



SCHOOL OF MEDICINE
Genetics



Zebrafish Atlas Phenotype Project (ZAPP): A Community-Driven Framework for Zebrafish Toxicology and Phenotype Data

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EHLC webinar
January 23rd, 2026

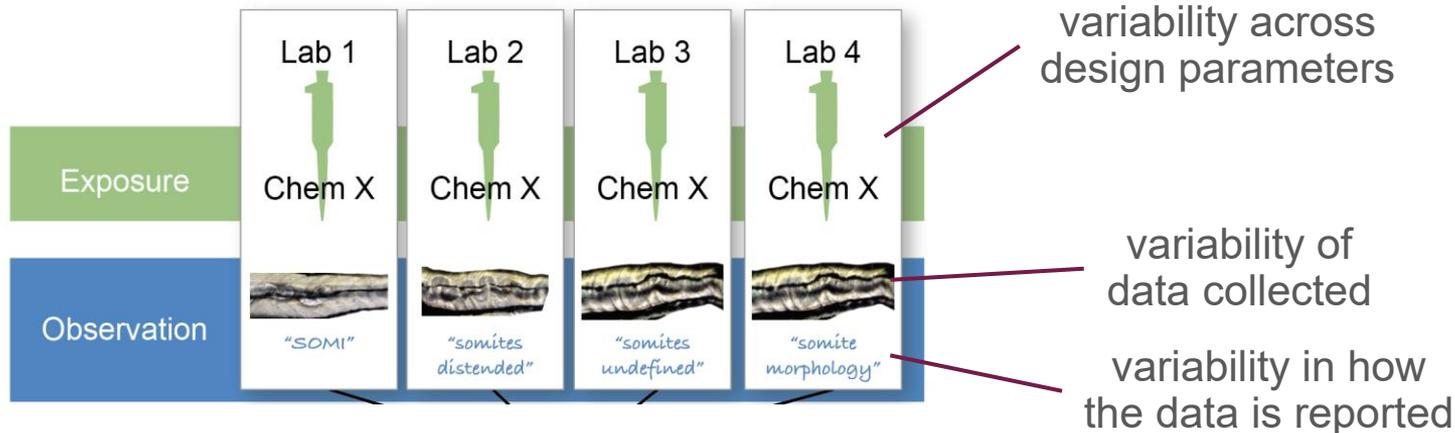
Lack of standards impedes data integration and reuse of zebrafish toxicology assay data

Systematic Evaluation of the Application of Zebrafish in Toxicology (SEAZIT) project

SEAZIT Goals:

- Provide scientific basis to decide whether or not zebrafish can be used for toxicological screening
- Provide fundamental knowledge on the use of zebrafish in toxicology

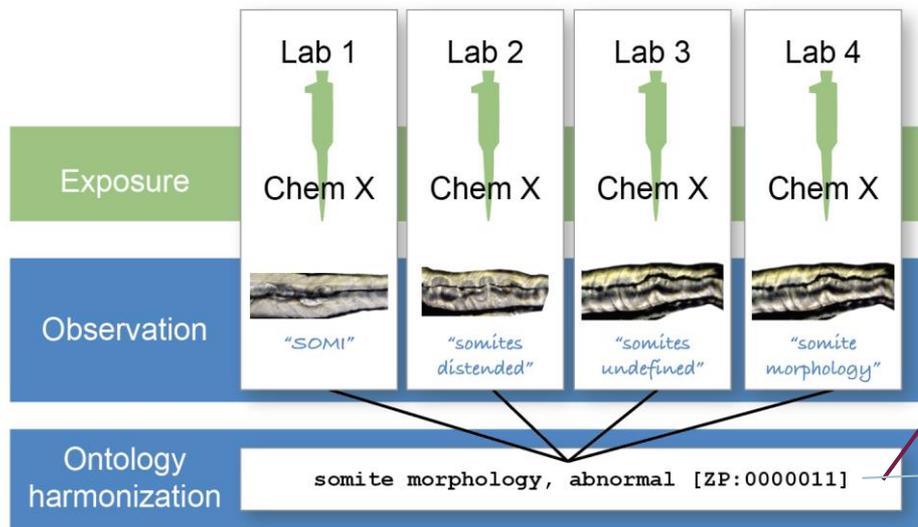
Inter-laboratory study



Lack of standards impedes data integration and reuse of zebrafish toxicology assay data

Systematic Evaluation of the Application of Zebrafish in Toxicology (SEAZIT) project

Inter-laboratory study



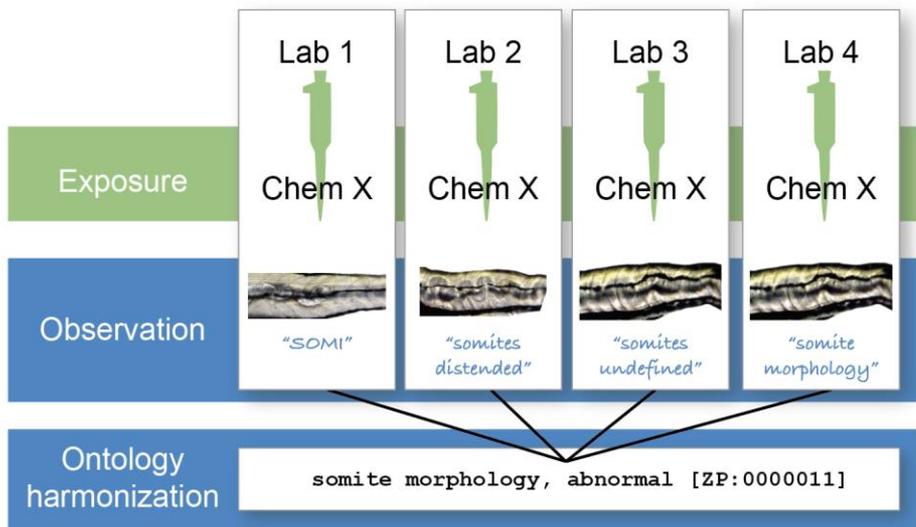
Use of standards (eg. ontology term to report phenotypes) reduces data variability

What does "abnormal somite morphology" phenotype look like?

Lack of standards impedes data integration and reuse of zebrafish toxicology assay data

Systematic Evaluation of the Application of Zebrafish in Toxicology (SEAZIT) project

Inter-laboratory study



Reporting phenotypic and toxicology data **using standards** would improve data interoperability, sharing, reuse, and comparison.

Adoption of standards is limited by practical use

The Research Community understands and agrees that the need for standards is crucial. However, they **don't know what to do in practice**.

- What are the standards and how to use them?
- Where to find the standards?
- What to do when what is needed is not available?
- How to use standards in existing research workflow?

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The Research Community understands and agrees that the need for standards is crucial. However, they **don't know what to do in practice**.

