

# Assessing sensitivity of intervention effects

MATHEMATICAL MODELING FOR INFECTIOUS DISEASE PLANNING IN  
AFRICA

60 minutes

# Questions: sensitivity analysis

**Question 1:** How sensitive are the intervention effects to changes in other parameter values?

**Question 2.** What parameters are most influential on the model outcomes?

# Learning objectives

At the end of this lecture, you would:

1. Be able to **assess** the sensitivity of intervention effects to changes in one or more key parameters
2. Be able to **interpret** results on the sensitivity of intervention effects

# Outline

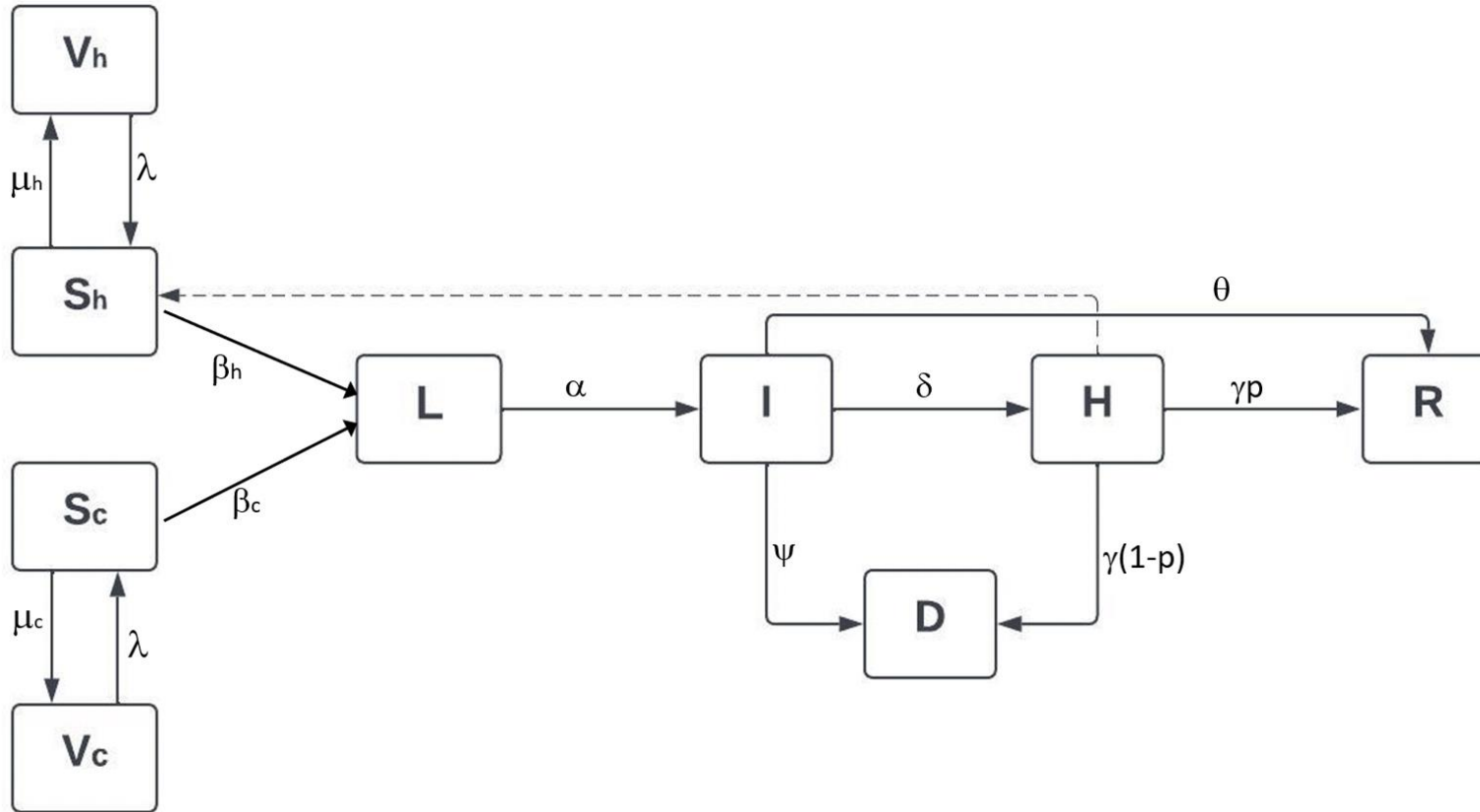
1. **Question 1:** How sensitive are the intervention effects to changes in other parameter values?

2. Demonstration with Berkeley Madonna

Slides on Question 2 are also provided (from slide 22) but will not be treated in this lecture.

**Question 1:** How sensitive are the intervention effects to changes in other parameter values?

# Intervention: Surveillance



Compartmental diagram for an Ebola transmission model

Relevant parameter:  $\delta$ ,  
rate at which infectious  
individuals are detected  
(and hospitalized)

Credit: Ebola Rwanda team

# Summary of surveillance effects

Total number of deaths by day 365	
$\delta = 1/4 \text{ day}^{-1}$	$\delta = 1/8 \text{ day}^{-1}$
1244	82159

$\delta$  = hospitalization rate for infectious individuals (per day)

**Conclusions based on results above:** The higher surveillance rate ( $\delta = 1/4$ ) leads to reduced mortality, compared to the lower surveillance rate ( $\delta = 1/8$ ).

