gemmell et al., 2014, 2020; Hoover and Miller, 2015; Costello et al., 2021). Yet, their hydrozoan relatives, *Physalia physalis* and *Velella velella*, use their pneumatophore to exploit the wind for their locomotion with a much higher propulsion efficiency (Ferrer and González, 2021; Lee et al., 2021). These two wind-propelled hydrozoans belong to the neuston.

Attention is being paid to the distribution and mass stranding of these two colonial organisms, *V. velella* and *P. physalis*, that travel the neuston (Canepa et al., 2020; Headlam et al., 2020; Fierro et al., 2021; Jones et al., 2021; Macías et al., 2021). The life cycle of *P. physalis* was recently revised (Munro et al., 2019), although *V. velella*'s life cycle requires further study (Schuchert, 2010; Duarte et al., 2019). Despite this recent interest, the metabolism of these wind-propelled hydrozoans has been meagerly studied. In this paper, the respiratory metabolism of these organisms is measured and compared with the pulsating scyphozoans *A. aurita* and *P. noctiluca*, thereby comparing between different types of locomotion.

Taking advantage of the chance-beaching of smacks of *P. physalis* and *V. velella* on the north coast of Gran Canaria Island (Canary Islands, Spain) and aquarium *A. aurita* and *P. noctiluca*, we were able to measure and examine the respiratory nature and biochemical composition in both hydrozoans and scyphozoans. Respiration was the metabolism proxy used to investigate the metabolic difference between these two types of locomotion (Webb, 1971; Cowen, 1996; Gemmell et al., 2013; Fu and Uye, 2021). In particular, we analyzed the oxygen consumption rate, i.e., physiological respiration (R), and the respiratory electron transport system (ETS) activity, i.e., potential respiration (Φ), which is the enzyme activity at the cellular level responsible for the physiological macroscopic response (Packard, 1985). Additionally, this work describes the application of enzymatic ETS analysis to study the respiratory metabolism of these fragile cnidarians that are often cumbersome to sample and incubate without damage (Raskoff et al., 2003). These respiratory results were also used to investigate the impact of these organisms as predators through the calculation of the associated carbon demand. On the other hand, measurements of the basic biochemical composition and energy content provided insight into the role of these species as prey. This work helps us to understand the feeding strategies of gelatinivore predators and further deconstruct the viewpoint that jellyfish are trophic-dead ends (Hamilton, 2016).

Therefore, we hypothesized that the biochemical and energy content per wet mass (WM) of the two hydrozoan species may be higher since these species have chitinous structures that are absent in other, more fragile gelatinous zooplankton. We also hypothesized that R, Φ , and derived heterotrophic energy transformation (described as heterotrophic energy production in Packard et al., 2015), as well as the R: Φ ratio (Hernández-León and Gómez, 1996), should be lower in *P. physalis* and *V. velella* than in pulsating hydrozoans or scyphozoans. The wind-driven hydrozoans would require less respiration-derived

energy to cover the same distance compared to their scyphozoan relatives. Our objective was to investigate the physiology of these understudied organisms resulting in the first measurements of the respiratory metabolism and biochemical composition (protein, lipid, carbohydrate, and associated energy content) in *P. physalis* and *V. veleva* and comparing them with measurements in *A. aurita* and *P. noctiluca* with differing locomotion. This information provides novel insight into the impact of hydrozoans as both predator and prey, i.e., insight on integral members of mostly overlooked food webs (Helm, 2021).

MATERIALS AND METHODS

Sampling and Storage

Samples of *A. aurita* and *P. noctiluca* were provided by the Loro Parque Foundation's in Tenerife (Canary Islands, Spain). The *A. aurita* culture traced its origins to a stock culture from the Madrid aquarium (Spain). Specimens of *P. noctiluca* were captured during a jellyfish bloom that occurred in the Canary Islands in 2012. Generations of these two species have been cultivated at Loro Parque since their capture. The culturing conditions included filtered seawater at 18–20°C, 33 PSU (‰), and a pH of 8.2. *A. aurita* was fed thrice daily with *Artemia* nauplii and *P. noctiluca* was fed both *A. aurita* portions and *Artemia* nauplii. Prior to the experiment, *A. aurita* was fed *Artemia* nauplii, *ad libitum*, while *P. noctiluca* was not provided with its feed due to the requirements of the aquarium.

In the case of *V. veleva* and *P. physalis*, both specimens were collected in the northeastern beaches of Gran Canaria (Canary Island, Spain). These organisms arrived naturally as smacks in January and February 2017. They were carefully collected by gloved hands as they beached by the tide, after which they were transported live in containers half full of seawater. The samples were separated to be used for respiratory analysis as well as wet mass (WM) and dry mass (DM) determination.

The physiological respiration was first measured in organisms used to study respiratory metabolism, followed by the determination of WM. Next, the samples were then stored at –80°C to prevent enzyme degradation until enzymatic analysis was performed (Ahmed et al., 1976; Thuesen and Childress, 1994).

To determine the relation between WM and DM in *V. veleva* and *P. physalis* individual colonies of each species were rinsed with distilled water, placed on several dried petri dishes to remove the excess water, weighed inside pre-weighed aluminum foil, and then left in the drying oven at 60°C until constant DM was observed (~7 days). The samples from respiration experiments followed a similar weighing procedure; however, they were stored in a bag at a temperature of –80°C instead of the drying oven.

Oxygen Consumption Measurement (Physiological Respiration)

Jellyfish R was determined using oxygen-sensitive optodes to monitor oxygen (O₂) consumption [Fibox-4 by Presens

(Regensburg, Germany) (Lilley and Lombard, 2015)]. The organisms, submerged in seawater, were carefully transferred to containers ready for incubation and O₂ monitoring. Great care was taken during manipulation to avoid harm and stress to the organisms. A pre-calibrated optode spot was already installed inside the containers. The calibration consisted of a two-point calibration of 0 and 100% oxygen saturation using a saturated solution of NaSO₃ and vortexed seawater at the incubation temperature, respectively. The container size was dependent on the organism's size and was selected to maintain a container volume (CV)-to-wet mass (WM) ratio above 30 mL g⁻¹. Such containers included 60 mL glass BOD bottles, 350 mL glass jars, 600 mL glass jars, and 1 L glass jars. In the case of *V. veleva* and *P. physalis*, their floating nature posed a challenge for the incubation. R was measured with the organism completely submerged. An identical-sized vessel, filled only with filtered seawater, was used as a control.

All incubations took place in the dark at 20°C for 3–6 h with measurements every 10–20 min, and bubbles were rigorously avoided throughout the incubations. The seawater in the container was gently mixed prior to measurement of O₂ concentration ([O₂]) in μmol L⁻¹.

Electron Transport System Analysis (Enzymatic Respiration)

The ETS assay used was performed kinetically (an absorbance time-course) as described by Packard and Christensen (2004). The chemistry was based on Packard's first experiments (Packard et al., 1974), which was improved by several authors (Owens and King, 1975; Gómez et al., 1996; Purcell et al., 2019). The *A. aurita* samples and smallest *P. noctiluca* were homogenized using an ultrasonic probe (Cole Parmer) with a Vibracell VCX 130 ultrasonic processor in cold 0.1 M phosphate buffer [0.1 M Na₂HPO₄, 0.1 mM KH₂PO₄, 75 μM MgSO₄·7H₂O, polyvinylpyrrolidone (1.5 mg mL⁻¹), TRITON X-100 (2 mL L⁻¹)] at 8.5 pH. *P. noctiluca* medusa, *V. veleva*, and *P. physalis* samples were homogenized with a motor-driven Teflon-glass tissue-grinder at 4,000 rpm for 2–3 min, in cold 0.1 M phosphate buffer. A glass microfiber filter (GF/F Circle, 25 mm) was inserted into the tissue grinder to serve as an abrasive (Packard, 1971). The largest specimens were first blended, in a mechanical liquefier, with a known volume of cold deionized water (Puranity TU3), that was, sufficient to macerate the sample (Packard, 1971; King et al., 1978). Then, a known aliquot of that blended homogenate was blended in cold buffer with a GF/F filter in the Teflon-glass tissue-grinder as described earlier. In the case of the hydrozoans, a known aliquot of that buffered homogenate was also diluted to achieve a measurable concentration. The maximum final volume-to-mass ratio for *A. aurita* was kept close to 10 mL homogenate per g sample, for *P. noctiluca* close to 15 mL homogenate per g sample, and for *V. veleva* and *P. physalis*, it was kept between 30 and 45 mL of homogenate per g sample. These homogenization steps were always performed at 0–4°C in ice baths to retard enzyme degradation (Packard, 1971). The crude homogenate was