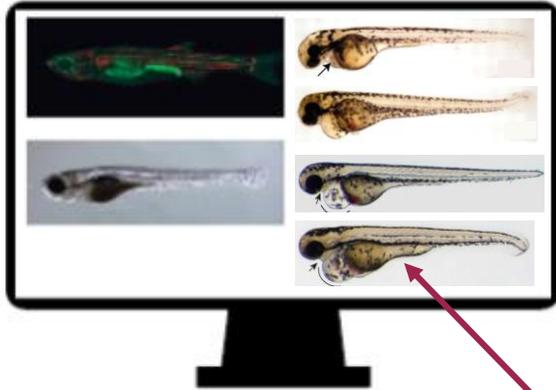
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Community-led Zebrafish Toxicology Phenotype Atlas (ZAPP) will

- **integrate existing standards, and develop new one when needed, for phenotypic and toxicology data**
- **make standard adoption approachable**
- **support community-driven improvement**

The Zebrafish Toxicology Phenotype Atlas (ZAPP)

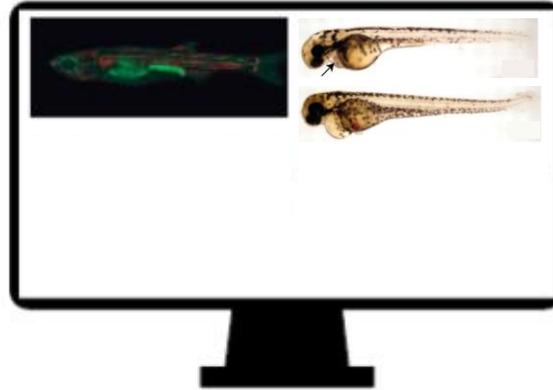
Zebrafish Phenotype Atlas



Explore

Search
Filter

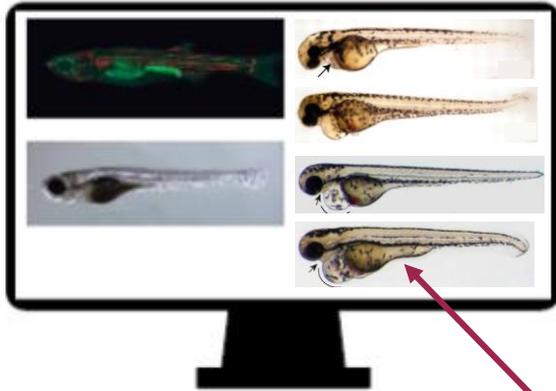
Zebrafish Phenotype Atlas



Visual definition
of phenotypes

The Zebrafish Toxicology Phenotype Atlas (ZAPP)

Zebrafish Phenotype Atlas

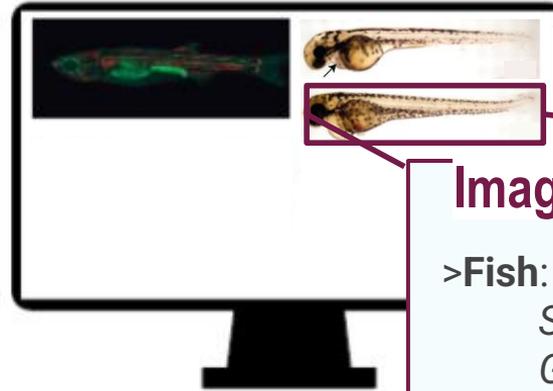


Visual definition
of phenotypes

Explore

Search
Filter

Zebrafish Phenotype Atlas



Images + Annotations

>Fish:

Strain: strain

Genotype: Genotype

Allele: Allele

...

>Experimental conditions

Toxicant: Toxicant

Concentration: mg/l

Timing of exposure: 2

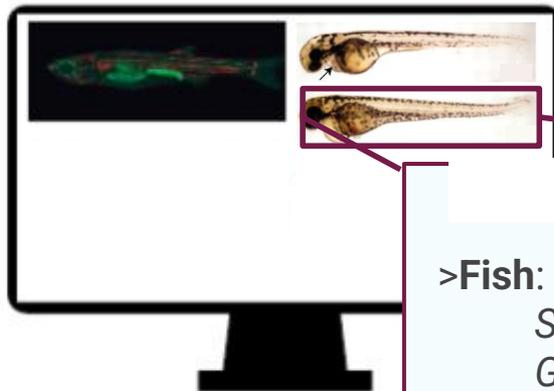
h

...



The Zebrafish Toxicology Phenotype Atlas (ZAPP)

Zebrafish Phenotype Atlas



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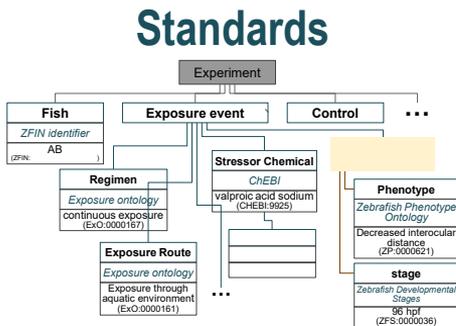
...

>Experimental conditions

Toxicant: Toxicant
Concentration: mg/l
Timing of exposure: 2

h

...

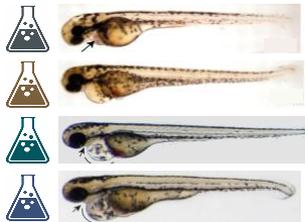


Same standards
for all data in the atlas

The Zebrafish Toxicology Phenotype Atlas (ZAPP)

Data Collection

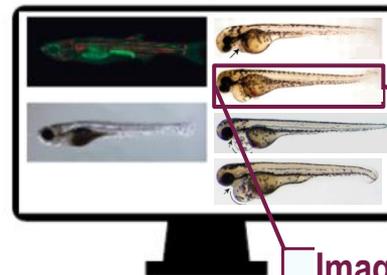
Zebrafish community



Submit to the Atlas



Zebrafish Phenotype Atlas



Images + Annotations

>Fish:

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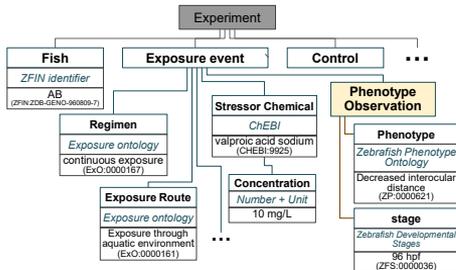
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>Phenotype: Phenotype



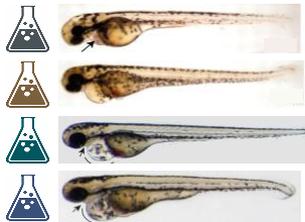
Standards



The Zebrafish Toxicology Phenotype Atlas (ZAPP)

Data Collection

Zebrafish community



Data Annotation (using Standards)

Provenance of Data

Control Information

Images

Control information

Experiment information

Fish

Exposure Event

Stressor CAS Identifier

Stressor concentration Unit μM mg/L Other

Exposure route

via environment (chemical in the water)

via diet (ingestion)

via gavage

via injection

Exposure regimen

Continuous exposure Repeated exposure

Stage at exposure start Hours

Stage at exposure end Hours

Phenotype Observation

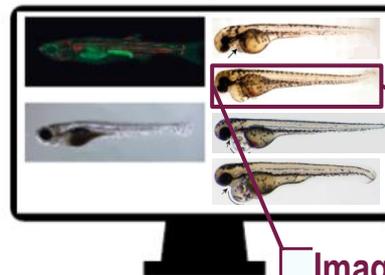
Images

Phenotypes shown in the image

Stage at observation Hours

Submit to the Atlas

Zebrafish Phenotype Atlas



Images + Annotations

>Fish:

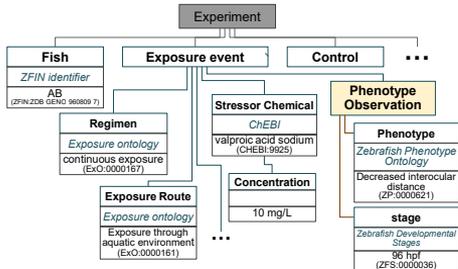
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Standards



Annotation Toolkit

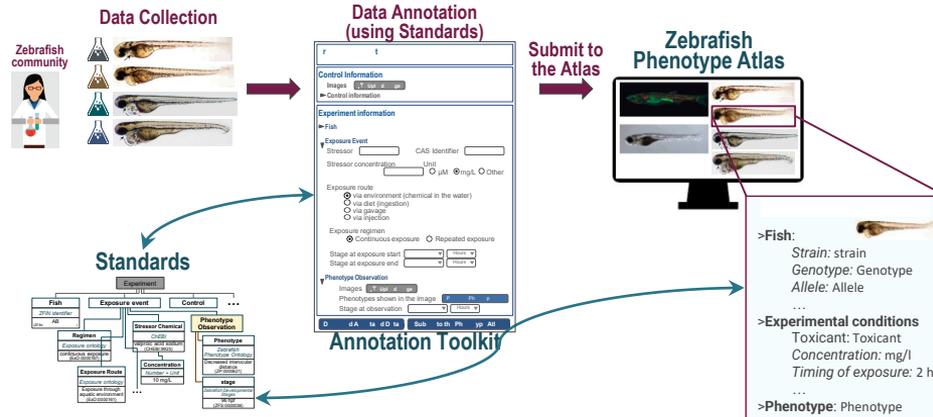
Community involvement is at the core of ZAPP

Community is involved in every part of the project

- Data annotation and submission
- Standards development
- Requirements gathering
- Governance

ZAPP community includes

- Zebrafish researchers
- Toxicologists
- Atlas experts
- Standards developers
- etc



A few caveats...

ZAPP is developed to be **species agnostic**

Though created to represent Zebrafish data, tools and standards can be adapted/reused, including

- Data model
- Annotation toolkit
- Ontology term picker

“Phenotypes? I am confused!”

Phenotype can be

- **Abnormal morphology** ← First iteration
- Change in gene expression
- Change in behavior

“Fish”

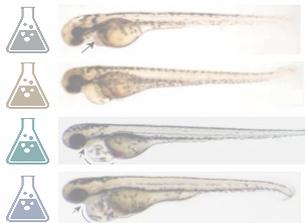
Fish can be

- **Wild-type** ← First iteration
- Transgenic
- Mutants

The Zebrafish Toxicology Phenotype Atlas (ZAPP)

Data Collection

Zebrafish community



Data Annotation (using Standards)

Provenance of Data

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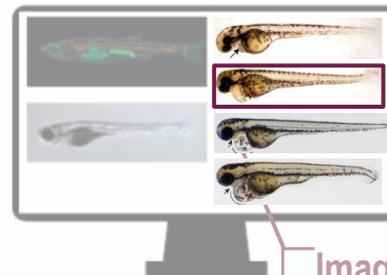
Images

Phenotypes shown in the image

Stage at observation Hours

Submit to the Atlas

Zebrafish Phenotype Atlas



Images + Annotations

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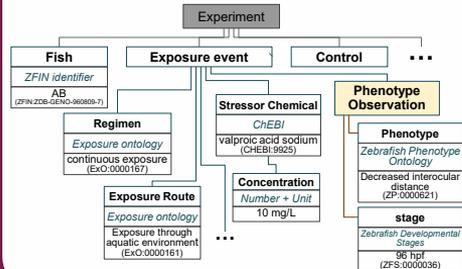
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Standards



Annotation Toolkit

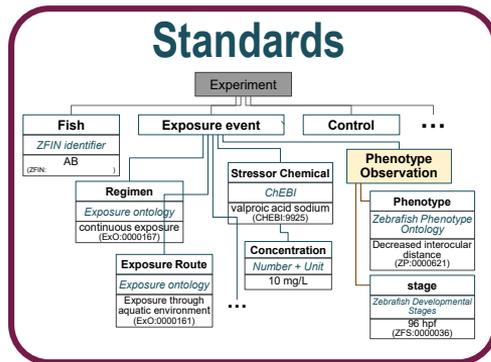
Standards for Toxicology Assay and Phenotype.

