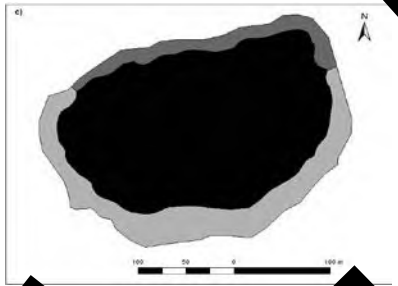




Predicting the effects of reintroducing a native predator (European eel, *Anguilla anguilla*) into a freshwater community dominated by alien species

Phillip J. Haubrock, Paride Balzani, Alberto Criado, Agustín P. Monteoliva & Elena Tricarico

Lake Arreo



Protected habitat

~ 136 ha

Max. depth: 24 m



Native species

Coleoptera



Gyrinus paykulli

Crayfish



Austropotamobius pallipes

Non-native species

Crayfish



Procambarus clarkii

Fish



Tinca tinca



Anguilla anguilla

Fish



Micropterus salmoides

Lepomis gibbosus

Cyprinus carpio

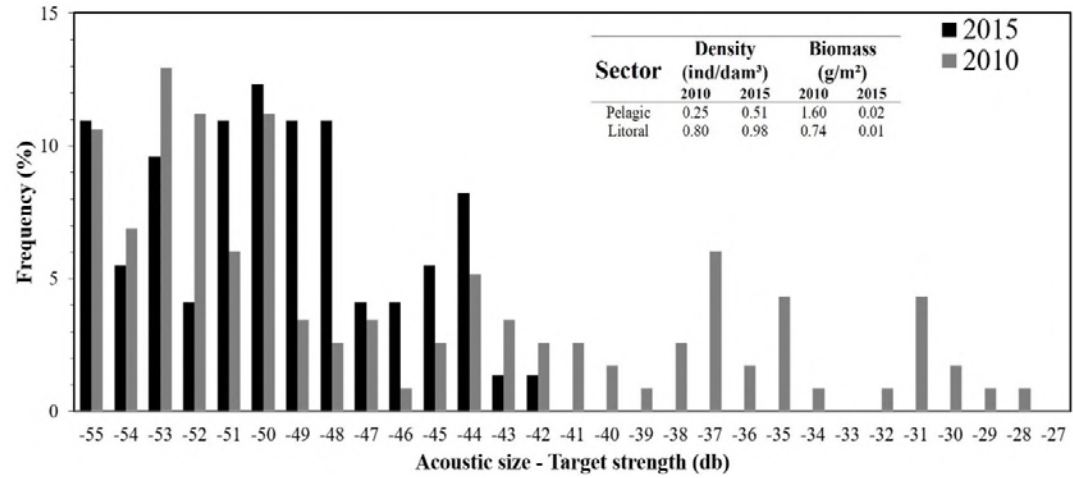
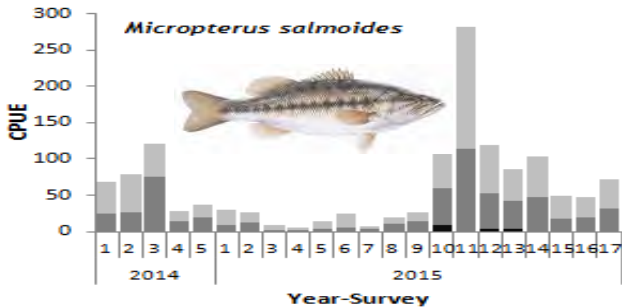
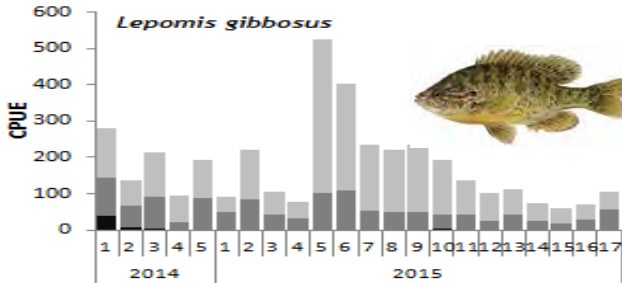
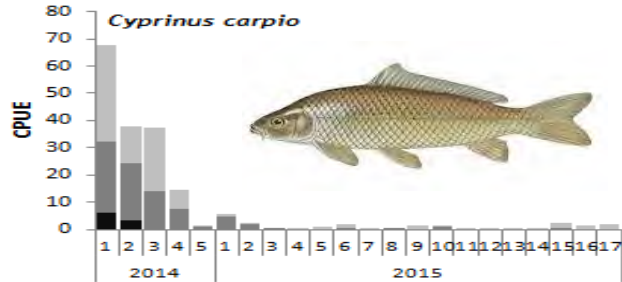
Vegetation



