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Title

Long-wavelength reflecting filters found in the larval retinas of one mantis shrimp family (Crustacea, Stomatopoda, Nannosquillidae)

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Summary

Animals are known to exploit either transmissive coloured filters or reflectors for adaptive visual benefits. Here we describe a new category of biological optical filtration that acts simultaneously as both a transmissive and reflective color filter. Discovered in the larval eyes of only one family of stomatopod crustaceans, each crystalline structure bisects the photoreceptive rhabdom into two tiers and contains an ordered array of membrane-bound vesicles with subphotonic diameters (152 nm). Axial illumination of these intrarhabdomal structures *in vivo* produces a narrow-band of yellow reflectance (mean peak 572nm). While analogous visual structures are not known in nature, the optical performance of these intrarhabdomal structural reflectors is similar to synthetic devices used in the optical industry, such as band gap filters, laser mirrors, or fiber Bragg gratings. The interaction of these structural filters with longwavelengths of light suggests these structures may have evolved for detecting pelagic bioluminescence in shallow water at night.

Key Words: invertebrate vision, stomatopod, larvae, photonic structure, bioluminescence

INTRODUCTION

Optical filters are commonly used in nature to modify the spectral or overall sensitivity of an animal's visual system. Many species of both vertebrates and invertebrates use a reflecting structure (tapetum) behind their photoreceptors to increase photon capture and enhance vision in dim light [1-4]. Others use coloured filters positioned lateral or distal to their photoreceptors to tune spectral sensitivity by transmitting specific wavelengths of light not absorbed or scattered by the filter [5-7]. We recently discovered an optical structure positioned within the waveguide pathway of photoreceptive rhabdoms in the eyes of mantis shrimp (stomatopod) larvae from a single family, the Nannosquillidae. These intrarhabdomal structural reflectors (ISRs) are specialized to simultaneously reflect and filter specific wavelengths of light onto rhabdomeric tiers above and below the structure, respectively. This optical performance is similar to several synthetic devices such as band gap filters, laser mirrors, and (in particular) fiber Bragg gratings used in optical sensors for a wide range of industries. To our knowledge, the mantis shrimp larval ISR is the first example of a naturally occurring analog to these human-made devices.

Specialized optical structures are well known in adult stomatopod eyes [5, 8, 9] that contain visual adaptations such as UV and visible pigmented filters [10-12], unsurpassed diversity in colour and polarization sensitivity [5, 10, 13, 14], as well as unique structural reflectors used for visual communication [15]. Stomatopod larvae, by comparison, lack all of these adult features and are understood to have comparatively simple apposition compound eyes. The larval stomatopod eye structure is understood to be universal among species and typical of most other planktonic crustaceans in the open-ocean habitats (epipelagic and mesopelagic) where they are found. Typical features of stomatopod larval visual ecology include compound eyes with a uniform array of ommatidia and a single photoreceptor type [16, 17]; morphological adaptations for hiding in open-water, such as highly transparent bodies and reflective eye camouflage [18]; and diel vertical migrations, where animals reside at depths during the day and travel to the surface at night. It should be noted that diel migratory behaviors (or any visually mediated behaviors) are poorly characterized in mantis shrimp larvae. Current understanding of these behaviors is deduced from a limited number of circumstantial studies (e.g. [19-21] as well as personal observations of nocturnal larval abundance in shallow water during specific moon phases relative to an absence of larvae at the same location during the day. Though the daytime depths of stomatopod larvae are not specifically known, one study collected *Squilla mantis* larvae at 10-200 m daytime depth in order to quantify geographic distribution of this species' abundance [22]. These depths are similar to what is known for the daytime depths and vertical migratory behavior of decapod crab zoea, which occur at about 12 m during the day, varying with larval stage [23].

Prior to this study it was understood that most species of stomatopod larvae possess similar large, apposition compound eyes for mediating visually guided behaviors such as predation and anti-predation responses, as well as migration [17, 24]. The gross examination of larval eyes from diverse taxa, however, reveals that a single family (Nannosquillidae) deviate from the typical eye structure by their possession of ISRs within their rhabdoms. In this study we characterize the anatomical and optical properties of nannosquillid larval ISR-containing retinas as well as consider how these structures may influence task specific behaviors in their nocturnal, dim light habitat.

RESULTS

Histology: In general, stomatopod larvae possess a pair of complex compound eyes composed of several hundred ommatidial units, each using transparent, apposition optics to focus light onto the photoreceptive rhabdom that is similar in structure to other crustacean larvae [25, 26]. The typical crustacean larval rhabdom is formed from a tightly packed column of visual pigment-expressing microvilli projected from a ring of seven retinular cells (R1-7). Each rhabdom is optically isolated by screening pigments in the retinular cells. In many crustacean larvae, including stomatopods, reflective structures lie on the surface of the retina, between each rhabdom, to camouflage the dark eye in open water [18]. While the majority of stomatopod larval retinas adhere to this typical arrangement, the retinas of nannosquillid species contain a conspicuous alteration: the intrarhabdomal structural reflector (ISR). The ISR is a 4-segmented, barrel-shaped structure that sits approximately one-third of the way down from the distal end of a rhabdom (Fig. 1). Electron and light microscopy reveal that these ISRs are not uniformly found across the eye but regionalized to the ventral and lateral ommatidia (Fig. 2). A subset of 30-50 ommatidia in the dorsal region of the eye are devoid of ISR structures and instead possess the typical untiered photoreceptor structure of other crustacean larval retinas (Fig. 1F, 2D). Two-photon microscopy further established the three-dimensional distribution of ISR-expressing and non-expressing ommatidia across the nannosquillid eye (Fig. 2A-B, Movie S1).

Diverse taxonomic sampling of stomatopod larvae for transmission electron microscopy (TEM) or two-photon microscopy revealed that ISRs may only occur in larvae from the family Nannosquillidae. At least four of the 13 genera described in the Nannosquillid family are represented by species in this study: *Pullosquilla thomassini*, *Pullosquilla litoralis*, *Alachosquilla vicina*, *Coronis scolopendra*, and two unknown nannosquillids (1 and 2) currently lacking adult DNA barcode reference sequences but nested within the nannosquillid clade (Fig. 3). The absence of ISRs from other stomatopod lineages suggests that this visual adaptation is unique to nannosquillid larval ecology. Additionally, ISRs were found at different developmental stages of each species, including the first and terminal larval stages, suggesting that their function serves the animal throughout the entire pelagic phase of life.

Within each ISR is an ordered lattice of spheroid, membranous vesicles, each an average of 152 ± 12 nm (mean $\pm$ 1 standard deviation; $n = 2074$) in diameter. TEM and electron tomography reveal that the ordered arrangement of these vesicles is preserved across the membranes of the four primary segments (Fig. 1D) and in three dimensions (Movie S2). Each ISR measures an average of 11.0 ± 2.0 μ m long by 4.8 ± 1.3 μ m wide and is positioned directly in the optical pathway of light, bisecting the retinular cells (R1-7) of the rhabdom into a proximal (R2, R3, R6, & R7) and distal tier (R1, R4, & R5, Fig. 1). This cellular arrangement is similar to that of the tiered rhabdoms found in the ommatidia of rows one to four of the midband of some adult stomatopod eyes [13].

