# $bigmouth buffalo Riv No Sal\ model\ summary$

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#### Overview

This document summarizes the results of a global sensitivity and uncertainty analysis for the **bigmouth-buffaloRivNoSal** habitat suitability index (HSI) model for *Ictiobus cyprinellus*. Metadata for the model is stored in the ecorest package in R.

The original documentation for this model can be found here<sup>1</sup>.

Sub-model: Riverine salinity SIV8 parameter not considered a potential problem in the study area

The bigmouthbuffaloRivNoSal model is comprised of 10 variables and 4 components.

#### Variables:

**Table 1.** SIV variables included in the bigmouthbuffaloRivNoSal model. Type indicates whether a variable is numeric or categorical and breakpoints indicates the number of distinct breakpoints in suitability graphs.

	Variable name	Type	Breakpoints
SIV1	pls.backwtrs.marsh.spring.summer.SIV	numeric	4
SIV2	avg.max.mthly.turb.summer.flow.SIV	numeric	12
SIV3	pH.level.SIV	numeric	5
SIV4	max.wtr.temp.summer.ad.SIV	numeric	14
SIV5	${\it avg.} max. wtr. temp. nursery. spring. summer. emb. SIV$	numeric	7
SIV6	min.DO.spring.summer.emb.ad.SIV	numeric	3
SIV7	avg.curr.vel.SIV	numeric	6
SIV9	dominant.subs.spwn.area.emb.SIV	categorical	4
SIV11	wtr.lvl.fluct.before.after.spwn.SIV	categorical	3
SIV13	${\it veg. cov. pls. backwtrs. marsh. proct. embayments. shoreline. SIV}$	numeric	4

 $<sup>^{1}</sup> https://ecolibrary.sec.usace.army.mil/resource/225f8c89-071b-438a-f0f8-b09bc5b8ffa2$ 

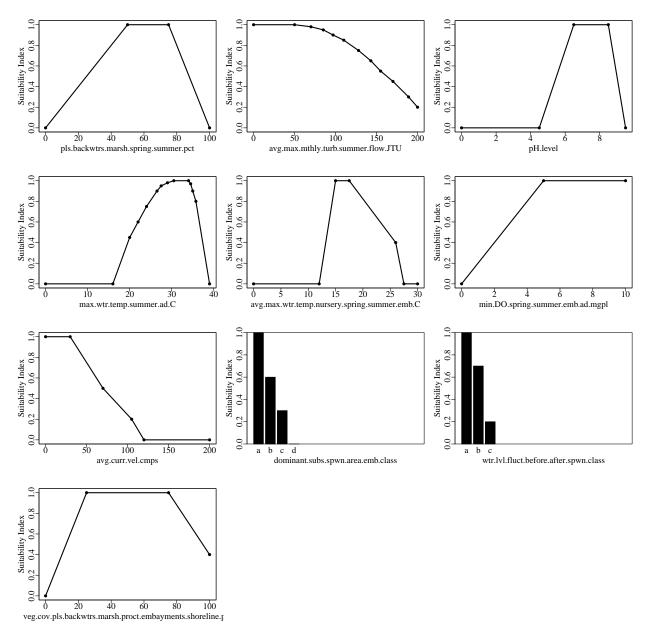


Figure 1. Suitability index graphs for variables included in the bigmouthbuffaloRivNoSal model in ecorest.

#### Components:

Table 2. Components included in the bigmouthbuffaloRivNoSal model in ecorest.

	Component	Equation
CCF	Food cover component	$(SIV1*SIV13)^(1/2)$
CWQ	Water quality component	$(SIV2*SIV3*(SIV4)^(2)*SIV6)^(1/5)$
CR	Reproduction component	ifelse(SIV5<=0.4 SIV9<=0.4,min(SIV5,SIV9,((SIV5^2)*SIV6*(SIV 9^2)*SIV11)^(1/6)),((SIV5^2)*SIV6*(SIV9^2)*SIV11)^(1/6))
COT	Other component	SIV7

# Model equation:

The equation to calculate an overall HSI index for the bigmouth buffaloRivNoSal model is: ifelse (CR<=0.4,min(CR,(CCF\*CWQ\*(CR^2)\*COT)^(1/5)),(CCF\*CWQ\*(CR^2)\*COT)^(1/5))

According to our classification, this model's format is: author-specified

#### Global sensitivity and uncertainty analysis:

We ran global sensitivity and uncertainty analyses on the bigmouthbuffaloRivNoSal model using the sensobol package in R (Puy et al. 2022). The following parameters were used for the sensobol analysis:

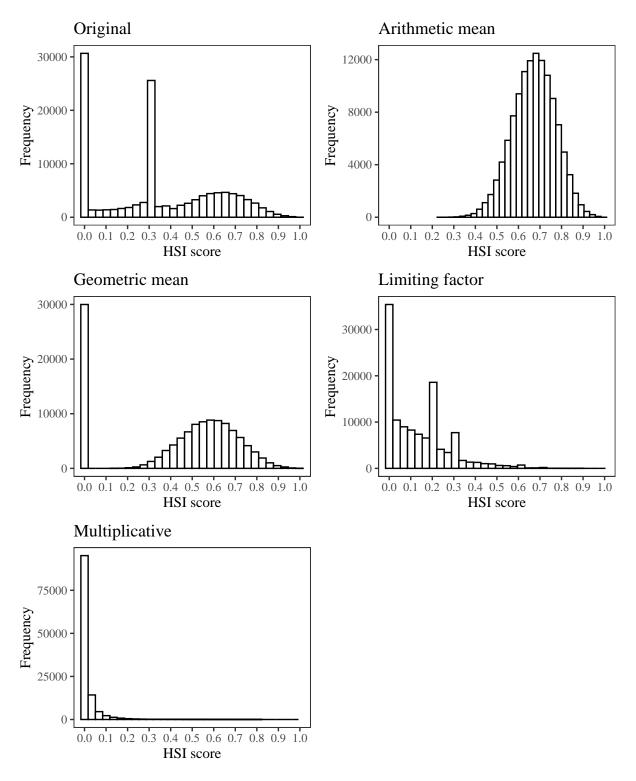
Table 3. Parameters and settings used for sensobol sensitivity and uncertainty analyses.

Parameter	Equation	Value
Number of input variables (M) Base sample size (n) Number of model evaluations (N) First order estimator Total order estimator	- - n*(M+2) See Puy et al. (2022) See Puy et al. (2022)	10 10000 120000 Saltelli Jansen
Number of bootstrap replications Sampling scheme Matrices	- - -	1000 Quasi-random A, B, AB

We ran a sensitivity and uncertainty analysis for the bigmouthbuffaloRivNoSal model using the original equation outlined in the documentation from Edwards (1983) and using arithmetic mean, geometric mean, limiting factor, and multiplicative equations to contrast the results across different equation structures.

#### Model uncertainty

We ran the bigmouthbuffaloRivNoSal model using 120000 combinations of its SIV variables, which were sampled from a uniform distribution spanning the range of possible values listed in the bigmouthbuffaloRivNoSal documentation. We limited the range of possible values for each parameter to the range in which the SIV values were greater than zero to prevent HSI score distributions with primarily zero values.



**Figure 2.** Empirical distributions of HSI scores for the bigmouthbuffaloRivNoSal model using the original author-specified model equation from Edwards (1983), and an arithmetic mean, geometric mean, limiting factor, and multiplicative structure incorporating all SIV variables. Note differences in the y axis.

We assumed a uniform distribution for all parameters because we evaluated all ecorest models in batch. Should you decide to run your own sensitivity analysis, this assumption should be evaluated independently for each parameter in the model.

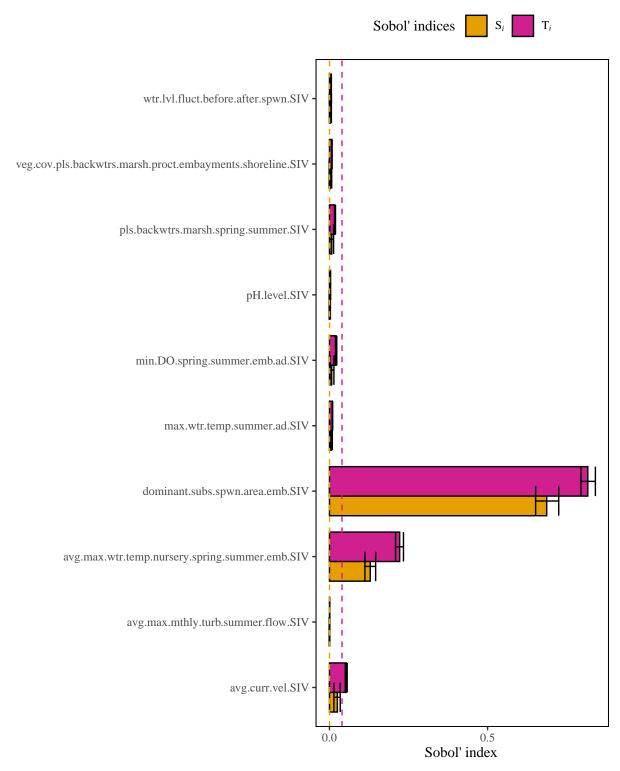
**Table 4.** Quantiles from the empirical distribution of HSI scores for the original bigmouthbuffaloRivNoSal model structure, an arithmetic mean equation, a geometric mean equation, a limiting factor equation, and a multiplicative equation structure.