Phragmites australis

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Management



Anguilla anguilla

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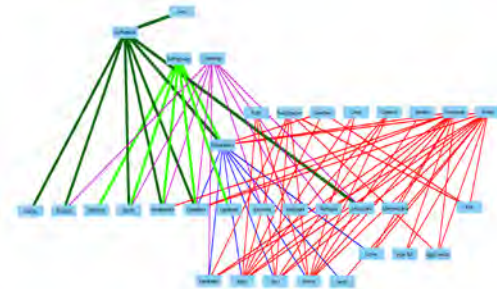
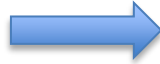
Sampling 2018

Species	Isotope Sample Size	Stomach Sample Size
<i>Micropterus salmoides</i>	15	50
<i>Lepomis gibbosus</i> *	15	50
<i>Cyprinus carpio</i>	11	11
<i>Procambarus clarkia</i>	15	50
<i>Phragmites australis</i>	5	



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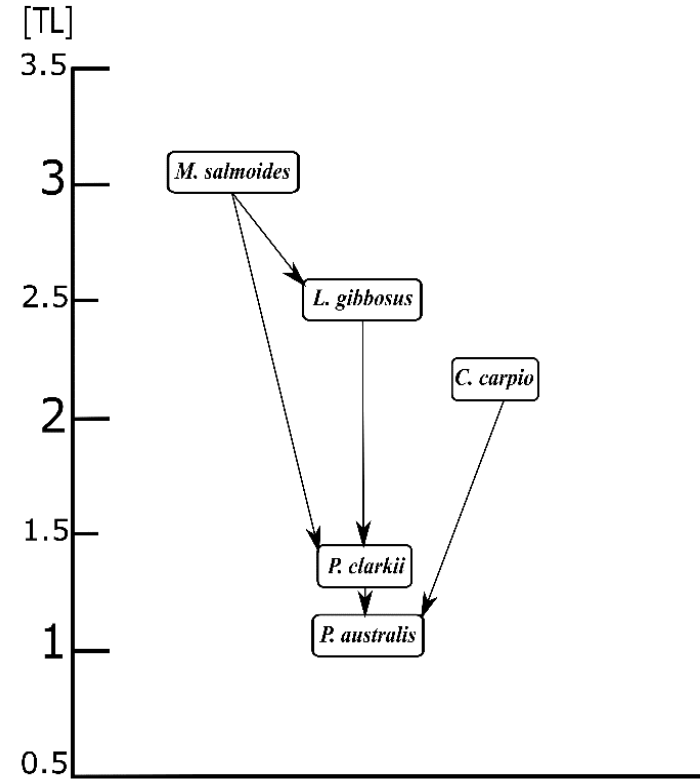
Laboratory procedure



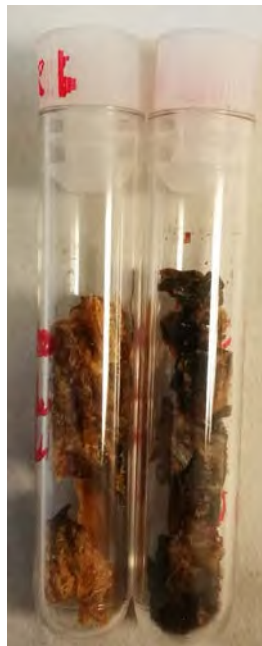
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Diet analysis