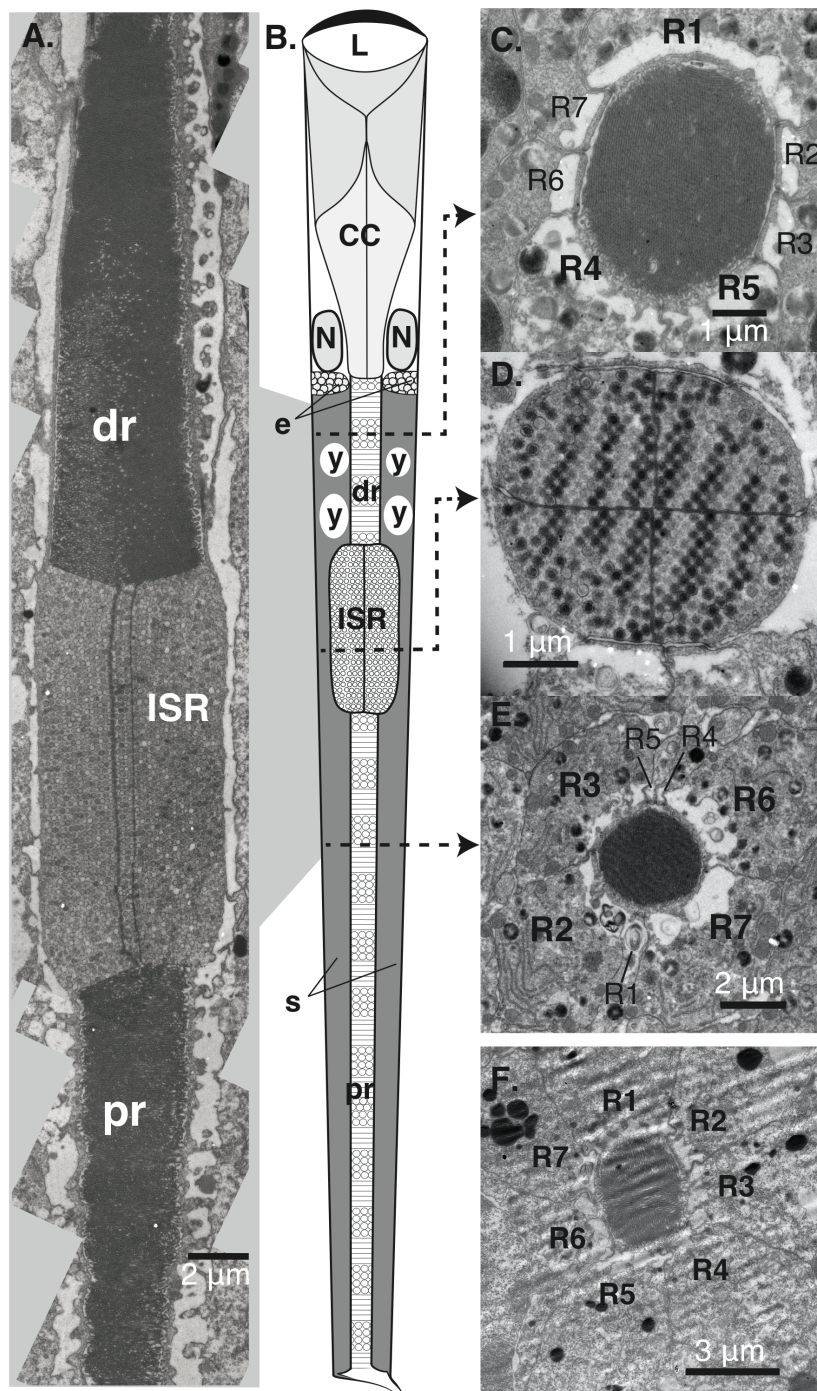


Fig. 1. Anatomy of stomatopod larval ommatidia containing intrarhabdomal structural reflectors (ISR) described by histology. (A) Composite TEM of longitudinal section through a lateral ommatidium. Orthogonal microvilli in the distal (dr) and proximal (pr) rhabdoms flank the ISR. (B) Diagram of an ISR-containing ommatidium. (C) TEM cross-section through the distal tier of the rhabdom, formed by microvilli from retinular cells R1, R4, and R5 (nomenclature from [10]). Like adult tiered rhabdoms, extensions of the remaining retinular cells are visible and do not contribute microvilli to the rhabdom. (D) TEM cross-section through ISR. (E) TEM cross-section through the proximal rhabdomeric tier, formed by microvillar projections from retinular cells R2, R3, R6 and R7. (F) TEM cross-section through a non-ISR expressing photoreceptor in the dorsal region of the eye. Note the equal contribution of microvilli from retinular cells R1-7. Lens, L; crystalline cone, CC; Retinular cell nucleus, N; reflective eye camouflage, e; yellow, long-pass screening pigments [27], y; lateral screening pigments, s.

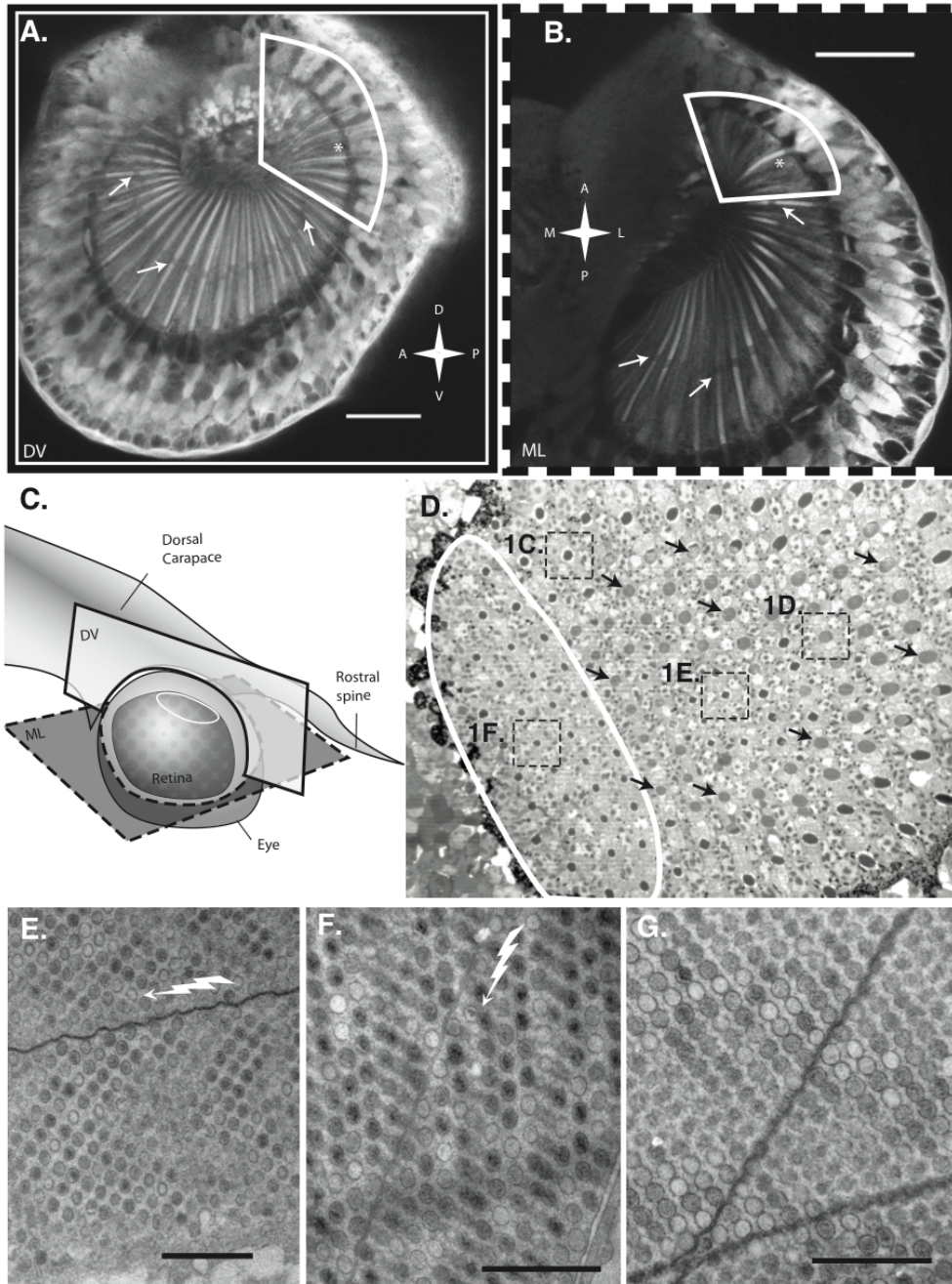


Fig. 2. Regional localization of ISRs across the retina and in the eyes of five different nannosquillid species. (A-B) Two-photon optical sections of *Coronis scolopendra* first stage larval eyes revealing a small region of dorsal pointing ommatidia (within white line) lacking ISR structures (denoted by *). Arrows indicate the dark, less autofluorescent ISRs separating rhabdoms into two distinct tiers. Compasses indicate anatomical directions: A, anterior, P, posterior, M, medial, L, lateral, D, dorsal, V, ventral. Scale bars, 50 μ m. (C) Diagram of dorsal-ventral (DV) and medial-lateral (ML) sections through eye in A (solid border) and B (dashed border), respectively. (D) Light micrograph of retina cross-section from early stage larva, unknown nannosquillid species 1. Boxes depict regions imaged via TEM in Fig. 1C-F. White line denotes dorsal region of untiered ommatidia lacking ISR expression, similar to zone identified in A-C. Arrows highlight subset of ISRs in the remainder of the eye. (E) TEM longitudinal section of ISR in last stage larva, *Pullosquilla thomassini* and (F) mid stage *Alachosquilla vicina*. Jagged arrows indicate incoming optical axis in longitudinal sections. (G) Oblique TEM section of ISR in early stage, *Pullosquilla litoralis* larva. E-G scale bars, 1 μ m.

***In vivo* Reflectance Spectroscopy:** The size and arrangement of vesicles within the ISR structures led us to hypothesize that certain wavelengths of light will be reflected when the structure is illuminated. To investigate how white light interacts with the ISR, we used a custom microscope system to illuminate, image, and measure reflectance spectra from the pseudopupils of larval ommatidia *in vivo* (Fig. S2; as in [18]). The pseudopupil is the dark spot that moves smoothly across the surface of a compound eye as it rotates, which is produced when the optical axes of a subset of ommatidia align with the optical viewing axis of an observer. Since the ISR lies in the optical axis, we predicted that a distinct reflection would be observed when light was imaged axially into ISR-containing rhabdoms.

Prior to this experiment, the phylogenetic distribution of species expressing ISRs was not known. Therefore, we tested a diverse range of species available at our field site (Lizard Island, Australia) both to measure the interaction of light with the ISR as well as to examine the distribution of species that express these structures. The experiment was conducted blind to species identity by using wild-captured larvae that were identified using DNA barcoding after light reflectance measurements were collected [28]. It was observed that some larvae reflected a band of yellow light (average peak of 572 ± 18 nm) from the ventral and lateral ommatidia, but only when illuminated on-axis (Fig. 3; Fig. S2). While side-illumination was sufficient to visualize and measure the blue and green camouflage structures that lie over the pigmented retina [18], only on-axis, epi-illumination could produce a yellow reflectance from the pseudopupil (Movie S3). After the experiment each animal was identified using DNA barcoding, which revealed that yellow pseudopupil reflectances were only found in nannosquillid species (Fig. 3H). These species represent three of the six nannosquillid species shown to contain ISRs histologically: *Pullosquilla thomassini*, *Alachosquilla vicina*, and unknown species 1 (Fig. 3H). While stochastic species sampling and post hoc identification allowed for unbiased data collection, these same methods also prevented our ability to collect reflectance measurements from the remaining species in this study. TEM was used to verify the presence of ISR structures in post-reflectance-measured specimens (Fig. 2D-F), confirming that ISR structures were the source of observed yellow reflectances. This conclusion was further supported by the agreement between patterns of ISR anatomical regionalization and the pattern of yellow-reflectance observed from the lateral and ventral ommatidia (Fig. 2A-D, Fig 3, Movie S1). Neither yellow reflectances, nor ISR structures were found in the dorsal-most ommatidia.