Assume we are uncertain about the transmission rate for frontliners,  $\beta_h$ .

What factors might influence this uncertainty?

- Uncertainty about per-contact infectivity
- Uncertainty about contact rates

*How do we interpret our intervention effects (surveillance) considering this uncertainty?*



# What if we are uncertain about $\beta_h$ ?

- Will these conclusions hold if the transmission rate ( $\beta_h$ ) is changed?
- Are the intervention effect results *sensitive* to changes in  $\beta_h$ ?

$\beta_h$	Total number of deaths by day 365	
	$\delta = 1/4 \text{ day}^{-1}$	$\delta = 1/8 \text{ day}^{-1}$
Baseline value: 2.4	1244	82159

$\delta$  = hospitalization rate for infectious individuals (per day)

$\beta_h$  = transmission rate for frontliners

→ **Conclusion:** Higher surveillance is better than lower surveillance for deaths.

*How may we go about answering the above question?*

# Assessing the sensitivity of surveillance effects conclusions to changes in $\beta_h$

$\beta_h$	Total number of deaths by day 365	
	$\delta = 1/4 \text{ day}^{-1}$	$\delta = 1/8 \text{ day}^{-1}$
Baseline value: 2.4	1244	82159
0		
0.25		
...		
3.25		
3.5		

Conclusion: Higher surveillance is better than lower surveillance for deaths.

Does the conclusion remain unchanged across different values of  $\beta_h$ ?

# Assessing the sensitivity of surveillance effects conclusions to changes in $\beta_h$

$\beta_h$	Total number of deaths by day 365	
	$\delta = 1/4 \text{ day}^{-1}$	$\delta = 1/8 \text{ day}^{-1}$
Baseline value: 2.4	1244	82159
0	54	1879
0.25	72	2755
...	...	...
3.25	4637	184859
3.5	6908	213921

Conclusion: Higher surveillance is better than lower surveillance for deaths.

Does the conclusion remain unchanged across different values of  $\beta_h$ ?

$\delta$  = hospitalization rate for infectious individuals (per day)

$\beta_h$  = transmission rate for frontliners

# Assessing the sensitivity of surveillance effects conclusions to changes in $\beta_h$

$\beta_h$	Total number of deaths by day 365	
	$\delta = 1/4 \text{ day}^{-1}$	$\delta = 1/8 \text{ day}^{-1}$
Baseline value: 2.4	1244	82159
0	54	1879
0.25	72	2755
...	...	...
3.25	4637	184859
3.5	6908	213921

**Baseline**

Do we arrive at the same conclusions with the **summary** as we do with the **baseline**?

<b>1436</b>	<b>64886</b>
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**Summary** (Column averages, excluding baseline)

$\delta$  = hospitalization rate for infectious individuals (per day)

$\beta_h$  = transmission rate for frontliners

# Conclusions

$\beta_h$	Total number of deaths by day 365	
	$\delta = 1/4 \text{ day}^{-1}$	$\delta = 1/8 \text{ day}^{-1}$
Baseline value: 2.4	1244	82159
0	54	1879
0.25	72	2755
...	...	...
3.25	4637	184859
3.5	6908	213921
	<b>1436</b>	<b>64886</b>

Surveillance effects are not sensitive to changes in  $\beta_h$ .

$\delta$  = hospitalization rate for infectious individuals (per day)

$\beta_h$  = transmission rate for frontliners

# Assessing the sensitivity of surveillance effects conclusions to changes in $\beta_h$

## Main idea:

1. We will compare the interventions across different values of  $\beta_h$  and summarize the intervention results for each intervention.
2. The summary result ( $Su$ ) will be compared to the baseline result ( $Ba$ ).
3. If  $Su$  agrees with  $Ba$ , intervention effects are not sensitive to changes in  $\beta_h$ . However, if  $Su$  does not agree with  $Ba$ , intervention effects are sensitive to changes in  $\beta_h$ .

A note: The same procedure may be applied if interested in only a few values of  $\beta_h$

$\beta_h$	Total number of deaths by day 365	
	$\delta = 1/4 \text{ day}^{-1}$	$\delta = 1/8 \text{ day}^{-1}$
Baseline value: 2.4	1244	82159
3.5	6908	213921

Baseline

Do we arrive at the same conclusions with **value 1** as we do with the **baseline**?