then centrifuged at 4,000 rpm (1,500 g) for 10 min at 0–4°C using a refrigerated microcentrifuge. The resulting supernatant contained the enzymes for analysis. The enzyme analysis consisted of mixing, in a cuvette, 0.1 mL of the supernatant, 0.3 mL of the buffered substrates [NADH (1.7 mM) and NADPH (1.7 mM)] and 0.1 mL of INT solution [INT tetrazolium salt 98%, 3-(4-Iodophenyl)-2-(4-nitrophenyl)-5-phenyl-2H-tetrazol-3-ium chloride (Acros Organics, Geel, Belgium) 2 mg per mL deionized H₂O (3.88 mM)]. The reaction in the cuvette was monitored in a Cary100 UV-Vis Spectrophotometer [Agilent Technologies (Santa Clara, CA, United States)] for 8 min at 490 nm until INT was reduced to a dark-red formazan. From the evolution of this absorbance, the rate of formazan production can be determined with the extinction coefficient of that batch of INT (13.76 mM⁻¹ cm⁻¹). The potential respiration (Φ) associated with this enzymatic system was calculated, knowing that 4 moles of electrons transferred down the ETS reduced 2 moles of INT and would normally reduce, at cytochrome oxidase, 1 mole of O₂. This implies that an ETS activity of 4 $\mu\text{mol e}^- \text{h}^{-1}$ is stoichiometrically equivalent to 1 $\mu\text{mol O}_2 \text{h}^{-1}$ of Φ (Packard, 1985). Blanks without homogenate were required to control any non-enzymatic INT reduction (Maldonado et al., 2012). The residual crude homogenate was stored at –20°C for biochemical determinations.

Biochemical Composition Measurements and Derived Energy Content

The protein content was measured in the samples homogenized for the ETS assay. These samples were diluted with buffer, when necessary, to reach a ratio of 120 mL of homogenate per g of sample. The analysis was based on the Biuret reaction as described by Lowry et al. (1951) and modified by Rutter (1967). It was further modified in this study by the addition of Dodecyl sulfate sodium salt (SDS) as suggested by Markwell et al. (1978) and Martínez et al. (2020). The lipid content was measured in samples homogenized previously without any further dilution. The content was determined based on the “index of total lipids,” as defined by Knight (Marsh and Weinstein, 1966; Knight et al., 1972; Barnes and Blackstock, 1973; De Coen and Janssen, 1997). We followed the methodology described by Bligh and Dyer (1959) for the extraction of lipids. A standard curve, ranging from 0 to 4.8 mg mL⁻¹ of olive oil dissolved in chloroform, was utilized for calculations. The carbohydrate content was measured in homogenized samples as previously described, without any further dilution. The content was studied using the method proposed by Dubois et al. (1956). A standard curve of glucose dissolved in homogenization buffer ranging between 0 and 1.5 mg mL⁻¹ was used. In the case of the smallest *A. aurita* and *P. noctiluca*, a more concentrated solution would have been desirable for the analysis, as in some cases, the biochemical components were below the limit of detection. The protein, lipid, and carbohydrate contents were used to calculate composition-based energy-contents and energy-densities (Ec) (Doyle et al., 2007) as proposed in Clarke et al. (1992). The average enthalpies of

combustion are 23.9, 39.5, and 17.5 KJ g⁻¹ for proteins, lipids, and carbohydrates, respectively.

Carbon Demand Calculations and Heterotrophic Energy Transformation

The daily respiratory carbon demand (RCD) in mg C d⁻¹ g WM⁻¹ can be calculated using the following equation (Uye and Shimauchi, 2005):

$$RCD = R \cdot RQ \cdot F \cdot t \cdot (AE \cdot 10^3)^{-1} \quad (1)$$

where: R represents respiration in $\mu\text{L O}_2 \text{h}^{-1} \text{g WM}^{-1}$, RQ represents the unitless ratio of CO₂-produced to O₂-consumed (0.85) (Uye and Shimauchi, 2005); F is the C:CO₂ conversion 0.536 $\mu\text{g C} (\mu\text{L CO}_2)^{-1}$, t is 24 to convert hours to days; and AE is the assimilation efficiency described by Schneider (1989). This is the percentage of assimilated carbon consumed, in this case, 0.8 for the scyphozoans (Schneider, 1989) and 0.9 for the hydrozoans (Purcell and Kremer, 1983). The equation is divided by 10³ to convert micrograms to milligrams. The same respiration rates were used to calculate the heterotrophic energy transformation (HET) in units of J d⁻¹ g WM⁻¹, using the associated energetics of the adenosine triphosphate (ATP) production (Packard et al., 2015; Azzaro et al., 2019). We prefer to use the term, “Heterotrophic Energy Transformation,” rather than the original term, “Heterotrophic Energy Production.” This is because according to the Law of Conservation of Energy (physics) and the First Law of Thermodynamics (chemistry), energy is neither produced nor destroyed. It is constant and can only be transformed from one form to another (Feynman et al., 1970). It would be calculated using R, in units of $\mu\text{mol O}_2 \text{h}^{-1} \text{g WM}^{-1}$, based on Equation (2):

$$HET = R \cdot Oe \cdot PO \cdot \Delta G_{ATP} \cdot t \quad (2)$$

where HET was reported as J d⁻¹ g WM⁻¹. Oe is 2, the number of electron pairs that participate in the reduction of one mole of O₂ to 2 moles of H₂O; PO equals 2.5 the modified P/O ratio (the ATP produced by the flow of one electron pair to atomic oxygen); ΔG_{ATP} equals to 0.048, the Gibbs free energy associated with ATP in J ($\mu\text{mol ATP}^{-1}$), and t is again 24 to convert hours to days. The original concept can be found in Ochoa’s work (Ochoa, 1943). The constants and ratios are explained by several authors (Alberty and Goldberg, 1992; Ferguson, 2010; Moran et al., 2012; Packard et al., 2015). To calculate potential HET (p-HET) from ETS activity, the calculation uses Φ instead of R.

Statistical Analysis

The boxplots were built using Microsoft excel (2016) (Microsoft-Corporation, 2016). R software was used to perform the statistical analysis (R Core Team, 2017) to compare the results for each species. The data for species have an unequal size and significantly unequal variances ($p < 0.05$) according to the Levene and Brown-Forsythe tests (Brown and Forsythe, 1974). Therefore, multiple comparisons were performed using Games-Howell *post-hoc* tests, which are known as a non-parametric approach based on Welch’s degrees of freedom correction and

do not require equal variances or equal sample size (Games and Howell, 1976; Lee and Lee, 2018).

RESULTS

Wet Mass

The wet masses of the samples and the number of samples (in parentheses) used in the respiration and composition studies ranged from 0.14 to 59.8 g for *A. aurita* ($n = 38$), 0.6–70.7 g for *P. noctiluca* ($n = 20$), 0.17–5.8 g for *P. physalis* ($n = 11$), and 0.19–1.1 g for *V. veleva* ($n = 5$).

Respiratory Metabolism

The respiration results (R & Φ) were normalized by WM and protein (Table 1 and Figure 1). R , normalized by WM, was observed to be higher in the hydrozoans than the scyphozoans (Table 1 and Figure 1A). Games-Howell statistical analysis showed a significant difference between the results of the scyphozoans and *P. physalis* ($p < 0.01$); however, it did not show a significant difference between the scyphozoans and *V. veleva*. The WM normalized Φ (Figure 1B) were even higher by a factor of 10. Here, the hydrozoans had a significant difference from the scyphozoans, but not among each other ($p < 0.0001$ for the test with *P. physalis* and $p < 0.05$ for the test with *V. veleva*). However, a higher water content was also observed for scyphozoans. Therefore, when the results were normalized by protein (P) (Figures 1C,D), R/P in *P. physalis* and *V. veleva* (1.81 ± 0.73 and $2.94 \pm 1.12 \mu\text{L O}_2 \text{ h}^{-1} \text{ mg P}^{-1}$, respectively) were below and significantly

different ($p < 0.0001$) from the P -specific R of *A. aurita* and *P. noctiluca* (15.82 ± 13.5 and $7.69 \pm 1.22 \mu\text{L O}_2 \text{ h}^{-1} \text{ mg P}^{-1}$, respectively). P -specific Φ (Figure 1D) was similar in all species with only *P. physalis* being significantly different from *A. aurita* with a $p = 0.041$.

