1 Title

2 Gadusol is a maternally provided sunscreen that protects fish embryos from DNA damage

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4 Authors

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13

14 Abstract

15 Ultraviolet radiation (UVR) and its deleterious effects on living cells selects for UVR-protective

16 mechanisms. Organisms across the tree of life evolved a variety of natural sunscreens to

17 prevent UVR-induced cellular damage and stress. However, in vertebrates, only melanin is

18 known to act as a sunscreen. Here we demonstrate that gadusol, a transparent compound

19 discovered over 40 years ago in fish eggs, is a maternally provided sunscreen required for

20 survival of embryonic and larval zebrafish exposed to UVR. Mutating an enzyme involved in

21 gadusol biosynthesis increases the formation of cyclobutane pyrimidine dimers, a hallmark of

22 UVB-induced DNA damage. Compared to the contributions of melanin and the chorion, gadusol

is the primary sunscreening mechanism in embryonic and larval fish. The gadusol biosynthetic
 pathway is retained in the vast majority of teleost genomes but is repeatedly lost in species

24 pathway is retained in the vast majority of teleost genomes but is repeatedly lost in species 25 whose young are no longer exposed to UVR. Our data demonstrate that gadusol is a maternally

provided sunscreen that is critical for early-life survival in the most species-rich branch of the

27 vertebrate phylogeny.

28

30 Introduction

31 Most life on earth relies on photosynthetic food webs for their energy source, which can result in extensive exposure to ultraviolet radiation (UVR)¹. UVR, especially UVB (280-320nm), can 32 damage proteins and DNA, leading to errors during DNA repair and replication. Excessive UVR 33 induces cellular death. Aquatic organisms risk UV exposure because biologically harmful levels 34 35 of UVB can penetrate >10 m in clear water². Organisms in diverse habitats adapt to avoid, ameliorate, or protect against the effects of UVR. Some of these adaptations include sun 36 37 avoidance behaviors (e.g., nocturnal lifestyle) and DNA repair machinery (e.g., nucleotide 38 excision repair)³. However, since sunlit habitats can have significant nutritive advantages over 39 dark environments and because no repair pathway is completely efficient, many organisms 40 employ sunscreens to avoid UVR damage from occurring in the first place⁴. 41 42 Sunscreens absorb UV photons before they penetrate vulnerable cells and dissipate this absorbed energy as less harmful heat. A wide variety of sunscreens are used by living 43 organisms, including flavonoids in plants, scytonemin in cyanobacteria, and melanin in 44 45 numerous organisms including vertebrates⁴. In fish and other aquatic vertebrates, melanin is produced in melanophores (homologous to melanocytes in mammals), which differentiate from 46

47 embryonic neural crest cells and migrate to cover aspects of the brain and body⁵. Recently, an

internal melanophore umbrella was shown to protect the hematopoietic stem and progenitor cell

⁴⁹ niche in developing zebrafish from UVR⁶. However, since melanophores emerge late in

50 embryonic development, they cannot protect early stages when the embryo is most sensitive⁷,

and some controversies remain about melanin's sunscreening role in fish^{8–10}. Thus, the

52 mechanisms that may protect the initial phases of development in externally fertilized vertebrate

embryos (e.g., the vast majority of fish) remain mysterious. Apart from melanin, no

54 endogenously produced sunscreen has been documented in vertebrates.

55

56 Mycosporine-like amino acids (MAAs) are a class of sunscreening compounds produced by 57 numerous algae and microbes⁸. Experiments indicate that depletion of MAAs causes UVR

58 sensitivity in cyanobacteria¹¹ and sea urchins^{12,13}. The eggs of many fish species also contain

⁵⁹ large quantities of an MAA-related UVR-absorbent compound called gadusol (first discovered in

eggs of the cod *Gadus morhua*)^{14,15}. Although the existence of gadusol in fish eggs and

embryos was discovered decades ago, its role as a sunscreen remains untested¹⁶. Gadusol in fish was originally thought to come from dietary sources^{1,2}. However, a two-gene cassette was

fish was originally thought to come from dietary sources^{1,2}. However, a two-gene cassette was recently discovered in many vertebrate genomes that enables the production of gadusol from

64 sedoheptulose-7-phosphate (an intermediate in the pentose phosphate pathway)¹⁷. Yeast

65 engineered to express the zebrafish biosynthetic pathway produced gadusol, which provided

66 protection against UVR in yeast¹⁷. With an understanding of the teleost genetic architecture

67 underlying gadusol production, and a genome-editing toolkit for zebrafish, it is now possible to 68 test the role of gadusol as a sunscreen *in vivo*.

69

Here, we test the role of gadusol in UVR protection of fish embryos and larvae by generating a

71 gadusol-deficient mutant zebrafish. We determine that gadusol is maternally provided in

72 embryos and provides protection from UVR throughout embryonic and larval development. We

find that gadusol is the primary sunscreen during fish development while melanin and other

74 mechanisms provide secondary protection. In a broader evolutionary context, we also find a

striking pattern of repeated loss of gadusol production in fish species whose embryos are not

responsed to sunlight. Together, our work provides evidence that gadusol is a widely distributed

and evolutionarily conserved sunscreen that protects vertebrate embryos in aquatic sun-lit

