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Data Article

Lipid quantitation and metabolomics data from vitamin E-deficient and -sufficient zebrafish embryos from 0 to 120 hours-post-fertilization

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ABSTRACT

The data herein is in support of our research article by McDougall et al. (2017) [1], in which we used our zebrafish model of embryonic vitamin E (VitE) deficiency to study the consequences of VitE deficiency during development. Adult 5D wild-type zebrafish (*Danio rerio*), fed defined diets without (E⁻) or with VitE (E⁺, 500 mg RRR- α -tocopherol/kg diet), were spawned to obtain E⁻ and E⁺ embryos that we evaluated using metabolomics and specific lipid analyses (each measure at 24, 48, 72, 120 hours-post-fertilization, hpf), neurobehavioral development (locomotor responses at 96 hpf), and rescue strategies. Rescues were attempted using micro-injection into the yolk sac using VitE (as a phospholipid emulsion containing d₆- α -tocopherol at 0 hpf) or D-glucose (in saline at 24 hpf).

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Specifications Table

Subject area	<i>Biology</i>
More specific subject area	<i>Nutrition; Antioxidants</i>
Type of data	<i>Graphs; Figures</i>
How data was acquired	<i>LC-MS (liquid chromatography-mass spectrometry) using a Shimadzu Nexera system (Shimadzu; Columbia, MD, USA) coupled to a high-resolution hybrid quadrupole–time-of-flight mass spectrometer (TripleTOF[®] 5600; SCIEX; Framingham, MA, USA); Embryos were assessed for viability, developmental progression and spontaneous movements (earliest behavior in zebrafish), using the zebrafish acquisition and analysis program (ZAAP). Locomotor Response Assay using a Viewpoint ZebraBox (software version 3.0, Viewpoint Life Sciences, Lyon, France).</i>
Data format	<i>Analyzed</i>
Experimental factors	<i>Adult 5D zebrafish fed defined diets without (E⁻) or with VitE (E⁺, 500 mg RRR-α-tocopheryl acetate/kg diet) were spawned to obtain E⁻ and E⁺ embryos that were evaluated up to 120 hours-post-fertilization (hpf)</i>
Experimental features	<i>Lipid and metabolomics analyses were performed in developing E⁻ and E⁺ zebrafish embryos collected daily from 24 to 120 hpf. Mortality and neurobehavioral outcomes were assessed at 96 hpf in two separate rescue experiments using E⁻ and E⁺ embryo populations micro-injected into the yolk (1) at 0 hpf with VitE or (2) at 24 hpf with D-glucose</i>
Data source location	<i>Oregon State University, Corvallis, OR 97330</i>
Data accessibility	<i>Data is within this article</i>

Value of the data

- Fatty acid quantification and peroxidation data during zebrafish embryonic development in E⁻ vs. E⁺ zebrafish embryos may be used by other researchers to investigate antioxidant effects of VitE with respect to specific lipids.
- The metabolomics dataset may be utilized by other researchers to investigate the secondary metabolic effects of VitE deficiency.
- Rescue studies using microinjection into the yolk sac may be compared to other methods of compound/nutrient delivery to developing zebrafish.

1. Data

Fig. 1. shows data from quantitative analyses of LA (linoleic acid, 18:2, omega-6); ARA (arachidonic acid, 20:4, omega-6); EPA (eicosapentaenoic acid, 20:5, omega-3); DHA (docosahexaenoic acid, 22:6, omega-3) in fatty acid extracts from samples with and without alcoholic saponification of E⁻ and E⁺ embryos collected at 24, 48, 72, and 120 hpf. Tables 1 and 2 provide detailed targeted metabolomics datasets for E⁻ and E⁺ embryos collected at 24, 48, 72, and 120 hpf. Relative response intensity metabolomics data for choline and methylation pathway intermediates in E⁻ and E⁺ embryos are shown in Fig. 2. Relative response intensities of antioxidant network components from metabolomic analyses, as well as quantification of α -tocopherol and ascorbic acid, in E⁻ and E⁺ embryos (pmol/embryo) are shown in Fig. 3. Relative response intensities of glycolytic and tricarboxylic acid cycle intermediates in E⁻ and E⁺ embryos are shown in Fig. 4. Relative response intensities of free saturated fatty acids and coenzyme A from metabolomics data in E⁻ and E⁺ embryo are shown in Fig. 5. Fig. 6 shows locomotor activity data from E⁻ and E⁺ embryos micro-injected into the yolk sac at 0 hpf

