



Potential interactions between UV filters and environmental microbial communities

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Patricia A. Holden, Ph.D., P.E.

*Professor, Bren School of Environmental Science & Management
Earth Research Institute, Univ. of Calif., Santa Barbara, CA 93106, U.S.A.*



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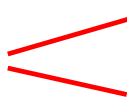
DBI-0830117
DBI-0216480



Approved active ingredients in sunscreen

Aminobenzoic acid
Avobenzone
Cinoxate
Dioxybenzone
Homosalate
Meradimate
Octocrylene
Octinoxate
Octisalate
Oxybenzone
Padimate O
Ensulizole
Sulisobenzene
Titanium dioxide
Trolamine salicylate
Zinc oxide

UV filters



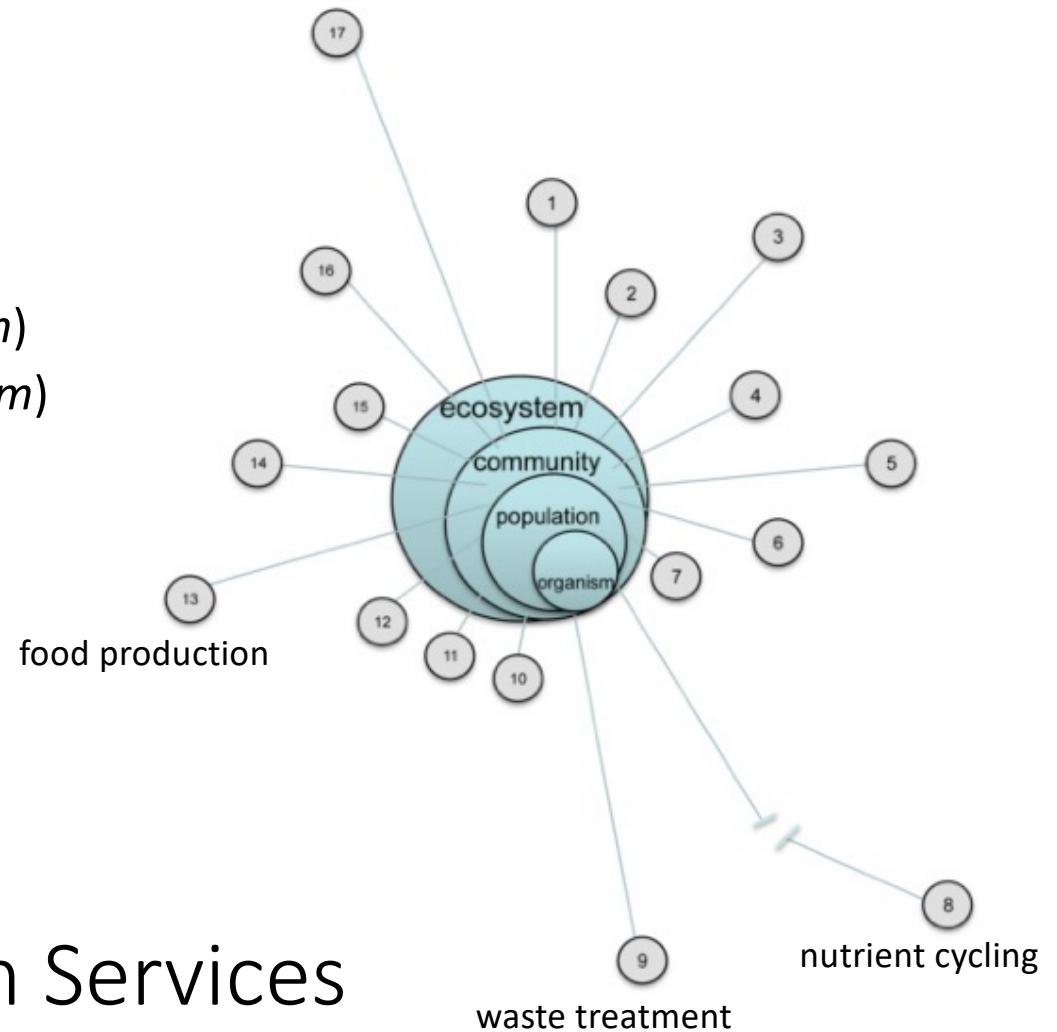
<https://www.fda.gov/drugs/understanding-over-counter-medicines/sunscreen-how-help-protect-your-skin-sun#ingredients>

questions

- How might microbial communities transform UV filters?
- How might UV filters affect microbial communities?