- 78 environments.
- 79
- 80 Results

81 Gadusol is maternally provided and protects embryos and larvae from UVR

82 To test if gadusol is a sunscreen in vertebrate embryos, we used CRISPR-Cas9 to delete most

- of exon 2 of zebrafish *eevs*, which encodes the enzyme essential for the first step in gadusol
- biosynthesis (**Fig. S1**). We chose zebrafish for these experiments because they live and spawn
- in shallow sunlit waters, they are known to produce gadusol¹⁷, and they are genetically
- tractable. Grown in our animal facility, where they are protected from UVR, homozygous *eevs*
- 87 mutant females and males survived to fertile adulthood like their wild-type peers. Using
- reciprocal crosses between homozygous mutant adults ($eevs^{-/-}$) and wild-type adults ($eevs^{+/+}$),
- 89 we generated heterozygous mutant embryos that lack maternal contribution of gadusol
- 90 (hereafter referred to as Meevs) and heterozygous mutant embryos that retain this maternal
- contribution (referred to as $eevs^{+/-}$) (**Fig. 1A**). Notably, Meevs and $eevs^{+/-}$ embryos have identical
- 92 genotypes but either lack or possess maternally provided gadusol, as judged by mass
- spectrometry (Fig. 1A) and UV-spectrophotometry (Fig. S2). We generated maternal-zygotic
 homozygous mutant embryos (referred to as MZeevs) from in-crosses of homozygous mutant
- 95 parents. Immediately after fertilization, gadusol was nearly absent in MZeevs embryos and
- 96 indistinguishable from Meevs (**Fig. 1B**).

97 We next asked how long maternally provided gadusol persisted in embryos and larvae. We

compared gadusol abundances from whole embryos and larvae with the following genotypes:

99 *eevs*^{+/+} (wild-type), Meevs, and MZeevs. We found only a modest increase in gadusol

abundance in Meevs relative to MZeevs at 5 days post-fertilization (dpf), suggesting that

101 maternally synthesized and deposited gadusol is the source of nearly all gadusol in the

- developing zebrafish (Fig. 1B). This is an example of a maternal effect, where disruption of the
- *eevs* gene in mothers eliminates gadusol presence in their embryos, regardless of embryogenotype.
- 105 To determine if gadusol protects zebrafish embryos against UVB, we developed an assay to
- deliver precise doses of UVB to embryos and measure the effect on swim bladder inflation at 5
- dpf (a hallmark of healthy development essential for survival, **Fig. S3**). We found that 450 joules
- $(J)/m^2$ of UVB (fluence rate: 2.5 W/m², see **Methods**) delivered at 24 hours post-fertilization
- (hpf) resulted in ~75% swim bladder inflation in wild-type and $eevs^{+/-}$ embryos, respectively, but
- did not result in gross developmental defects (**Fig. S4**). In stark contrast, MZeevs and Meevs
- embryos were extremely vulnerable to the same dose of UVB; all embryos failed to inflate their
- swim bladders (**Fig. 1C**).
- 113 Since zygotic production of gadusol was still minimal at 5 dpf (**Fig. 1B**), we hypothesized that
- 114 larvae lacking maternal gadusol should be highly sensitive to UVB at this later stage. We

- repeated UVB dosage curves on 5 dpf larvae and identified 2.5 kJ for a small but significant
- impact on wild-type larvae survival (**Fig. S5**). We grew UV-exposed and control larvae in our
- 117 fish facility nursery to 28 dpf, which requires developing animals to forage for food to survive.
- 118 We found that only 2% of exposed Meevs larvae survived, compared to ~50% of controls
- exposed to the same dose of UVB (**Fig. 1D**). Together, these data demonstrate that maternally
- 120 provided gadusol provides powerful UVB protection to early embryos and older larvae.

121 Gadusol prevents DNA damage and apoptosis

- 122 Next, we sought to understand the mechanism by which gadusol protects embryos from UVB. In
- 123 other species, gadusol and related molecules were hypothesized to function as antioxidants as
- 124 well as sunscreens^{14,17,18}. To test if gadusol serves as an antioxidant in zebrafish embryos, we
- exposed 24 hpf embryos to hydrogen peroxide to induce oxidative stress. At 5 dpf, gadusol-
- depleted Meevs and control eevs^{+/-} embryos had similar responses to increasing doses of
- 127 oxidative stress, suggesting that gadusol does not function as an antioxidant *in vivo* (Fig. S6).
- 128 To test if gadusol serves as a sunscreen by absorbing UVB, we measured the production of
- 129 cyclobutane pyrimidine dimers (CPDs), a signature of UVB-induced DNA damage¹⁹. If gadusol
- acts as a sunscreen, then it would absorb UVB photons and shield the underlying DNA from
- 131 CPD formation. We exposed 24 hpf embryos to UVB and used immunohistochemistry to detect
- 132 CPDs and quantify fluorescence intensity. Embryos that lacked gadusol had significantly higher
- 133 levels of CPD formation after UVB exposure compared to controls containing gadusol (Fig.
- 134 **2A,B**). CPDs are cytotoxic and at high abundance induce apoptosis. We used
- immunohistochemistry to detect a fast-acting apoptotic marker (activated caspase-3) in embryos
- exposed to UVB²⁰ (Fig. 2C, Fig. S7). We found that embryos lacking gadusol had increased
- levels of apoptotic nuclei, relative to controls (**Fig. 2C**), supporting a role for gadusol in
- absorbing UVB and preventing DNA damage.
- 139 To characterize transcriptional responses to UVR in the absence of gadusol, we performed
- 140 RNAseq comparing gadusol-depleted Meevs and wild-type embryos. Five hours after exposure
- 141 to UVB, embryos lacking gadusol had significantly higher expression of many key stress
- response genes (*tp53, gadd45aa, ddb2, & cdkn1a*) relative to UVB-treated controls (**Fig. 2D**).
- 143 GO terms enriched in UV-exposed gadusol-depleted embryos included response to UV,
- response to DNA damage, response to light, and other stress response terms (**Fig. S8, Table**
- 145 **S1**). Together, our imaging and gene expression data confirm that gadusol in zebrafish embryos
- acts as a true sunscreen to provide efficient protection against UV-induced DNA damage,
- 147 cellular stress, and cell death.
- 148 Gadusol is the primary sunscreen in early fish development
- 149 In light of our finding that gadusol acts as a sunscreen, we compared the relative sunscreening
- potency of gadusol compared to other potential UV-blocking/absorbing mechanisms in larval
- zebrafish. Melanin is a well-known sunscreen in many organisms including humans. In
- zebrafish, melanophores become pigmented around 36 hpf, ultimately forming stripes that
- partially cover the larval brain and body, a pattern that is stable until ~14 dpf^{21,22}. Melanophores
- protect the hematopoietic niche in larval zebrafish⁶, but their role as a whole-body sunscreen
- remains untested. The *nacre/mitfa* mutant disrupts a key melanophore master regulator and