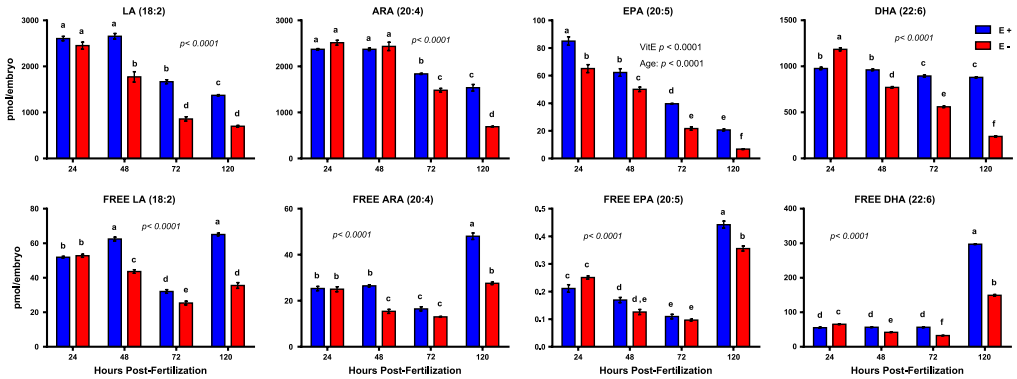


Fig. 1. Quantified levels of total and free (unesterified) fatty acids in E- vs. E+ embryos. Area counts normalized using internal standards ($n=3$ samples/group, with $n=10$ –15 embryos/sample for total lipids; $n=4$ samples/group with $n=15$ –30 embryo/sample for free fatty acids). Shown are saponified (upper row) or extracted only (lower row) samples, means \pm SEM; p -values are for VitE \times Age interactions, unless main effects (VitE or Age) are indicated (Tukey's post-test, $p < 0.05$ for bars bearing different letters). Abbreviations: LA (linoleic acid); ARA (arachidonic acid); EPA (eicosapentaenoic acid); DHA (docosahexaenoic acid).

with either saline or a VitE-emulsion. Fig. 7 shows locomotor activity data from E- and E+ embryos micro-injected into the yolk sac at 24 hpf with either saline or *D*-glucose.

2. Experimental design, materials and methods

2.1. Study design

All experiments (*i.e.* lipid quantifications, targeted metabolomics analyses, and micro-injection rescue studies) were performed in duplicate and have been reported in detail [1].

2.2. Zebrafish husbandry and diets

The Institutional Animal Care and Use Committee of Oregon State University approved this protocol (ACUP Number: 4344). Tropical 5D strain zebrafish were housed in the Sinnhuber Aquatic Research Laboratory and complete details of the housing and husbandry have been reported [1].

2.3. Analyses

Diet and embryo α -tocopherol [2] and ascorbic acid [3] were determined using high-pressure liquid chromatography with electrochemical detection as reported [1].

Extraction and sample preparation for metabolomic analysis were performed following 24, 48, 72, and 120 hpf, embryos ($n=15$ per replicate, $n=4$ replicates per group), as described [1]. Chromatography was performed with a Shimadzu Nexera system (Shimadzu; Columbia, MD, USA) coupled to a high-resolution hybrid quadrupole-time-of-flight mass spectrometer (TripleTOF[®] 5600; SCIEX; Framingham, MA, USA). Two different LC analyses using reverse phase and HILIC columns were used, as described [1].

Analysis of total DHA, EPA, ARA, and LA were performed as described [2] with modifications, as described [1]. Chromatographic separations were carried out on 4.6×250 mm J'sphere ODS-H80 ($4 \mu\text{m}$, YMC Co, Kyoto, Japan) for negative ion analysis. TOF-MS and TOF-MS/MS were operated with