Data Model

= all the pieces of information required
AND how they should be reported

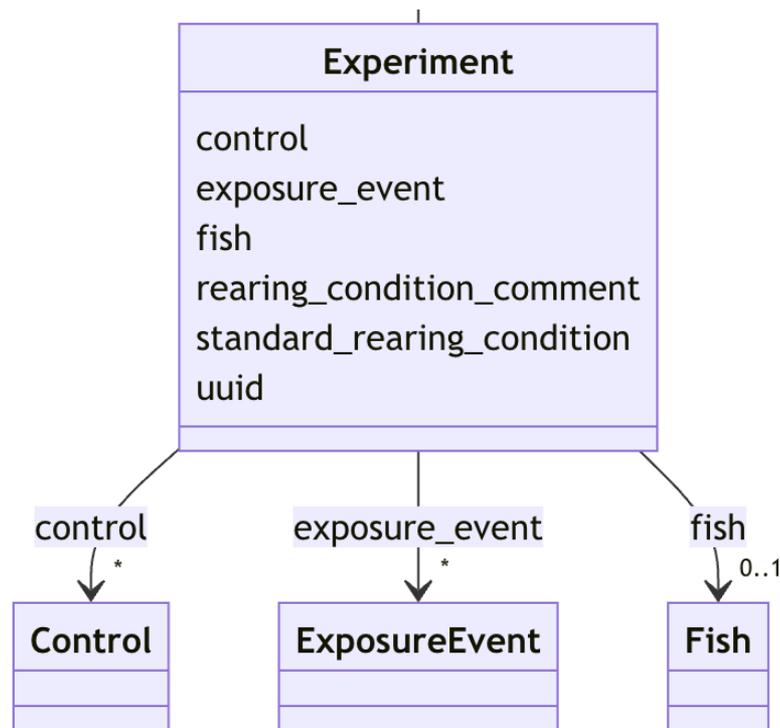
Ontologies and standard vocabularies

= standards to use to report the information



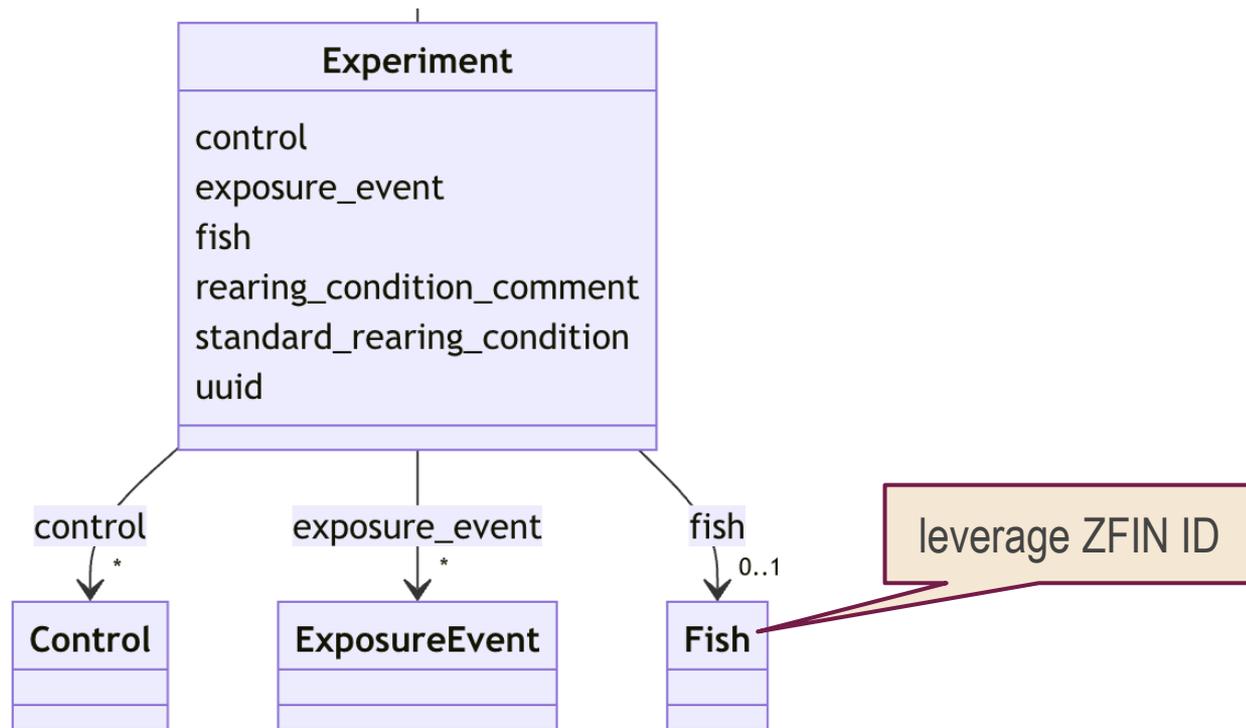
Zebrafish Toxicological Exposure Data Model

- Created based on community input
- Refined based on community feedback
- Created for Zebrafish toxicological assay, but can be reused and/or expanded for other species/model and other toxicological assay



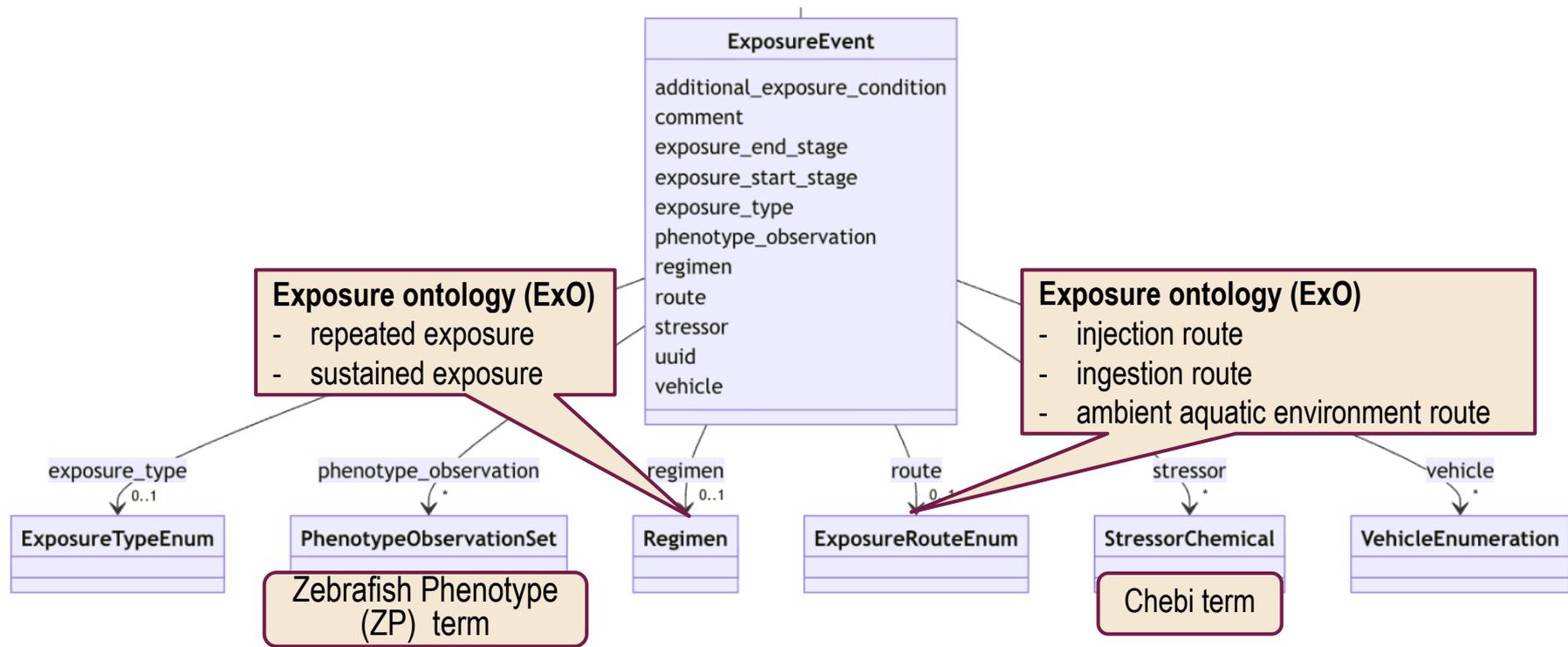
LinkML-based ; <https://github.com/zappfish/zebrafish-toxicology-atlas-schema>