Of the three nannosquillid species measured for *in vivo* pseudopupil reflectance, the wavelength of peak reflectance did not vary significantly among these species (ANOVA, $n_{\text{UKNan1}} = 40$, $n_{\text{Puth}} = 3$, $n_{\text{Alvi}} = 2$, $F_{1,43} = 0.01$, $p = 0.923$), though different spectral shapes were observed from each species (Fig. S2). Variation in spectral shape may be attributed to differences in eye size, ommatidial diameter, or subtle differences among species in their ISR anatomical assemblies (Fig. S1). Variations in the amount of reflectivity (seen in Fig S2B) were also common due to a variety of factors, including subtle alignment deviations of the sample probe with the optical axis; imaging through multiple layers of dioptrics and rhabdomeric material; and different sizes of the eyes and ommatidia aperture among individuals of the same species but different stages.

Several ISR reflectances contain short-wavelength components we attribute as possible contamination within our reflectance set-up from the surrounding blue-reflective material (eyeshine) that lies on the surface of the retina between ommatidia (Fig S2B [18]). ISR-containing ommatidia correspond to the lateral and ventral areas of the eye that express this blue eyeshine (Movie S1; Fig. 3A). Ommatidia with green eye shine in the dorsal region of the

eye lack ISRs. At present, a pure reflectance from the ISR alone cannot be obtained until methods are devised to adequately remove the crystal from within the center of the eye.

Photonic Modelling: To understand the wavelength-selective reflectance from the ISR (Fig. 3G), we developed a semi-analytical optical model. This mathematical model uses a combination of Bragg's law [29] with finite distance time domain numerical simulations. By using these methods we were able to account for the findings of the *in vivo* experiment and test for any morphological disorder of the ISR potentially induced by TEM. The amount of disorder and the periodicity of the vesicles was determined by calculating the structure factor estimated from TEM micrographs (S.I.). The structure factor of the vesicles within the ISR is related to a face centered cubic geometry of the vesicles with an estimated lattice constant of 392 ± 7 nm (mean $\pm$ 1 standard deviation; Fig. 4A), which takes into account any structural deformations from TEM embedding and slicing of the tissue. Since the materials contained within the ISR vesicles are not currently known, values for two vesicle refractive indices (n_v) were tested in the model. A high-refractive index material, pteridine (n_v 1.70), and a low-refractive index material, lipid (n_v 1.48), were used to model the unknown contents of the ISR vesicles (further details in S.I.). Pteridine was selected as the high-refractive index material due to the consistency of our TEM results with membrane-bound pteridine granules found in other crustacean photonic structures [30-32]. Our model resulted in good agreement with the mean experimental reflectivity spectrum for lipidic vesicles with a histological expansion factor (ψ) of 8% (further details in Supplemental Information, Fig S2F, Fig4B). This is further evidence to suggest that the observed yellow reflectance measured from the larval pseudopupil is produced by the ISR. The residual difference between the reflectivity of our predicted and observed reflectances can be attributed to several factors including potential histological distortion of ISR dimensions (i.e. ψ); unknown identities (refractive indices) of vesicle and matrix materials; and additional disorder in the orientation and periodicity of the 3D lattice across large volumes outside the range of TEM tomography (Fig S2F-J).

Though we were able to collect sufficient data to describe the reflection properties of the ISR, we were unable to measure how light is transmitted through the structure. The location of the ISR within the rhabdom poses a logistical problem for collecting light transmission data. Frozen, fresh, or fixed sectioning methods all distort the vesicle lattice, affecting spectral transmission. Though transmission data could not be collected, gross dissection and visual inspection of fixed retinas revealed that ISR structures lack colorful pigments (Movie S4) suggesting that any transmitted wavelengths are likely the result of scattering or reflection from the photonic structure, rather than absorption by photostable pigments. The banded staining pattern of the ISR vesicles in TEM is also consistent with other micrographs of paracrystalline assemblies of clear, layered vesicles in snail retinas [33].

corresponds to the Nannosquillidae family. ‡ indicate barcoded species where reflectance was not measured, but ISR absence or presence was determined by histology only.

DISCUSSION

The discovery of ISR structures raises three major questions. First, what source of light in the natural habitat of nannosquillid larvae interacts with the ISRs? We have established that the ISR maximally reflects light centered at a wavelength of 572 nm. Stomatopod larvae encounter few sources of light in this range of the visible spectrum since, like many other larval crustaceans, they live at depths up to 200 m by day and come towards the surface at night. The broad irradiance spectrum of moonlight is abundant close to the water's surface and during periods of the full moon. However, it is unlikely that nannosquillid visual systems would be tuned to detect the long-wavelength components of moonlight since this light source is highly variable with lunar phase. During the half and dark phases of the moon the total irradiance is 1/10 and 1/100 that of full moonlight, respectively [34]. Nannosquillid larvae have only been observed to rise in the water column to within one meter of the surface during the quarter and dark phases of the moon (pers. observ; pers comm. RL Caldwell). The only other source of long-wavelength light in the nannosquillid larval habitat is bioluminescence (BL). BL has evolved multiple times in diverse taxa inhabiting the pelagic environment, resulting in a range of peak wavelengths of emission that vary with depth. The majority of bioluminescent species live in the deep sea and emit spectra that peak between 460 nm and 500 nm [35]. While there are less than half as many bioluminescent species characterized from shallow water (where mantis shrimp larvae reside), the emission spectra of these species are red-shifted, with peaks between 460 nm and 520 nm [35]. Though BL is dim, since nannosquillid larvae are generally much smaller than other mantis shrimp (2 mm to 12 mm total length) it may be a salient signal in the close range of larval interactions.

A second question is raised after consideration of the type of light nannosquillid larvae encounter in their habitat: Does the ISR improve detection of these natural light sources (moonlight or BL)? One known mechanism for improving the contrast detection of BL is the use of yellow, transmissive filters in the lenses of many deep sea fish. In these fish species, only the long-wavelengths of BL emission spectra are able to reach these fishes' photoreceptors, improving the contrast of a BL target against the short-wavelength dominant background of the deep sea (for example [36, 37]. The ISR, however is not a short-wavelength absorbing, long-pass filter, but rather a long-wavelength reflective, short-pass filter. The convergence of selection towards similar wavelengths onto photoreceptors in these two systems led us to hypothesize that the ISR may have evolved for improving sensitivity to BL. Since many of the variables surrounding nannosquillid ecology and visual physiology have yet to be described (including a potential BL source), we developed a model to test the potential effect of ISRs on nannosquillid retinas viewing BL light sources that peak at different wavelengths. The model estimates the relative proportion of photons captured by the rhabdomic tiers from an incoming light source (quantal catch, QC) with and without the reflective or filtering effect of the ISR (full details in Supplementary Materials, [based on 38].

The model incorporates experimental morphological measurements, the mean measured ISR spectral reflectivity (estimated at 50% reflectance), an estimation of ISR transmission (1 – IRS reflectance), and two visual pigments known to be expressed in different species of nannosquillid larvae (450 nm and 500 nm lambda max, Fig. S3A). The spectral forms of BL emission spectra were obtained from the literature (Fig S4B, [39, 40] and modelled as a 66 nm

FWHM Gaussian spectral distribution (black trace, Fig. S4C). This estimation allowed us to test how the ISR impacts relative absorbance changes for different BL emission spectra that peak at each wavelength in a range from 450 nm to 650 nm (Fig 4D-G). Moonlight irradiances at different phases were also tested, though the effect was negligible during the quarter phase (1.8%) when nannosquillid larvae are found near the surface (see Supplemental Materials for further details and discussion).

Though all visual pigment scenarios of the model predict an increase in QC, the maximum increase from ISR reflectance onto the distal tier arises when the visual pigment $\lambda_{\max}$ is 500 nm. This conservative estimate (at 50% ISR reflectivity) predicts up to an 11% boost in photon capture for BL emissions peaking between 475 nm and 520 nm, the range of BL emissions in the shallow pelagic habitat (Fig 4; [35]). The proximal tier receives a decrease in QC, regardless of visual pigment expression. This is due to a baseline reflectance of 20% for all wavelengths in our experimental data. Use of a modeled ISR reflectance/transmission spectrum (Gaussian curve, Fig. S3A) produced similar patterns of increased QC in the distal tier and decreased QC in the proximal tier, though to a lesser extent than with the experimental data (Fig S3E-H). This is a result of the Gaussian model's baseline being set to zero. The model predicts that for both the modeled and experimental reflectance data, the maximum capture of photons occurs when the proximal tier $\lambda_{\max} = 450$ nm and the distal tier $\lambda_{\max} = 500$ nm (Fig 4F, Fig S3G). Though stomatopod larvae are typically understood to express one spectral type of photoreceptor [16, 17], the discovery of the ISR and the results of this study provide evidence for a reasonable hypothesis that nannosquillids may possess a dichromatic retina.

These calculations demonstrate a potential for photonic interaction between the ISR and common bioluminescent sources in the shallow pelagic environment. To properly test the hypothetical interactions with BL light, future research must target characterization of the chemical nature of the ISR components and the optical properties of the overlying materials (lens & crystalline cone), as well as assess the spectral sensitivity of each rhabdomeric tier in a nannosquillid retina. Incorporation of precise measurements of these currently unavailable factors may reveal that the increase in QC of the distal rhabdom is indeed much greater than our conservative estimates.

Finally, we questioned which ecological pressures might have led ISRs to localize to specific regions of the eye? In the lateral and ventral regions of the eye, ISRs would be beneficial for imaging point sources of bioluminescent light around and below [26] the animal in order to aid predation of potential food sources. Nannosquillids may have evolved a selective advantage by gaining more visual information about a specific food resource relative to other larval species of stomatopod. In contrast, the dorsal region of the eye lacks the ISRs. As previously mentioned, some mesopelagic fish, use yellow filters in upward looking visual fields to break BL camouflage used on the ventral surfaces of many prey species of fish. The absence of yellow-reflecting ISRs in the dorsal retina matches that fact that this is not be a behaviorally relevant task for stomatopod larvae, with evolutionary selection instead for more typical pelagic type photoreceptors in the dorsal regions each eye. In comparison to other species of stomatopod larvae, nannosquillid larvae are much smaller and faster swimmers (pers. observations). An evaluation of nannosquillid larval behavior may reveal that they use different predation strategies relative to other, often sympatric species of mantis shrimp larvae.