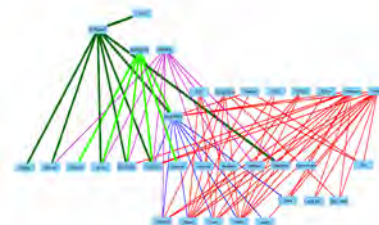
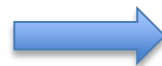
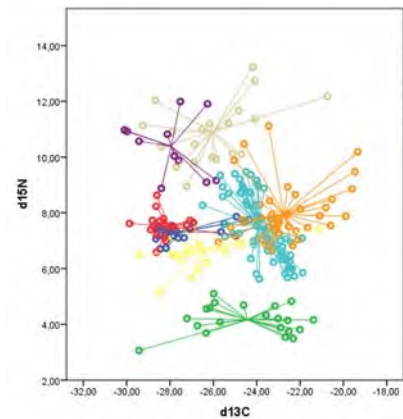
Prey	<i>L. gibbosus</i>	<i>M. salmoides</i>	<i>C. carpio</i>
<i>P. clarkii</i>	5.2	78.7	0.0
<i>L. gibbosus</i>	0.0	5.6	0.0
Odonata	0.0	4.0	0.0
Mollusks	1.5	0.0	0.0
Hymenoptera	10.3	0.5	0.0
Heteroptera	4.5	0.0	0.0
Snails	0.0	0.0	0.0
Araneae	0.5	9.7	0.0
Diptera	1.3	0.0	0.0
Odonata	2.1	0.0	0.0
Orthoptera	5.2	0.0	0.0
Coleoptera	49.7	0.0	0.0
Formicidae	0.5	0.0	0.0
unid. Insects	44.1	0.0	0.0
detritus	32.6	0.5	45.5
fisheggs	3.7	0.0	0.0
plant material	1.6	0.0	0.0



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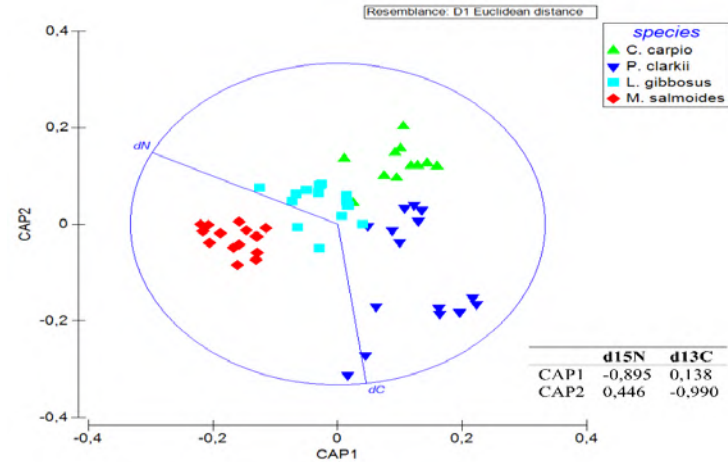
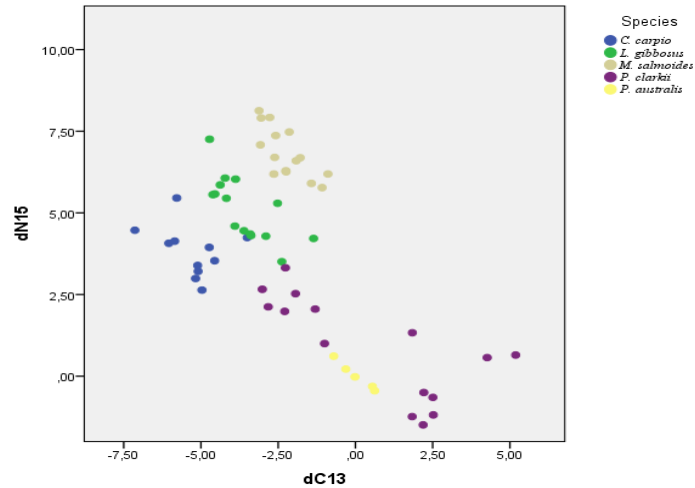