	1%	2.5%	5%	25%	50%	75%	95%	97.5%	99%	100%
Original	0.00	0.00	0.0	0.00	0.30	0.58	0.78	0.82	0.87	1.00
Arithmetic	0.43	0.47	0.5	0.61	0.67	0.74	0.83	0.86	0.89	1.00
Geometric	0.00	0.00	0.0	0.16	0.52	0.64	0.78	0.82	0.86	1.00
Limiting	0.00	0.00	0.0	0.00	0.11	0.20	0.39	0.49	0.59	0.99
Multiplicative	0.00	0.00	0.0	0.00	0.00	0.01	0.08	0.14	0.23	0.98

The empirical distribution of the original bigmouthbuffaloRivNoSal model has a coefficient of variation (CV) of **0.796**, while the arithmetic mean model has a CV of **0.148**, the geometric mean model has a CV of **0.634**, the limiting factor model has a CV of **1.018**, and the multiplicative model has a CV of **2.763**. Hence, the **Multiplicative** model is the most uncertain, while the **Arithmetic mean** model is the least uncertain.

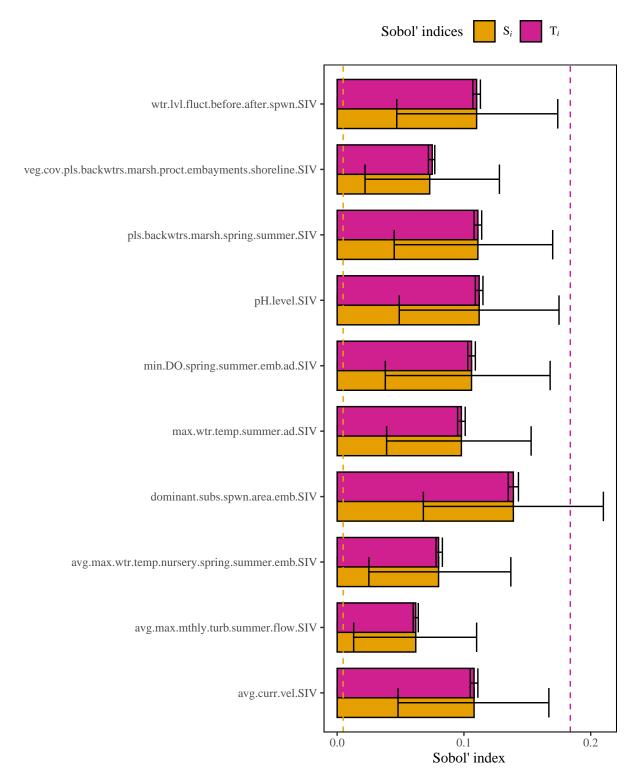
# Model sensitivity

Below are the results of the global sensitivity analysis for the bigmouthbuffaloRivNoSal model using the original equation, an arithmetic mean, a geometric mean, a limiting factor, and a multiplicative model structure. The sensobol package uses variance-based sensitivity metrics, so the model's sensitivity to a given parameter is a measure of how much variance in the HSI score decreases in response to that parameter being fixed (Puy et al. 2022). For each parameter, the observed changes in the variance of the HSI score can be described with a first order sensitivity index ( $S_i$ ) that accounts for the influence of a single parameter of interest on variance in HSI, or with a total order index ( $T_i$ ) that accounts for the influence of a single parameter on its own and in combination with all other parameters (*i.e.*, interactions) (Puy et al. 2022). We can compare the 95% confidence intervals for the first and total order indices to a dummy parameter, which represents a parameter that has no influence on the variance in a model's output. While an uninfluential variable should theoretically have an  $S_i$  and  $T_i$  of zero, small approximation errors can lead variables to have a non-zero influence on a model's output (Puy et al. 2022). If the confidence interval of the  $S_i$  and  $T_i$  index for a given parameter overlaps the confidence interval of the dummy parameter, we can deduce that the parameter has a negligible effect on variance in HSI scores, both on its own and in combination with all other variables.