Stomatopods, at all life history stages, continue to provide a compelling system for understanding visual adaptation in the ocean. We have described a new type of visual structure

that functions concurrently as both a transmissive and reflective colour filter in stomatopod larval eyes. Looking forward, the homology between these ventral and lateral larval nannosquillid ISR ommatidia and tiered adult stomatopod rhabdoms may provide insight into the developmental mechanisms that led to the rise of the elaborate colour vision system found in many adult stomatopod eyes.

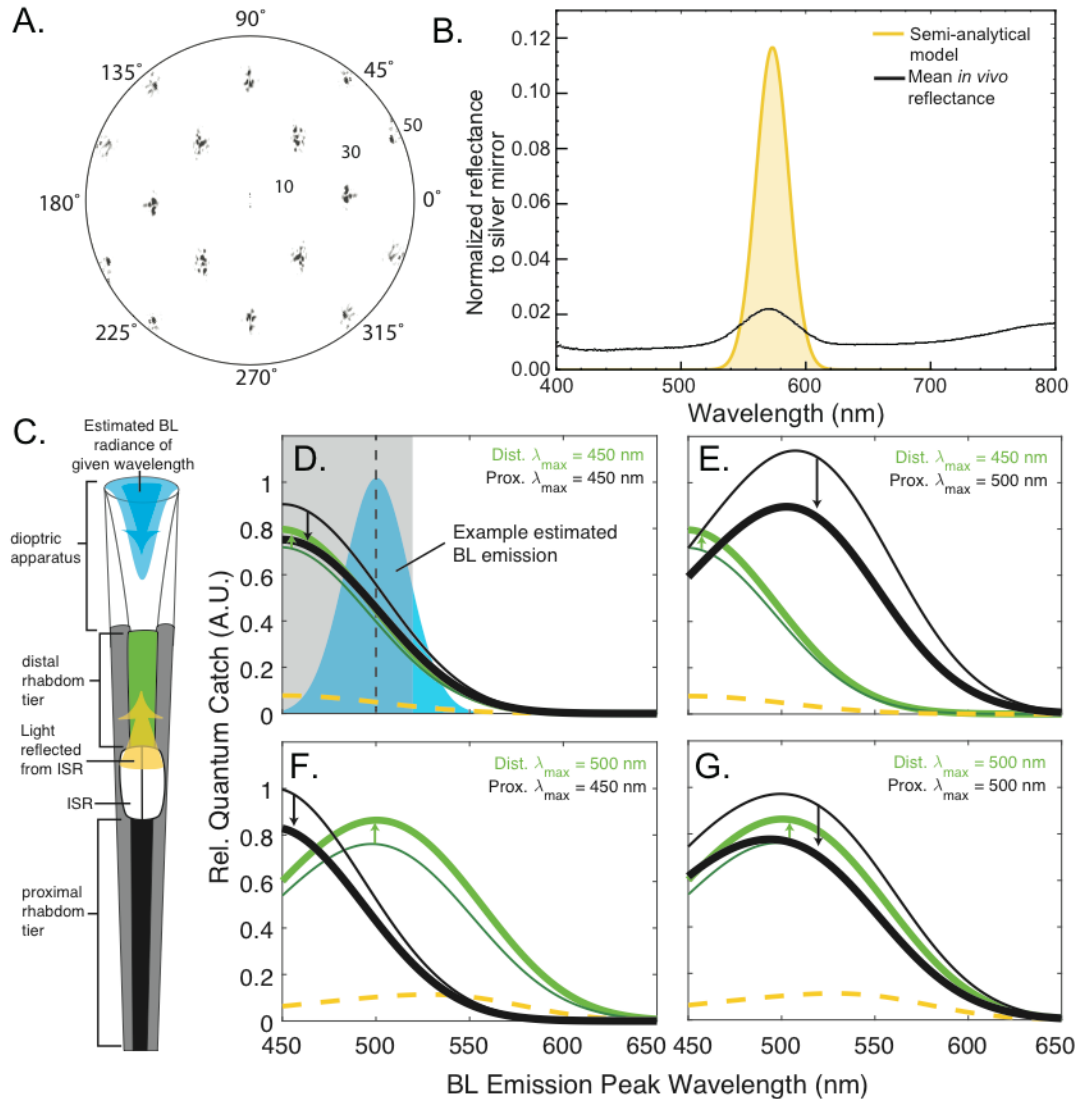


Fig. 4. Results of two optical models (A) Two-dimensional structure factor unveiling the FCC packing of the vesicles in the ISR, measured from TEMs of unknown nannosquillid 1 retina. (B) Comparison between the optical response of the ISR predicted by our semi-analytical model and the experimental data, yellow and black curve, respectively. The model indicates an exact correspondence between the predicted wavelength of maximum reflection and the observed wavelength-selective response. Both calculated and measured reflectances were mathematically normalized to a silver mirror. (C) Diagram representing different components of QC model to BL light sources modeled with and without experimental ISR reflectance. Colors correspond to traces in D-G (D-G) Relative proportion of photons captured from different BL emission spectra. Calculations for distal and proximal tier contain different

combinations of visual pigments with peak absorbances (λ_{max}) of 450 nm and 500 nm (combinations indicated top right of each panel). Left two panels (D & F) contain one visual pigment; Right two panels (E & G) contain two different visual pigments. Green traces, QC calculated for distal (Dist.) tier rhabdom; blue traces, QC calculated for proximal (Prox.) tier rhabdom. Thin line, QC calculated for ommatidium without ISR; thick line, QC calculated for ommatidium with ISR; yellow dashed line, amount of light reflected from ISR that is absorbed by the receptor. Example BL emission spectrum with peak emission at 500 nm provided in blue; intersection of black dashed line with green curves represents QC calculated to the 500 nm peaking BL spectrum. Grey shading denotes range of biologically relevant estimated BL emission spectra in the shallow pelagic environment reviewed in [35] (475 nm – 520 nm).

ACKNOWLEDGEMENTS

General: We would like to thank the following people for help with this work: Animal collection and identification: Megan L Porter, Roy L Caldwell, Elliott P. Steele, Sitara Palecanda, Lindsley DeMelo, the Marine Resources Center at the Marine Biological Laboratory, and Australian Museum Lizard Island Research Station, run by Lyle Vale and Anne Hoggart; Spectral measurements and analysis: Tom Jordan, Justin Marshall, John Cataldi, Sonke Johnsen; Statistical analyses: Jenny Gumm, Rachael Feord; Manuscript feedback: Paloma Gonzalez-Bellido, Alice Chiou, Camilla Sharkey.

Funding: US Air Force Office of Scientific Research (FA8655-12-2112 and FA9550-12-1-0321) and a travel grant from the Australian Microscopy and Microanalysis Research Facility Travel and Access Program (2010)

Author contributions: KDF made the initial discovery and performed experiments. KDF, TWC, NWR designed the experiments and analyzed the data and wrote and edited the manuscript. TEM by KDF and JM; photonic modelling by GJ & SV; BL modeling by DW; and 2-photon microscopy prep by KDF and imaging by TJW. Figures created by KDF with additions by DW, GJ, and revisions by co-authors

Competing interests: Authors declare there are no competing interests.

SUPPLEMENTAL MATERIALS

Materials and Methods

Animals & DNA Barcode Identification: All animals were captured at night while wading in shallow water (1-2 m depth) with dip nets and underwater lights at Lizard Island Research Station (August 2015; Queensland, Australia). Stomatopod larvae were sorted from the total plankton catch and stored in cups of seawater until completion of in vivo measurements. Identification was accomplished using previously published methods for DNA barcoding [28]. In brief, telson tissue was fixed in absolute ethanol and transported to the University of Bristol where total DNA was isolated (DNA XS, Machery-Nagel) and DNA amplicons of the cytochrome oxidase 1 mitochondrial gene were generated. Amplicons were sequenced (Eurofins

Genomics, UK) and aligned to a database of manually curated stomatopod reference sequences (Geneious Pro 5.5.6) to generate a maximum likelihood phylogenetic tree (PHYML, [41]. Positive identification of a sample to a reference was given if the sequences exhibited either reciprocal monophyly or less than 3% sequence divergence. Several larval specimens were able to be identified to family (Nannosquilidae) due to their nesting within references of this taxonomic unit. These species were thus referred to as “unknown species 1” and “unknown species 2” since a reference barcode does not currently exist to the species level for this group of samples. The two different species of Unknown Nannosquillid found, represent one that is unique to this study and one that corresponds to a nannosquillid larval species found in a previous study of larval photoreceptors (KM982429, [16]. The same assignments were also done for two larval sequences nested within the Squilloid superfamily, referred to as “unknown squilloids.” A complete list of sample sequences and their Genbank accession numbers is provided in Supplemental Table S1. Reference stomatopod DNA barcode sequences compiled from Genbank. One sequenced larval species, *Pullosquillia litoralis*, lacks a published adult reference CO1 sequence and was identified via direct collection from hatched clutches collected at the University of California Berkeley Gump Station (Moorea, French Polynesia). *Coronis scolopendra* larvae were hatched from morphologically identified and isolated adults, therefore they did not necessitate the use of DNA barcoding methods for identification.