The ratio R/Φ representing the fraction of the maximum enzymatic activity used by the physiological respiration varied as seen while comparing the hydrozoans and the scyphozoans. *A. aurita* and *P. noctiluca* showed an R/Φ of 0.61 ± 0.38 and 0.40 ± 0.15 , respectively, which were significantly different from each other ($p < 0.05$), whereas the ratios in *P. physalis* and *V. veleva* were significantly lower ($p < 0.0001$) than those of the scyphozoans by about a factor of 5 (0.11 ± 0.08 and 0.09 ± 0.04 , respectively). Hence, the wind-driven hydrozoans from this study used less of their potential metabolism, as hypothesized. Earlier zooplankton studies found ratios around 0.5 (Packard, 1985; Hernández-León and Gómez, 1996), which was the expected ratio between the *in vivo* enzyme activity and the maximum enzyme activity (Cleland, 1967).

Composition

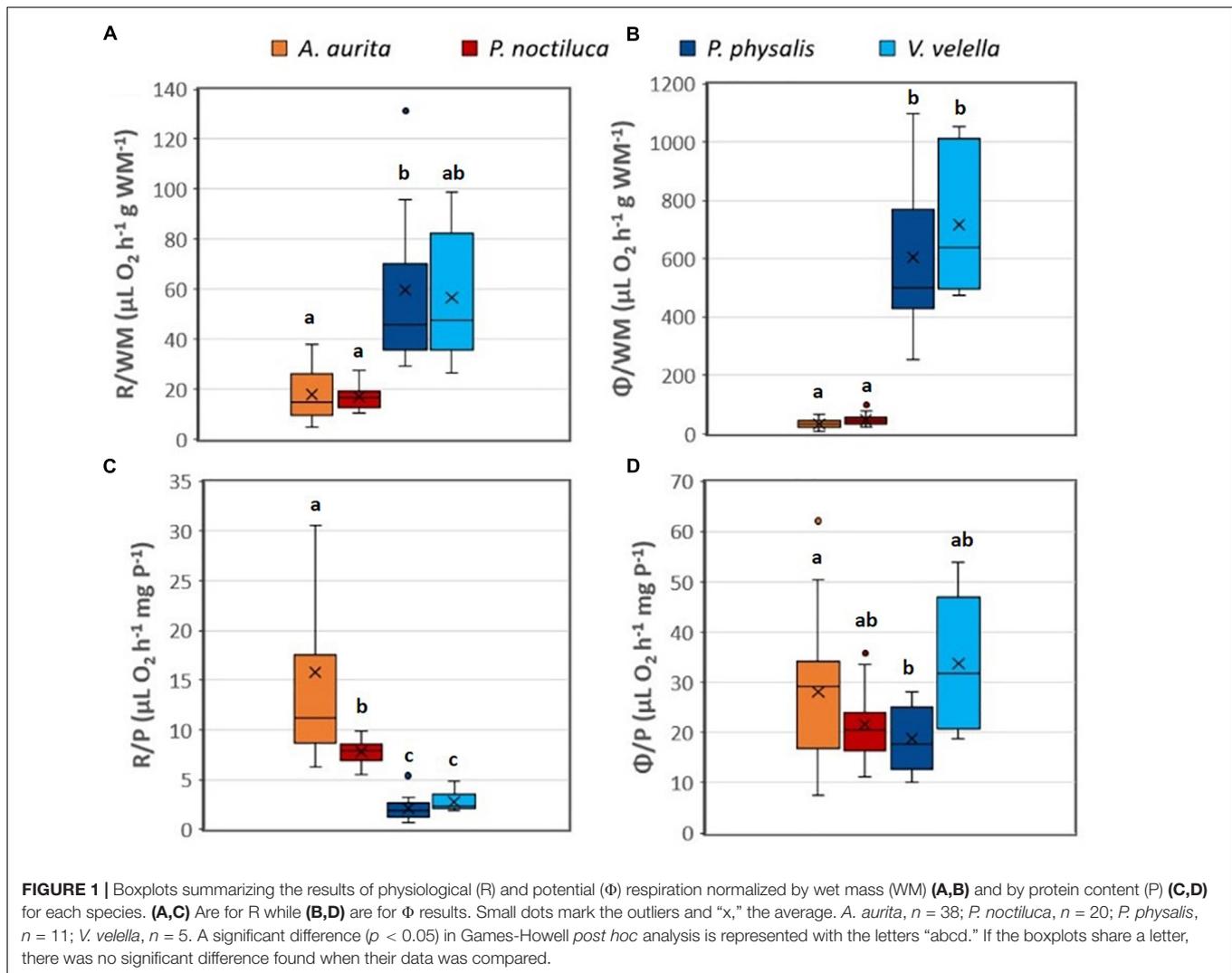
The protein (P), lipid (L), and carbohydrate (K) reported in this study, are the first such measurements in *P. physalis* and *V. veleva*. These results are showcased per WM in Figure 2. The average content per WM for *A. aurita*, *P. noctiluca*, *P. physalis*, and *V. veleva* for P were $0.12 \pm 0.04\%$, $0.21 \pm 0.07\%$, $3.72 \pm 0.97\%$, $2.39 \pm 0.76\%$ (Figure 2), showing significant differences from each other ($p < 0.005$). The results for L/WM were $0.11 \pm 0.14\%$, $0.07 \pm 0.06\%$, $0.62 \pm 0.19\%$, $0.24 \pm 0.15\%$ (Figure 2), where only *P. physalis* varied from the other species ($p < 0.05$). In the case of K , the scyphozoans demonstrated a few cases below the detection limit, showing the resulting values of 0.05 ± 0.05 , 0.02 ± 0.01 , 0.3 ± 0.05 , 0.17 ± 0.08 , respectively (Figure 2). The K content in scyphozoans was significantly different from hydrozoans ($p < 0.05$) with the exception of *A. aurita* and *V. veleva* ($p = 0.053$). The resulting WM-specific E_c in the hydrozoans was significantly higher than in the scyphozoans ($p < 0.0001$) and significantly different between the two hydrozoan colonial species ($p < 0.0001$).

The P , L , and K measurements were normalized by dry mass (DM) based on the $DM:WM$ ratio of 51 samples of *P. physalis* (0.04–13.1 g) and 24 of *V. veleva* (0.02–0.6 g). The resulting percentages of DM per WM were $11.7 \pm 1.6\%$ and $9.3 \pm 1.1\%$ for *P. physalis* and *V. veleva*, respectively. These values were higher than those of *A. aurita* and *P. noctiluca* (3.6 and 6%, respectively) and other gelatinous organisms (Morand et al., 1987; Malej et al., 1993; Lucas, 1994; Uye and Shimauchi, 2005). This is consistent with the water content in these two hydrozoans (89.3 and 90.7%) below the range expected for gelatinous zooplankton (94–97.6%), including pyrosomes (92.8%) (Bailey et al., 1995; Lucas et al., 2011; Ikeda, 2014). With respect to WM , the P , L , and K content of every component were higher in the two hydrozoans since the water content in these two is lower. However, for DM (Table 2), the P content was higher in the hydrozoans compared to scyphozoans measured, while lipid and carbohydrate contents in

TABLE 1 | Wet mass (WM) specific physiological respiration (R) results for each of the species studied here in the format of mean \pm SD.

Species	Source	R/WM	
		($\mu\text{L O}_2 \text{ h}^{-1} \text{ g WM}^{-1}$)	
<i>A. aurita</i>	This study	17.72 \pm 9.64	
	Uye and Shimauchi (2005)	2.96–3.72	↓
	Møller and Riisgård (2007)	15–180	≡↑
	Shimauchi and Uye (2007)	42.24–87.38	↑
	Han et al. (2012)	4.07–6.22	↓
<i>P. noctiluca</i>	This study	16.8 \pm 5.04	
	Malej (1989b)	0.58–5	↓
	Lilley et al. (2014)	6.61–9.06	↓
	Lilley and Lombard (2015)	0.43–0.55	↓
	<i>P. physalis</i>	This study	59.48 \pm 31.53
<i>V. veleva</i>	Larimer and Ashby (1962)	75–106	↑≡
	This study	56.62 \pm 27.43	

For each species ranges of WM specific R in the literature are also reported. Adjacent to each value is a symbol that represents how these ranges compare to the result in this study for the same species, i.e., ↑ if the range is above the mean, ↓ if the range is below the mean, ≡ if the range contains the mean. When ≡ is to the right of an arrow, it indicates that the mean \pm SD falls partially inside the range. When arrows are to the right of ≡, it is indicated that most of the range is either above or below the mean \pm SD. The results of this study are highlighted in bold to differentiate from the others published.



scyphozoans were closer to the content measured in hydrozoans, yet still lower.