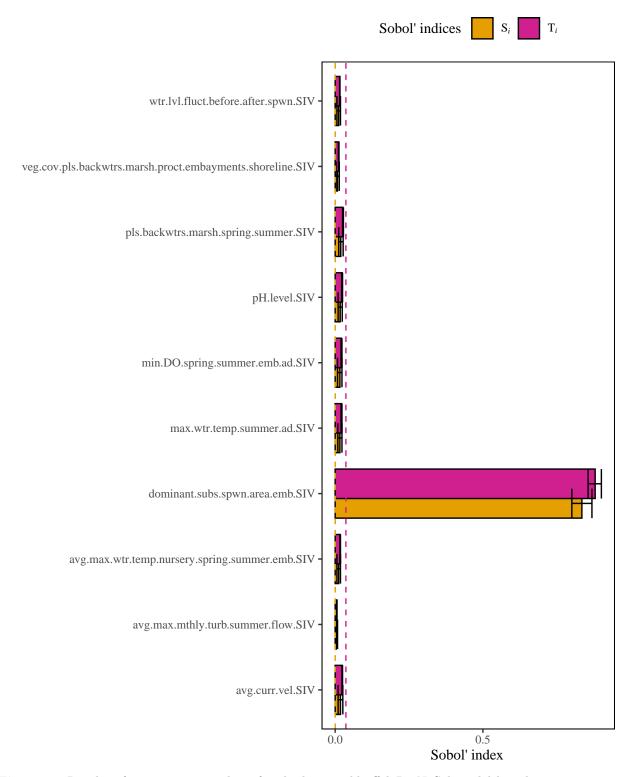
**Figure 3.** Results of a sensitivity analysis for the bigmouthbuffaloRivNoSal model based on the original author-specified model outlined in Edwards (1983). Dashed lines represent baseline numerical approximation error for  $S_i$  and  $T_i$  (*i.e.*, dummy variables).

## Arithmetic mean model



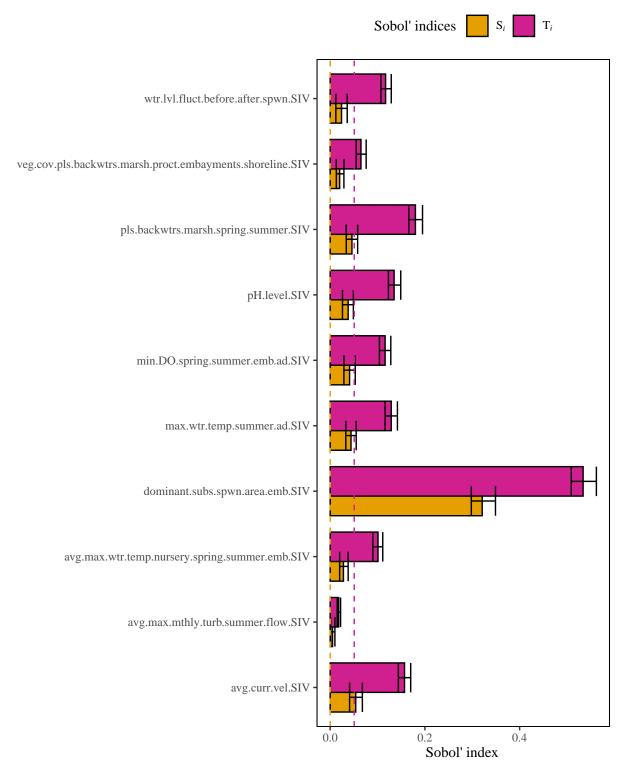
**Figure 4.** Results of a sensitivity analysis for the bigmouthbuffaloRivNoSal model based on an arithmetic mean structure. Dashed lines represent baseline numerical approximation error for  $S_i$  and  $T_i$  (*i.e.*, dummy variables).

## Geometric mean model



**Figure 5.** Results of a sensitivity analysis for the bigmouth buffaloRivNoSal model based on a geometric mean structure. Dashed lines represent baseline numerical approximation error for  $S_i$  and  $T_i$  (*i.e.*, dummy variables).

# Limiting factor model



**Figure 6.** Results of a sensitivity analysis for the bigmouthbuffaloRivNoSal model based on a limiting factor structure. Dashed lines represent baseline numerical approximation error for  $S_i$  and  $T_i$  (*i.e.*, dummy variables).