Value 1

For example,

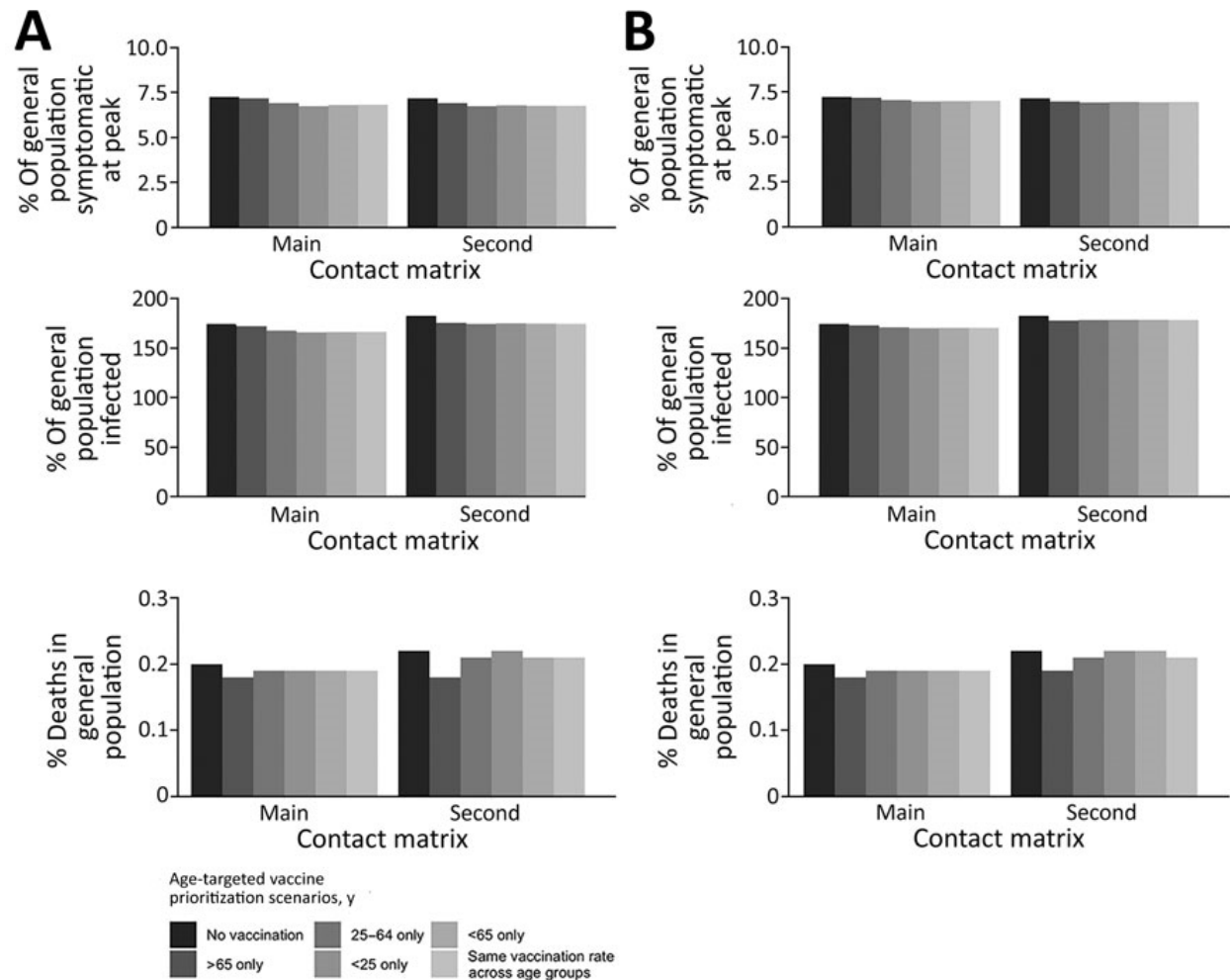


Figure 1. Effects of various vaccination scenarios on symptomatic infections at peak (upper panels), cumulative infections (middle panels), and deaths (lower panels) as a percentage of the general population, Ghana. The assessment used 2 different contact matrices in the main analysis and an effective reproductive number of 3.13 for the initial strain. A) Results assuming 1 million persons were vaccinated in 3 months. B) Results assuming 1 million persons were vaccinated in 6 months. Percentage of cumulative infections is  $\geq 100\%$  because of waning immunity from natural infection and vaccination.



A procedure for assessing the sensitivity of  
intervention effects to changes in key parameters

# Assessing the sensitivity of surveillance effects conclusions to changes in an uncertain parameter

## Main idea:

1. We will compare the interventions across different values of the uncertain parameter ( $p$ ) and summarize the intervention results for each intervention.
2. The summary result ( $Su$ ) will be compared to the baseline result ( $Ba$ ).
3. If  $Su$  agrees with  $Ba$ , intervention effects are not sensitive to changes in  $p$ . However, if  $Su$  does not agree with  $Ba$ , intervention effects are sensitive to changes in  $p$ .

# Assessing the sensitivity of surveillance effects conclusions to changes in $\beta_h$

## Steps:

1. Write down your research question for this analysis: How sensitive are the results on surveillance effects to changes in  $\beta_h$ ?
2. Decide on the model outcome for this analysis.
3. Write down the values of  $\beta_h$  to be assessed. (Refer to activity 1 where you defined bounds for key parameters)
4. Decide on the surveillance levels to be compared.

# Assessing the sensitivity of surveillance effects conclusions to changes in $\beta_h$

## Steps:

5. For each surveillance level, compute the model outcome for all values of  $\beta_h$ .

6. With the results from step 5, complete the table below.

<div>Uncertain parameter</div>	$\beta_h$	Total number of deaths by day 365		<div>Model outcome</div>
		$\delta = 1/4 \text{ day}^{-1}$	$\delta = 1/8 \text{ day}^{-1}$	
<div>Values of interest for uncertain parameter</div>	Baseline value: 2.4			<div>Intervention levels of interest</div>
	0			
	0.25			
	...			
	3.25			
	3.5			

# Assessing the sensitivity of surveillance effects conclusions to changes in $\beta_h$

## Steps:

7. Summarize the results from step 6 separately for each surveillance level: compute an average of the model outcomes for each surveillance level.
8. Compare the summary results ( $Su$ ) from step 7 to the baseline ( $Ba$ ). If  $Su$  agrees with  $Ba$ , we may conclude that surveillance effects are not sensitive to changes in  $\beta_h$ . If  $Su$  does not agree with  $Ba$ , we may conclude that surveillance effects are sensitive to changes in  $\beta_h$ .