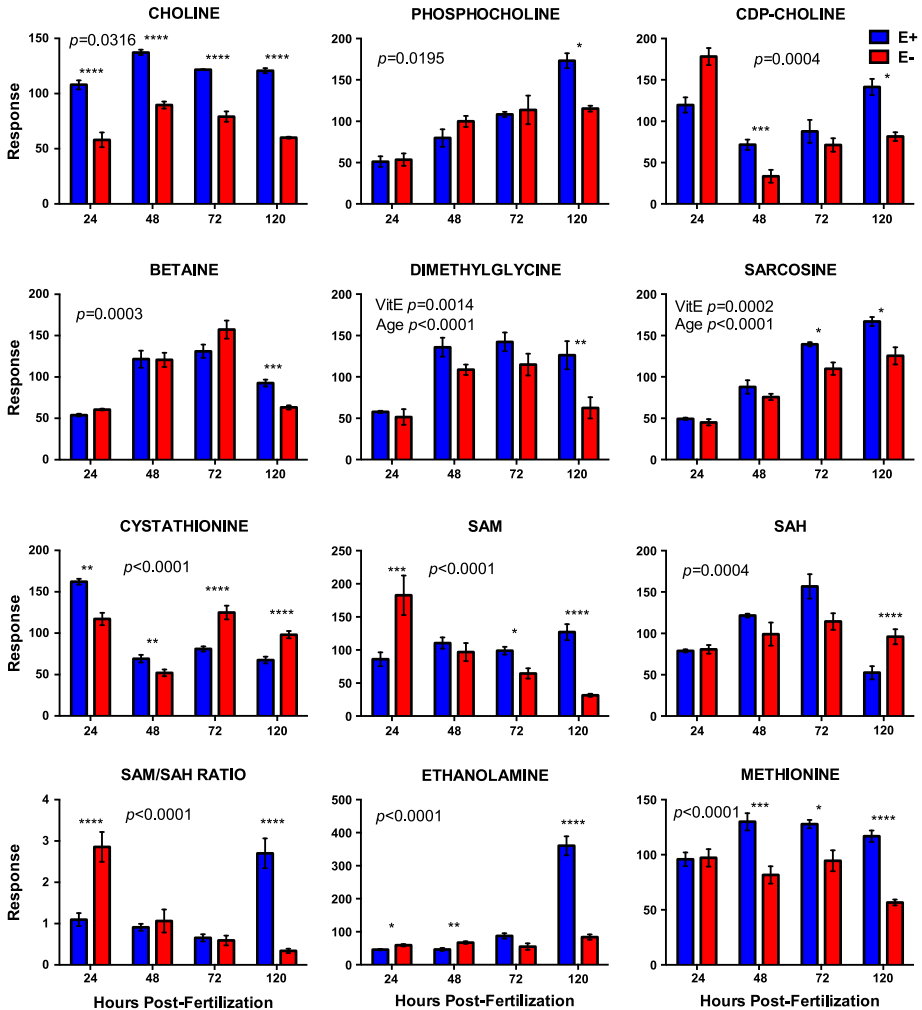


Fig. 2. Relative response intensities of choline and methylation pathway intermediates. E- and E+ embryo ($n=15$ /sample; 4 samples/group) data were normalized against QC sample intensities ($n=4$) for each individual metabolite. Statistical significance ($p < 0.05$) was calculated using 2-way ANOVA with Sidak's post-test for multiple comparisons of normalized and natural log-transformed intensity values. Shown are means \pm SEM; p -Values are for VitE x Age interactions, unless indicated otherwise. Paired comparison, p -values are indicated as * < 0.05 , ** < 0.005 , *** < 0.001 , **** < 0.0001 .

same parameters as for metabolomics, as described [1].

2.4. Microinjection rescue studies

Embryos were microinjected as described and criteria used to assess supplementation tolerance of zebrafish embryos using ZAAP at 24, 48, and 120 hpf, as described [1].