Zebrafish Toxicological Exposure Data Model



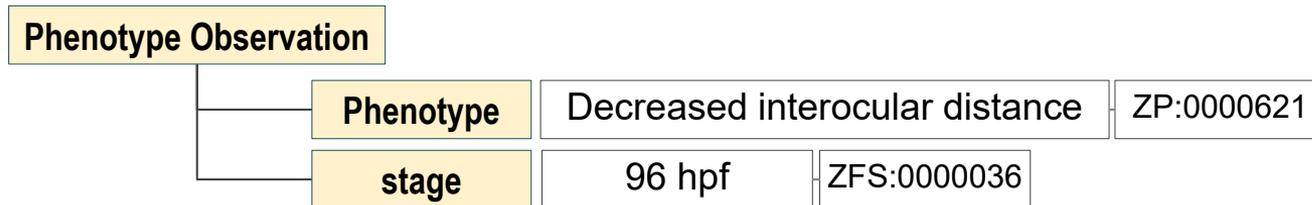
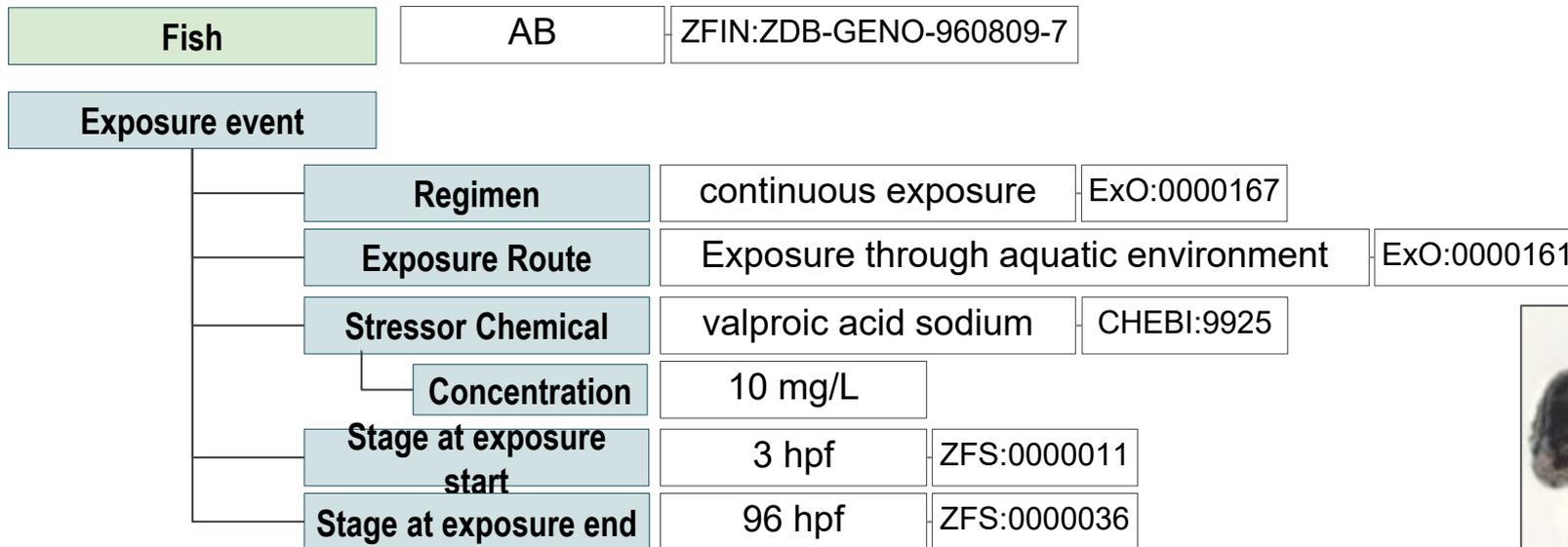
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Zebrafish Toxicological Exposure Data Model



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Annotations following the data model: example



Part of the model omitted for clarity.

Ontologies and Standard Vocabularies

We **reuse existing** ontologies/terminologies, and **improve/enhance them** as needed.

- **Exposure ontology (ExO)**
 - Species non-specific
 - We created new terms to represent exposure route and exposure regimen
- **Environment Exposure Ontology (ECTO)**
 - Species non-specific
 - We currently use the term “exposure to chemical” (ECTO:0000231)
 - ECTO allows for the representation of other exposure type (eg: exposure to smoke - ECTO:7000059)
- **Zebrafish Phenotype Ontology (ZP)**
 - Specific to Zebrafish, but already integrated into uPheno (species non-specific phenotype ontology)
 - Computationally created from ZFIN phenotype annotations
 - Represented by concatenation of different ontology terms
 - Terms names are cumbersome and non-intuitive
 - We are improving ZP by:
 - creating user friendly names
 - adding definition to terms

Standards for Toxicology Assay and Phenotype.

Data Model

LinkML-based Zebrafish
Toxicological Exposure Data Model

<https://github.com/zappfish/zebrafish-toxicology-atlas-schema>

Ontologies

Exposure ontology (ExO)

<https://github.com/CTDbase/exposure-ontology>

Environment Exposure Ontology (ECTO)

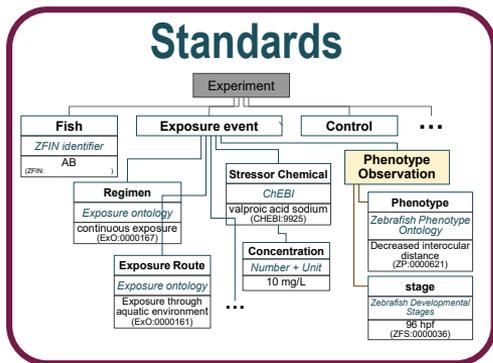
<https://github.com/EnvironmentOntology/environmental-exposure-ontology>

Zebrafish Phenotype Ontology (ZP)

<https://github.com/obophenotype/zebrafish-phenotype-ontology>

Zebrafish Anatomy Ontology (ZFA)

<https://github.com/ZFIN/zebrafish-anatomical-ontology>

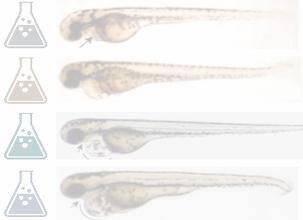


Standards are created for Zebrafish toxicological assay, but planned to be reusable or extendable to other species/model and other toxicological assay

Standards adoption

Data Collection

Zebrafish community



Data Annotation (using Standards)

Provenance of Data

Control Information

Images

Control information

Experiment information

Fish

Exposure Event

Stressor CAS Identifier

Stressor concentration Unit μM mg/L Other

Exposure route

via environment (chemical in the water)

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Exposure regimen

Continuous exposure Repeated exposure

Stage at exposure start Hours

Stage at exposure end Hours

Phenotype Observation

Images

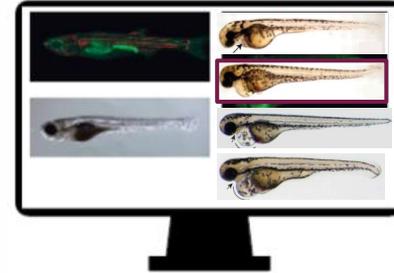
Phenotypes shown in the image

Stage at observation Hours

Submit to the Atlas



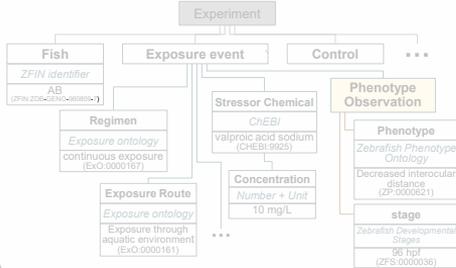
Zebrafish Phenotype Atlas



Fish + Annotations

- >Fish:
 - Strain: strain
 - Genotype: Genotype
 - Allele: Allele
 - ...
- >Experimental conditions
 - Toxicant: Toxicant
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 - ...
- >Phenotype: Phenotype

Standards

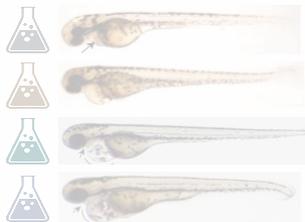


Annotation Toolkit

Standards adoption

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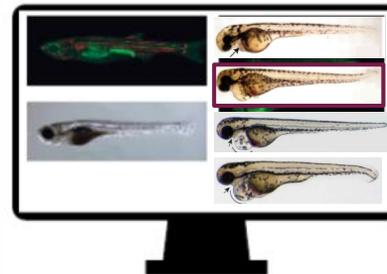
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Zebrafish Phenotype Atlas