Respiratory Carbon Demand and Heterotrophic Energy Transformation

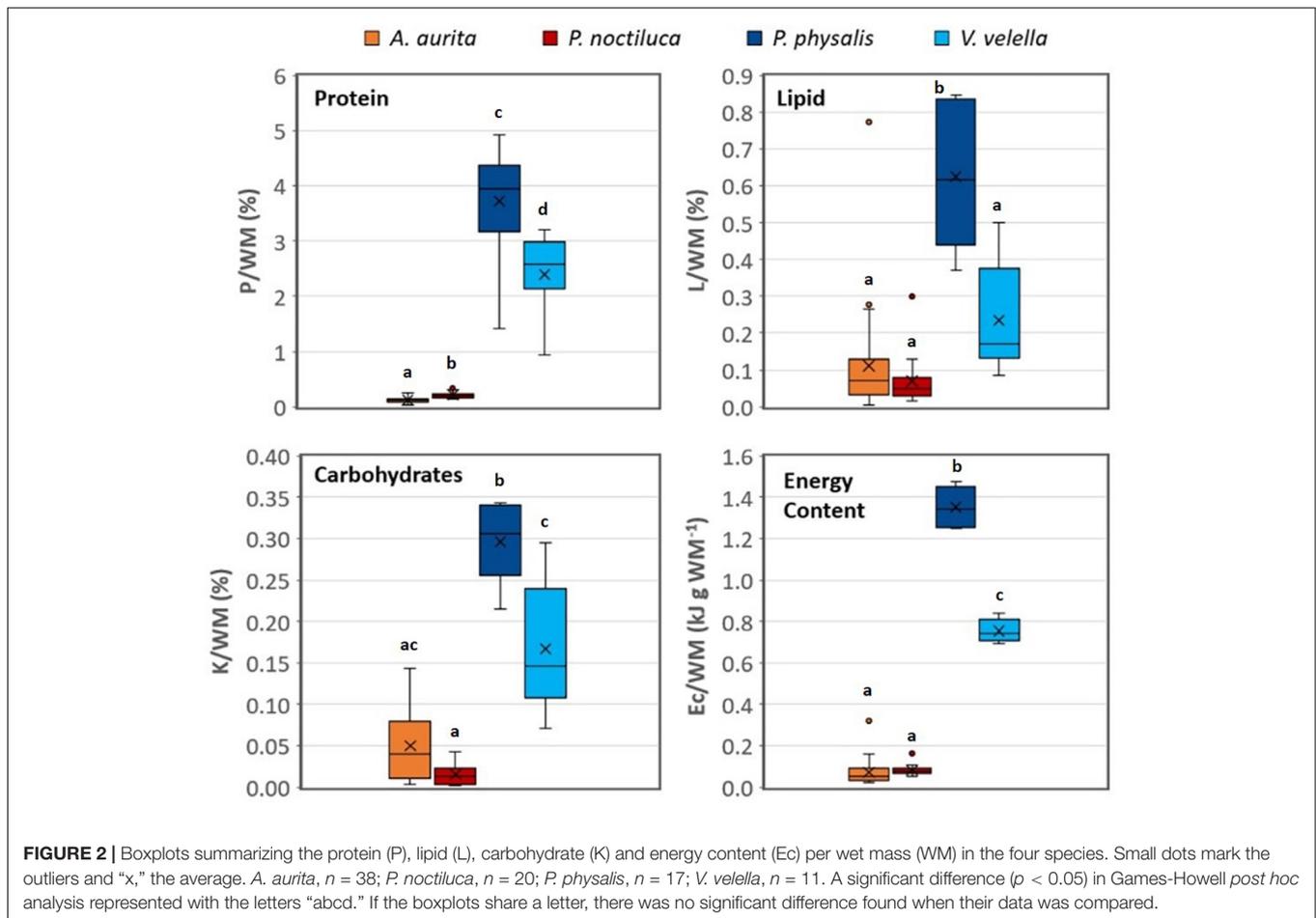
The results of respiratory carbon demand (RCD) and heterotrophic energy transformation (HET), calculated from R and Φ , are shown in **Tables 3, 4**. RCD and HET per WM were higher in the hydrozoans than in the scyphozoans, particularly the potential parameters derived from Φ .

The RCD in **Table 3** was further normalized by carbon (C) to showcase the percentage of each organism's C content required daily for survival. The resulting C-specific RCD illustrates that the daily respiratory C requirements fall below 5% of their own body-C mass except for *A. aurita*, which required an estimated 17% d^{-1} . This high percentage in *A. aurita* has been described previously (**Table 3**; Uye and Shimauchi, 2005). C ingested can sometimes even exceed its own body-C mass as reported by ingestion experiments with different types of preys

(Båmstedt et al., 1994). In the case of *P. noctiluca*, the results presented (**Table 3**) are fairly close to the percent of body-C respired reported by Malej (1989a). However, Lilley et al. (2014) described this percentage as closer to 7%. These values fit into the range found in other gelatinous zooplankton (Bailey et al., 1995; **Table 3**).

Similarly, HET was normalized by the Ec calculated from the P, L, K composition to determine the daily percentage of its own Ec required by the organism to survive. Here, all organisms required less than 10% of their own Ec daily with *P. physalis* and *V. verella* requiring less than 2% (**Table 4**).

The inverse of C-specific RCD and Ec-specific HET serve to estimate the number of days that the organism can survive with only its own carbon or Ec as sources to satisfy only their respiratory requirements. In this case, based on R-derived RCD, *A. aurita* would last for almost 5 days, *P. noctiluca* for 29, *P. physalis* for almost 50, and *V. verella* around 25. Based on Ec-specific HET derived from R, *A. aurita* and *P. noctiluca* would last 13 and 18 days, respectively, while *P. physalis* and *V. verella* would survive, 88 and 51 days, respectively. It must be noted



that these are rough estimations. Nevertheless, these estimations help in achieving a better understanding of both carbon- and energy-based metabolism.

DISCUSSION

Comparison and Implications of Derived Respiratory Carbon Demand and Heterotrophic Energy Transformation

The RCD reported here can be compared with estimations of carbon (C) ingestion rates to determine the percentage of C ingested that is approximately destined for respiratory metabolism. Uye and Shimauchi (2005) published an ingestion rate between 0.06 and 0.30 mg C d⁻¹ gWM⁻¹, whereas Møller and Riisgård (2007) reported a range between 0.58 and 0.69 mg C d⁻¹ gWM⁻¹ for *A. aurita*. Lilley et al. (2014) estimated the necessary carbon ingestion for an 8 cm *P. noctiluca* to be 0.57 mg C d⁻¹ g WM⁻¹. Based on these values and our RCD results (Table 3), we can infer that respiration in the scyphozoans requires a significant percentage of the carbon ingested with the potential of requiring further carbon. Uye and Shimauchi (2005) concluded that around one-half of the carbon ingested was used

for respiration, which implies an even higher percentage of the carbon assimilated.

Fish larvae have been identified as the main prey of *P. physalis* (Purcell, 1984). It was documented that *P. physalis*, with floats between 1 and 20 cm, captured fish larvae between 2 and 20 mm in length (Purcell, 1984). It was estimated that about 120 fish larvae were consumed daily by each colony (Purcell, 1984). The carbon content (g C) per fish larva can be estimated using the equation, $C = 0.44 \cdot 10^{-3} \cdot L^{3.272}$ (where C is carbon in grams and L is the length, in cm, for each larva) (Harris et al., 1986) for *Gadus chalcogramma* from the Gadidae family, one of the fish families sampled by Purcell (1984). This was followed by a calculation of the range of carbon content based on the fish length ranges published by Purcell (1984) (0.002–4.25 mg C) and determination of the associated C demand based on the amounts of fish consumed (about 120 individuals). Hence, it was found the C demand per individual colony associated with those predation rates, ranged between 0.27 and 510.04 mg C d⁻¹ ind⁻¹.