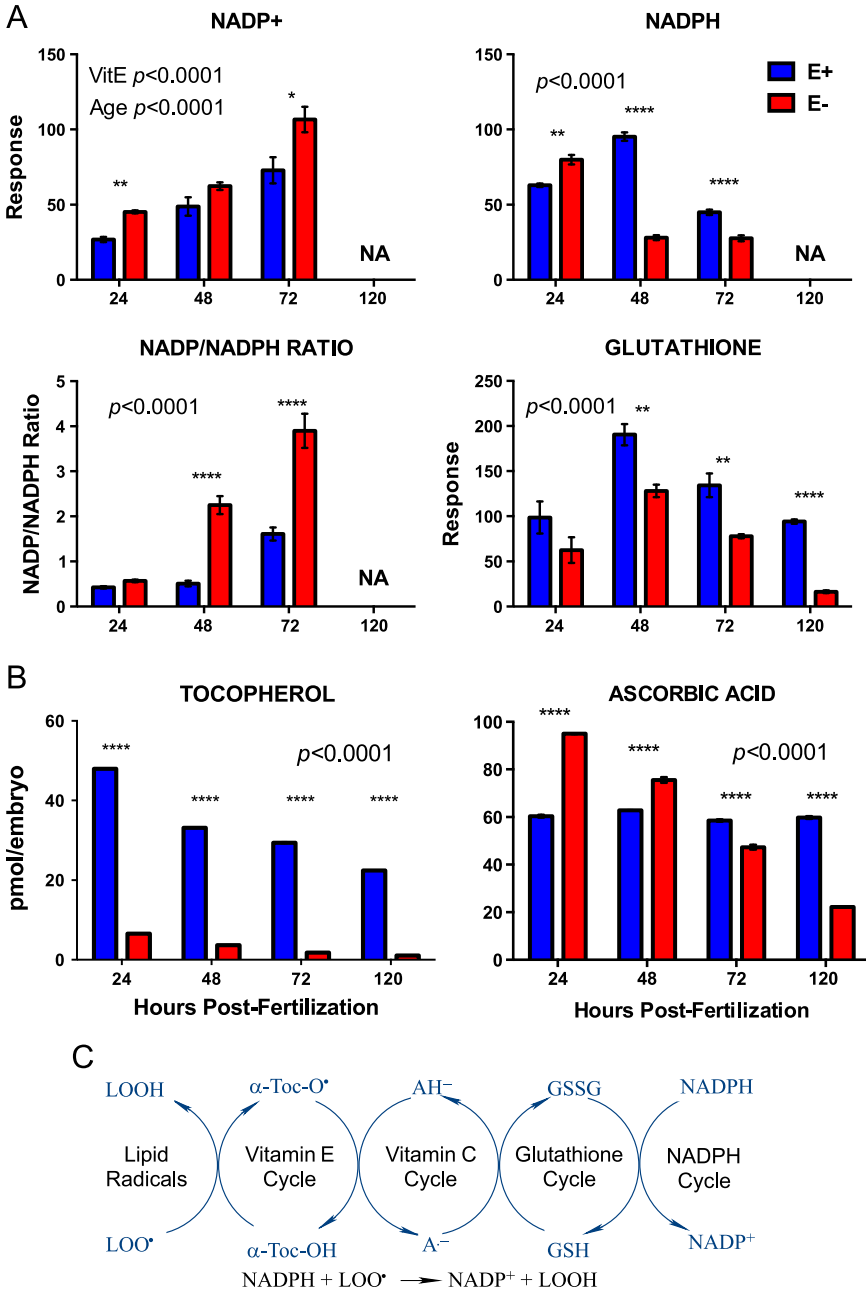


Fig. 3. Relative response intensities of antioxidant network components from metabolomics and quantification of α -tocopherol and ascorbic acid. A. E- and E+ embryo ($n=15$ /sample; 4 samples/group) relative response data was normalized against QC sample intensities ($n=4$) for each individual metabolite. B. Quantified levels of α -tocopherol and ascorbic acid, according to established protocols (31) and (33), respectively. Statistical significance ($p < 0.05$) was calculated using 2-way ANOVA with Sidak's post-test for multiple comparisons of normalized and natural log-transformed intensity values. Shown are means \pm SEM; p -Values are for VitE \times Age interactions, unless indicated otherwise. Paired comparisons p -values are indicated as * < 0.05 , ** < 0.005 , *** < 0.001 , **** < 0.0001 . C. Antioxidant network scheme showing interaction of antioxidants with lipid radicals and consumption or NADPH.