- 156 lacks melanophores. We generated two groups of larvae, each with pigmented and
- unpigmented siblings. One group contained no maternal gadusol, while the other group
- 158 contained gadusol (**Fig. 3A**). We treated all 5 dpf larvae with 2.5 kJ of UVB and assessed
- survival in the nursery at 28 dpf. Larvae with gadusol were highly resistant to UVB stress,
- regardless of pigmentation status (**Fig. 3B**). All larvae that lacked gadusol were highly sensitive
- to UVB, and larvae that lacked both gadusol and melanin were slightly more sensitive to UVB
- than their pigmented siblings. At a lower UVB dose (1.5 KJ), we also found a modest but
- significant effect of melanophores in protecting against UVB (Fig. S9). We conclude that while
- melanin plays a minor role in UVR protection, gadusol is the primary sunscreen in early fish
- 165 development.
- 166 Another potential UV-protective mechanism is the chorion, the nearly transparent eggshell that
- 167 contains perivitelline fluid and the embryo from fertilization until 2-3 dpf. We tested the
- 168 sunscreening role of the chorion by mechanically removing it with forceps and exposing these
- 169 embryos, and sibling controls that retained the chorion, to 450 J of UVB at 24 hpf. We found that
- the chorion does provides significant protection from UVB as ~60% of dechorionated embryos
- failed to inflate their swim bladders, significantly less than sibling controls (**Fig. 3C**). We
- examined if gadusol was present in the chorion or in the fluid within the chorion but found little to
- none (**Fig. S10**). These results suggest that the chorion structure itself can shield some
- incoming UVB. However, we conclude that the chorion provides less UV protection than
- 175 gadusol, as gadusol-depleted embryos even with intact chorions all failed to inflate their
- swim bladders when challenged with the same dose of UVB (**Fig. 1C**).
- 177 Together, our findings support a model where embryonic and larval fish are protected by
- multiple layers of UVB protection that span early development (**Fig. 3D**). The egg is maternally
- 179 loaded with gadusol, which provides the primary and most important layer of UV protection from
- 180 fertilization until at least 5 dpf. The chorion and melanophores are secondary, and less effective,
- 181 means of UVR protection. The chorion protects the developing embryo between fertilization and
- hatching (2-3 dpf), when pigmented melanophores emerge and modestly protect the growing
- 183 larval fish.
- 184 Gadusol has been repeatedly lost in fish species whose embryos are no longer exposed to185 sunlight
- 186 The two-enzyme biosynthetic pathway necessary for gadusol production (Eevs and MT-Ox) is
- 187 encoded in numerous vertebrate genomes, including fish, birds, reptiles, and amphibians¹⁷.
- 188 Osborn et al. identified the loss of the gadusol biosynthetic pathway in the coelacanth genome,
- and suggested the loss might be attributable to lack of UV penetration in the deep sea habitat of
- this species¹⁷. To test for broader patterns of conservation and loss among fish, we surveyed
- additional genomes, including many species that live in habitats not exposed to UVR. We
- hypothesized that gadusol pathway genes would not be required in species that live in deep
- waters, caves, are live bearers, or use electroreception to navigate habitats with poor light
 penetrance²³. To test this hypothesis, we searched 136 teleost genomes for inactivation or loss
- 194 of either eevs or MT-Ox. In all species, we identified a syntenic genomic region demarcated by
- 196 highly conserved flanking genes and assessed the presence or absence of intact ORFs
- 190 Inging conserved national copies of cove and MT-Ox. Our approach largely confirmed that the view of the second sec
- 197 encoding functional copies of *eevs* and MT-Ox. Our approach largely confirmed that the vast

majority of teleosts have functional copies of eevs and MT-Ox¹⁷. However, our survey identified 198 199 16 independent losses of either the eevs or MT-Ox genes across the teleost phylogeny (Fig. 4, 200 red species). Most of these genomes had lost orthologs of both eevs and MT-Ox, while others had lost only one gene or had pseudogene remnants. The loss of genes involved in gadusol 201 production was significantly correlated with lifestyle traits that identified species that live or 202 spawn in habitats protected from the sun (p = 0.012) (see Fig. S12, Methods and Table S2 for 203 204 details). To corroborate the link between loss of eevs or MT-Ox and loss of gadusol, we 205 measured gadusol levels in medaka embryos, which have intact eevs and MT-Ox genes, and 206 ovaries of channel catfish, which have lost eevs and MT-Ox. We found a strict correlation 207 between the presence of intact genes and maternally provided gadusol (Fig. S11). We conclude 208 that gadusol production has been repeatedly lost during evolution in teleost species whose 209 lifestyles protect them from UVR.