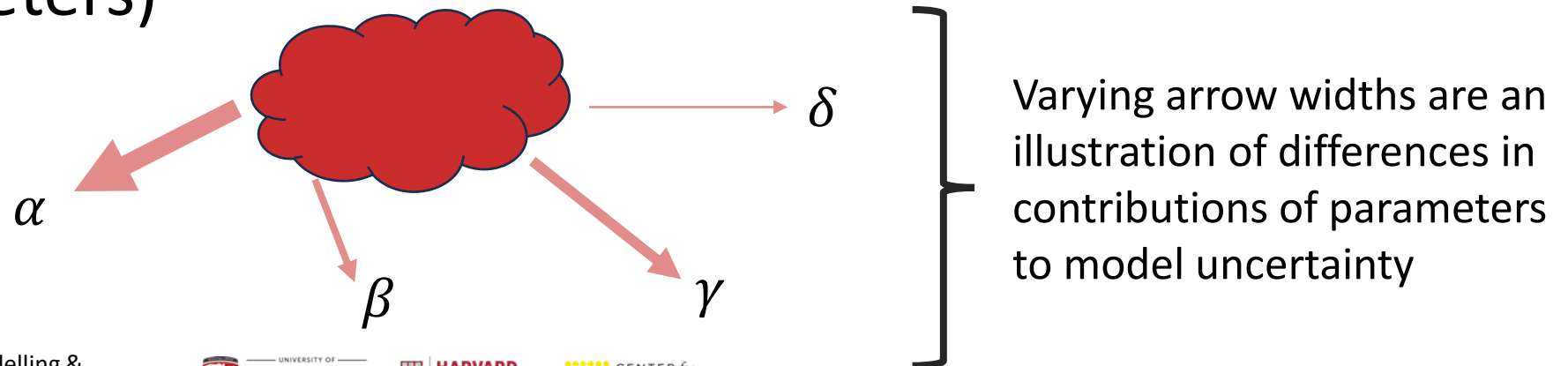
# Assessing the sensitivity of surveillance effects conclusions to changes in $\beta_h$

**A demonstration in Berkeley Madonna  
using the outlined steps**

Question 2. What parameters are most influential on the model outcomes?

# A closer look at Question 2

- Could be interpreted as: *Where is the uncertainty in the model outcome coming from?*
- Analysis on question 2 quantifies how variation in the model outputs can be apportioned to the various inputs (or parameters)



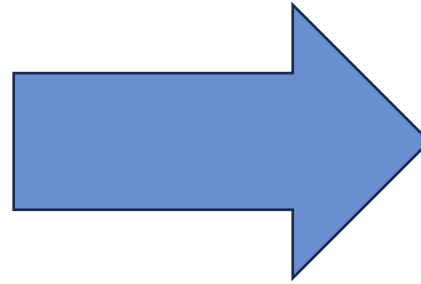
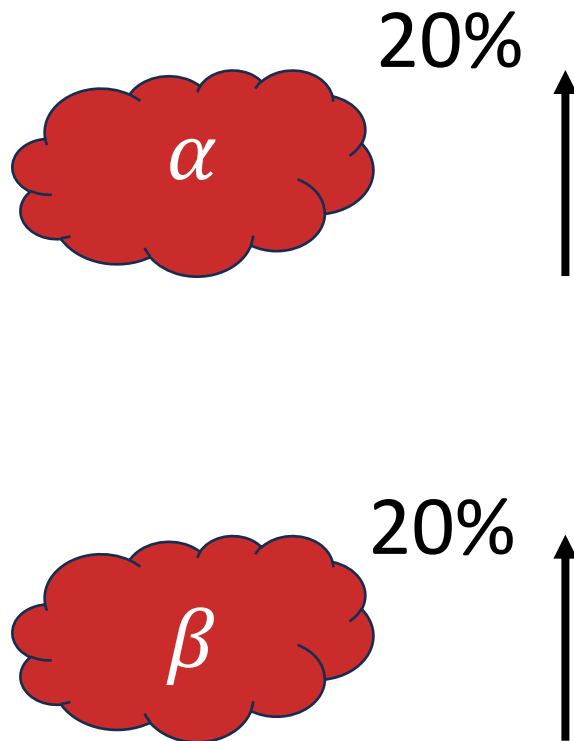