***In vivo* Reflectance Measurements:** In the field, a Leitz Dialux 22 compound microscope (Wetzlar, Germany) was modified to fit two custom machined optical fiber adaptors that were mounted in place of the eye pieces. An optical fiber (1000 um UV-Vis, Ocean Optics) was attached to each adaptor and connected to either a white light source (tungsten LS-1, Ocean Optics) or a spectrometer (QE6500, Ocean Optics) connected to a computer (MacBook Pro, Apple) running SpectraSuite software (Ocean Optics). The z-plane of each optical fiber adaptor was adjustable so that the focal point of each optical fiber could be fixed through the objective. This was accomplished by connecting the light source to one adaptor and focusing the beam onto a white standard while looking down the eye piece, then doing the same procedure in the reverse position for the second optical fiber adaptor. Once both optical fibers were focused, all imaging took place through the digital camera mount (Canon D-series). Live larvae were fixed to sticks with cyanoacrylate adhesive and mounted in a calibrated angle holder. The mounted larva was then immersed in a tray of seawater and imaged via a submerged 10× microscope objective (Carl Zeiss 4820813, West Germany) protected with clear plastic film (Fig. S2A). A goose-neck laboratory lamp was used to illuminate the specimen from the side for focusing purposes but was turned off for all spectral measurements. A Spectralon diffuse white standard (Labsphere; North Sutton, NH, USA) was affixed near the larva to generate a 100% reflectance measurement from the epi-illumination light source through the microscope. Dark (0% reflectance) measurements were collected by shunting the light to the camera viewing mount. In this way, all internal reflections through the microscope system were removed from the reflectance measurement. Once dark and white standards were generated, the pseudopupil of the larval compound eye was

brought into focus and spectral reflectances recorded. Off-optical axis, epi-illumination measurements of regions of the eye outside the pseudopupil produced similar reflectance spectra from the eye camouflage as were previously recorded using 45° side-illumination [18], though reflectivity was greater in the epi-illumination system than when measured using side-illumination (i.e. Fig. 3G vs. [18]). Epi-illuminated reflectances of the pseudopupil often contained contributions from the off-axis reflective camouflage due to the size of the illumination spot and reflectance sample area exceeding the size of the pseudopupil, which contained four to six ommatidia. On-axis pseudopupil and off-axis reflectance measurements were collected from the ventral, lateral and dorsal region of each eye. Photographs were also collected of each specimen in on- and off-illumination conditions by setting a partial shunt between the camera and the eyepiece connected to the light source (Fig. 3; Movie S3). Spectra were plotted and analysed using Matlab2015b to generate total ISR reflectance averages as well as an average pseudopupil reflectances from each ISR-expressing species measured (Fig. S2B-E)

Electron & Two-Photon Microscopy: Eyes were dissected from live animals in the field immediately post reflectance measurements and fixed with 2.5% glutaraldehyde in PEMS buffer [0.05 M PIPES, 0.05 M PIPES 2K, 10 mM EGTA, 0.5 mM MgCl₂, and 3.8% sucrose, pH 7.0; based on [42] for 60 min with intermediate gentle agitation. Remaining tissue was fixed in absolute ethanol for DNA barcoding identification. Eye tissue was then washed 3 x 10 min in PEMS buffer and stored in fresh buffer at 4°C for transport to University of Maryland Baltimore County (UMBC) Porter Imaging Facility or University of Bristol Wolfson Imaging Facility. The remaining body tissue was fixed in absolute ethanol for DNA barcoding for identification post hoc. Eyes were postfixes for 60 min in 1.0% osmium tetroxide in water, washed 3 x 10 min in megapure water, then gently agitated in a second postfix of 2% uranyl acetate aqueous solution for 60 min. Samples were dehydrated in an ethanol series before being transferred to propylene oxide for slow infiltration with Epon resin (Taab). Each infiltration step occurred for a minimum of three hours followed by a final embedding in 100% resin overnight. Tissue was then transferred to fresh resin and polymerized 7-8 hours at 70 °C. Ultrathin sections (70-80 nm thickness) were cut using a diamond knife, and imaged via a Zeiss 10 CA transmission electron microscope at 60 kV (UMBC) or a FEI Tecnai 12 120kV TEM (Bristol).

TEM micrographs were analysed using ImageJ Fiji software [43] to measure dimensions of ISR structures, vesicles, and rhabdoms. Since vesicles were often observed as ellipsoid in TEM micrographs, the average diameter (d) of each ISR vesicle was determined by measuring the area (A) of an ellipse drawn along the membrane of the the two-dimensional image of each vesicle and using the equation

$$d = 2 \sqrt{\frac{A}{\pi}}$$

The nature of this method for determining diameter resulted in the banded pattern of recorded values observed in Fig S1A, especially in images where the scale was set in microns.

Two unknown nannosquillid species 1 specimens were used for conducting TEM tomography experiment (LI15001 and LI15008). Serial sections of 300 nm thickness, were cut from Epon embedded tissue and mounted on open-slot copper grids coated with Formvar film. Prior to visualization, 15nm gold fiducials (Aurion, NL) were applied to each side of the grid. The sample was then mounted in a Fischione tomography sample holder and the images recorded on an FEI 4k x 4k Eagle camera using FEI Tomography acquisition software in an FEI Tecnai 20, fitted with a LaB6 filament and operated at 200kV. Tilt series were then reconstructed using IMOD software (Boulder, Colorado) and segmented using AMIRA software (FEI Visualization Sciences Group, Bordeaux).

An individual from unknown nannosquillid species 1 (LI15030) and a first stage *Coronis scolopendra* larva hatched from captive adults in the laboratory (Marine Biological Laboratory, Woods Hole, MA) were fixed, dehydrated, and cleared for two-photon microscopy using previously published methods [44]. Imaging of eye structures was completed using an Olympus XLSLPLN25XGMP objective, exciting autofluorescence using a Newport Spectra-Physics InSight® DS+™ laser at 810 nm, and generating average images (from 32 frames) with a resonant scanner as part of a Bruker (Prairie Technologies) Ultima IV *In Vivo* Microscope using GFP and RFP detection channels. The voxel resolution was 0.2 μm in X, Y and Z. Images were compiled and animated using Vaa3D [45] and Fiji software.

Photonic Model: To understand the origin of the wavelength-selective response observed from the pseudopupil we developed a semi-analytical model. Our model combines an analytical approach (Bragg's law [29], and numerical simulations (Finite distance time domain, FDTD) to describe the optical properties of the ISR.

First, to estimate the structural parameters of the ISR we measured its structure factor starting from TEM images. All TEM and reflectance data used in the model are from unknown nannosquillid species 1, which provided the most complete set of data. The structure factor ($S(q)$) is formally defined as

$$S(q) = \frac{1}{N} \left\langle \sum_{i,j=1}^N e^{-iq \cdot (r_i - r_j)} \right\rangle,$$

where q is the wave vector, N the total number of particles, $r_{i,j}$ is the position of the vesicles labeled i and j , and the summation equation $\langle \dots \rangle$ denotes ensemble average. The structure factor allowed us to quantify the positional disorder of the ISR (Fig. 4A). From the structural factor we then integrated $S(q)$ along the direction connecting to vesicles to estimated the lattice constant (α), which was determined to be 392 ± 7 nm.

Our analytical approach used Bragg's law to predict wavelength position (λ) of the optical response of the periodic crystal to incident light

$$\lambda = a' * n_{eff},$$

where n_{eff} is the effective refractive index, defined by $n_{eff} = \sqrt{\phi * n_v^2 + (1 - \phi) * n_m^2}$ with the refractive index of the vesicles (n_v) and the refractive index of the surrounding matrix (n_m) are estimated to be 1.48 (membrane) and 1.35 (cytoplasm), respectively. The constant ϕ is the filling fraction of the FCC crystal (i.e. the fraction of the crystal structure occupied by the vesicles) which is defined as $\phi = \frac{\frac{16}{3}\pi*(r')^3}{(a')^3}$. The variables a and r are the size of the crystal cell and the radius of the vesicles, respectively, adjusted such that $a' = a * \psi$ and $r' = r * \psi$. The variable ψ is a fitting parameter that accounts for artificial compression during the sample preparation for the TEM imaging [46]. In these experiments ψ was equal to 1.08, which translates to an estimation of 8% compression, or shrinkage, of the tissue.

Though Bragg's law is powerful tool for analytical analysis of the interaction between light and a material, it is not sufficient for determining intensity or spectral width of the reflection. For this reason we also performed numerical FDTD simulations of the optical response of the ISR using the commercial software LUMERICAL 8.18 (Lumerical 4 Solutions Inc., Vancouver, BC, Canada). These calculations incorporate values of vesicle periodicity (or distribution within the ISR), vesicle size, and the dimensions of the entire ISR structure. Using this approach, we also took into account the finite numerical aperture (NA) of the objective used during the *in vivo* reflectance experiments (NA=0.22), which strongly affects the intensity and width of the response from a periodic system (see Fig. S2H).

To incorporate positional disorder and particle size distribution of vesicles within the ISR structure, we then combined the information generated from the FDTD simulations on predicted reflectance spectral shape with the predicted peak locations determined from Bragg's Law (Fig. S2I-J). To do this, we calculated the peak position for each value of a and r allowed by their distributions, with mean and standard deviation of 392 ± 7 nm and 82 ± 5 nm, respectively. Then, we weighted the numerical line shape with the probability of each value of a and r , such as the value of the associated normalized probability density function. The resulting line shapes (grey lines in Fig. S2I-J) were then fitted with a Gaussian curve that conserves the area of the starting numerical line shape. An unexpected finding in our calculations was that numerical aperture has a larger impact on the calculated optical response than the vesicle size distribution in the structure (Fig. S2H & I). In contrast to this finding, when we account for positional disorder in the calculations, the result is a broader and less intense reflection peak (Fig. S2J).

Experimental reflectance spectra data collected from live larvae (Fig. 3G) used a white diffuser to generate white light reference standards. These experimental data thus needed to be normalized against the total incident radiation (silver mirror) to allow for direct comparison with the theoretical line shape. To do this, we calculated the fraction of light reflected by a white

diffuser with a numerical aperture (NA) equivalent to the experimental condition (NA=0.22) using Lambert's Law

$$R(\theta_e) = \frac{\int_0^{\theta_e} \cos(\theta) \sin(\theta) d\theta}{\int_0^{90} \cos(\theta) \sin(\theta) d\theta}.$$

Where θ_e is the angle of a cone of reflectance off the standard white diffuser, and $R(\theta_e)$ is the fraction of light reflected by a standard white diffuser within the defined cone. For a NA of 0.22, θ_e was 9.8° and $R(\theta_e)$ was 0.03. The experimental data were then multiplied by $R(\theta_e)$ to obtain values of reflectance referenced to the total incident radiation.