210 Discussion

Plants and microorganisms use numerous UV-absorbing compounds as sunscreens^{4,24}. 211 However, other than melanin, the repertoire of vertebrate sunscreens - especially compounds 212 213 that protect the most vulnerable early stages of development - remain essentially unknown. 214 Here, we provide experimental and phylogenomic evidence that gadusol is an ancient 215 sunscreen essential for protecting fish embryos from UVR. First, we use a CRISPR mutant that 216 disrupts gadusol biosynthesis to show that gadusol is produced during oogenesis and persists 217 in the embryo until at least 5 dpf. Second, we demonstrate that maternally deposited gadusol 218 safeguards embryonic and larval development by preventing UV-induced developmental defects and improving survival. Third, we find that gadusol acts as a true sunscreen preventing the 219 220 formation of CPDs, a signature of UVB-induced DNA damage, and consequently reducing 221 levels of cell and organismal death. Gadusol does not have any obvious functions beyond 222 protecting against UVR, as mutants survive to adulthood and are fertile. Together, these data 223 demonstrate that gadusol is a maternally provided sunscreen employed during early fish 224 development.

- Que work explores two alternative mechanisms of LIV protection during early d
- 225 Our work explores two alternative mechanisms of UV protection during early development. We 226 find that the chorion, a transparent eggshell that shields the developing embryo, also provides
- modest UV protection during embryogenesis. This protection is short lived (zebrafish hatch by
- 228 2-3 dpf) but may provide secondary protection during the most vulnerable stages of
- development. Melanin pigmentation emerges around embryo hatching and serves a relatively
- modest role as a whole-body sunscreen in 5 dpf larvae. Together, our results show that gadusol
- is the primary sunscreen across embryonic and larval development, while melanin and the
- 232 chorion play secondary roles during distinct phases of development.
- Finally, our phylogenetic analysis of gadusol biosynthetic genes, building on a previous study¹⁷,
- suggest that gadusol is an ancient sunscreen conserved broadly to protect teleost embryos.
- However, gadusol production has been repeatedly lost during teleost evolution. Intriguingly,
- these genes are absent in many fish species whose embryos are not exposed to UVR, including
- 237 deep sea-dwelling and electroreceptive fish. We suggest that similar to our protected fish facility
- environment, gadusol is also dispensable for embryonic development in natural environments
- that lack UVR. In microorganisms, the production of sunscreening compounds have been

- estimated to require >10% of all metabolic activity²⁴. Perhaps the loss of gadusol production in
- nutrient-poor dark habitats provides some evolutionary advantage, analogous to the energy
- conservation hypothesis invoked to explain the repeated loss of eyes in Mexican cavefish^{26,27}.
- 243 However, once these genes have been lost, descendent species may enter an evolutionary
- fitness trap where they are confined to breeding environments lacking UVR.

245 It remains unclear what role gadusol might play in other tetrapods. Functional copies of *eevs*

- and MT-Ox have been found in numerous vertebrate genomes¹⁷, but to our knowledge the
- 247 presence of gadusol has never been reported in vertebrates other than fish. Gadusol has been
- detected in the eggs or embryos of several aquatic invertebrates, including sponge²⁸, starfish²⁸,
- sea urchin²⁹, and brine shrimp³⁰. We hypothesize that gadusol may also protect early
- 250 development in these diverse aquatic organisms.
- 251 Here, we show that aquatic vertebrates produce and employ an additional sunscreen to
- 252 melanin. Melanin and gadusol both absorb well in the UVB spectrum. However, melanin also
- absorbs most wavelengths in the visible light spectrum, making it opaque and conspicuous
- while gadusol is transparent and invisible. Transparency as camouflage is a common trait in
- aquatic animals, especially in the open ocean where there is nothing to hide behind³¹. To date,
- 256 gadusol has only been detected in aquatic organisms. We speculate that gadusol has been
- 257 particularly advantageous to these animals as it offers protection from UVR, enabling an
- organism to stay in nutrient-rich sunlit areas, while remaining optically inconspicuous. We
- 259 propose that aquatic ecosystems exhibit unique ecological challenges that have selected for the
- 260 use of a transparent sunscreen.
- 261

262 Materials and Methods

263 Zebrafish Husbandry

All zebrafish work was performed at University of Utah's CBRZ zebrafish facility. This study was conducted under the approval of the Office of Institutional Animal Care and Use Committee (IACUC no. 18-2008) of the University of Utah's animal care and use program.

267 Generation of eevs mutant line

To generate a stable gadusol-depleted mutant line, eevs was targeted using CRISPR-Cas9 268 mutagenesis. Four gRNAs (**Table S3**) were designed using ChopChopV2³², targeting exon 2 269 (Fig. S1) due to the lack of suitable target sites within the small exon 1. Guide RNAs were 270 271 synthesized from DNA oligos using standard protocols³³. Freshly laid wild-type TU-strain embryos were injected with SpCas9 protein (NEB) mixed with gRNAs (~300 ng/ul), KCI, and 272 phenol red. 1-2 nanoliters were injected into each embryo. Mosaic mutant embryos were raised 273 to adulthood and outcrossed to wild-type Tübingen strain. Primers designed from ChopchopV2³² 274 were used to amplify the region targeted for CRISPR editing and to select for edited alleles with 275 large deletions. A compound deletion allele was identified by Sanger sequencing that removes 276 379bp and shifts the eevs open reading frame (Fig. S1, sequences in Table S3) (Genewiz). 277 278 This eevs mutant allele was given the designation zj2 and can be genotyped using PCR with 279 allele specific primers (Table S3). Sibling fish with the zj2 allele were crossed to produce

homozygous KO eevs^{zj2/zj2} fish, labeled as eevs^{-/-} throughout the text. Lack of gadusol was

determined using UPLC MS/MS and spectrophotometry (see below for details). Embryos

resulting from crosses of *eevs*^{-/-} mothers had little to no gadusol compared to wild-type embryos,

confirming successful generation of a gadusol-depleted line.