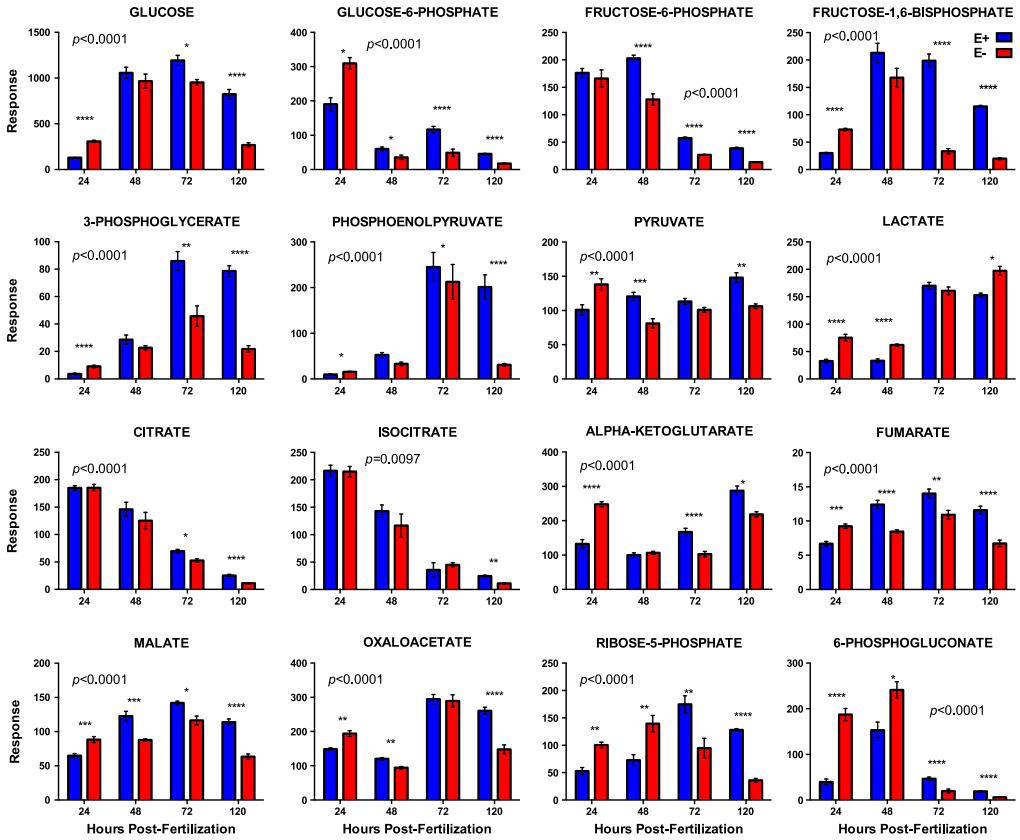


Fig. 4. Relative response intensities of glycolytic and tricarboxylic acid cycle intermediates. E- and E+ embryo ($n=15$ /sample; 4 samples/group) data were normalized against QC sample intensities ($n=4$) for each individual metabolite. Statistical significance ($p < 0.05$) was calculated using 2-way ANOVA with Sidak's post-test for multiple comparisons of normalized and natural log-transformed intensity values. Shown are means \pm SEM; p-values are for VitE \times Age interactions. Paired comparisons p-values are indicated as * < 0.05, ** < 0.005, *** < 0.001, **** < 0.0001.

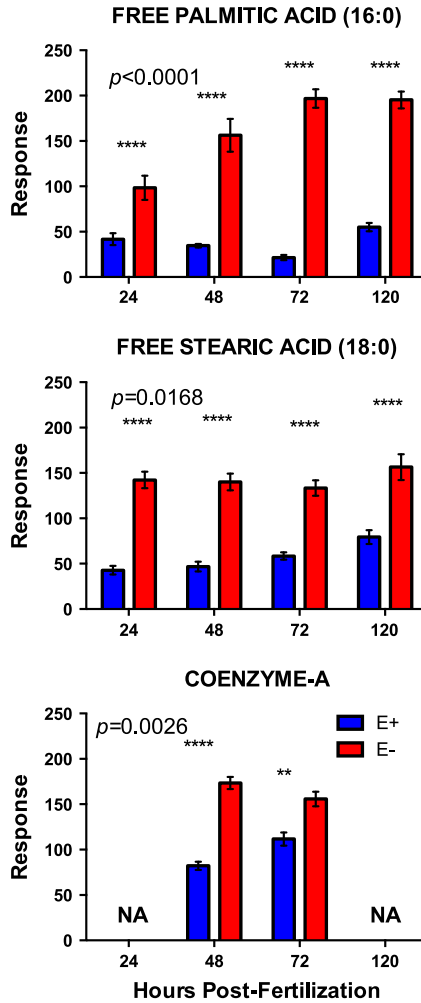


Fig. 5. Relative response intensities of free saturated fatty acids and coenzyme A from metabolomic analyses. E- and E+ embryo ($n=15$ /sample; 4 samples/group) data were normalized against QC sample intensities ($n=4$) for each individual metabolite. Statistical significance ($p < 0.05$) was calculated using 2-way ANOVA with Sidak's post-test for multiple comparisons of normalized and natural log-transformed intensity values. Shown are means \pm SEM; p -values are for VitE x Age interactions. Paired comparison p -values are indicated as * < 0.05 , ** < 0.005 , *** < 0.001 , **** < 0.0001 .