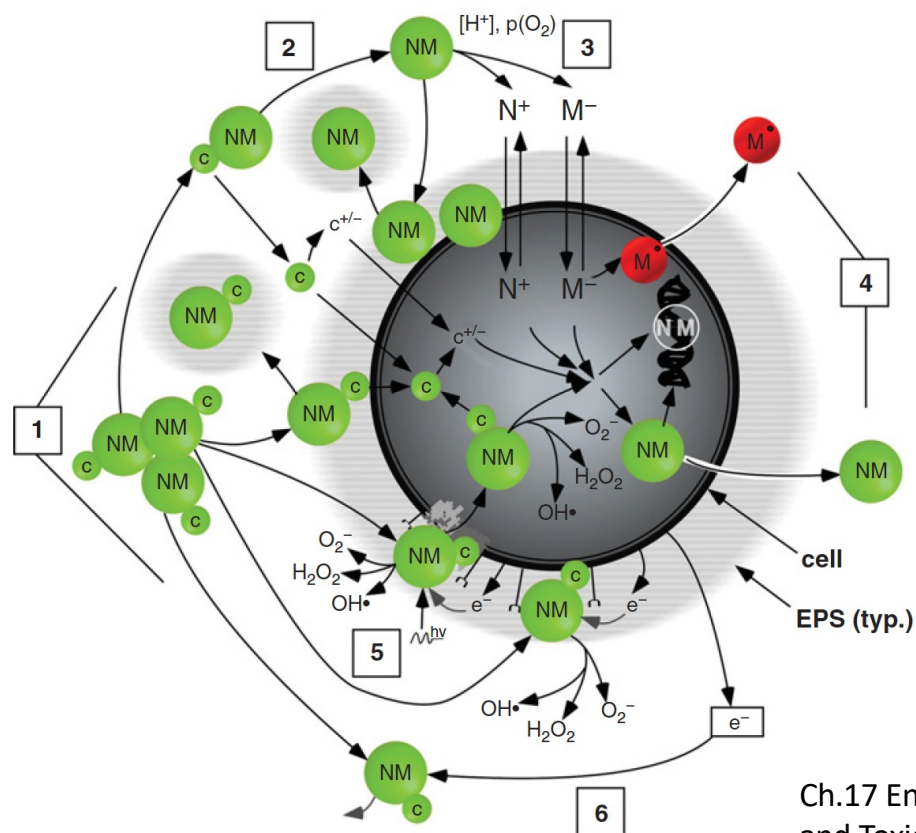
Microbial Communities

- Microbes
 - bacteria (*Bacteria* and *Archaea*; $1\ \mu\text{m}$)
 - fungi, algae, protozoa (*Eukarya* $10\ \mu\text{m}$)
 - viruses ($1\ \text{nm}$)
 - symbioses