Even after normalizing the data and accounting for the measured disorder, there was a substantial difference in the intensity between the experimental data and our model (Fig. 3B), which may arise as the result of several possibilities which are not mutually exclusive. First, due to distortion of the tissue during histological preparation, there may be error in the measured size of the ISR itself. Increasing the thickness of the ISR produces an increase in modeled reflectance (Fig. S2G). It may be that *in vivo* the crystals are much shorter in length than was observed in histological sections, producing a less intense reflection (Fig. S2G). A second potential source of the difference between theory and experiments is the uncertainty surrounding the refractive indices of the ISR materials (vesicles and matrix). If the ISR is composed of materials differing in refractive index from the predicted membrane and cytoplasm, the optical performance of the material would change.

Since the chemical identity of the membrane vesicles or their contents are not presently known, it was necessary to consider a range of refractive index values for potential biological materials. Fig S2G depicts the predicted optical response of the ISR with two different values for vesicle refractive index (n_v), 1.48 and 1.70, which model lipidic and pteridine-based systems, respectively. Pteridine was selected as a candidate vesicle material (alternate to lipidic) due to its presence in other crustacean photonic systems [30-32]. Changing n_v from 1.48 to 1.70 increases the effective refractive index of the system from 1.35 to 1.39, resulting in both a red-shift of the peak position and an increase in reflectivity. A slight decrease in the shrinkage constant (ψ) from 8% to 5% was applied to offset the red-shift in the higher n_v model so as to align both modeled and experimental reflectance spectra to the same peak for spectral shape comparison (Fig S2G). Even when the wavelength of peak reflectance is the same, use of $n_v=1.70$ creates a greater deviation of the model width and intensity from the observed reflectance (Fig. S3E), leading to the conclusion that 1.48 more closely mimics the actual refractive index of the material within the vesicles.

A third reason for reflectivity differences between the observed and predicted data is related to the true level of disorder in the orientation and periodicity of the 3D lattice of the structure. This information was neither accessible by two-dimensional image data, such as TEM, nor the limited thickness able to be sampled using TEM-tomography. Thus, the true values of disorder among the 3D lattice is currently unknown. The final reason for differences in the calculated and measured reflectances is that we do not know the true NA of the light returning from the ISR. In the measured reflection, a fraction of the incident light is absorbed into the rhabdom as well as

refracted through the animal's light focusing units (crystalline cones and lens). Since our calculations reveal that NA has a substantial impact on the calculated reflectance (Fig S2H), this may be a major source of disparity between observed and predicted spectra.

Quantal Catch Modeling: Mathematical model scripts were compiled in Matlab2017b and are provide as supplemental, annotated documents (visualModelfinal.m; a1SSHtemplate.m). In the visual model, the change in light flux, I , as a function of distance from the start of the rhabdom, z , was calculated by

$$\frac{dI}{dz} = -\{\eta(\lambda)f(z)\kappa_v(z)\alpha_v(\lambda) + [1 - \eta(\lambda)]\kappa_s(z)\alpha_s(\lambda)\}I(z, \lambda),$$

from Arikawa et al. (1999, [38]. Where η is the fraction of the light flux which is inside the rhabdom boundary as a function of wavelength, λ ; f is the fraction of the rhabdom cross-section taken up by a single photoreceptor (in this case $f = 1$); κ_v and κ_s are the peak absorption coefficients of the visual pigment and screening pigments respectively; and α_v and α_s are the normalized absorption spectra. $\alpha_v(\lambda)$ was modelled using a visual pigment absorption template1 [47]. The fraction of light within the rhabdom therefore was approximated by: $\eta(V) = a - be^{-cV}$ with $a=0.96$, $b=2.82$, $c=1.27$ and the waveguide parameter, $V = \pi d(n_1^2 - n_2^2)^{1/2}/\lambda$; the rhabdom internal refractive index, $n_1=1.36$ and the surrounding refractive index $n_2=1.34$. Further model parameters are summarized in Fig. S3 and Supplemental Table 3.

To test the impact of the ISR on light detection, we used a mathematical model of light absorption in rhabdomeric photoreceptors surrounded by screening pigments [38]. Previously, yellow, long-pass lateral screening pigments were described in the distal retinas of Nannosquillid larvae (Fig. 1, [27]. Since the transmission spectrum of these pigments is similar to the reflectance spectrum of the ISR, we incorporated these data into our model of light absorption in the photoreceptors (for transmission spectrum see [27]. The light source used to pass along the photoreceptor in this model was an estimated either the irradiance spectrum of moonlight at 3 m depth (digitized from [48]) and normalized to 1 for full moon conditions, or as bioluminescent radiance. The bioluminescent (BL) emission spectrum defined by a Gaussian curve with a full-width at half-maximum of 66 nm (Fig. S3C). This estimated BL was determined by examining the spectral width of common bioluminescent emission spectra in the literature [39, 49]. Since the specific bioluminescent target viewed by nannosquillid retinas is not known, we calculated the quantal catch (QC) for each estimated BL emission spectrum with a peak between 450 nm and 650 nm. Since the range of peak BL emissions in the shallow, pelagic environment range from 450 nm to 520 nm (gray shading Fig. 4D & E, [35], the QC calculations outside this range are theoretical. Waveguide properties of the rhabdoms and values used in the optical model are reported in Fig S3D and Table S2, respectively.

The light absorbed by the visual pigment and lateral screening pigments was calculated per unit length of the rhabdom along with the resultant attenuation of the light. Light reaching the ISR was either reflected back along the distal tier, with a model reflection spectrum, $R(\lambda)$, calculated

for a conservative 50% reflectivity (Fig. 4) or transmitted through to the proximal tier, with transmission spectrum = 1-, since the absorption spectrum of the IPC was assumed to be nil. The light absorbed by the visual pigment was integrated along the length of the photoreceptor and taken as a measure of relative quantum catch (QC). QC was calculated as a function of the peak wavelength of the target Gaussian light source (each estimated BL). Two visual pigments have been measured from different species of Nannosquillid larvae, with peak absorption wavelengths of 450 nm and 500nm [16]. Calculations were thus performed using all potential combinations of opsin templates for these two visual pigments in the proximal and distal rhabdomeric tiers (Fig. S3A).

For the BL radiance calculations, the ISR reflection produced little to no effect on the QC of the distal tier in the 450 nm λ_{max} visual pigment condition due to comparatively minimal overlap between the reflectance and absorption spectra (Fig. 4D). Contrastingly, a rhabdom containing a 500 nm λ_{max} visual pigment is more strongly affected by the ISR (Fig. 4E). In this visual pigment condition, QC begins to increase to occur at estimated BL emission spectra that peak at 475 nm and longer, which corresponds to the range of BL emissions in the nannosquillid larval habitat (shallow pelagic zone). For animals who are active in dim light, this demonstrates that the ISR has the potential to increase nannosquillid larval capture of biologically relevant photons in the distal rhabdomeric tier. The effect on the proximal tier was different, due to the predicted filtering effect of long wavelengths by ISR reflectance. When the proximal tier expressed the visual pigment with λ_{max} 500 nm, the QC was decreased, while expression of the λ_{max} 450 nm was unaffected, regardless of the distal tier visual pigment expression. Though all marine crustacean larvae (including mantis shrimp) are understood to have a single class of photoreceptor in their retinas [16, 17, 24], were this the case in the tiered nannosquillia retina then the animal would experience a loss of photon, and decreased sensitivity of the proximal rhabdom. If, however, the spectral sensitivity of each tier were different, with a λ_{max} 500 nm expressed distally and λ_{max} 450 nm expressed proximally, then there would be no net loss of photons by the eye.

The results of the calculations with moonlight irradiance revealed that during the full-moon at 3 m depth, there is only a 1.2% increase in QC when the distal tier visual pigment absorbs maximally at 450 nm. In the presence of a 500 nm λ_{max} visual pigment, however, the ISR improves the relative QC in the distal tier by up to 18%, though this gain would be diminished to 1.8% during the quarter phase and further to 0.18% during the new moon. While a 18% gain in signal would be substantial for improving sensitivity to the background light near the surface, this is only during the few days when the moon is at its maximum face each month. While we cannot reject the hypothesis that the ISR evolved for improving sensitivity to moonlight, the fact that larvae do not rise to the surface during the full moon and the coincidence of ISR reflectance wavelengths with known BL tuning systems led us to conclude that BL is the more likely visual

target. These results provide concrete hypotheses with which to probe the visual physiology of
nannosquillid eyes, using behavioral, electrophysiological, and/or molecular methods.

Statistical Analysis: All statistical analyses were conducted in Matlab2015b using custom
scripts. One-way ANOVA with Tukey HSD post hoc statistical analyses were conducted
measurements of ISR vesicle size ($n_{\text{Puli}} = 629$, $n_{\text{Alvi}} = 315$, $n_{\text{Puth}} = 631$, $n_{\text{UKNan1}} = 499$, $F_{3,2069} =$
 127.43 , $p = 9.77\text{e-}76$), ISR total length ($n_{\text{Puli}} = 23$, $n_{\text{Alvi}} = 28$, $n_{\text{Puth}} = 53$, $n_{\text{UKNan1}} = 43$, $F_{3,146}$
 $= 53.81$, $p = 2.41\text{e-}23$), and ISR total width ($n_{\text{Puli}} = 85$, $n_{\text{Alvi}} = 46$, $n_{\text{Puth}} = 43$, $n_{\text{UKNan1}} = 60$, $F_{3,233}$
 $= 63.18$, $p = 7.85\text{e-}30$) among species. Results of post hoc tests, original data points, mean,
standard deviation, and standard error of the mean are reported in Fig S1A-C using notBoxPlot
Matlab function (version 1.31.0.0, Rob Cambell). Histograms of combined values measured
from all species for each ISR variable (vesicle diameter, ISR length, and ISR width) were plotted
with Matlab2015b to show distribution of total anatomical measurement data. Similar methods
were used to test for significant differences in wavelength of peak reflectance from ISR-
containing ommatidia among species (ANOVA, $n_{\text{UKNan1}} = 40$, $n_{\text{Puth}} = 3$, $n_{\text{Alvi}} = 2$, $F_{1,43} = 0.01$, $p =$
 0.923).