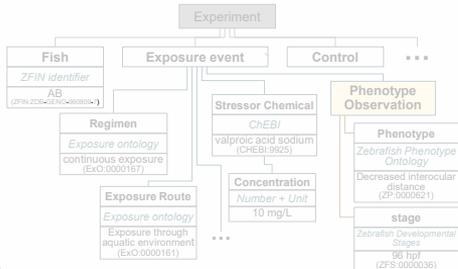
Publish standardized data



Fish + Annotations

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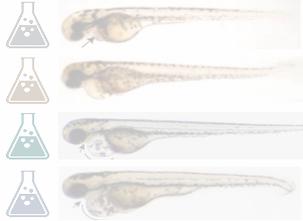


Annotation Toolkit

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Data Annotation (using Standards)

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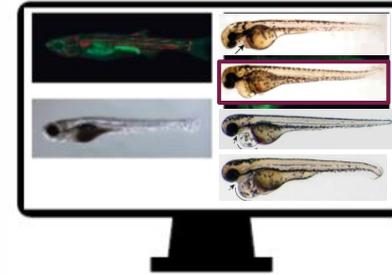
Phenotypes shown in the image

Stage at observation Hours

Submit to
the Atlas



Zebrafish Phenotype Atlas



Publish
standardized
data



The ultimate goal is for the reported data to be unambiguous, comparable, reproducible, and reusable.

Inputs + Annotations

>Fish: 
Strain: strain
Genotype: Genotype
Allele: Allele
...
>Experimental conditions
Toxicant: Toxicant
Concentration: mg/l
Timing of exposure: 2 h
...
>Phenotype: Phenotype

The Annotation Toolkit simplifies standards adoption

The annotation toolkit allows the use of standards, without requiring expertise in ontologies or standards development.

The research community can use the tool to **annotate** their data using standards, and

- **submit** their annotated data to the Atlas
- **download** their annotated data for publication submission (e.g. supplementary data of a manuscript)

Provenance of Data

▶

Control Information

Images

▶ Control information

Experiment information

▶ Fish

Exposure Event

▼

Stressor CAS Identifier

Stressor concentration Unit μM mg/L Other

Exposure route

via environment (chemical in the water)

via diet (ingestion)

via gavage

via injection

Exposure regimen

Continuous exposure Repeated exposure

Stage at exposure start

Stage at exposure end

Phenotype Observation

▼

Images

Phenotypes shown in the image

Stage at observation

The Annotation Toolkit simplifies standards adoption

Provenance of Data

Control Information
Images
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Experiment information

Fish

Exposure Event
Stressor CAS Id
Stressor concentration Unit μM mg/L Other

Exposure route
 via environment (chemical in the water)
 via diet (ingestion)
 via gavage
 via injection

Exposure regimen
 Continuous exposure Repeated exposure

Stage at exposure start
Stage at exposure end

Phenotype Observation
Images
Phenotypes shown in the image
Stage at observation

Fish Information
Strain/Background (WT) ⓘ

AB	ZDB-GENO-960809-7 • ZDB-FISH-150901-27842
AB/C32	ZDB-GENO-070425-3 • ZDB-FISH-150901-19012
AB/EKW	ZDB-GENO-091223-1 • ZDB-FISH-150901-18519
AB/TL	ZDB-GENO-031202-1 • ZDB-FISH-150901-29235
AB/TU	ZDB-GENO-010924-10 • ZDB-FISH-150901-29084

- Common names found with **autocomplete**
- IDs are shown to bring awareness

The Annotation Toolkit simplifies standards adoption

Provenance of Data

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Continuous exposure Repeated exposure

Stage at exposure start Hours

Stage at exposure end Hours

Phenotype Observation

Images

Phenotypes shown in the image

Stage at observation Hours

Limited options when constraints in data model

User friendly names

Translation to ID behind the scenes

Exposure route

- via environment (chemical in the water)
- via diet (ingestion)
- via gavage
- via injection

ExO:0000161
ambient aquatic environment route

Exposure regimen

- Continuous exposure
- Repeated exposure

ExO:0000167
sustained exposure

The Annotation Toolkit simplifies standards adoption

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- via environment (chemical in the water)
- via diet (ingestion)
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- via injection

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Continuous exposure Repeated exposure

Stage at exposure start

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Phenotype Observation

Images

Phenotypes shown in the image

Stage at observation

Pick a term from the Zebrafish Phenotype Ontology

Ontology Term Picker Tool



The **Ontology Term Picker** allows users to find phenotype terms without the need to understand how ontologies work.

Phenotype Picker

Use this tool to find phenotype terms for your zebrafish research. Browse the **Zebrafish Anatomy (ZFA)** hierarchy on the left, then select associated **phenotype terms** from the Zebrafish Phenotype Ontology (ZP) on the right.

Filter by Anatomy

- zebrafish anatomical entity
 - anatomical line

Phenotype Results



microencephaly



Filter by Anatomy

- zebrafish anatomical entity
 - anatomical line
 - anatomical space
 - anatomical structure
 - anatomical surface
 - portion of organism substance
 - unspecified

Clear search to
browse by
anatomy



Search results for "microencephaly"

Filter within results...

1 phenotype

microencephaly

ZP_0000711 • 307 ZF

microencephaly

Synonyms

- abnormally small brain
- brain decreased size, abnormal

Definition

An abnormal reduction in the size of the brain.

http://purl.obolibrary.org/obo/ZP_0000711

ZFin usages: 307

Request changes



Filter by Anatomy

- nervous system
 - autonomic nervous system
 - central nervous system
 - brain
 - brain nucleus
 - brain vasculature
 - cephalic flexure
 - choroid plexus
 - choroidal fissure
 - forebrain
 - forebrain midbrain boun...
 - granule cell
 - hindbrain
 - midbrain
 - midbrain hindbrain bou...
 - supraoptic tract
 - brainstem
 - floor plate
 - grey matter
 - neuraxis flexure
 - neuromere

Find term
using
anatomy

Phenotypes for brain

Filter within results...

