

Examining the effect of low vs high
doses of TCDD and Mercury both alone
and in combination on *Microgadus*
tomcod

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Why is my project important/interesting ?

- Diamond Alkali
 - TCDD
 - Not an intended product
 - Effects AHR pathway
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- Effects
- Hackensack River
 - Mercury
- Effects in human are well known
- How does it affect organisms in waterways?

What is the experimental question and what I am hoping to discover from my analysis ?

- How do certain chemicals affect organisms differently at low doses versus high doses and therefore this should also be taken into account.
 - TCDD
 - Mercury
- Look at difference in expression particularly for genes associated with the Atlantic tomcod heart

What is the experimental design (how many factors, how many levels , how many replicates) ?

- Organism: *Microgadus tomcod*
 - Why?
 - Resistance
- Chemicals:
 - TCDD
 - Mercury
 - Low vs. High doses
 - In combination
- Control
 - Acetone
 - Water
- Averaging each group

What am I planning to use?

- Salmon
- EdgeR & DESeq2

- Visualization
 - Java TreeView 3.0

- Clustering
 - ClustVis

Are there any obstacles?

- Setting a cut off for fold change
 - Suggested: 1.5
 - Arbitrary
 - Initially useful
- Lack of high quality reference genome
 - Perhaps, another fish genome can be used as reference.
- Limitation
 - Not like a waterway

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