Laboratory procedure



SENCKENBERG

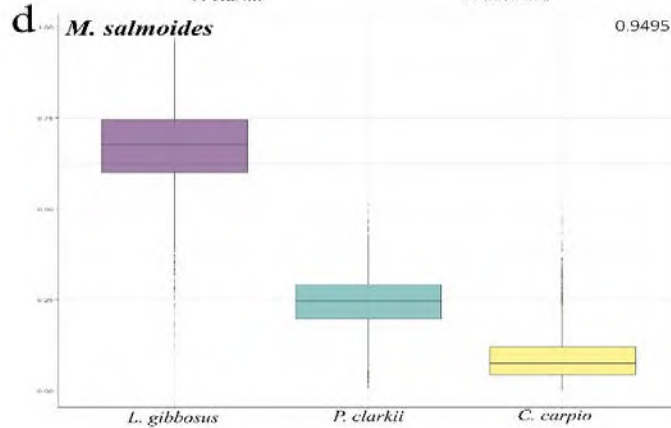
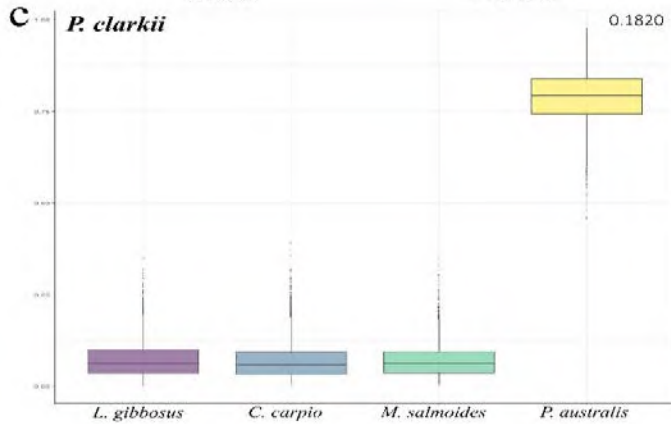
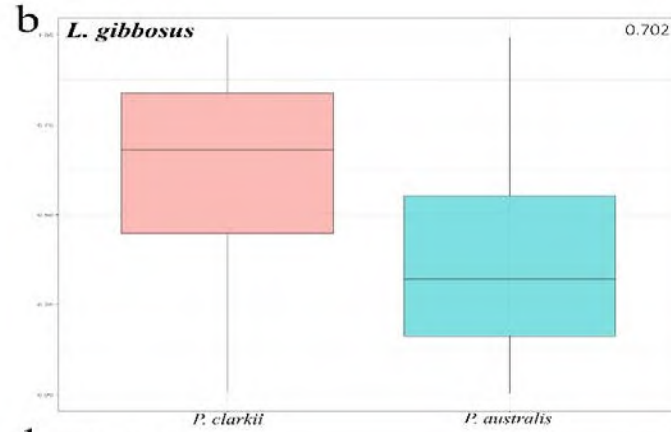
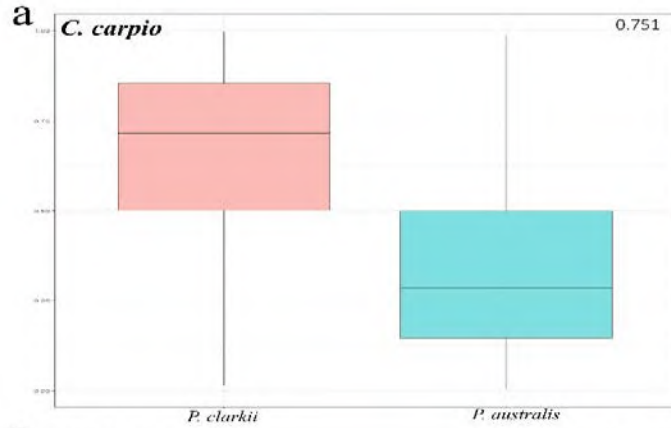
Stable Isotope Analysis



Group	Layman metrics and Stable Isotope Analysis results								
	Mean $\delta^{15}\text{N}$	Mean $\delta^{13}\text{C}$	NR	CR	TA	CD	MNND	SDNND	SEAc
<i>Cyprinus carpio</i>	12.2	-33.9	2.8	3.8	5.4	1.0	0.6	0.5	2.5
<i>Lepomis gibbosus</i>	13.5	-32.2	3.8	3.4	4.9	1.2	0.5	0.4	2.2
<i>Micropterus salmoides</i>	15.2	-30.8	2.2	2.3	2.6	0.9	0.3	0.2	2.3
<i>Procambarus clarkii</i>	9.4	-28.2	4.8	9.6	16.3	3.2	0.8	0.7	10.2
<i>Phragmites australis</i>	8.4	-28.6	1.0	1.4	0.1	0.6	0.3	0.2	0.1
Community	na	na	8.6	12.3	53.4	3.0	0.4	0.4	17.1