# Multiplicative model

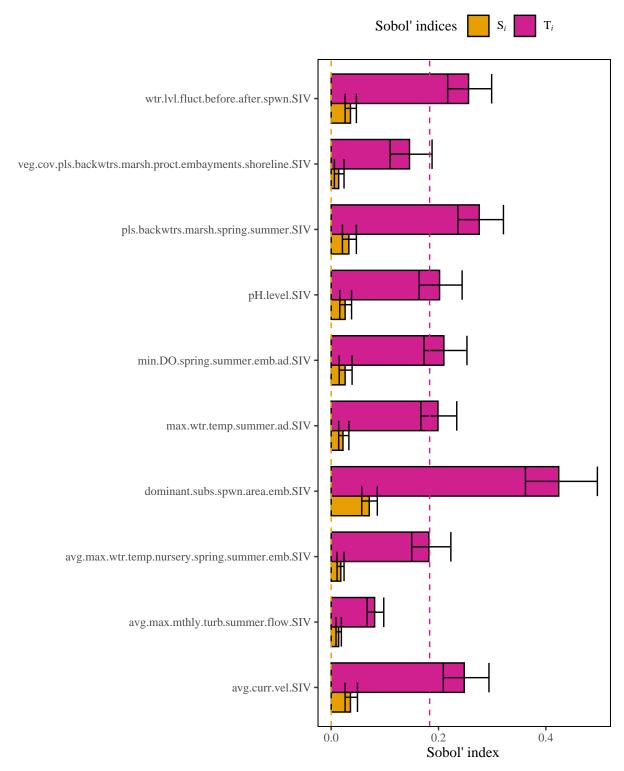


Figure 7. Results of a sensitivity analysis for the bigmouth buffaloRivNoSal model based on a multiplicative mean structure. Dashed lines represent baseline numerical approximation error for  $S_i$  and  $T_i$  (*i.e.*, dummy variables).

#### Summary of influential variables

Original model In the original bigmouthbuffaloRivNoSal model, 6 of 10 variables are influential and dominant.subs.spwn.area.emb.SIV has the highest first order sensitivity. In addition, dominant.subs.spwn.area.emb.SIV has the highest total order sensitivity.

# Un-influential variables in original model:

```
avg.max.mthly.turb.summer.flow.SIV
pH.level.SIV
wtr.lvl.fluct.before.after.spwn.SIV
veg.cov.pls.backwtrs.marsh.proct.embayments.shoreline.SIV
```

Arithmetic mean model In the arithmetic mean bigmouthbuffaloRivNoSal model, 10 of 10 variables are influential and dominant.subs.spwn.area.emb.SIV has the highest first order sensitivity. In addition, dominant.subs.spwn.area.emb.SIV has the highest total order sensitivity.

Un-influential variables in arithmetic mean model:

None

Geometric mean model In the geometric mean bigmouthbuffaloRivNoSal model, 10 of 10 variables are influential and dominant.subs.spwn.area.emb.SIV has the highest first order sensitivity. In addition, dominant.subs.spwn.area.emb.SIV has the highest total order sensitivity.

Un-influential variables in geometric mean model:

None

Limiting factor model In the limiting factor bigmouthbuffaloRivNoSal model, 10 of 10 variables are influential and dominant.subs.spwn.area.emb.SIV has the highest first order sensitivity. In addition, dominant.subs.spwn.area.emb.SIV has the highest total order sensitivity.

Un-influential variables in limiting factor mean model:

None

Multiplicative model In the multiplicative mean bigmouthbuffaloRivNoSal model, 10 of 10 variables are influential and dominant.subs.spwn.area.emb.SIV has the highest first order sensitivity. In addition, dominant.subs.spwn.area.emb.SIV has the highest total order sensitivity.

Un-influential variables in multiplicative model:

None

#### References

- 1. Edwards, EA. 1983. Habitat suitability index models: Bigmouth buffalo. U.S. Dept. Int., Fish Wildl. Serv. FWS/OBS-82/10/34. 23 pp.
- 2. McKay S, D Hernandez-Abrams, and K Cushway. 2024. ecorest: conducts analyses informing ecosystem restoration decisions. R package version 2.0.0, https://CRAN.R-project.org/package=ecorest.
- 3. Puy, A, S Lo Piano, A Saltelli, and SA Levin. 2022. sensobol: an R package to compute variance based sensitivity indices. Journal of Statistical Software 102(5):1-37. doi: 10.18637/jss.v102.i05