284 Gadusol extraction and UPLC MS/MS detection

285 Gadusol was extracted twice from embryos (7.5mg of vacuum dried egg material, crushed with

a microfuge pestle) using 150 ul of a (80:20, v/v) methanol:water solution. The extraction

287 supernatant was analyzed using ultraperformance liquid chromatography (Waters Acquity I-

Class, 2.1 x 100 mm BEH Amide column) and mass spectrometry (Waters Xevo G2 QToF)

289 (UPLC-MS) in negative ionization mode (detector range of 50-2000 Da). We used a regular

phase chromatography method starting with 95% acetonitrile (+0.1% formic acid) and 5 % water
 (+0.1% formic acid) following a linear gradient over 12 minutes ending with 30% acetonitrile

(+0.1% formic acid) following a linear gradient over 12 minutes ending with 50% accountine
 (+0.1% formic acid). Analytical standards of pure gadusol were run during the same acquisition

- run to match the retention time and observed mass between embryo samples and the pure
- 294 standard.

295 Gadusol detection via Nanodrop

296 To monitor gadusol production the UV-vis spectrometry on a Nanodrop was employed to

297 determine relative gadusol concentrations. Briefly, 25 embryos/larvae were placed in a

298 microfuge tube. All excess water was removed with a Pasteur pipette. 100 ul of 80:20

299 methanol:water was added to embryos. Embryos were mashed with a microfuge pestle for 15

- seconds. Samples were left to extract for at least 15 minutes, and then centrifuged at 12,000 g.
- 301 Clear supernatant, containing polar compounds such as gadusol, was separated and analyzed
- 302 on the nanodrop.

303 UV exposure, swim bladder inflation, and survival assays

24 hpf embryos were exposed to 450 J of UVB as measured on a radiometer (Solarmeter UVB)

at a fluence rate of 2.5 W/m² in 30ml of clear E3 media. This is a conservative estimate of a

³⁰⁶ physiologically relevant UVB dose that fish embryos would routinely experience in the wild⁶. A

raised and inverted UVP transilluminator with 306 nm broadband UVB bulbs was used (Ushio
 30000318) on the "low" setting (see Fig. S13). Embryos were returned to the incubator and kept

in the dark after mock or UV exposure. Swim bladder inflation was scored at 5 dpf by adding ice

- to the petri dish to stun the larvae, followed by manual counting on a dissection scope. A
- standard dose curve was conducted to determine that 450 J was an appropriate dose (Fig. S3).

5 dpf larvae were exposed to a dose curve to determine that 2.5 kJ was an appropriate dose

(**Fig. S5**). After mock or UV exposure, larvae were placed in an incubator for 1 day (dark) and

then placed in the nursery at 6 dpf. Survival was scored at 28 days post-fertilization to ensure

that all living juveniles could feed on their own and were not being sustained on maternal yolk.

- 316 Determination of CPDs in 24 hpf embryos
- 24 hpf embryos were dechorionated to obtain more consistent UV exposure. Embryos were
- exposed to 450 J of UVB and then immediately fixed after exposure in 4% PFA for 1 hour at

- 319 25°C. Fixed embryos were then washed in PBST. Embryos were exposed to 2 M HCl for 1 hour
- to break apart dsDNA and expose CPD epitopes. Samples were blocked in 5% NGS + PBST.
- 321 Mouse anti-CPD primary antibody (TDM-2, Cosmo Bio) was used to stain for CPDs. Goat anti-
- mouse AF546 secondary antibody (Invitrogen) was used to visualize CPDs. Embryos were also
- 323 stained with DAPI to visualize nuclei. Prior to imaging on a confocal microscope, tails were
- removed from embryos and placed on a flat glass slide with a small drop of PBST. A cover slip
- was mounted over the tails and sealed with nail polish. Tails were then imaged on an inverted
- 326 confocal microscope with a 20x objective (Zeiss 880). Images were analyzed using ImageJ³⁴ to
- 327 determine mean fluorescence intensity / tail area using the DAPI channel to create a mask for
- 328 the tail.

329 Apoptosis assay

- 24 hpf embryos within chorions were exposed to 450 J of UVB and then placed in the incubator
- for 5 hours. Chorions were removed and embryos were fixed for 1 hour in 4% PFA. Embryos
- 332 were stained with an activated caspase-3 antibody (BD Biosciences, anti:Rabbit) to mark
- apoptotic cells. Goat anti-rabbit AF594 secondary antibody (Invitrogen) was used to visualize
- apoptotic cells. Embryo tails were removed, processed, and imaged as above. ImageJ was
- used to process images and count the number of activated caspase-3 positive nuclei/mm².
- 336 RNAseq sample prep, library prep, sequencing, and analysis
- After 5 or 24hrs post UV exposure embryos were smashed with a microfuge pestle (MTC Bio)
- and RNA extracted using TRI Reagent (Zymo) and purified via Direct-zol RNA Miniprep Plus
- 339 (Zymo). Library prepared using NEBNext Ultra II Directional RNA Library Prep with poly(A)
- 340 mRNA Isolation. Samples then sequenced with Total RNA (eukaryote) NovaSeq SP Reagent
- Kit v1.5_50x50 bp. Each sample sequenced to a depth of 25 million reads. Reads aligned using $\frac{1}{2}$
- 342 STAR³⁵ and zebrafish reference genome (GRCz11). Optical duplicates removed and adapters
- trimmed. Differential expression analysis conducted with $DESeq2^{36}$ and specifically the
- Bioconductor package³⁷.
- 345 Generating embryos that lack melanin and gadusol
- To generate embryos that lacked melanin, *mitfa*^{w2/w2} fish were crossed with *mitfa*^{+/w2} fish to</sup>
- 347 produce clutches of 1:1 pigmented:unpigmented siblings, all with gadusol (Fig. 3A). To
- 348 generate embryos that lack both melanin and gadusol, *mitfa*^{+/w2}; *eevs*^{-/-} females were crossed to
- 349 $mitfa^{w2/w2}$; $eevs^{-/-}$ males to produce 1:1 pigmented:unpigmented siblings that all lacked gadusol.
- 350 Chorion UV protection assay
- 24 hpf wild-type TU-strain embryos were manually dechorionated with forceps in a dish with a
- thin film of 0.5% agar on the base of the dish. Embryos were moved with a fire-smoothened
- 353 Pasteur pipette. Embryos were exposed to 450 J of UVB as described above and then placed in
- incubator and swim bladder inflation was scored at 5 dpf.
- 355 Phylogenetic analysis of eevs and MT-Ox presence
- 123 genomes were gathered from the UCSC genome ark (GenArk) and additional 11 genomes
- 357 for deep sea and electro-receptive fish were gathered NCBI genomes for all except the Yap