# Why sensitivity analyses?

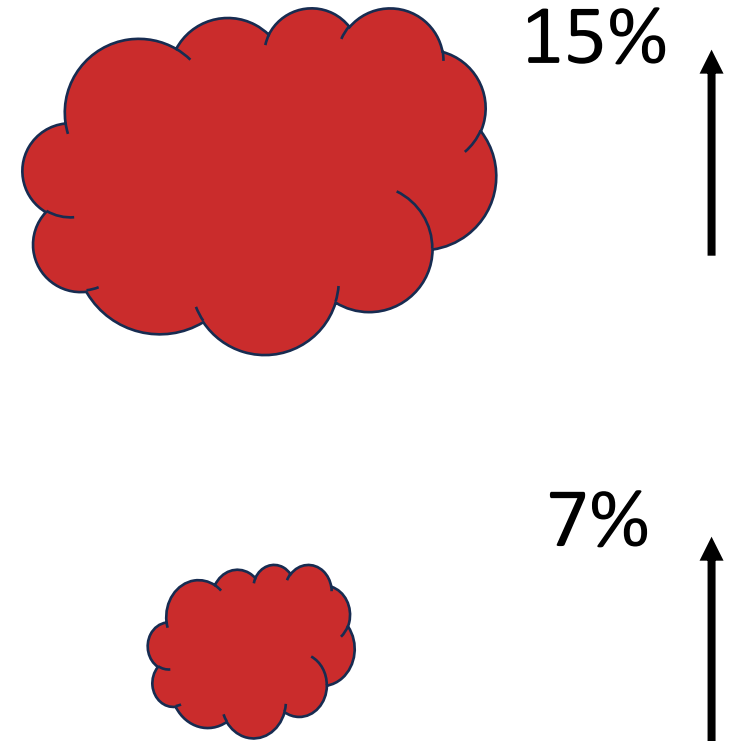
- To investigate how much variation in a parameter influences the variation in the model output
  - *How much of the uncertainty in model incidence is explained by the transmission rate parameter?*
  - Investigate parameter importance
  - If parameter is of biological/other significance, SA results allow us to make statements about the **connection between the biological/other factors and the transmission process/outcomes of interest**

# Why sensitivity analyses?

## Parameter uncertainty



## Model output uncertainty



# Sensitivity analyses vs uncertainty analyses

## Sensitivity analyses (SA)

Question: **Where is the uncertainty in the model inference coming from?**

Quantifies how variation in the model outputs can be apportioned to the various parameters

## Uncertainty analyses (UA)

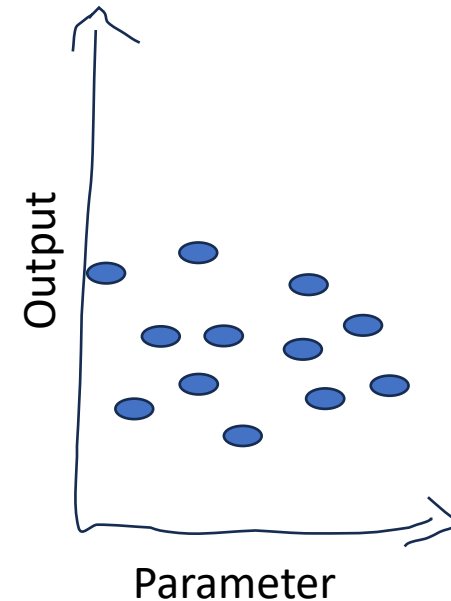
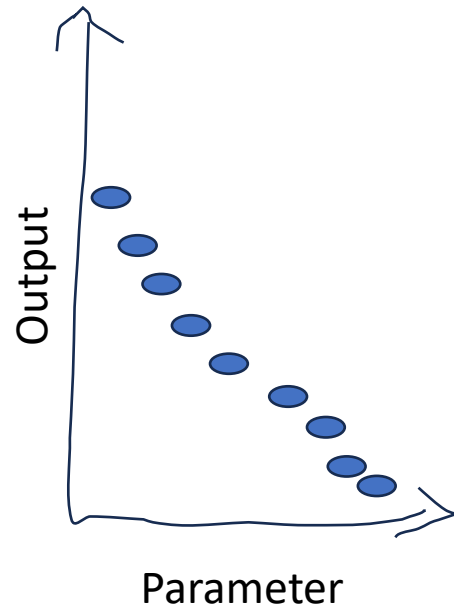
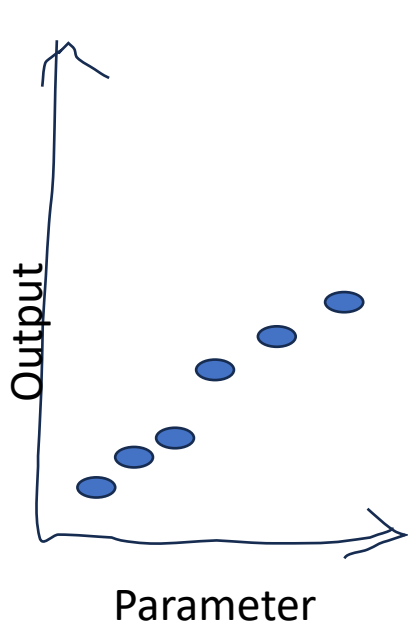
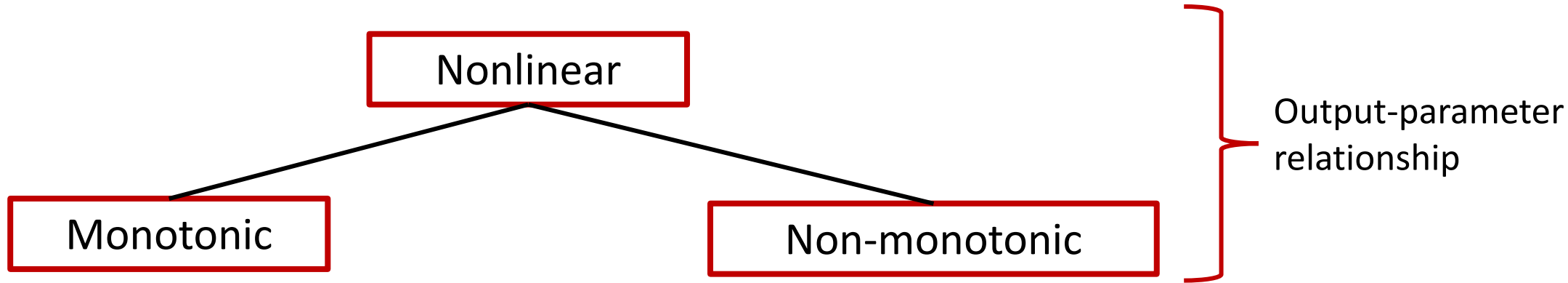
Question: **How uncertain is the model inference (or estimate)?**