In the *P. physalis* colonies, the RCD per colony and potential RCD per colony, in mg C d⁻¹ ind⁻¹, ranged from 0.19 to 3.51 and 0.90–50.49, respectively (Table 3). Considering the fact that the float length of our study's siphonophores ranged from 2.39 to 6.77 cm, it is reasonable to propose that our results are closer to the lower values stated by Purcell (1984). Our

TABLE 2 | Protein (P), Lipid (L), Carbohydrate (K) and energy content (Ec) per dry mass (DM) for each species as mean ± SD.

Species	Source	P/DM (%)		L/DM (%)		K/DM (%)		Ec/DM (kJ g DM ⁻¹)	
<i>A. aurita</i>	This study (n = 38)	3.45 ± 1.22		3.1 ± 3.82		1.39 ± 1.31		2.02 ± 1.56	
	Lucas (1994)	2.07–28.6	≡↑	3.5–11.5	↑≡	0.35–1.12	↓≡	–	
	Arai (1997)	–		–		–		2.3–3.6	↑≡
<i>P. noctiluca</i>	This study (n = 20)	3.51 ± 1.18		1.11 ± 1.04		0.24 ± 0.2		1.29 ± 0.41	
	Malej et al. (1993)	11–17.6	↑	1.6–2.8	↑≡	0.3–0.5	↑≡	–	
	Arai (1997)	–		–		–		3.1–4.1	↑
<i>P. physalis</i>	This study (n = 17)	31.8 ± 8.27		5.34 ± 1.64		2.53 ± 0.43		11.6 ± 0.91	
	Wissing et al. (1973)	–		–		–		15–16.3	↑
<i>V. veleva</i>	This study (n = 11)	25.7 ± 8.16		2.53 ± 1.65		1.79 ± 0.86		8.12 ± 0.62	

For each species, ranges of composition and energy content per DM in the literature are reported. Next to each value is a symbol representing how these ranges compare to the result in this study for the same species. ↑ if the range is above the mean, ↓ if the range is below the mean, ≡ if the range contains the mean. When ≡ is to the right of an arrow, it indicates that the mean ± SD falls partially inside the range. When arrows are to the right of ≡, it indicates that the majority of the range is either above or below the mean ± SD. The results of this study are highlighted in bold to differentiate from the others published.

TABLE 3 | Compilation of the RCD (mean ± SD) for our four species of planktonic cnidarians normalized by wet mass (WM) and by carbon content (C).

Source	Description	RCD/WM		RCD/C	
		(mg C d ⁻¹ g WM ⁻¹)		(% d ⁻¹)	
		From R	From Φ	RCD (R)	RCD (Φ)
This study	<i>A. aurita</i> (20°C)	0.24 ± 0.13	0.45 ± 0.22	17.25 ± 9.38	32.35 ± 15.31
This study	<i>P. noctiluca</i> (20°C)	0.23 ± 0.07	0.63 ± 0.27	3.36 ± 1.01	9.23 ± 3.91
This study	<i>P. physalis</i> (20°C)	0.72 ± 0.38	7.83 ± 3.59	1.97 ± 1.04	21.34 ± 9.78
This study	<i>V. veleva</i> (20°C)	0.69 ± 0.33	7.89 ± 2.55	3.86 ± 1.87	44.26 ± 14.31

Malej (1989a)	<i>P. noctiluca</i> (19–23°C)	–	–	0.6–4.2	–
Bailey et al. (1995)	Medusae and ctenophores (6°C)	–	–	0.6–18.28	–
Uye and Shimauchi (2005)	<i>A. aurita</i> (15–25°C)	–	–	2.5–7.1	–
Lilley et al. (2014)	<i>P. noctiluca</i> (20°C)	–	–	0.4–7	–

C/WM: 0.14% for *A. aurita* (Uye and Shimauchi, 2005; Purcell et al., 2010), 0.68% for *P. noctiluca* (Morand et al., 1987), 3.67% for *P. physalis* (Beers, 1966) and 1.78% for *V. veleva* (Kemp, 1986). Other gelatinous zooplankton carbon specific respiratory carbon demand results in the literature are reported below the dashed line.

results are consistent with predation observations, considering that the amount of the ingested C destined for respiration will be lower than the total C ingested. Hence, RCD would be the minimum C requirement. For example, Purcell and Kremer (1983) found that the siphonophore, *Sphaeronectes gracilis*, at low food levels, ingested 5.5 times more C than necessary to meet metabolic demands and even 9 times more at high food

levels. This observation agrees with the low metabolic strategy hypothesized but begs to question if the rest of the assimilated C is invested in growth, reproduction, or the generation of specific organic substances in these colonial pelagic hydrozoans.

Regarding *V. veleva*, there has been some discussion regarding its main prey. According to the most recent findings (Bieri, 1966; Purcell et al., 2012, 2015; Zeman et al., 2018), its main prey include cladocerans, euphausid larvae, copepods, and fish eggs. For example, it has been found that *V. veleva* gut contained 43% cladocerans, 19% copepods (6% cyclopoida, 13% calanoid), 28% fish eggs, and 5% invertebrate eggs (Zeman et al., 2018). In other examples (Purcell et al., 2015), the prey found in *V. veleva*'s gastrozooids mainly included calanoid copepods and euphausid larvae (calycopsis and furcilia). By converting the ingestion rates that appear in these articles n° of prey consumed colony⁻¹ d⁻¹ to carbon mass consumed colony⁻¹ d⁻¹, we can compare the values obtained from ingestion studies with our metabolic results. To achieve this, we needed the approximate carbon content for the different types of prey ingested by *V. veleva* (µg C ind⁻¹). These carbon content values were for the calanoid copepod, 1.55

TABLE 4 | Compilation of the HET (mean ± SD) for our four species of planktonic cnidarians normalized by wet mass and by energy content (Ec).

Species	HET/WM		HET/Ec	
	(J d ⁻¹ g WM ⁻¹)		(% d ⁻¹)	
	From R	From Φ	HET (R)	HET (Φ)
<i>A. aurita</i>	4.56 ± 2.48	8.55 ± 4.05	7.55 ± 3.82	14.36 ± 6.58
<i>P. noctiluca</i>	4.32 ± 1.3	11.87 ± 5.03	5.47 ± 1.3	15.59 ± 6.62
<i>P. physalis</i>	15.29 ± 8.11	165.78 ± 75.97	1.13 ± 0.6	12.27 ± 5.62
<i>V. veleva</i>	14.56 ± 7.05	167.06 ± 54.01	1.93 ± 0.93	22.13 ± 7.16

and for the cladoceran, 0.72 (Walve and Larsson, 1999); for the fish eggs, 30 (Pereira et al., 2014); for the euphausiid larvae it depended on the stage, 10 for calycopsis, 984 for furcilia (Lindley, 1999; Guglielmo et al., 2015). Based on the ingestion rates we obtained the range 4.10–13.09 mg C ind⁻¹ d⁻¹ for 94.9% of the prey ingested (Purcell et al., 2015) as well as the range 0.02–7.62 mg C ind⁻¹ d⁻¹ for 84% of the prey ingested (Zeman et al., 2018). From our results (Table 3), the RCD per colony ranged from 0.21 to 0.35 mg C ind⁻¹ d⁻¹ and potential RCD per colony ranged from 1.4 to 6.2 mg C ind⁻¹ d⁻¹. It must also be considered that *V. velella* colonies used for our research had a sail length between 1.16 and 3.39 cm. Sail lengths from Purcell et al. (2015) ranged between 1.7 and 5.2 cm and between 1.9 and 3.3 cm from Zeman et al. (2018). Our results are in the lower end of the range compared to previous ingestion studies. These observations are similar to the *P. physalis* comparison. The ingestion may be higher to fulfill the demand for biological processes other than respiration, such as *V. velella*'s 0.5 mm daily growth (in length) (Bieri, 1977). Low carbon destined for respiratory metabolism is in line with our hypothesis of the low metabolic strategy of these neuston wanderers.