Ecosystem Services

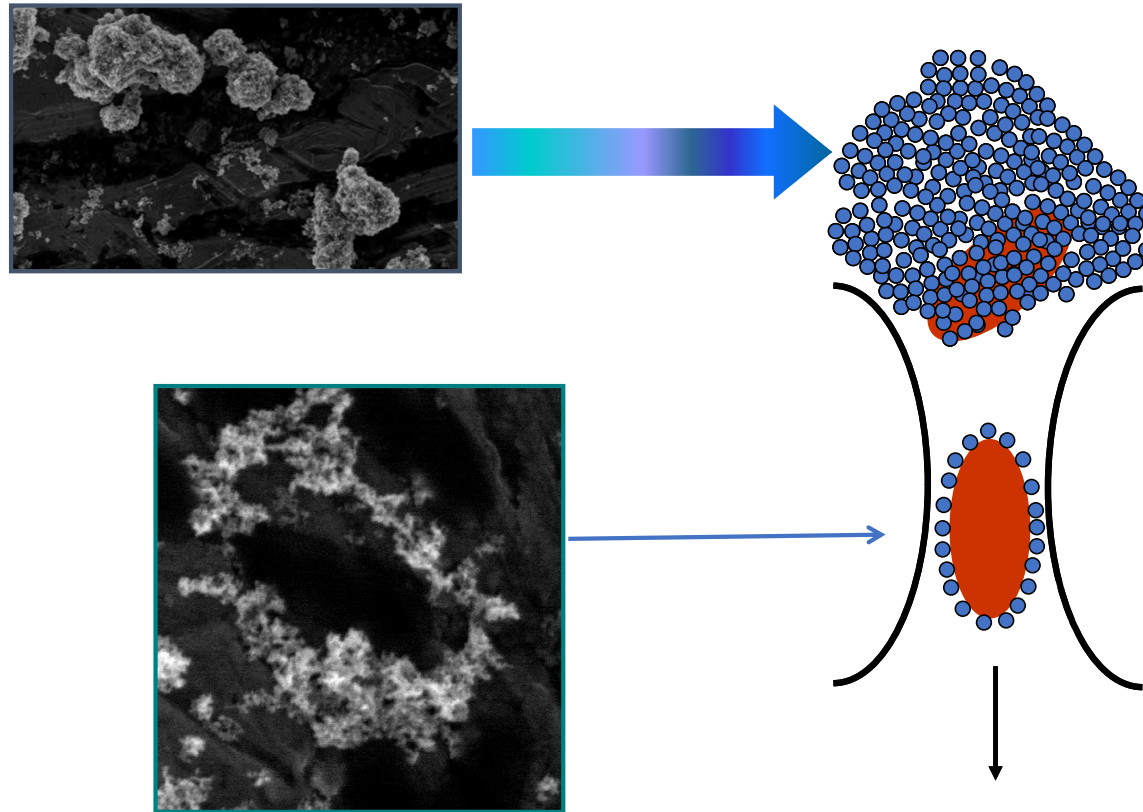
Potential Cell-NM Interactions



1. Sorption / desorption
2. Cap or coating loss
3. Decomposition
4. Expulsion
5. Membrane interactions
6. Extracellular e⁻ transfer

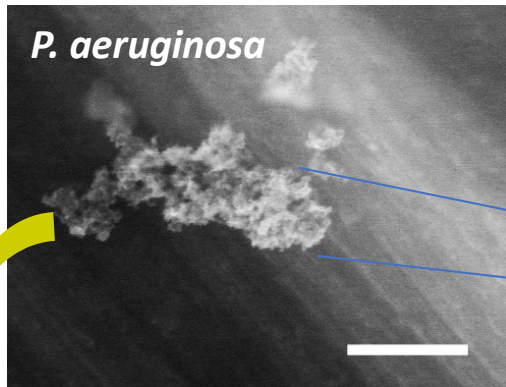
Ch.17 Engineered Nanoparticles and the Environment: Biophysicochemical Processes and Toxicity, First Edition. Edited by Baoshan Xing, Chad D. Vecitis and Nicola Senesi. © 2016 John Wiley & Sons, Inc. Published 2016 by John Wiley & Sons, Inc.

1. nTiO_2 sorbs to bacteria; disagglomerates

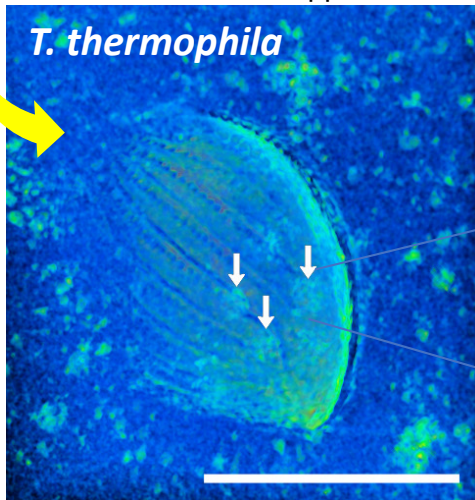
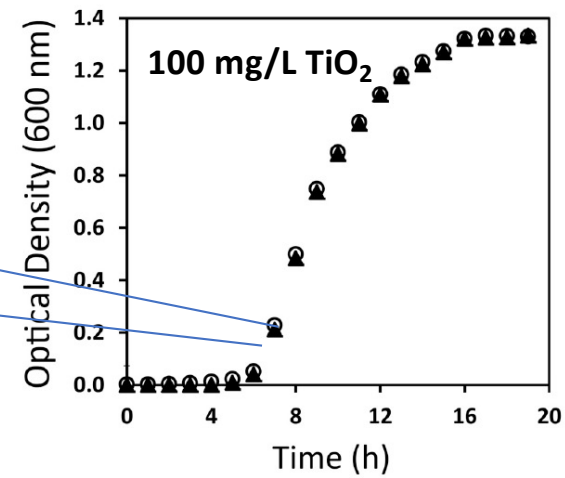


Horst *et al.* 2010. Appl. Environ. Microbiol.