Showing 50 of 118 phenotypes

abnormal brain morphology

ZP_0000100 • 356 ZFIN references



abnormal brain

ZP_0000032 • 343 ZFIN references



microencephaly

ZP_0000711 • 307 ZFIN references



abnormally necrotic brain

ZP_0002272 • 294 ZFIN references



abnormally increased number of brain

ZP_0010501 • 252 ZFIN references



Filter by Anatomy

- nervous system
 - autonomic nervous system
 - central nervous system
 - brain
 - brain nucleus
 - brain vasculature
 - cephalic flexure
 - choroid plexus
 - choroidal fissure
 - forebrain
 - forebrain midbrain boun...
 - granule cell
 - hindbrain
 - midbrain
 - midbrain hindbrain bou...
 - supraoptic tract
 - brainstem
 - floor plate
 - grey matter
 - neuraxis flexure
 - neuromere

Showing the ontology tree suggests a more specific terms

Phenotypes for brain

increased size

4 phenotypes matching "increased size"

increased size of the macrophage in brain

ZP_0148851

• 2 ZFIN references

increased size of the proliferative region in brain

ZP_0007309

• 1 ZFIN references

macroencephaly

ZP_0003410

• 16 ZFIN references

brain hyperplasia

ZP_0015592

• 1 ZFIN references

Filter by Anatomy

- nervous system
 - autonomic nervous system
 - central nervous system
 - brain
 - brain nucleus
 - brain vasculature
 - cephalic flexure
 - choroid plexus
 - choroidal fissure
 - forebrain
 - forebrain midbrain boun...
 - granule cell
 - hindbrain
 - midbrain
 - midbrain hindbrain bou...
 - supraoptic tract
 - brainstem
 - floor plate
 - grey matter
 - neuraxis flexure
 - neuromere

Phenotypes for brain

increased size

4 phenotypes matching "increased size"

I cannot find my phenotype

increased size of the macrophage in brain

ZP_0148851 • 2 ZFIN references

increased size of the proliferative region in brain

ZP_0007309 • 1 ZFIN references

macroencephaly

ZP_0003410 • 16 ZFIN references

brain hyperplasia

ZP_0015592 • 1 ZFIN references

Ontology Term Picker Tool



The **Ontology Term Picker** allows users to find phenotype terms without the need to understand how ontologies work.

This tool is developed to **accommodate 'any' ontologies**, especially ontologies using 2 types of concepts

Examples: phenotype and anatomy
breed and taxon
disease and gene

The Annotation Toolkit simplifies standards adoption



The annotation toolkit allows the use of standards, without requiring expertise in ontologies or standards development.

The research community can use the tool to **annotate** their data using standards, and

- **submit** their annotated data to the Atlas
- **download** their annotated data for publication submission (e.g. supplementary data of a manuscript)

Provenance of Data

Control Information

Images

Control information

Experiment information

Fish

Exposure Event

Stressor CAS Identifier

Stressor concentration Unit μM mg/L Other

Exposure route

via environment (chemical in the water)

via diet (ingestion)

via gavage

via injection

Exposure regimen

Continuous exposure Repeated exposure

Stage at exposure start

Stage at exposure end

Phenotype Observation

Images

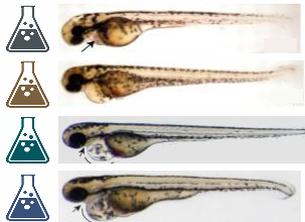
Phenotypes shown in the image

Stage at observation

Zebrafish Phenotype Web Application

Data Collection

Zebrafish community



Data Annotation (using Standards)

Provenance of Data

Control Information

Images

Control information

Experiment information

Fish

Exposure Event

Stressor CAS Identifier

Stressor concentration Unit μM mg/L Other

Exposure route

via environment (chemical in the water)

via diet (ingestion)

via gavage

via injection

Exposure regimen

Continuous exposure Repeated exposure

Stage at exposure start Hours

Stage at exposure end Hours

Phenotype Observation

Images

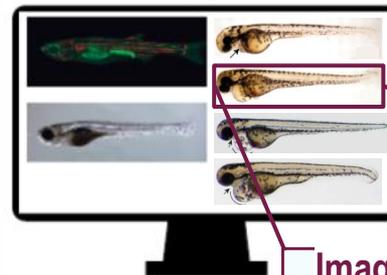
Phenotypes shown in the image

Stage at observation Hours

Submit to the Atlas



Zebrafish Phenotype Atlas



Images + Annotations

>Fish:

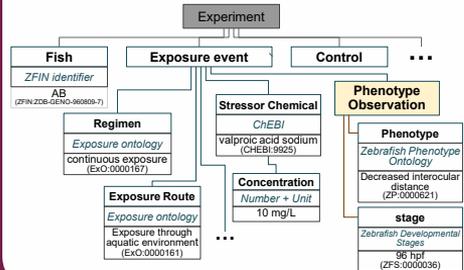
Strain: strain
Genotype: Genotype
Allele: Allele