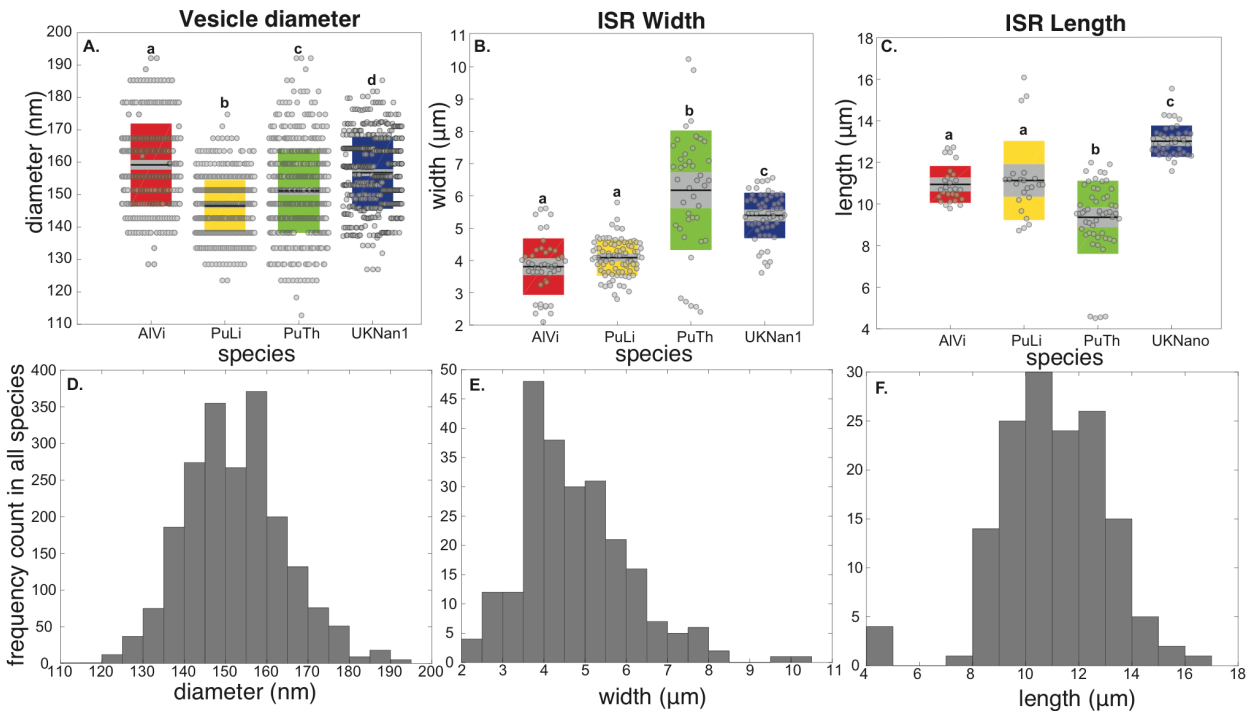


Fig. S1. Dimensions of intrarhabdomal photonic structure (ISR) features of different species measured from TEM micrographs. (A) ISR vesicle diameter measurements by species. Among species, means ranged from 146.5 nm to 159.2 nm. Banding pattern of data points are a result of the method used to calculate diameter. (B) The total width of ISRs in each species. Means range among species from 3.8 μm to 6.2 μm . (C) The total length of ISR structures by species. Means range among species from 9.4 μm to 13.0 μm among species. (A-C) Colors correspond to each species: Red, AlVi, *Alachosquilla vicina*; yellow, PuLi, *Pullosquilla litoralis*; green, PuTh, *Pullosquilla thomassini*; blue, UKNan1, unknown nannosquillid species 1. Same for B-C. Lowercase letters indicate significance groups determined by one way ANOVA with Tukey HSD post hoc analysis. Black, solid lines represent mean value. Red, yellow, green and blue colored patches depict 1 standard deviation from mean. Grey patches, 95% confidence interval (or 1.96 standard error of the mean). Grey dots, individual data points. (D-F) Histograms showing the combined distribution of vesicle diameter, ISR diameter, and ISR length from all species.

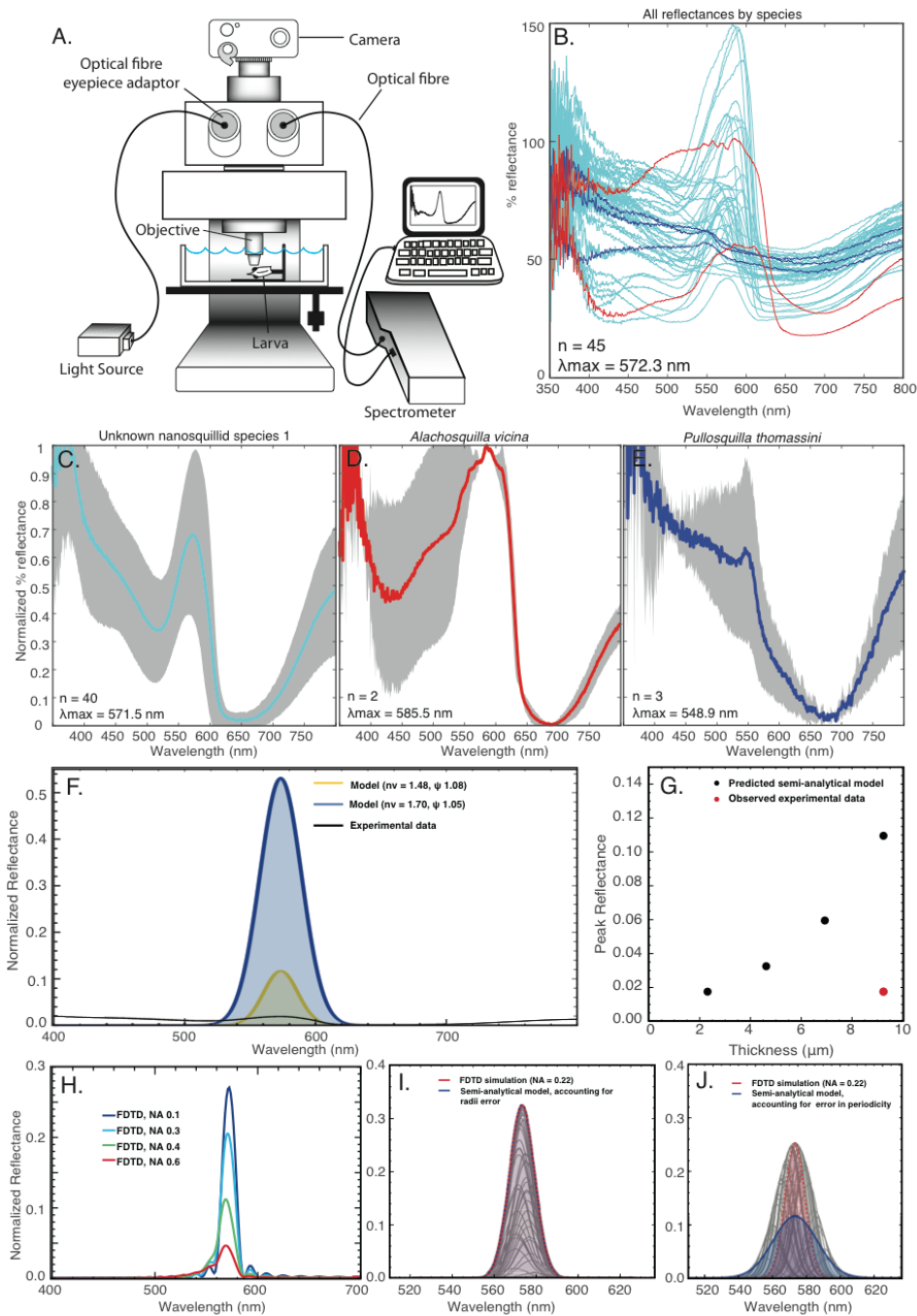


Fig. S2. Measured and modeled light reflected from the ISR (A) diagram of epi-reflectance microscope set-up. Objective has numerical aperture of 0.22, which is relevant to reflectance model in H. (B) All reflectances measured from illuminated pseudopupil of Nannosquillid larvae. Variation in reflectivity are likely the result of difference in ommatidial size, objective alignment with pseudopupil, contributions from the blue reflective camouflage that surrounds each ommatidium, as well as variables evaluated in F-J. Trace colors correspond to species in C-E. (C) Average pseudopupil reflectance from unknown Nannosquillid species 1 (D) *Alachosquilla vicina* and (E) *Pullosquilla thomassini*. n values and average wavelength of peak

reflectance ($\lambda_{\max}$) reported in the bottom left corner of each figure. **(F)** Peak intensity calculated as a function of the thickness of the ISR. To retrieve a peak reflectance comparable to that of the experimental data it is necessary to reduce the thickness of the ISR. Thus, the difference in observed and predicted thickness may related to distortion of the tissue during histological preparation. **(G)** Results of modeled reflectance using two different refractive index values (n_v) and associated expansion factors (ψ) versus experimental reflectance (black trace). **(H)** Simulated reflectance as a function of the numerical aperture (NA). Note how reflectivity changes as a function of numerical aperature of the objective used to collect the reflected light. **(I)** Comparison of semi-analytical curves for different values of vescicles' radii **(J)** and ISR periodicity.