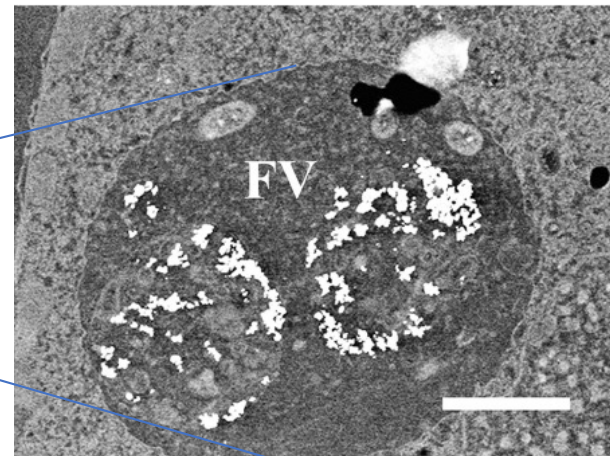
1. nTiO₂ Trophic Transfer to Protozoans



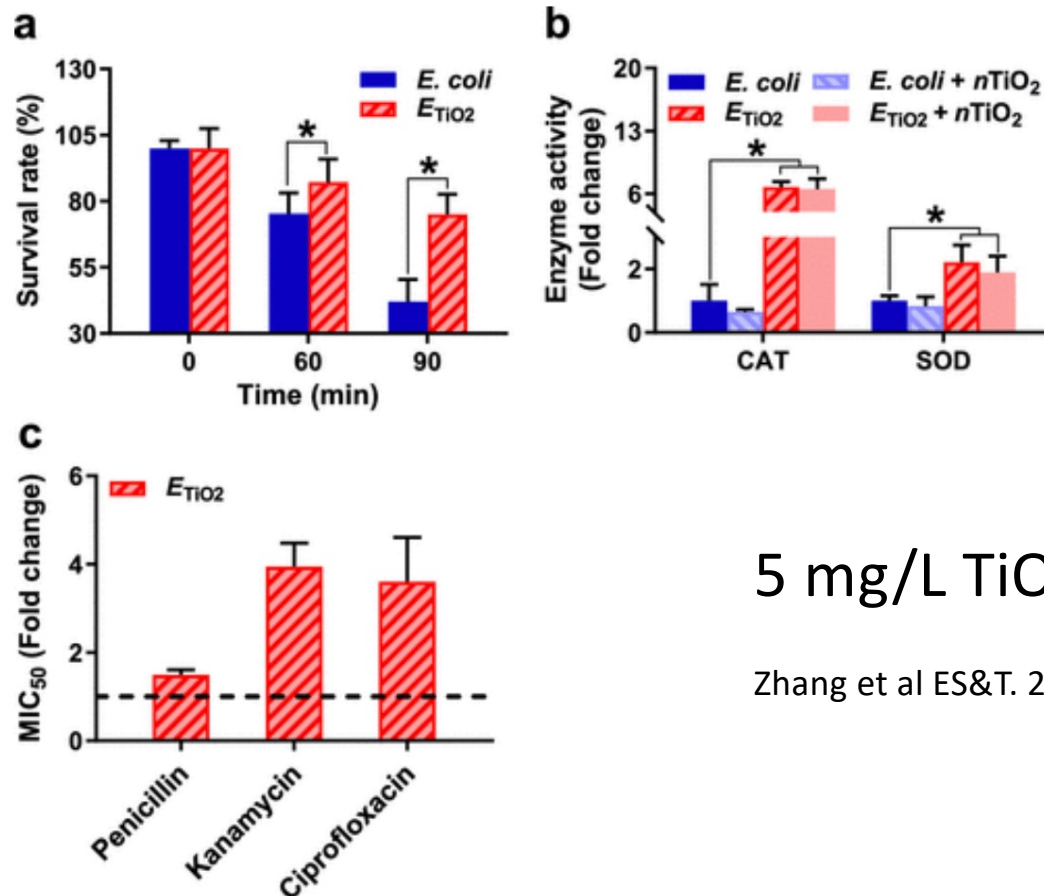
Horst *et al.* 2010. Appl. Environ. Microbiol.



Mielke *et al.* 2013. Appl. Environ. Microbiol.



2. Commensal to Pathogen Transition in *E. coli*



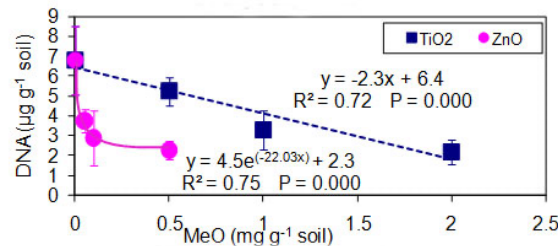
5 mg/L TiO₂

Zhang et al ES&T. 2020

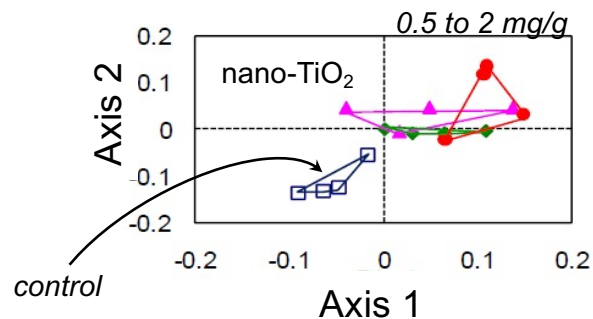
2. Evidence for ZnO and TiO₂ MNMs Impacting Soil Bacterial Communities