The measurement of HET presents great potential. If HET were measured in other organisms, we may be able to assess and compare their energy transforming capabilities and study energy circulation in the future (Karl, 2014). This may also be useful to address Margalef's suggested unifying principle in ecology regarding complexity and energy expenditure: "The energy required to maintain an ecosystem is inversely related to complexity, with the natural trend toward decreasing flow of energy per unit biomass; that is, increased maturity" (Margalef, 1963). Furthermore, HET also allows for calculations such as the one in Lane (2015) which described that the energy release, in terms of ATP synthesis, for an average person per gram was greater than that of the sun by a factor of 10,000. This inspired our calculation where we compared the HET by scyphozoan and hydrozoan mitochondria with the mass-specific energy-release rate of the sun at 0.017 J d⁻¹ g⁻¹ (Williams, 2018). From Table 4, the resulting ranges of average HET were between 4 and 15 J d⁻¹ g WM⁻¹, approximately. The calculation based on potential-HET was approximately between 8 and 170 J d⁻¹ g WM⁻¹ considering the four species in this study. Consequently, the WM-specific heterotrophic energy transformation in these jellyfish is between 500 and 10,000 times greater, on a per mass basis, than the sun's energy release rate. If jellyfish dry-mass is used in this calculation, the difference would be at least more than a thousand-fold greater. This is consistent with the findings presented by Lane (2015).

Composition

Regarding the scyphozoan composition measurements, our results for *A. aurita* can be compared with those of Lucas (1994). Protein and lipid content per DM in this study (Table 2) were found to be within the ranges presented by Lucas, but on the lower side. Carbohydrate measurements per DM were slightly above those illustrated by Lucas. For *P. noctiluca*, we compared our results to the results by Malej et al. (1993), which were generally higher than the percentages reported in this study (Table 2).

There is much less information about the general biochemical composition of *P. physalis* and *V. velella*. The explanation behind the high DM per WM percentage in the two hydrozoans compared to the other gelatinous organism could be related to the presence of chitin in the formation of the pneumatophore (Kirkpatrick and Pugh, 1984). The chitin gives both *P. physalis* and *V. velella* a more rigid structure that supports the changing interior gas pressure of the pneumatophore used to maintain the colony floating through the neuston. These harder structures are absent in other gelatinous zooplankton. However, the DM/WM of *V. velella* in this study ($9.3 \pm 1.1\%$) is higher than previously described $5.93 \pm 1.56\%$ (Kemp, 1986).

Comparing the P, L, K content per WM in this study (Figure 2) with the gelatinous zooplankton composition analyses by several authors (Arai et al., 1989; Clarke et al., 1992; Malej et al., 1993; Lucas, 1994, 2009; Bailey et al., 1995; Dubischar et al., 2006; De Barba et al., 2016), we can observe that the content of these components is higher in *P. physalis* and *V. velella* with certain minor exceptions. Hence, for a similar WM, these two could be a better source of some of these components due to their low water content.

When comparing the biochemical content per DM (Table 5) we observed that the range in the literature for protein was generally below the mean content determined in *P. physalis* and *V. velella* (Arai et al., 1989; Clarke et al., 1992; Malej et al., 1993; Lucas, 1994, 2009; Bailey et al., 1995; Dubischar et al., 2006; De Barba et al., 2016). Table 5 also depicts several exceptions, for example, the two cubozoans, *Chiropsalmus quadrumanus* and *Tamoya haplonema* (De Barba et al., 2016); as well as a few arctic species (Percy and Fife, 1981); *Aglantha digitale* (Ikeda, 1972); certain edible species (Khong et al., 2016); and the gonad tissue of *A. aurita* and *Cyanea capillata* (Lucas, 1994; Doyle et al., 2007). The higher presence of protein in the neustonic hydrozoans may be due to the protein-based blue pigment, protein-based venoms, chitin-protein complexes associated with the pneumatophore, or the protein in extracellular fluids (Lane et al., 1965; Gainey, 1972; Zagalsky and Herring, 1977; Tamkun and Hessinger, 1981; Roberts, 1992).

The gelatinous zooplankton L/DM range in the literature varied widely (Table 5; Arai et al., 1989; Clarke et al., 1992; Malej et al., 1993; Lucas, 1994, 2009; Bailey et al., 1995; Dubischar et al., 2006; De Barba et al., 2016). *P. physalis* presents above most measurements in the other species of this article (Figure 2). This could be due to their different diets (Lopes et al., 2016), which, in the case of *P. physalis*, is mainly fish-focused, or it could be related to the lipid content associated with nematocyst capsules and toxins (Stillway, 1974, 1976; Joseph, 1979). The exceptions to the lipid content in gelatinous species were the higher values in the arctic species described by Percy and Fife (1981; Table 5). These values will have implications on the energy content explored later. The arachidonic acid (20:4 ω 6), eicosapentaenoic acid (20:5 ω 3), and docosahexaenoic acid (22:6 ω 3) are proposed to be essential fatty acids (EFAs) (Cater et al., 2021), which may explain the reason jellyfish are in demand by some predators (Arai et al., 1989; Stenvers et al., 2020). These EFAs have often been found at high levels in both hydrozoans and scyphozoans (Stillway, 1976; Lopes et al., 2016; Tilves et al., 2016; Stenvers et al., 2020).

TABLE 5 | Protein (P), Lipid (L), Carbohydrate (K) and energy content (Ec) per dry mass (DM) for *P. physalis* and *V. velella* presented as mean \pm SD.

Source	Description	P/DM (%)		L/DM (%)		K/DM (%)		Ec/DM	
								(kJ g DM ⁻¹)	
This study	<i>P. physalis</i>	31.8 \pm 8.3		5.34 \pm 1.64		2.53 \pm 0.43		11.6 \pm 0.91	
This study	<i>V. velella</i>	25.7 \pm 8.16		2.53 \pm 1.65		1.79 \pm 0.86		8.12 \pm 0.62	
<hr/>									
This study	Scyphozoans (<i>A. aurita</i> , <i>P. noctiluca</i>)	0.94–6.88	↓	0.1–7.69	≡	0.03–4	≡↓	0.59–8.91	↓≡
Ikeda (1972)	<i>Aglantha digitale</i>	56.5	↑	3	≡	0.8	↓	–	
Percy and Fife (1981)	Arctic species	7.7–31	≡↓	4.4–22.1	≡↑	0.4–1	↓≡	4.35–16.7	≡↑
Arai et al. (1989)	<i>Aequorea victoria</i>	6.56	↓	2.17	↓≡	0.57	↓≡	2.05	↓
Clarke et al. (1992)	Southern Ocean gelatinous zooplankton	6.77–22.6	↓≡	0.72–8.41	≡	0.26–1.95	≡↓	2.56–6	↓
Malej et al. (1993)	<i>P. noctiluca</i>	11–17.6	↓	1.6–2.8	↓≡	0.3–0.5	↓≡	–	
Lucas (1994)	<i>A. aurita</i>	2.07–28.5	≡	3.5–11.5	≡↑	0.35–1.12	↓≡	–	
Bailey et al. (1995)	Medusae and ctenophores	0.21–7.77	↓	0.1–3.06	≡↓	0.06–0.8	↓≡	–	
	<i>C. sericeum</i> , <i>A. tortugensis</i>	11.6–21.4	↓≡	0.7–9.97	≡↑	0.99–1.6	↓≡	–	
Dubischar et al. (2006)	<i>Salpa thompsoni</i>	2.8–9.5	↓	1.3–9.1	≡	0.1–1.7	≡↓	–	
Doyle et al. (2007)	Compilation hydrozoans, scyphozoans and ctenophores	–		–		–		0.47–6.52	↓
Lucas (2009)	<i>Periphylla periphylla</i>	3.41–10.8	↓	1.15–4.81	≡	0.51–1.44	≡↓	–	
De Barba et al. (2016)	Cubozoans, hydrozoans and scyphozoans	10.6–27.7	≡↓	1.31–3.74	≡↓	0.94–5.92	≡	–	
Khong et al. (2016)	Edible species	19.2–39.2	≡	0.18–0.67	↓	8.87–18.2*	↑	3.9–9.08	≡↓

*In this article the K content was determined subtracting the percentage of moisture, protein, fat and ash from 100% instead of measuring it.