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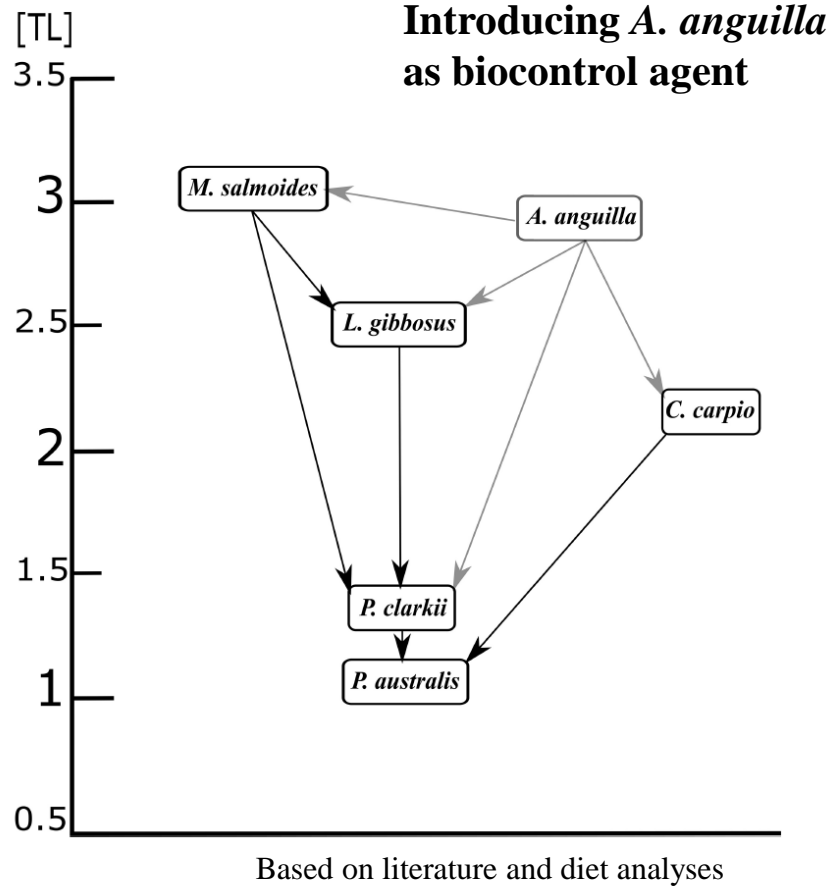
Stable Isotope Analysis Mixing Models



Idea: introduce 1.000 – 2.000 grown individuals



Anguilla anguilla



**Introducing *A. anguilla*
as biocontrol agent**

1. Identifying suitable isotope data

Limited availability of data

Dörner et al., (2009) *Journal of Fish Biology* 74

Similar community:

- *Perca fluviatilis*
- *Rutilus rutilus*
- *Tinca tinca*
- *Orconectes limosus*
- *Phragmites australis*