Decrease Soil DNA

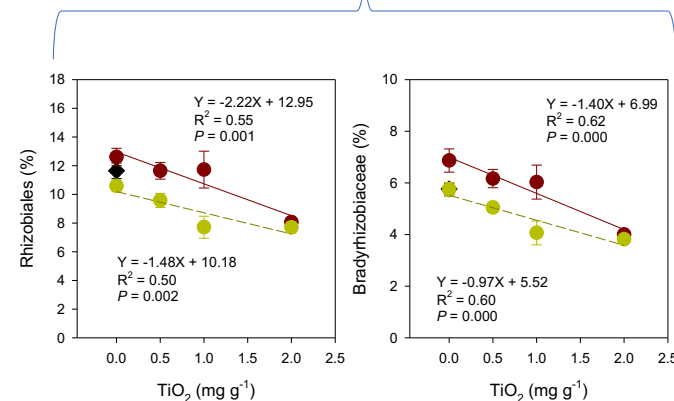


Change bacterial community



Ge et al. 2011. ES&T.

Alter select taxa abundance



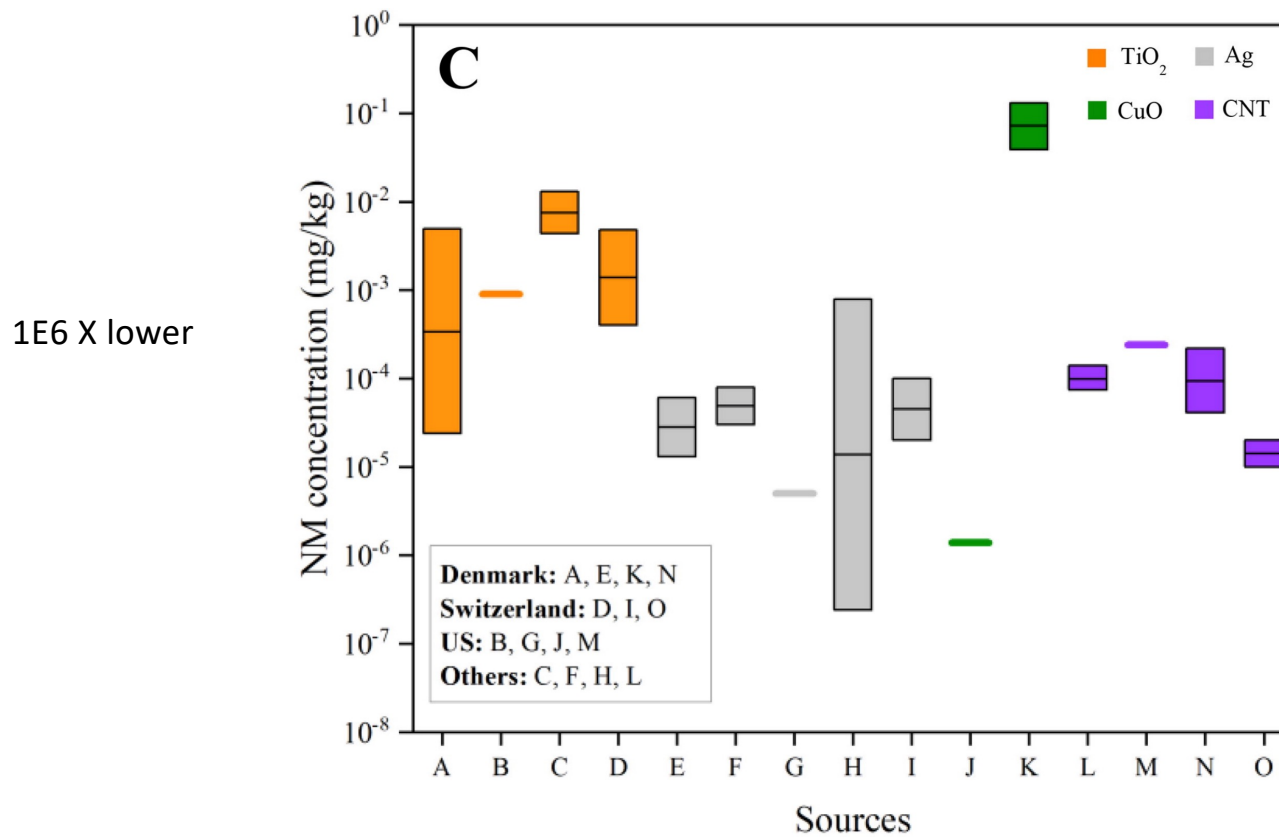