Below the dashed line, ranges other gelatinous zooplankton examples in this study and in the literature are displayed for comparison. Next to each value a symbol represents how these ranges compare to the results in this study for either hydrozoan. ↑ if the range is above the mean, ↓ if the range is below the mean, ≡ if the range contains the mean. When ≡ is to the right of an arrow, it indicates that the mean \pm SD falls partially inside the range. When arrows are to the right of ≡, it indicates that the majority of the range is either above or below the mean \pm SD. The results of this study are highlighted in bold to differentiate from the others published.

The K/DM range in the literature was normally between 0.1 and 1.7% (Table 5; Larson and Harbison, 1989; Clarke et al., 1992; Malej et al., 1993; Lucas, 1994, 2009; Bailey et al., 1995; Doyle et al., 2007). The results here for *A. aurita* and *P. noctiluca* were within this range while the K/DM ratios for the two hydrozoans were, in general, above this range (Table 5). The only exceptions were the K content described in the previously mentioned cubozoans, in the hydrozoan, *Olindias sambaquiensis* (De Barba et al., 2016), and in edible jellyfish (Khong et al., 2016). However, this was mainly due to the K content being calculated from other components instead of being measured. Carbohydrates are, in general, the least abundant of the three biochemical components (Lucas, 2009). Chitin intermediates present in both *V. velella* and *P. physalis*, as well as hexose associated with the float in *P. physalis* (Gross et al., 1958), were likely to be responsible for the K concentrations in their tissue.

Based on composition, the resulting Ec/WM (Figure 2) in *A. aurita* (0.07 \pm 0.06 kJ g WM⁻¹) and *P. noctiluca* (0.08 \pm 0.02 kJ g WM⁻¹) were within the range of the compiled observations in gelatinous zooplankton, as shown by Doyle et al. (2007), at 0.07–0.3 kJ g WM⁻¹, excluding the arctic species. Since arctic species such as those analyzed by Percy and Fife (1981), have a high lipid content, their maximum Ec/WM would be 0.85 kJ g WM⁻¹. The results for *P. physalis* (1.35 \pm 0.11 kJ g WM⁻¹) and *V. velella* (0.75 \pm 0.06 kJ g WM⁻¹) were close or above this maximum Ec/WM. Hence, for the same WM, the Ec in *P. physalis* was the highest. The Ec/DM for the hydrozoans (Tables 2, 4) was slightly above the results compiled by Doyle et al. (2007), except for the Ec/DM in high lipid arctic hydrozoans, which was double the maximum Ec/DM reported (Percy and Fife, 1981;

Table 5). The Ec/DM values in *A. aurita* and *P. noctiluca* were slightly below the ranges for these species published by Arai (1997; Table 2). In the tables of Ec compiled by Arai (1988), the Ec/WM for siphonophores was 0.67 kJ g WM⁻¹ (Musayeva and Sokolova, 1979), and the Ec/DM in *P. physalis*, based on bomb calorimetry, was 15–16.3 kJ g DM⁻¹ (Wissing et al., 1973). The first value agrees with our observations while the second, based on calorimetry, suggests a higher Ec than the one calculated here based on composition (Table 2).

Respiratory Metabolism

Due to logistical problems associated with incubations, we encourage research on enzyme-based alternatives. Enzyme-based techniques are also useful when working with fragile gelatinous zooplankton species that are often damaged during sampling, hindering live measurements (Purcell et al., 2010).

Comparing our WM-specific R results with similar studies in the literature (Table 1), we observe that, with regards to *A. aurita*, the average R/WM presented in this article (Table 1) was below some of the recently published results (Møller and Riisgård, 2007; Shimauchi and Uye, 2007) and above some others (Uye and Shimauchi, 2005; Han et al., 2012; Iguchi et al., 2017).

Iguchi et al. (2017) reported ETS activities in 10 *A. aurita* medusae (Table 6). They used this information to estimate R in the giant jellyfish *Nemopilema nomurai*, observing great potential in the application of the ETS assay. However, they were cautious about their results and advocated further studies in their final paragraphs. The results in this work for the 6 *A. aurita* medusae were closest in size to *A. aurita* in their study (Table 6), showing a considerable difference. The difference between studies may

TABLE 6 | Comparison between ETS results (mean \pm SD) on *A. aurita* medusae from Iguchi et al. (2017) and from our study.

	Iguchi et al. (2017)	This study
N	10	6
Size range (g WM)	0.7–1.9	0.7–1.5
R ($\mu\text{L O}_2 \text{ h}^{-1} \text{ g WM}^{-1}$)	5.9 \pm 1.4	28.6 \pm 8
Φ ($\mu\text{L O}_2 \text{ h}^{-1} \text{ g WM}^{-1}$)	7.1 \pm 2.6	52.1 \pm 6.8
R/ Φ	0.9 \pm 0.3	0.57 \pm 0.2

Note that in the literature, the ratio, R/ Φ , is often referred to as R/ETS (Filella et al., 2018).

have several explanations. We measured R with optodes, whereas the researchers used the Winkler technique that only measured the initial and final O_2 concentrations from the incubations. This technique is known to be an endpoint-type of analysis. Furthermore, their long (24 h) incubations may have led to O_2 -limited respiration (Gnaiger, 2001; Purcell et al., 2010).

Apart from the deterioration mentioned in their article, there are other factors in the study (Iguchi et al., 2017) that may have led to a loss in enzymatic activity. They describe freezing the homogenates, instead of the samples, with liquid nitrogen at -196°C , followed by storing the frozen homogenate at -80°C . From our experience, this leads to a significant loss in enzyme activity within 24 h (Ahmed et al., 1976). Furthermore, they also used the previous substrate concentrations (Owens and King, 1975), which might have slightly underestimated the ETS activity in scyphozoans. They also used the extinction coefficient of INT,

15.9 $\text{mM}^{-1} \text{ cm}^{-1}$, from Packard and Williams (1981) rather than determining the INT-formazan extinction coefficient for their own solutions. Most research teams do the same; however, we have found that this practice can lead to underestimations of ETS activity. In summary, Iguchi et al. (2017) inadvertently, may have underestimated both respiration and ETS activity. To resolve this type of discrepancy in the literature, an international intercalibration workshop or program should be organized for ETS analysis.

In the case of *P. noctiluca*, the R/WM results here (Figure 1 and Table 1) are higher than those reported in the literature (Table 1; Malej, 1989b; Lilley et al., 2014; Lilley and Lombard, 2015). For these two hydrozoans, the results are one of the first regarding their respiratory metabolism. No other study of R in *V. veleva* was found. R measurements in *P. physalis* tissue were reported in Larimer and Ashby (1962). Their results at 25°C were slightly above but close to our results (Table 1).

While comparing *P. physalis* and *V. veleva* with other jellyfish, WM-specific results here (Figure 1 and Table 1) were generally higher due to the lower water content compared to other gelatinous zooplankton. Therefore, the following comparisons are DM- or P-specific (Table 7). Compared to the study from Bailey et al. (1995), the R/DM results here (Table 7) are higher than in most gelatinous organisms except the Narcomedusa, *Aeginura grimaldii* and *in situ* incubations of other hydrozoan and scyphozoan species (Table 7). However, considering that these measurements were at temperatures between 4 and 7°C it is reasonable to conclude that the R/DM results here could be closer to or below those in the literature (Table 7).

TABLE 7 | Dry mass (DM) specific and protein (P) specific physiological respiration (R) results for *P. physalis* and *V. veleva* studied are presented as mean \pm SD.