Characterizes the confidence bounds for a model output

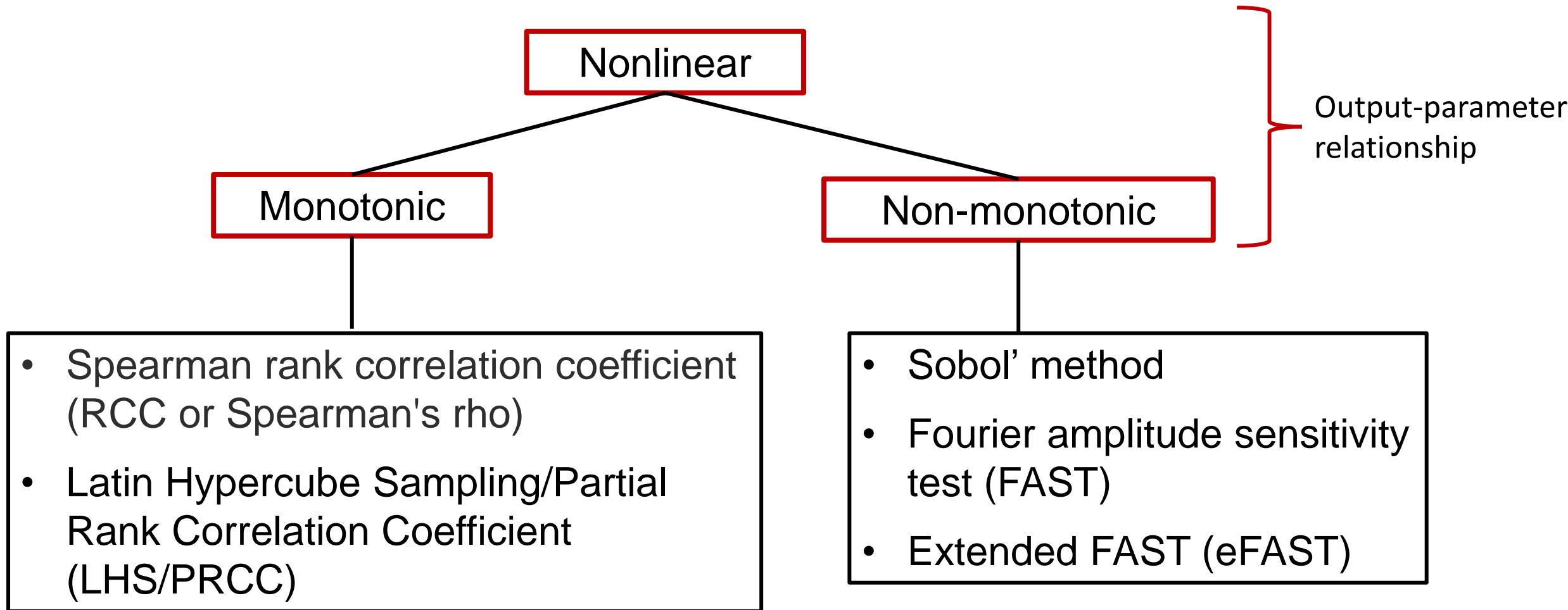
# In practice...

UA and SA are coupled and termed "sensitivity analysis", although they have different objectives.

# Examples of SA methods



# Examples of SA methods



# One-at-a-time SA (OAT-SA) may not be reliable

- OAT-SA involves changing the value of a parameter OAT while keeping the others constant
- Why OAT-SA will not work with the models we are studying:
  - Assumes model is *linear* – in many cases, epidemic models are not
  - Does not consider interaction effects between parameters (simultaneous change of parameters is needed for interactions to be detected)
  - Does not sufficiently explore the parameter space (this problem is worse in higher dimensions)
- See Saltelli and Annoni (2010) for proof