Upper: 15d
Lower: 60d

Incl.: N₂ fixation, C cycling, CH₄ oxidation

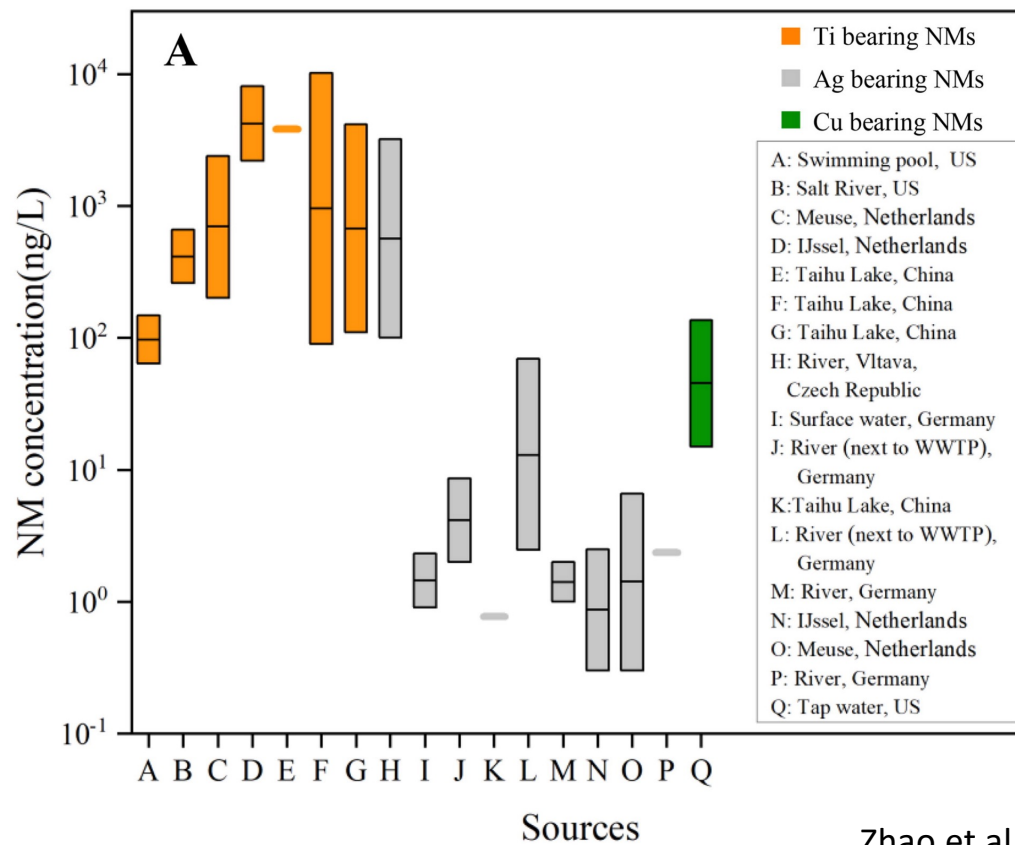
Potential Ecosystem Effects

Ge et al. 2012. Appl. Environ. Microbiol.