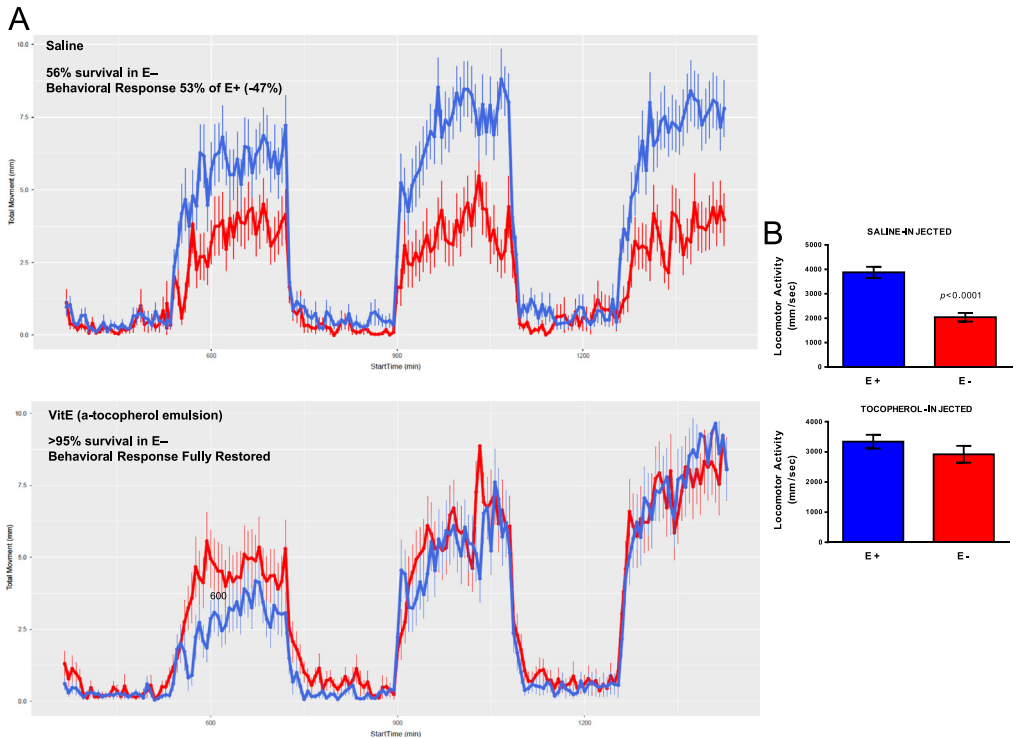


Fig. 6. E- compared with E+ embryos have impaired behavior when injected with saline (upper panel), but restored responses when injected with VitE (lower panel). A. Embryos were analyzed in 96-well plates (128 embryos per group). Locomotor activities following a series of light stimuli (a stimulus every 6 for 24 min) were measured as distance moved (mm) over time (seconds). At 96 hpf, E- (red) embryos treated with saline (upper panel) were 47% less responsive to light than were E+ embryos (E- area-under-curve, AUC: 2040 ± 178 ; E+ AUC: 3877 ± 228 ; $p < 0.0001$). Embryos with morphological defects were not included in data analysis. E- behavior was restored using VitE injection into the yolk at the 1 cell stage (lower panel E- AUC: 2970 ± 280 ; E+ AUC: 3340 ± 226 , not significantly different). B. Bar chart comparisons of respective time-course data. VitE (tocopherol)-injected E- and E+ embryo locomotor activities were not significantly different.