# Latin Hypercube Sampling/Partial Rank Correlation Coefficient (LHS/PRCC)

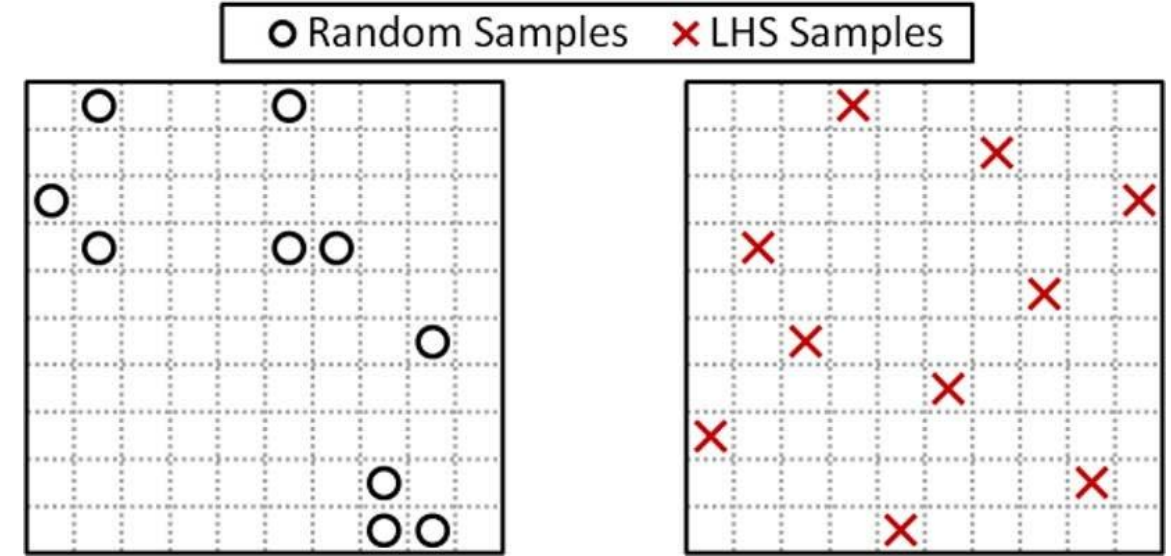
## General idea

- Efficiently explore parameter space (**LHS**)
- Identify and rank key parameters whose uncertainties contribute to model output uncertainty (**PRCC**)



# LHS

- Sampling scheme



1. Divide parameter range/space\* into equally probable intervals
  2. Sample n times without replacement from each interval
- Explores the parameter space more efficiently than simple random sampling

\*The figure assumes a two-dimensional parameter space.

# Partial Rank Correlation Coefficient

- **Correlation**
  - Strength of linear association between parameter and output
- **Partial correlation**
  - Correlation between a parameter and an output while discounting the linear effects of other parameters on the output
- **Partial rank correlation**
  - Partial correlation computed on *rank-transformed data* (why transform?)

Condition: Little to no correlation between parameters

# LHS meets PRCC

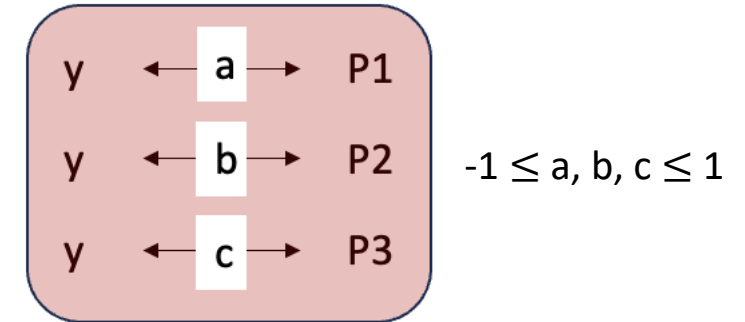
1. Set ranges for each parameter and get LHS samples.

Parameter		
P1	P2	P3
0.5	0.7	0.02
...	...	0.05
...	...	...

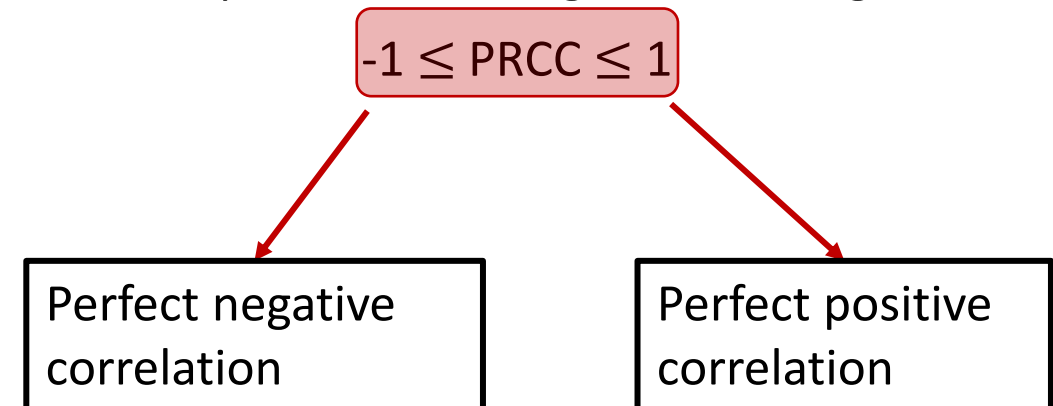
2. Compute model output (y) for each LHS sample and rank transform all results.

P1	P2	P3	y
0.5	0.7	0.02	1
...	...	0.05	3
...	...	...	...