>Experimental conditions

Toxicant: Toxicant
Concentration: mg/l
Timing of exposure: 2 h

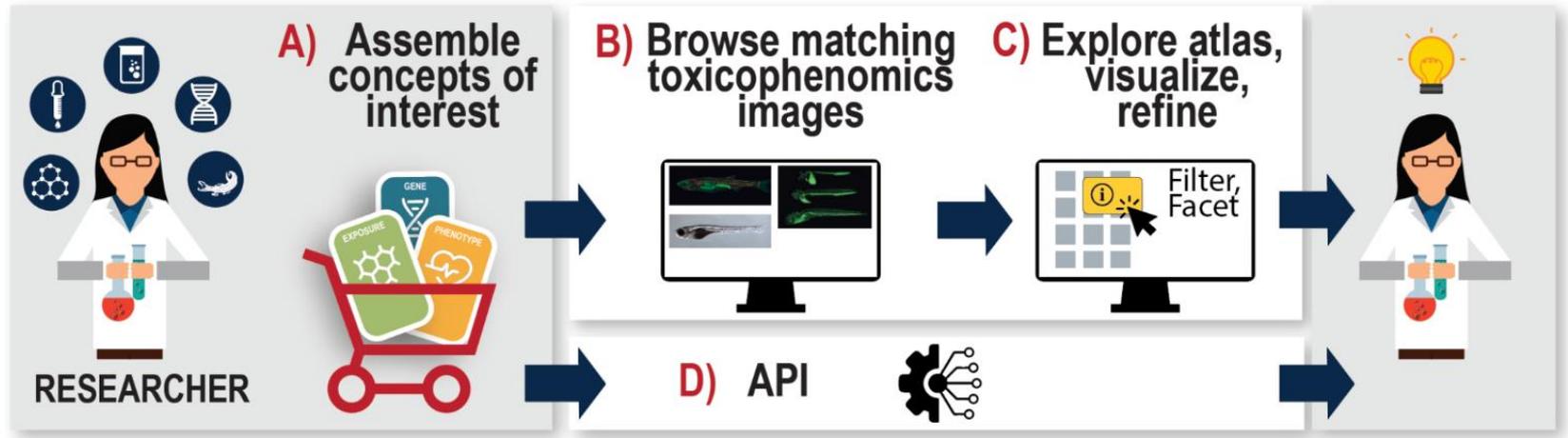
>Phenotype: Phenotype

Standards



Annotation Toolkit

Zebrafish Phenotype Web Application



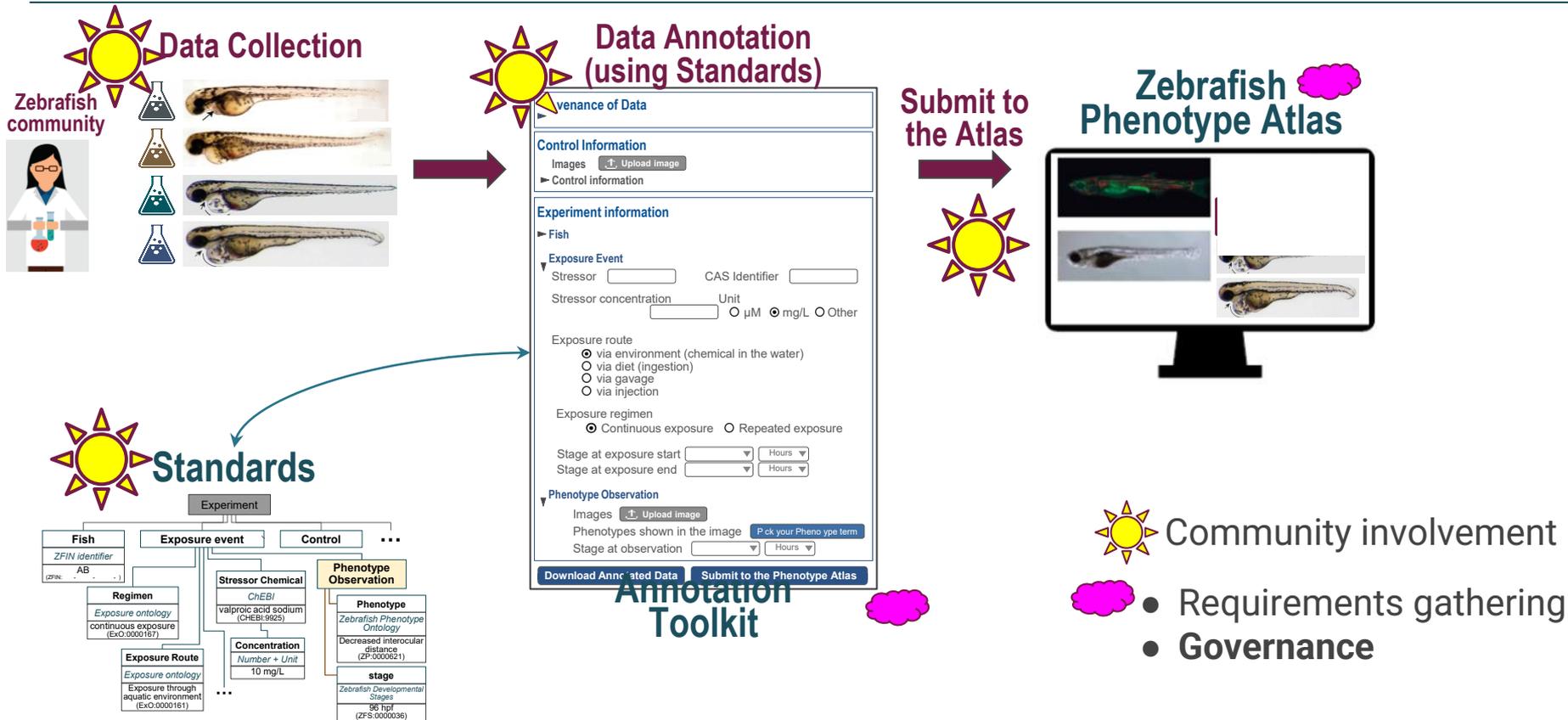
Functionality

- Browsing
- Filtering
- *TBD based on community requirements*

Features

- Open access
- All data attributed and provenanced
- Content submitted by the community
- All data standardized
- Data “peer reviewed” - *process TBD*

Community involvement is at the core of ZAPP



The Zebrafish Research Community is our primary community

The Research Community understands and agrees that the need for standards is crucial. However, they **don't know what to do in practice**.

- How to use standards in existing research workflow?
- What are the standards and how to use them?
- Where to find the standards?
- What to do when what is needed is not available?

We meet the community where they are, and we “nudge them” into understanding standards and adopting them.



Engaging with the community: Meeting the community where it is

- Speak “their” language: communication style and vocabulary
 - *e.g avoid using the words “data model”*
 - *e.g workshops and hands-on exercises to gather requirements*
- Develop tools that allow standards adoption without the need to understand what standards are and how they are developed

Engaging with the community: Fostering community awareness

- “Standards” is not a commonly discuss topic in the research community.
- We bring awareness through: workshops, posters, presentations at conferences.
- We partner with conference organizers** to create working groups and dedicated sessions to
 - highlight standards and community projects (which are rarely discussed at zebrafish research conferences)
 - promote how (and why) the research community can (and should) get involved

*** This has been possible thanks to Alexa Burger’s connections with the Zebrafish research community and position on the International Zebrafish Society (IZFS) board of directors*

Engaging with the community:

Clearly communicating how people can participate

- Join the ZAPP community:
 - tislabs.org/ZebraPhenoAtlas-signup
 - Slack channel on Zebrafish Rock!: #zapp-fish
- Participate in the workshops
 - Gather information about data standards, all information required
 - Gather feedback, community needs, etc
- Send us feedback
 - About what you need, what you want
 - About how we can make the use of standard easier (ie with minimal disruption of every day research)
 - Error, misrepresentation, changes
- Use the tools** (annotation tools and atlas) and submit your data

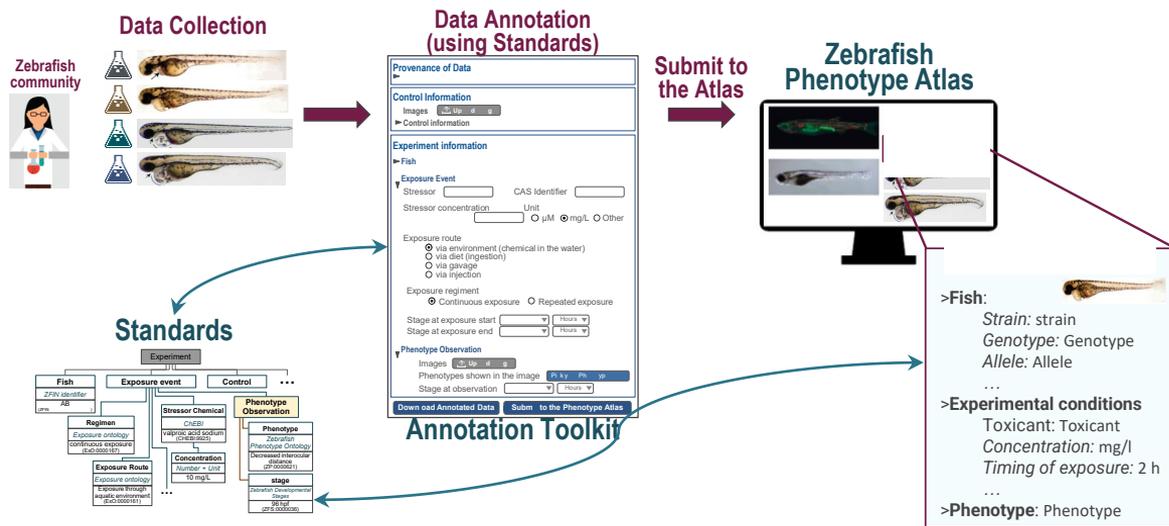


** when available

The Zebrafish Toxicology Phenotype Atlas (ZAPP)

Community-led Zebrafish Toxicology Phenotype Atlas (ZAPP) will

- integrate existing standards, and develop new one when needed, for phenotypic and toxicology data
- make standard adoption approachable
- support community-driven improvement



The ultimate goal is for the reported data to be unambiguous, comparable, reproducible, and reusable.

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tislabs.org/ZebraPhenoAtlas-signup