2. Concentrations (Ti NMs) in soil



2. Concentrations (Ti NMs) in waters

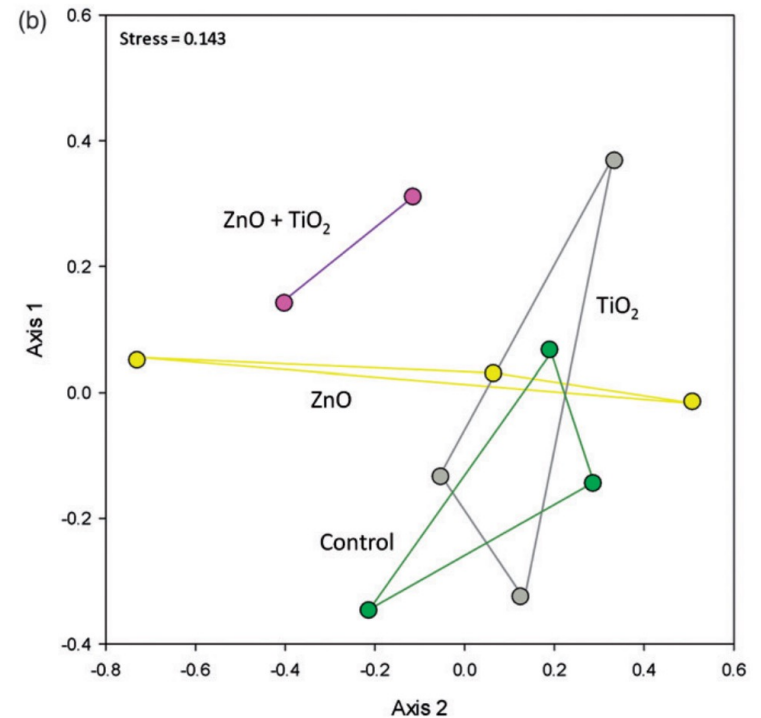
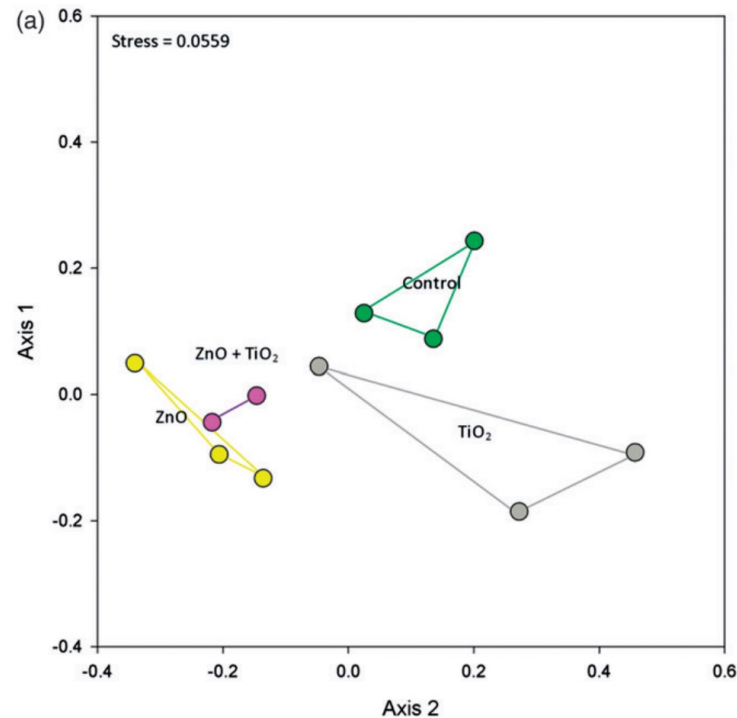


Zhao et al. 2020. Crit. Rev. Environ. Sci. Technol.

2. Effects on microbial communities

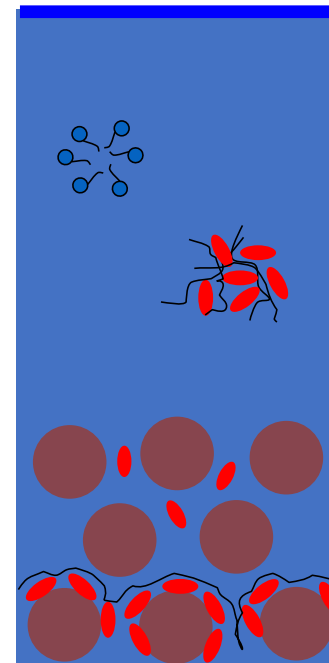
5 mg/L TiO₂: river water bacterial community membrane damage (light > dark; free living > biofilm) Battin et al. ES&T 2009.