Hadal snailfish³⁸ and *pseudoliparis swirei*³⁹. A BLAST database for each species was created by using the zebrafish sequence spanning from FRMD4B to FOXP1 to find the same region in all

360 curated genomes. If there was no BLAST hit for FOXP1 or MITF then the genome was dropped

361 for low quality. We then performed a tBLASTn search on the created databases for the

362 remaining genomes, using the zebrafish EEVS and MtOX translated nucleotide sequence as

the query. If there was no hit for EEVS or Mt-OX in the tblastn search that species was labeled

as not having gadusol.

To correlate the presence/absence of gadusol with life history traits we first collected life history data for all species (**Table S2**). The life history traits that we annotated were deep-sea, livebearing, electro-reception, and cave dwelling. We then built a species tree using fishtree⁴⁰ and added the Yap hadal snailfish and *Pseudolapris swirei* using the phylogenetic relationship determined in Mu et. al ³⁸. Due to gene loss in sister species not being independent, we used Bayestraits⁴¹ to perform the correlation test. We used discrete model testing and a likelihood ratios test comparing each of the five life-history traits to loss of gadusol (**Table S2**).

372 When running Bayestraits the loss of gadusol (parameter beta1 in the independent model and

q31 and q42 in dependent model) was set as trait one and the various life history traits were set

as trait two. The rate at which gadusol can be regained after loss was constrained to zero

because we were scoring for loss of the gene, and assumed it is nearly impossible to regain the

376 gene, especially in the short time span we are investigating. The parameters that estimate the

- 377 rate of life history traits changing from absent to present (q12,q34,q21,q43) were constrained to
- equal to each other, under the assumption that it is unreasonable that a fish would change its

life-style after loss of gadusol. When comparing the cave life history to gadusol loss, the

parameter that estimates the rate of moving from cave to surface (q21 and q43) was

constrained to zero under the assumption that species don't re-emerge from a cave after

adapting to that lifestyle.

383 The significance of the correlation between life-history trait and loss of gadusol was determined

using a likelihood ratio test which is calculated by 2*((dependent model likelihood)-(dependent

model likelihood)). The significance is then determined using a chi-square distribution with 2

degrees of freedom.

All parameters and code to re-run these models can be found in https://github.com/nclark-lab/gadusol

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397 with Native Nations and Urban Indian communities.

- Author Contributions: MCR and JAG conceived of the study. MCR created *eevs* knockouts.
- 399 MCR carried out and designed UV experiments. JM and MCR performed RNAseq experiments.
- 400 MCR analyzed RNAseq data with U of U bioinformatics core. DLF analyzed and ran samples on
- 401 UPLC MS/MS. JHL explored fish phylogeny and determined gene loss with input from NLC and
- 402 MCR. MCR and JAG wrote the manuscript with input from all authors.
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Figure 1. Gadusol is maternally provided and protects zebrafish embryos and larvae from UVB.

417 **A.** Experimental diagram for generating heterozygous mutant *eevs*^{+/-} embryos and larvae with

418 identical genotypes but containing maternal contribution of gadusol (top), or depleted of

419 maternally provided gadusol (bottom). On the right, UPLC mass spectra of 0hpf egg extracts

420 from each genetic cross; arrow indicates gadusol mass.

- B. Absorption values at 296nm from the indicated genotypes at the indicated timepoints. All
 absorption values normalized to wild type. Error bars indicate standard deviation from biological
 replicates.
- C. Distribution of swimbladder inflation scored in 5 dpf larvae, with genotypes and gadusol
 presence indicated, after mock exposure (grey) or UVB exposure (dark grey) at 24 hpf stage. All
 embryos resulted from crosses between TU and AB strain parents, except the TU in-cross that
 generated MZeevs embryos. From left to right, n = 50, 50, 75, 75 100, 97, 50, 50; N = 2, 2, 3, 3,
 4, 4, 2, 2.
- D. Survival distribution scored at 28 dpf, with genotypes and gadusol presence indicated, after
 mock exposure (grey) or UVB exposure (dark grey) at 5 dpf. From L-R n = 100, 95, 100, 97, N =
 4 for all groups.
- 432 n = embryos/larvae. N = clutches. statistics: student t test b, Fisher's Exact t-test c, d, *p<0.05; 433 **p<0.01; ***p<0.0001.