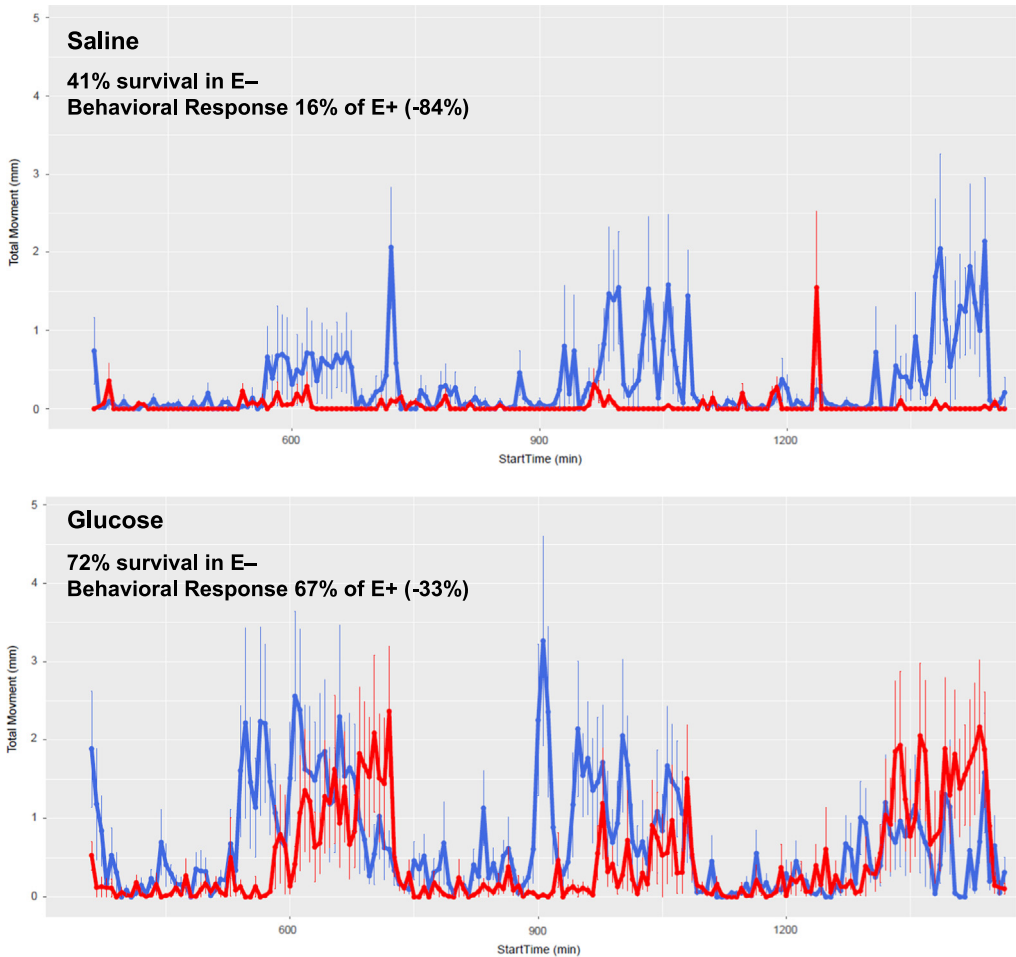


Fig. 7. Locomotor response assay activity data showing neurobehavioral impairment. E⁻ and E⁺ embryos (96 hpf) were analyzed in 96-well plates (128 embryos per group). Locomotor activities following a series of light stimuli (every 6 for 24 min) were measured as distance moved (mm) over time (seconds). E⁻ (red) embryos treated with saline (upper panel) were 84% less responsive to light than were E⁺ (blue) embryos (E⁻ area-under-curve, AUC: 572 ± 72 E⁺ AUC: 3580 ± 387 ; $p < 0.0001$). Embryos with morphological defects were not included in data analysis. E⁻ behavior was partially restored by approximately 50% following glucose injection into the yolk at 24 hpf (lower panel; E⁻ AUC: 2502 ± 150 ; E⁺ AUC: 3734 ± 359 ; $p < 0.0001$). Statistical significance was determined using a Kolmogorov–Smirnov test ($p < 0.01$).

2.5. Behavioral assessments

Locomotor activity was measured in a total of $n=128$ embryos per group using Viewpoint Zebrabox [4,5], as described [1].

2.6. Data processing and statistical analyses

All data processing and statistical analyses were performed as described in [4–6], with modification made as reported [1].

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Transparency document. Supporting information

Transparency data associated with this article can be found in the online version at <http://dx.doi.org/10.1016/j.dib.2017.02.046>.

Appendix A. Supplementary material

Supplementary data associated with this article can be found in the online version at <http://dx.doi.org/10.1016/j.dib.2017.02.046>.

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