700 µg/L TiO₂
70 µg/L ZnO
In river water;
Londono et al.
Nanotox. 2017



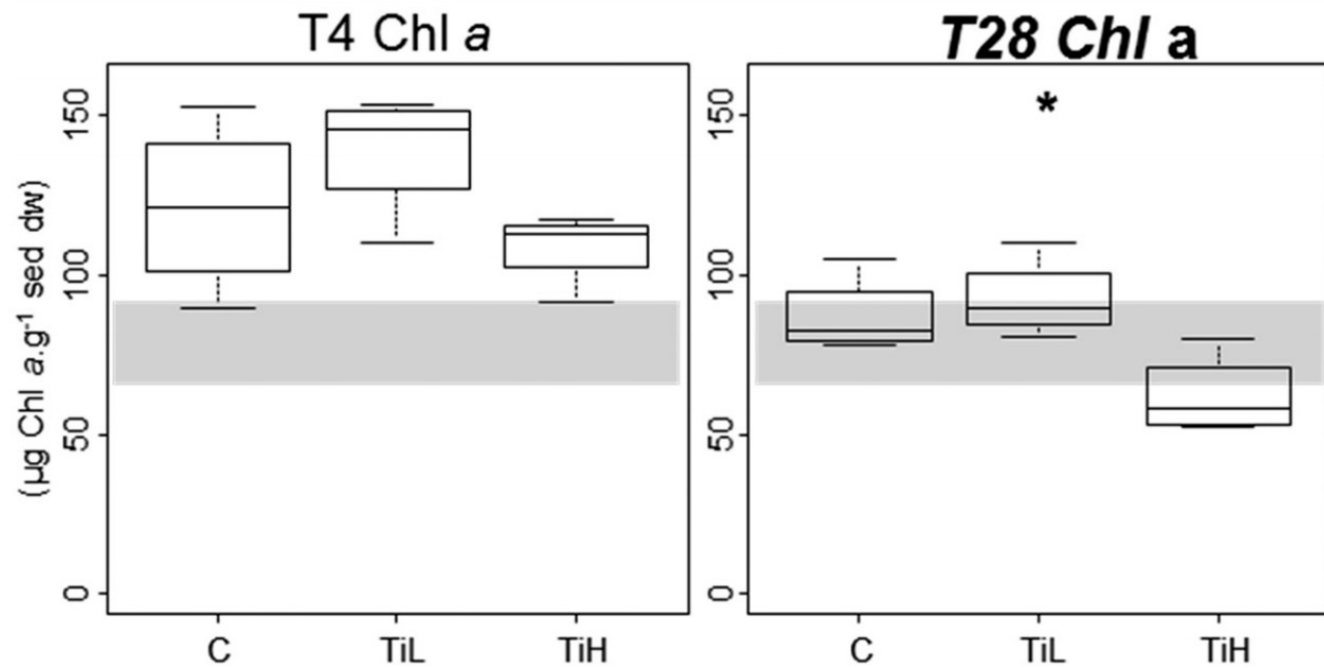
Microbes: aquatic Sub-compartments

- Air/ water interface (neuston)
- Column
 - Micelles
 - Flocs
 - Pore water
- Solid/water interface
 - biofilms



2. Primary production: coastal benthic algae

Winter

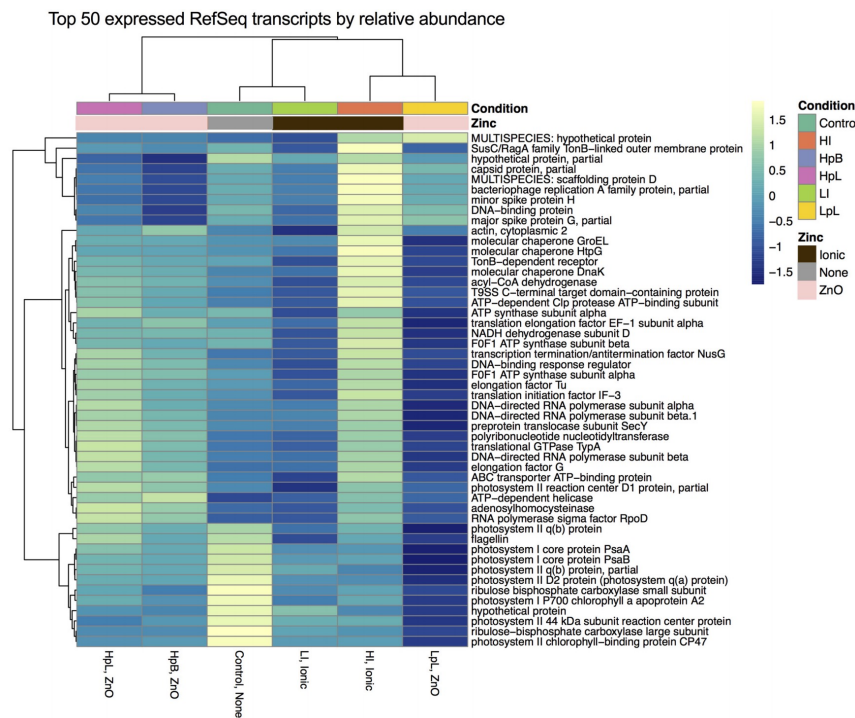


Passarelli et al. 2020. Environ. Sci. Nano.

25 $\mu\text{g/L}$

25 mg/L

2. Aquatic (riverine) biofilms responses to ZnO w/ coatings, and Zn ions



- Exposures on order of 100 $\mu\text{g/L}$
 - Biofilms concentrated Zn
- Gene abundances altered
 - Photosynthesis, N cycling, lipid degradation and synthesis, antibiotic resistance, xenobiotic degradation
- Yet overall function maintained through compensation
- Forms similar, although coatings matter
- Biofilm structure (‘dissolution’) changed (motility).

Summary Points

- *How might microbial communities transform UV filters?*
 - Physical association:
 - disagglomeration → dispersion → distribution
 - trophic transfer → food webs
 - EPS/ biomacromolecules: distribution, toxicological properties (corona)
- *How might UV filters affect microbial communities?*
 - community → composition → diversity → functional groups
 - f(concentration, media, physical state, compartment)
 - functional implications inferred
 - transcription
 - function (primary production, N cycling, waste treatment)
 - Pathogenicity
- Factors:
 - NM coating, concentration
 - Environmental compartment
 - Growth habit: biofilm vs floc or planktonic
- Methods: "omics" revealing process-related effects w/o necessarily large changes in diversity or stress response. Ecosystem-relevant microbial functional changes evidenced.