- 436 **A.** Immunohistochemistry, using an antibody that recognizes CPDs (TDM-2), on 24 hpf
- 437 immediately after mock or UVB exposure. Representative images shown.
- 438 **B.** Quantification of CPD labeling normalized to tail area (mm²). From left to right, n = 19, 19, 19, 439 21; N = 2 for all groups.
- 440 **C.** Quantification of immunohistochemistry, using an antibody that recognizes activated
- 441 caspase-3. n = 14, 16, 16, 20. N = 2 for all groups.
- 442 **D.** Significant upregulation of select UVR response and DNA damage GO term-associated
- genes measured from the indicated conditions and genotypes using RNAseq on 24 hpf embryos

- 444 after mock exposure or UVB exposure. RNA was collected 5 hours post mock or UVB exposure.
- 445 Gene expression is scaled by rows. Significance determined via Fishenricher⁴².
- 446 Student's T-test P*<0.05; P**<0.01; P***<0.0001. n = number of embryos. N = number of 447 clutches.

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451 Figure 3. Melanin and the chorion serve as secondary UV-shielding mechanisms in

452 embryonic and larval fish.

- **A.** Experimental diagram for generating embryos that lack either melanin, maternally providedgadusol, or both.
- B. Survival distribution scored at 28 dpf, with presence of melanin and gadusol indicated, after
 mock exposure (grey) or UVB exposure (dark grey) at 5 dpf. From left to right, n = 48, 48, 48, 48, 48, 48, 36, 60, 36; N = 2 for each group.
- 458 **C.** Distribution of swimbladder inflation scored in 5 dpf larvae after mock exposure (grey) or
- 459 UVB exposure (dark grey) at 24 hpf stage, with or without chorions. n = 100 for each group. N = 3 for each group.
- 461 **D.** Model illustrating the relative importance and timing of multiple UV-shielding mechanisms
 462 used in early zebrafish development.
- Student's T-test *p<0.05 ** p<0.01 ***p<0.0001. n = total number of individual embryos/larvae.
 N = total number of clutches.

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468 Figure 4. Gadusol production has been lost in several species no longer exposed to UVR.

- For each of 136 teleost species (full tree in **Fig. S12**), we assessed various life history traits that
- identify habitats that may not require embryonic protection from UVR, including electroreception,
- 471 live-bearing, cave dwelling, and deep-sea dwelling, indicated with colors in the legend to the left
- of the phylogeny. For each species, we identified the presence of intact open reading frames for
- 473 eevs and/or MT-Ox. Species that have lost the genes required for gadusol production are
- indicated in red. We found 16 independent losses across this phylogeny. We found that fish with
- these traits are more likely than by chance to lose gadusol (p=0.012).
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479 **REFERENCES**

- Williamson, C. E. What role does UV-B radiation play in freshwater ecosystems? *Limnology and Oceanography* 40, 386–392 (1995).
- Karentz, D. & Lutze, L. H. Evaluation ofbiologically harmful ultraviolet radiation in Antarctica
 with a biological dosimeter designed for aquatic environments. *Limnology and Oceanography* 35, 549–561 (1990).
- 3. Dahms, H.-U. & Lee, J.-S. UV radiation in marine ectotherms: Molecular effects and
 responses. *Aquatic Toxicology* 97, 3–14 (2010).
- 488 4. Cockell, C. S. & Knowland, J. Ultraviolet radiation screening compounds. *Biological Reviews*489 74, 311–345 (1999).
- 490 5. Schartl, M. *et al.* What is a vertebrate pigment cell? *Pigment Cell & Melanoma Research* 29,
 491 8–14 (2016).
- 492 6. Kapp, F. G. *et al.* Protection from UV light is an evolutionarily conserved feature of the
 493 haematopoietic niche. *Nature* 558, 445–448 (2018).
- 494 7. Dong, Q., Svoboda, K., Tiersch, T. R. & Todd Monroe, W. Photobiological effects of UVA
 495 and UVB light in zebrafish embryos: Evidence for a competent photorepair system. *Journal*496 of Photochemistry and Photobiology B: Biology 88, 137–146 (2007).
- Armstrong, T. N., Reimschuessel, R. & Bradley, B. P. DNA damage, histologial changes
 and DNA repair in larval Japanese medaka (Oryzias latipes) exposed to ultraviolet-B
 radiation. *Aquatic Toxicology* 58, 1–14 (2002).
- Fabacher, D. L. & Little, E. E. Skin component may protect fishes from ultraviolet-B
 radiation. *Environmental Science and Pollution Research* 2, 30–32 (1995).
- Fabacher, D. L., Little, E. E. & Ostrander, G. K. Tolerance of an albino fish to ultraviolet-B
 radiation. *Environmental Science and Pollution Research* 6, 69–71 (1999).
- 11. Garcia-Pichel, F. & Castenholz, R. W. Occurrence of UV-absorbing, mycosporine-like
 compounds among cyanobacterial isolates and an estimate of their screening capacity.
 Applied and Environmental Microbiology 59, 163–169 (1993).
- 12. Adams, N. L. & Shick, J. M. Mycosporine-like amino acids prevent UVB-induced
 abnormalities during early development of the green sea urchin Strongylocentrotus
 droebachiensis. *Marine Biology* 138, 267–280 (2001).
- Adams, N. L. & Shick, J. M. Mycosporine-like Amino Acids provide protection against
 ultraviolet radiation in eggs of the green sea urchin Strongylocentrotus droebachiensis.
 Photochemistry and Photobiology 64, 149–158 (1996).
- 14. Plack, P. A. *et al.* Gadusol, an enolic derivative of cyclohexane-1,3-dione present in the roes
 of cod and other marine fish. Isolation, properties and occurrence compared with ascorbic
 acid. *Biochem J* 199, 741–747 (1981).
- 516 15. Chioccara, F., Delia Gala, A., De Rosa, M., Novellino, E. & Prota, G. Mycosporine
 517 aminoacids and related compounds from the eggs of fishes. *Bulletin Des Societes*518 *Chimiques Belges* 89, 1101–1106 (1980).
- 16. Osborn, A. R. & Mahmud, T. Interkingdom Genetic Mix-and-Match To Produce Novel
 Sunscreens. ACS Synth. Biol. 8, 2464–2471 (2019).
- 521 17. Osborn, A. R. *et al.* De novo synthesis of a sunscreen compound in vertebrates. *Elife* **4**, e05919 (2015).