2. Extract species of interest and potential baseline organism

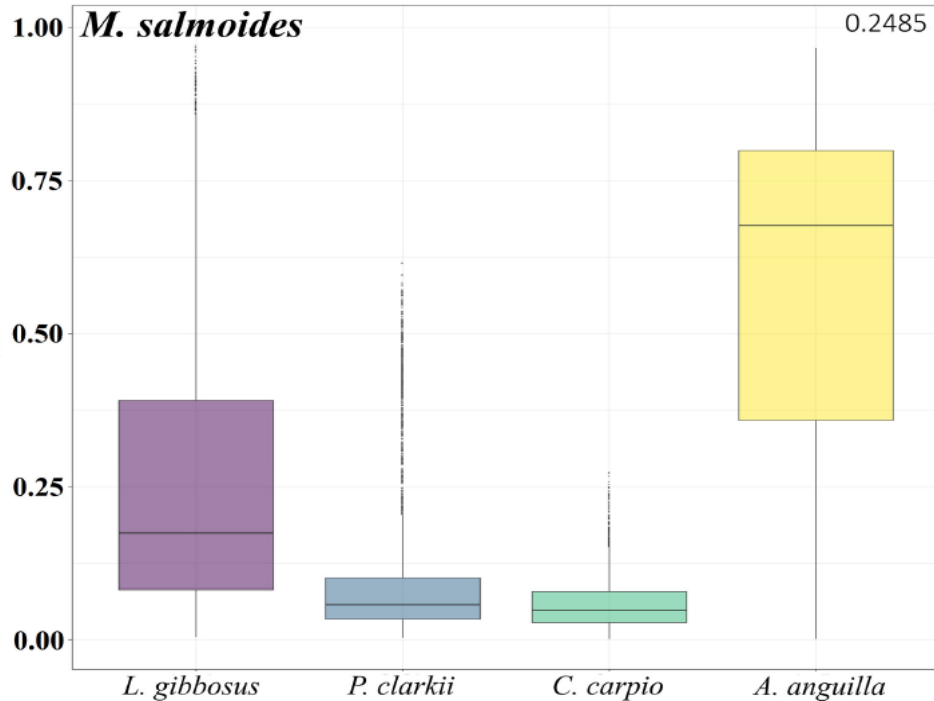
Anguilla anguilla & *Phragmites australis*

3. Standardize species of interest and community by the common baseline organism

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Assumption:

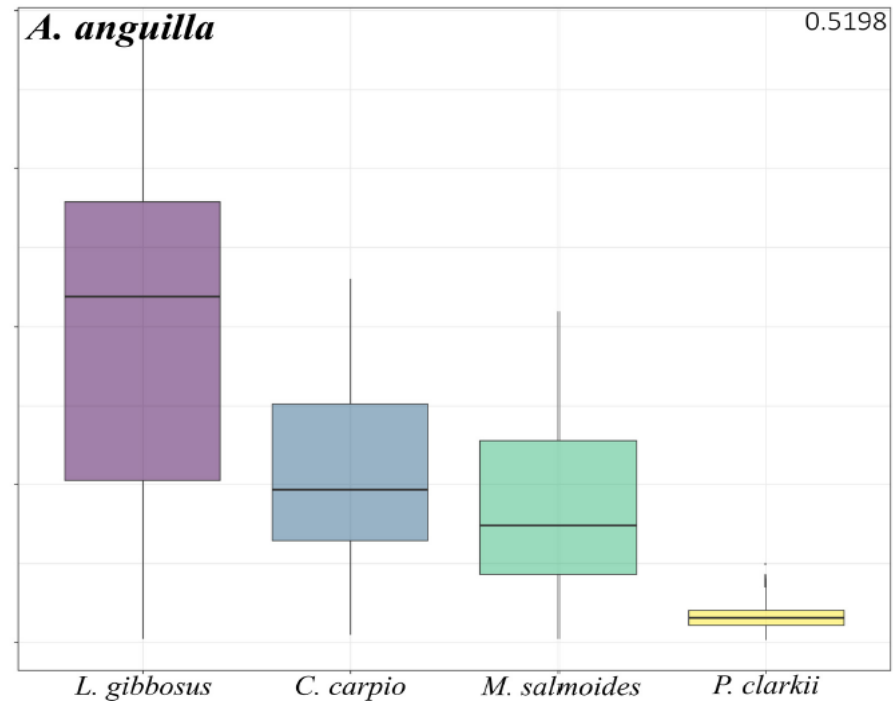
1. potential prey abundances reflecting abundances from last monitoring
2. introduction of 1000 individuals of *A. anguilla*