Source	Description	R/DM		R/P	
		$(\mu\text{L O}_2 \text{ h}^{-1} \text{ mg DM}^{-1})$		$(\mu\text{L O}_2 \text{ h}^{-1} \text{ mg P}^{-1})$	
This study	<i>P. physalis</i> 20°C	0.45 \pm 0.24		1.81 \pm 0.73	
	<i>V. veleva</i> 20°C	0.61 \pm 0.29		2.94 \pm 1.12	

	Scyphozoans 20°C (<i>A. aurita</i> , <i>P. noctiluca</i>)	–		5.4–50.22	↑
Nival et al. (1972)	Medusae, ctenophores 10 – 29°C	0.54–1.8	↑=	–	
	Siphonophores 15°C	0.14–0.42	↓=	–	
	Tunicates 15°C	0.12–8.3	≡↑	–	
Biggs (1977b)	Suborder Physonectae 23 – 29°C	–		5–40	↑
	Suborder Cystonectae 23 – 29°C	–		4–6.9	↑=
	Suborder Calycophorae 23 – 29°C	–		5–75	↑
Larson (1987)	Hydromedusae 10 – 15°C	0.09–0.93	≡	–	
	Scyphomedusae 15°C	0.24–0.88	≡	–	
Barangé et al. (1989)	Sessile hydrozoan 21°C	0.5–3.9	↑=	–	
Malej (1989b)	<i>P. noctiluca</i> 12 – 25°C	0.05–0.68	≡	–	
Bailey et al. (1995)	Shipboard incubations 6°C	0.061–0.136	↓	–	
	Shipboard Incubations 6°C (<i>Aeginura grimaldii</i>)	0.472–0.754	≡	–	
	<i>In situ</i> incubations 4.6 – 7.2°C	0.01–0.81	≡	–	
Møller and Riisgård (2007)	<i>A. aurita</i> 7 – 22°C	0.13–0.58	≡	–	

Below the dashed line, ranges of other gelatinous zooplankton examples in this study and in the literature are displayed for comparison. Next to each value, a symbol demonstrates how these ranges compare to the results in this study for the two neustonic hydrozoans. ↑ if the range is above the mean, ↓ if the range is below the mean, ≡ if the range contains the mean. When ≡ is to the right of an arrow, it indicates that the mean \pm SD falls partially inside the range. When arrows are to the right of ≡, it indicates that most of the range is either above or below the mean \pm SD. The results of this study are highlighted in bold to differentiate from the others published.

On the contrary, our protein-specific results (**Figure 1** and **Table 7**) are lower than similar results from a variety of hydrozoan species, including several siphonophores (**Table 7**; Biggs, 1977b). However, these measurements taken by Biggs (1977b) were performed at $26 \pm 3^\circ\text{C}$, thus, it is also reasonable to assume that our R results are low but within the range. A detailed comparison between the siphonophore species present in Biggs (1977b) showed that individuals of the suborder Cystonectae, *Bathypphysa sibogae*, and *Rhizophysa filiformis* had the lowest respiratory rates. Both species are from the same suborder as *P. physalis*, and neither are observed to pulsate predominantly. This was consistent with our hypothesis that using wind (or currents) for movement through the ocean permits life at low levels of respiration, which can be detected by a low R: Φ ratio. Meanwhile, the genus from the suborder Physonectae, such as *Forskalia*, *Agalma*, or *Cordagalma* and suborder Calycophorae *Sulculeolaria* or *Stephanophyes* had the highest respiratory values, and were observed to pulsate frequently (Bone and Trueman, 1982; Costello et al., 2015). However, exceptions, such as the results from species of the genus *Diphyes* and *Abyla*, showed low respiration regardless of their pulse. This was not in line with our hypothesis. Similarly, our respiration results were lower than those compiled by Barangé et al. (1989) for sessile hydrozoans. In addition, Nival et al. (1972) presented very similar respiration rates in pulsating hydrozoans and much higher ranges in medusae and tunicates at 15°C (**Table 7**).

Low R: Φ ratios may support our hypothesis that, from a respiratory standpoint, these neuston dwellers should have a low R: Φ , due to their drifting nature. Few studies have measured both R and Φ in a cnidarian. In this study, *P. noctiluca* and *A. aurita* showed R: Φ ratios of 0.40 ± 0.15 and 0.61 ± 0.38 , respectively. Iguchi et al. (2017) observed a high R: Φ ratio of 0.9 ± 0.3 in *A. aurita*. Three pulsating hydrozoans (*Leuckartiara octona*, *Clytia gregaria*, and *Stomatoca atra*) showed R: Φ ratios between 0.5 and 0.92 (King and Packard, 1975). The exception amongst pulsating jellyfish may be the results in *Cassiopea* sp., which presented ratios R: Φ of approximately 0.17 ± 0.02 (Aljbour et al., 2017). These upside-down jellyfish have a more benthic lifestyle with a strong reliance for food on its symbiotic dinoflagellate (Lampert, 2016). Hence, their pulsation is less locomotive and the presence of these autotrophs may increase the ETS activity measured, leading to a higher Φ (Bondyale-Juez et al., 2017). A low R: Φ (0.19 ± 0.10) was also found in sessile polyps of *A. aurita* (Purcell et al., 2019). The low R: Φ in this study (0.11 ± 0.08 and 0.09 ± 0.04 in *P. physalis* and *V. velella*, respectively) resembled the low ratio of *A. aurita* polyps. Since both hydrozoans in this study are wind-propelled polyp colonies, there may be a low ratio amongst sessile life forms that do not require ATP for locomotion as readily. In the case of the free-floating, wind-driven hydrozoans in this study, a low metabolism may allow them to survive prolonged periods without prey consumption as they sail along the surface of pelagic oligotrophic waters (Biggs, 1977a).

In the case of *V. velella*, the presence of endosymbiotic photosynthetic dinoflagellates (Blank and Trench, 1986) may allow it to survive without prey and live with an even lower R: Φ ratio (Bondyale-Juez et al., 2017). However, we realize that our measurements need confirmation and replication in order to

strengthen them (Schmidt, 2009). For example, the hydrozoan samples were close to being stranded, and there is a possibility that the colonies near the shore did not feed as much as in the open ocean. Lane et al. (1965) suggested that *P. physalis* could not provide reliable physiological data after being maintained for more than a few hours due to problems with interaction with aquariums (Lane et al., 1965). In our experiments, no specimen that clouded the water or deflated abnormally was used.

However, gases from the floats from both organisms could cause an increase in O_2 concentration during incubation. Munro et al. (2019) described gas release in young *Physalia* and the unlikely release of gas naturally in mature colonies. Carter (1931) suggested there is gas exchange between pneumatophores of *Physalia*, *Veleva*, and *Porpita* species with the exterior. However, Larimer and Ashby (1962) tested that CO was the main gas secreted, while O_2 and N_2 were diffused inward. The possibility of pneumatophores with a respiratory function has been speculated (Carter, 1931), which could also reduce the consumption of the O_2 dissolved measured in the water if it was obtained from the gases in their pneumatophore. The idea of gas exchange between the pneumatophore and the exterior with a production of bubbles in the case of vertical migrating siphonophores has also been proposed (Pickwell, 1966), which is presumably not the case for *P. physalis*. No abnormal deflation, production of bubbles, or sudden increase in O_2 was observed. Nevertheless, these issues should be considered. Research on enzymatic alternatives is also encouraged, and hence, we tentatively considered low R: Φ ratios to support our hypothesis; however, we realize that future research addressing alternative explanations for the high Φ is needed.

In both cases, it is also possible that nearshore samples may not exhibit the same healthy metabolism demonstrated by open ocean samples. Regardless, these novel respiration and composition measurements bring us closer to understanding trophic exchanges and physiological strategies occurring in the neuston ecosystem.

CONCLUSION

The composition of *P. physalis* and *V. velella* varies from other gelatinous zooplankton. Their lower water content (approximately between 85 and 91%) leads to a higher concentration of protein, lipid, carbohydrate, and energy content per wet mass.

Enzymatic ETS analysis serves as a complementary or alternative measurement to study the metabolism of fragile and difficult-to-incubate zooplankton. In this study, we propose the optimum conditions for assaying ETS activity in cnidarian zooplankton. These respiratory metabolism measurements facilitate RCD and HET calculations.

We hypothesized that the respiratory metabolism of wind-driven, sail-based, *V. velella* and *P. physalis* would be lower compared to that of *A. aurita* or *P. noctiluca* with their muscular-based pulsation-driven propulsion. This hypothesis was supported by the respiratory results in these four jellyfish. A lower R: Φ ratio (~ 0.1) was found in these hydrozoans. A lower protein-specific R (1.81 and $2.94 \mu\text{L O}_2 \text{ h}^{-1} \text{ mg P}^{-1}$,

respectively) in *P. physalis* and *V. veleva* was measured compared to results here and in the literature. The respiratory carbon demand compared to their carbon ingestion rates in the literature was below that of scyphozoans. The novel HET/EC was approximately 2%. All these are consistent with our hypothesis.

DATA AVAILABILITY STATEMENT

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

AUTHOR CONTRIBUTIONS

JP, TP, and MG conceived of the presented idea. DB-J, VR-K, and IM designed and performed the experimental analysis supervised by TP and MG. DB-J and VR-K processed the data in consultation with TP and IM. DB-J and TP wrote the manuscript in consultation with all the other co-authors. All authors contributed to the article and approved the submitted version.

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