18. Arbeloa, E. M., Uez, M. J., Bertolotti, S. G. & Churio, M. S. Antioxidant activity of gadusol 523 524 and occurrence in fish roes from Argentine Sea. Food Chemistry **119**, 586–591 (2010). 525 19. Mori, T. et al. Simultaneous establishment of monoclonal antibodies specific for either cyclobutane pyrimidine dimer or (6-4) photoproduct from the same mouse immunized with 526 527 ultraviolet-irradiated DNA. Photochemistry and photobiology 54, 225–232 (1991). 528 20. Yamashita, M. Apoptosis in zebrafish development, Comparative Biochemistry and 529 Physiology Part B: Biochemistry and Molecular Biology 136, 731–742 (2003). 530 21. Kelsh, R. N. Genetics and Evolution of Pigment Patterns in Fish. *Pigment Cell Research* 17, 326-336 (2004). 531 22. Kelsh, R. N., Harris, M. L., Colanesi, S. & Erickson, C. A. Stripes and belly-spots -- a review 532 533 of pigment cell morphogenesis in vertebrates. Semin. Cell Dev. Biol. 20, 90-104 (2009). 534 23. Schwassmann, H. O. Ecological aspects of electroreception. in Sensory ecology 521-533 535 (Springer, 1978). 536 24. Gao, Q. & Garcia-Pichel, F. Microbial ultraviolet sunscreens. Nature Reviews Microbiology 537 **9**, 791–802 (2011). 538 25. Summers, C. G. Vision in albinism. Transactions of the American Ophthalmological Society 539 **94**, 1095 (1996). 540 26. Krishnan, J. & Rohner, N. Cavefish and the basis for eye loss. Philosophical Transactions of 541 the Royal Society B: Biological Sciences 372, 20150487 (2017). 542 27. Moran, D., Softley, R. & Warrant, E. J. The energetic cost of vision and the evolution of 543 eyeless Mexican cavefish. Science Advances 1, e1500363 (2015). 544 28. Bandaranayake, W. M., Bourne, D. J. & Sim, R. G. Chemical Composition during Maturing 545 and Spawning of the Sponge Dysidea herbacea (Porifera: Demospongiae). Comparative 546 Biochemistry and Physiology Part B: Biochemistry and Molecular Biology 118, 851-859 547 (1997). 548 29. Chioccara, F., Zeuli, L. & Novellino, E. Occurrence of mycosporine related compounds in 549 sea urchin eggs. Comp. Biochem. Physiol. 85, 459-461 (1986). 550 30. Grant, P. T., Middleton, C., Plack, P. A. & Thomson, R. H. The isolation of 4 551 aminocyclohexenimines (mycosporines) and a structurally related derivative of cyclohexane-1-3-dione (gadusol) from the brine shrimp, Artemia. Comparative Biochemistry and 552 Physiology B-Biochemistry & Molecular Biology 80, 755–759 (1985). 553 554 31. Johnsen, S. Hidden in plain sight: the ecology and physiology of organismal transparency. 555 The Biological Bulletin 201, 301–318 (2001). 556 32. Labun, K., Montague, T. G., Gagnon, J. A., Thyme, S. B. & Valen, E. CHOPCHOP v2: a 557 web tool for the next generation of CRISPR genome engineering. Nucleic acids research 558 44, W272–W276 (2016). 33. Takasugi, P. R. et al. Orthogonal CRISPR-Cas tools for genome editing, inhibition, and 559 CRISPR recording in zebrafish embryos. Genetics 220, iyab196 (2022). 560 34. Abràmoff, M. D., Magalhães, P. J. & Ram, S. J. Image processing with ImageJ. 561 562 Biophotonics international 11, 36–42 (2004). 35. Dobin, A. et al. STAR: ultrafast universal RNA-seg aligner. Bioinformatics 29, 15-21 (2013). 563 564 36. Love, M. I., Huber, W. & Anders, S. Moderated estimation of fold change and dispersion for RNA-seq data with DESeq2. Genome biology 15, 1-21 (2014). 565

- 37. Gentleman, R. C. *et al.* Bioconductor: open software development for computational biology
 and bioinformatics. *Genome biology* 5, 1–16 (2004).
- 38. Mu, Y. *et al.* Whole genome sequencing of a snailfish from the Yap Trench (~ 7,000 m)
 clarifies the molecular mechanisms underlying adaptation to the deep sea. *PLoS genetics* 17, e1009530 (2021).
- 39. Wang, K. *et al.* Morphology and genome of a snailfish from the Mariana Trench provide
 insights into deep-sea adaptation. *Nature ecology & evolution* 3, 823–833 (2019).
- 40. Rabosky, D. L. *et al.* An inverse latitudinal gradient in speciation rate for marine fishes. *Nature* 559, 392–395 (2018).
- 41. Pagel, M., Meade, A. & Barker, D. Bayesian estimation of ancestral character states on
 phylogenies. *Systematic biology* 53, 673–684 (2004).
- 42. Kuleshov, M. V. *et al.* Enrichr: a comprehensive gene set enrichment analysis web server 2016 update. *Nucleic acids research* **44**, W90–W97 (2016).
- 579