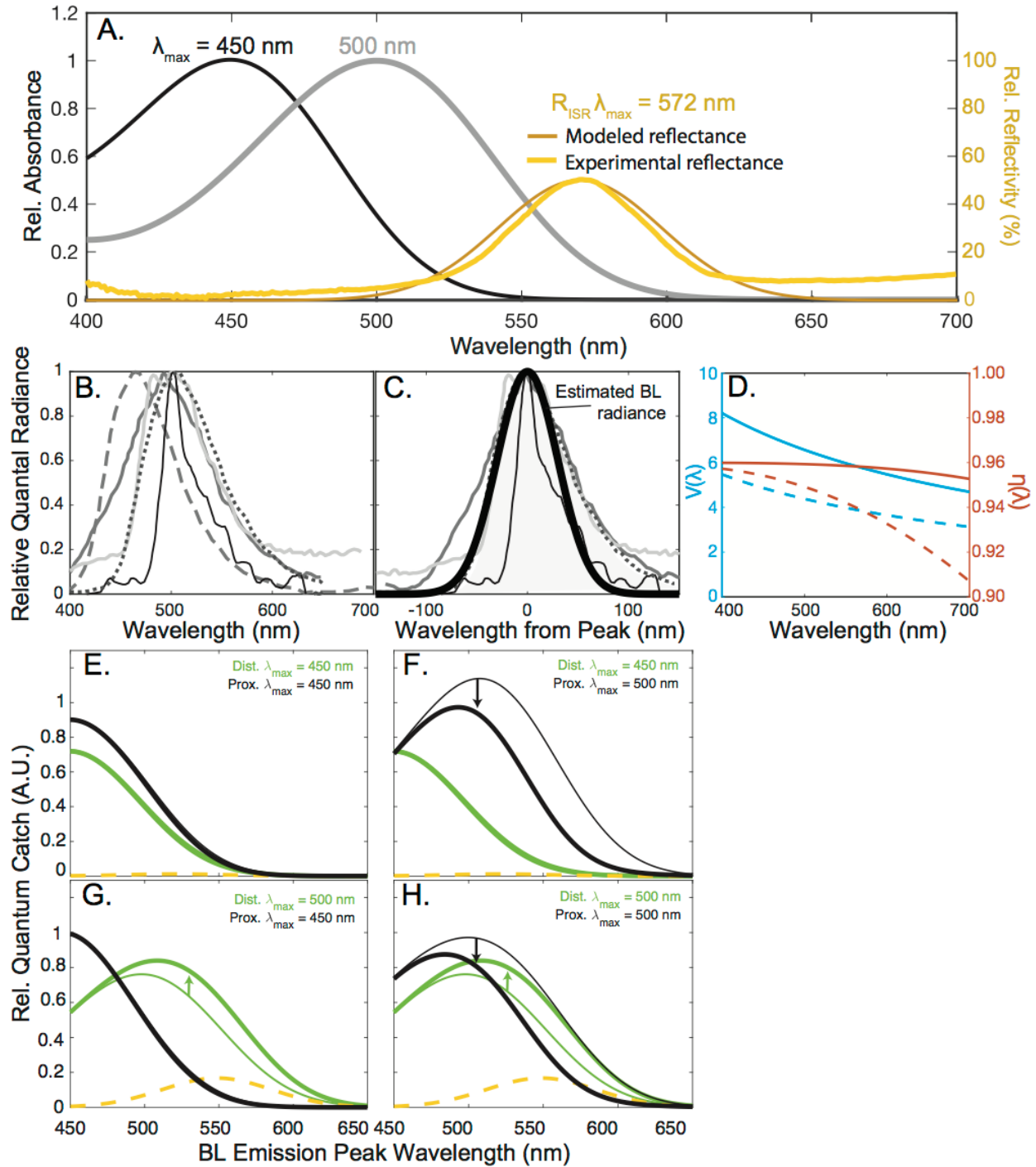


Fig. S3. QC model spectra, waveguide properties, and results calculated using Gaussian model of ISR reflectance. (A) 450 nm and 500 nm peak absorbing opsin template used to describe two possible visual pigment absorptions in the distal tier. These absorption values were selected based on nannosquid visual pigment data from the literature [16, 27]. Thin, dark yellow curve, Gaussian model of ISR reflectance used to calculate QC in E-H. Thick, light yellow curve, average reflectance measurements reported in this paper. Both modeled and measured reflectances calculated for 50% reflectivity (B) Sample spectra of bioluminescent sources as a function of wavelength digitized from the literature: dark grey long dash line – Ostracod,

Vargula hilgendorf [39]; dark grey solid line – Siphonophore, Bargmannia elongata [49]; light grey solid line – Midshipman, Porichthys motatus [39]; thin black line – Medusa, Clytia hemisphaericum [49]; dark grey short dash line – Ctenophore, mertensiid [49]. **(C)** The same sample spectra from (B) plotted relative to peak wavelength. Thick black curve shows a 66 nm FWHM Gaussian used in the quantum catch model to approximate different bioluminescent emission spectra. **(D)** The waveguide parameter V (shown in blue), for the distal (solid) and proximal (dashed) rhabdom tiers. For the parameters used here, the rhabdom is multi-mode ($V > 2.405$) for all visible wavelengths. The fraction of light inside the rhabdom, (shown in red) is between 0.9 and 0.96 across the spectrum for both the distal (solid) and proximal (dashed) rhabdom tiers.

Table S1. Larval stomatopod specimens used in TEM, 2-photon, and *in vivo* reflectance experiments. Most larvae were captured from the wild and identified post hoc by DNA barcoding, with the exception of *Coronis scolopendra*, whose larvae were hatched from adults captive in the laboratory. Pseudopupil reflectances were observed in all nannosquillid larvae tested, with the exception of one *P. thomassini* postlarva, likely due to the degeneration of the larval retina after metamorphosis.

Species	Larval stage	Genbank #	TEM	2Photon	Reflectance
<i>Alima pacifica</i>	last stage	MK397440	x		
<i>Gonodactylellus affinis</i> (Feller & Cronin 2016)	mid stage	KM982428.1	x		
<i>Unknown nannosquillid 2</i> (Feller & Cronin 2016)	early stage	KM982429			
<i>Gonodactylus childi</i>	early stage	MK397441	x		
<i>Pullosquilla thomassini</i>	last stage	MK397442	x		
<i>Unknown nannosquillid 1</i>	mid stage	MK397443	x		+
<i>Unknown nannosquillid 1</i>	mid stage	MK397444	x		+
<i>Alachosquilla vicina</i>	early stage	MK397445	x		+
<i>Unknown squilloid</i>	early stage	MK397446			-
<i>Unknown nannosquillid 1</i>	mid stage	MK397447			+
<i>Gonodactylus smithii</i>	mid stage	MK397448			-
<i>Gonodactylellus affinis</i>	mid stage	MK397449			-

<i>Unknown nannosquillid 1</i>	early stage	MK397450	x	+
<i>Pullosquilla thomassini</i>	early stage	MK397451		+
<i>Unknown nannosquillid 1</i>	mid stage	MK397452		+
<i>Unknown nannosquillid 2</i>	mid stage	MK397453		+
<i>Pullosquilla thomassini</i>	post larva	MK397454	x	-
<i>Alima pacifica</i>	last stage	MK397455		-
<i>Unknown squilloid</i>	mid stage	MK397456		-
<i>Haptosquilla trispinosa</i>	mid stage	MK397457		-
<i>Chorisquilla hystrix</i>	mid stage	MK397458		-
<i>Unknown nannosquillid 1</i>	mid stage	MK397459		+
<i>Unknown nannosquillid 1</i>	early stage	MK397460		+
<i>Pullosquilla thomassini</i>	last stage	MK397461		+
<i>Pullosquilla thomassini</i>	last stage	MK397462		+
<i>Chorisquilla hystrix</i>	mid stage	MK397463		-
<i>Chorisquilla hystrix</i>	mid stage	MK397464		-
<i>Unknown nannosquillid 1</i>	mid stage	MK397465		+
<i>Odontodactylus cultrifer</i>	mid stage	MK397466		-
<i>Unknown nannosquillid 1</i>	early stage	MK397467	x	+
<i>Unknown nannosquillid 1</i>	mid stage	MK397468		+
<i>Unknown nannosquillid 1</i>	mid stage	MK397469		+
<i>Unknown nannosquillid 1</i>	early stage	MK397470		+
<i>Unknown nannosquillid 1</i>	mid stage	MK397471		+

Unknown nannosquillid 1	early stage	MK397472		+
<i>Pullosquilla litoralis</i>	early stage	MK397473	x	
<i>Pullosquilla litoralis</i>	early stage	MK397474	x	
<i>Pullosquilla litoralis</i>	early stage	MK397475		
<i>Coronis scolopendra</i>	first stage	hatch ID		x

Table S2. Summary of parameters used in visual optical model. Data is sourced from this investigation unless otherwise stated.

Parameter	Value	Source
Length, distal rhabdom	34 μm	-
Length, proximal rhabdom	72 μm	-
Diameter, distal rhabdom	4.5 μm	-
Diameter, proximal rhabdom	3.0 μm	-
	0.01 μm^{-1}	Cronin & Marshall (1989a,b), Warrant & Nilsson (1998)
	0.131 μm^{-1}	Jutte <i>et al.</i> (1998)
Peak wavelength of IPC reflection	572 nm	-

Movie S1: Z-stack of the eye of *Coronis scolopendra* first larval stage, measured using two-photon microscopy using autofluorescence of the fixed tissue. ISR structures are visible as slightly darkened sections of the rhabdoms, which appear bright. Untiered, non-ISR expressing ommatidia are visible in the dorsal region of the eye.

Movie S2: Three-dimensional arrangement of vesicles (yellow) within intrarhabdomal photonic structures (ISR), segmented from TEM tomography data. This order is preserved across the membranes of the four primary ISR cells (pink). Arrow indicates the direction of incoming, on-axis light.

Movie S3: Yellow reflection is produced from on-axis illumination of ISR containing ommatidia. This video was captured during the *in vivo* reflectance measurement experiments and demonstrates how the pseudopupil moves as the animal rotates the eye in response to changing illuminations under the objective.

Movie S4: Light microscopy of fixed, disassociated retinas reveal that the ISRs do not contain colorful, photostable pigments, which is unlike adult mantis shrimp intrarhabdomal filters. Scale bar, 10 μm

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