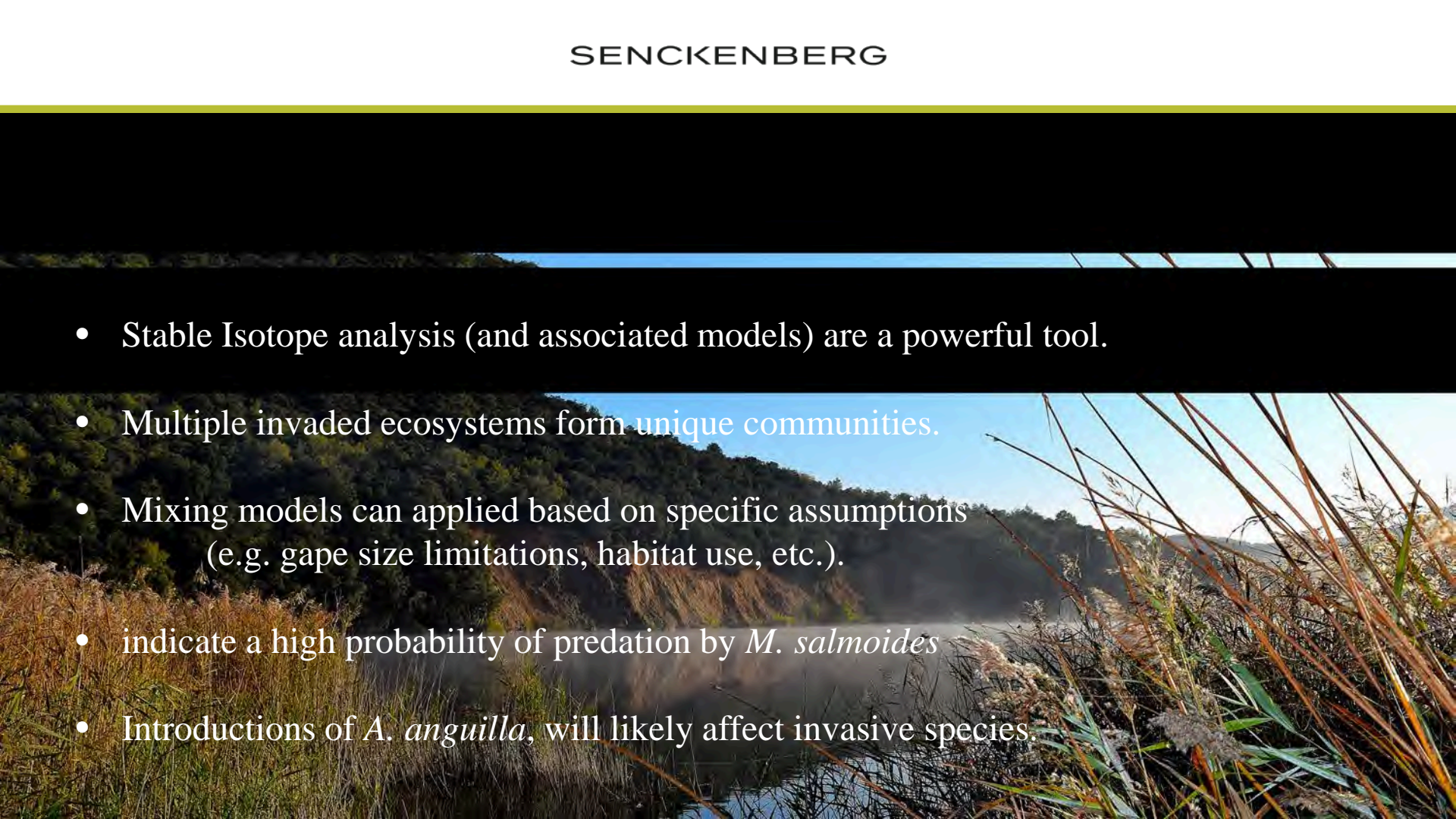
Limited effect due to predation by *M. salmoides*

Assumption:

1. potential prey abundances reflecting abundances from last monitoring
2. *A. anguilla* preying on only juv. fish



Highly variable due to prey availability?

- 
- Stable Isotope analysis (and associated models) are a powerful tool.
 - Multiple invaded ecosystems form unique communities.
 - Mixing models can applied based on specific assumptions (e.g. gape size limitations, habitat use, etc.).
 - indicate a high probability of predation by *M. salmoides*
 - Introductions of *A. anguilla*, will likely affect invasive species.



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**Thank you
for your
Attention**



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