



# MULTIPLE CONTAMINANTS IN COMPLEX COMMUNITIES EVALUATING NON-ADDITIVE EFFECTS OF MULTIPLE SIMULTANEOUS STRESSORS ON BIOMASS FLUX AND ECOSYSTEM FUNCTIONING



Top consumer

Producer

Intermediate consume

# Variation in…

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Current approaches to **environmental risk assessment** (ERA) are often limited to assessing the effect of single contaminants on single species, **CHALLENGE**<br>Current approaches to environmental risk assessment (ERA)<br>overlooking the effects that can occur at higher ecological scales.

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REVETIMING ERA<br>
How do multiple contaminants<br>
impact biodiversity, stability<br>
and ecosystem function through. impact biodiversity, stability and ecosystem function through…

Can we develop tools for ERA that allow us to evaluate how **multiple contaminants**  acting on **multiple traits** among species in communities impact **biodiversity, stability and ecosystem function**?

# RESULTS RESCALED DEVIATION FROM ADDITIVITY of Community Biomass

Top consumer

Intermediate consumer

Target trait (*mode of action*) Target species (*contaminant specificity*) Target trophic level Community size, structure and complexity Environmental conditions

## Food Web modelling

**Biomass dynamics in a community are determined by species traits [1]** 



# METHODS

FUTURE WORK

**Rescaled DA for Community Biomass** 

> 2.0  $1.3$  $0.5$

> > $-9.5$

**Community biomass** is differentially impacted depending on the **trait and trophic level** targeted by the pesticide

**Additivity, antagonistic buffering and antagonistic suppression** are the most commonly observed interaction types, with **no synergy** observed

**Antagonistic suppression** is observed when pesticides target **foraging** in the **intermediate consumer**, due to compensatory biomass dynamics inherent to the model resulting in reallocation of biomass among trophic levels 3

1. Explore buffering and suppression of total biomass by exploring biomass re-allocation across trophic levels 2. Apply method to stability 3. Expand community complexity

#### References

#### metabolism Metabolism Foraging Trait Rate Foraging  $QQQQQQQQQQ$ Growth  $\boldsymbol{C}$ Stressor intensity **GROWTH** 3 species case study Measuring Biomass Varying target trait Pesticide Contro top Metabolism of TOP  $0.8$ Foraging of top

**[1] Williams, Brose & Martinez (2007) Homage to Yodzis and Innes 1992: Scaling up feeding-based population dynamics to complex ecological networks [2] Tekin** *et al.* **(2020) Using a newly introduced framework to measure ecological stressor interactions**

### In silico experimental design

- Use differential equation **food web model** to simulate dynamics of plausible **tri-trophic** food chain
- Specify inhibitory **contaminant effects** on populations via linear reductions or increases of trait rates
- Generate 2-contaminant scenarios with
	- **1 herbicide targeting growth** and
	- **1 pesticide targeting either metabolism or foraging**
- Measure **community biomass**
- **Classify interactions** by calculating deviation from additivity of community biomass**[3]**





Using Tekin *et al.*'s framework for measuring **ecological stressor interactions [2]** , which incorporates;

- **Standardisation** of effect sizes (rescaling)
- **Categorisation** of interaction types
- Measures **Deviation from Additivity** (DA) - Synergy, Buffering and Suppression

1 2



## Contaminant Effects

## Classifying Interactions



- Synergy joint effect greater than sum of its parts
- Additive no interaction
- Antagonistic buffering joint effect smaller than sum of its parts
- Antagonistic suppression (negative) effect of negative stressor masks that of the positive stressor
- Antagonistic suppression (positive) effect of positive stressor masks that of the negative stressor