3. Compute the partial correlation between output and each parameter.



4. Interpret, considering statistical significance:



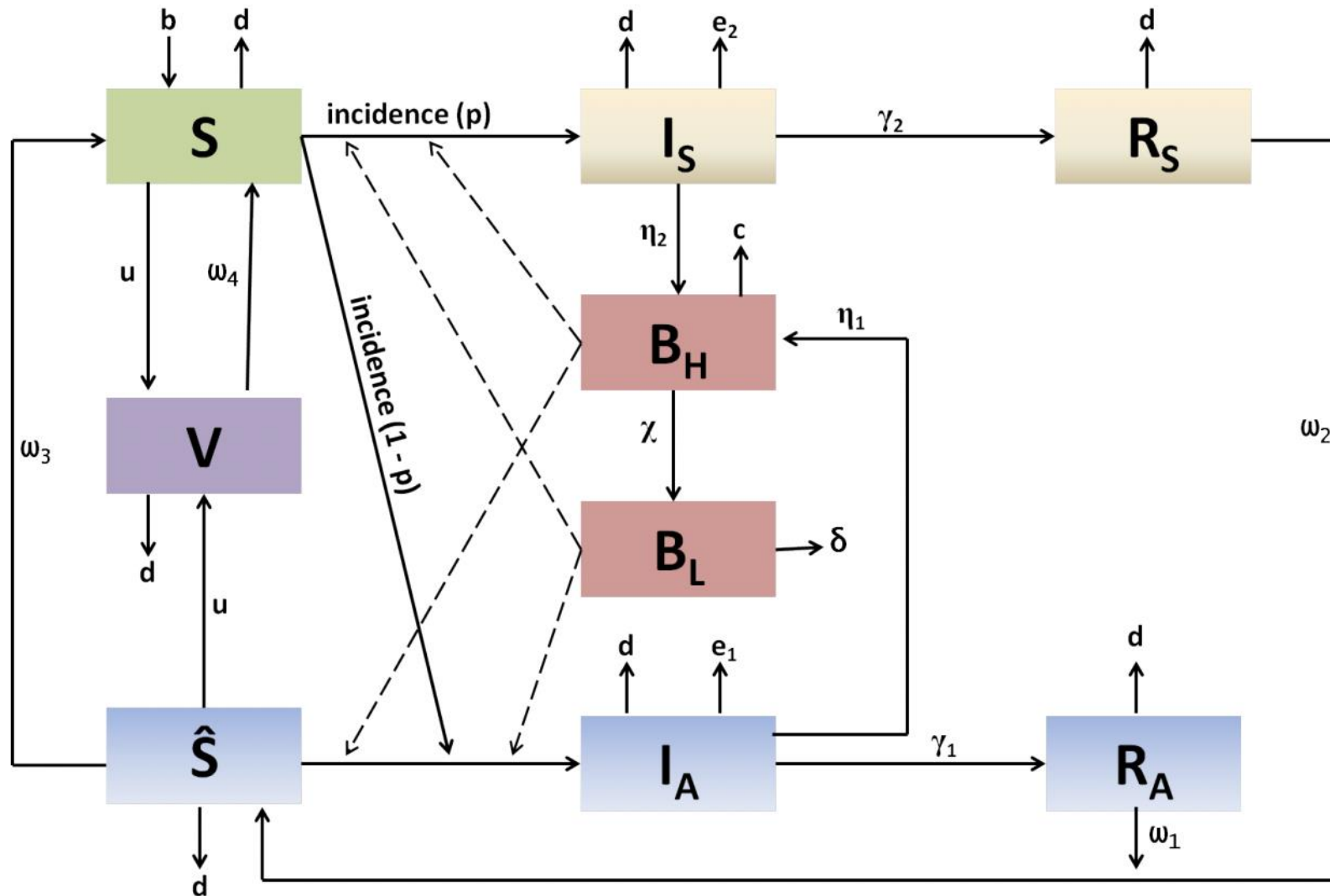
Activity: Discuss an application of LHS/PRCC to a cholera model

# Application of LHS/PRCC to a cholera model

## Outline

1. Model structure
2. Outcomes of interest
3. Testing monotonicity assumptions
4. LHS parameter ranges
5. LHS/PRCC results

# Cholera transmission model



# Parameter table

**Table 3.1:** Parameter List for Cholera Epidemic Model.

Symbol	Description	Value
$\hat{S}_0$	Initial # susceptible humans with partial immunity	3000
$S_0$	Initial # susceptible humans without partial immunity	$10,000 - \hat{S}_0$
$I_{A0}$	Initial # asymptomatic infecteds	0
$I_{S0}$	Initial # symptomatic infecteds	0
$R_{A0}$	Initial # recovered humans (asymptomatic)	0
$R_{S0}$	Initial # recovered humans (symptomatic)	0
$V_0$	Initial # humans with vaccinated immunity	0
$B_{H0}$	Initial concentration of highly infectious (HI) vibrios in environment	0
$B_{L0}$	Initial concentration of non-highly infectious (non-HI) vibrios in environment	$\kappa_L/2$
$p$	Probability of infecteds moving from symptomatic class to infected class without partial immunity	0.6
$r$	Scaling factor used to compute $\beta_H$ from $\beta_L$ .	0.1
$\beta_L$	Ingestion rate of non-HI vibrio from environment	$0.008 \text{ day}^{-1}$
$\beta_H$	Ingestion rate of HI vibrio from environment.	$r * \beta_L \text{ day}^{-1}$
$\kappa_L$	Half saturation constant of non-HI vibrios	103 cells/ml
$\kappa_H$	Half saturation constant of HI vibrios	$\kappa_L/700$ cells/ml



# Parameter table continued

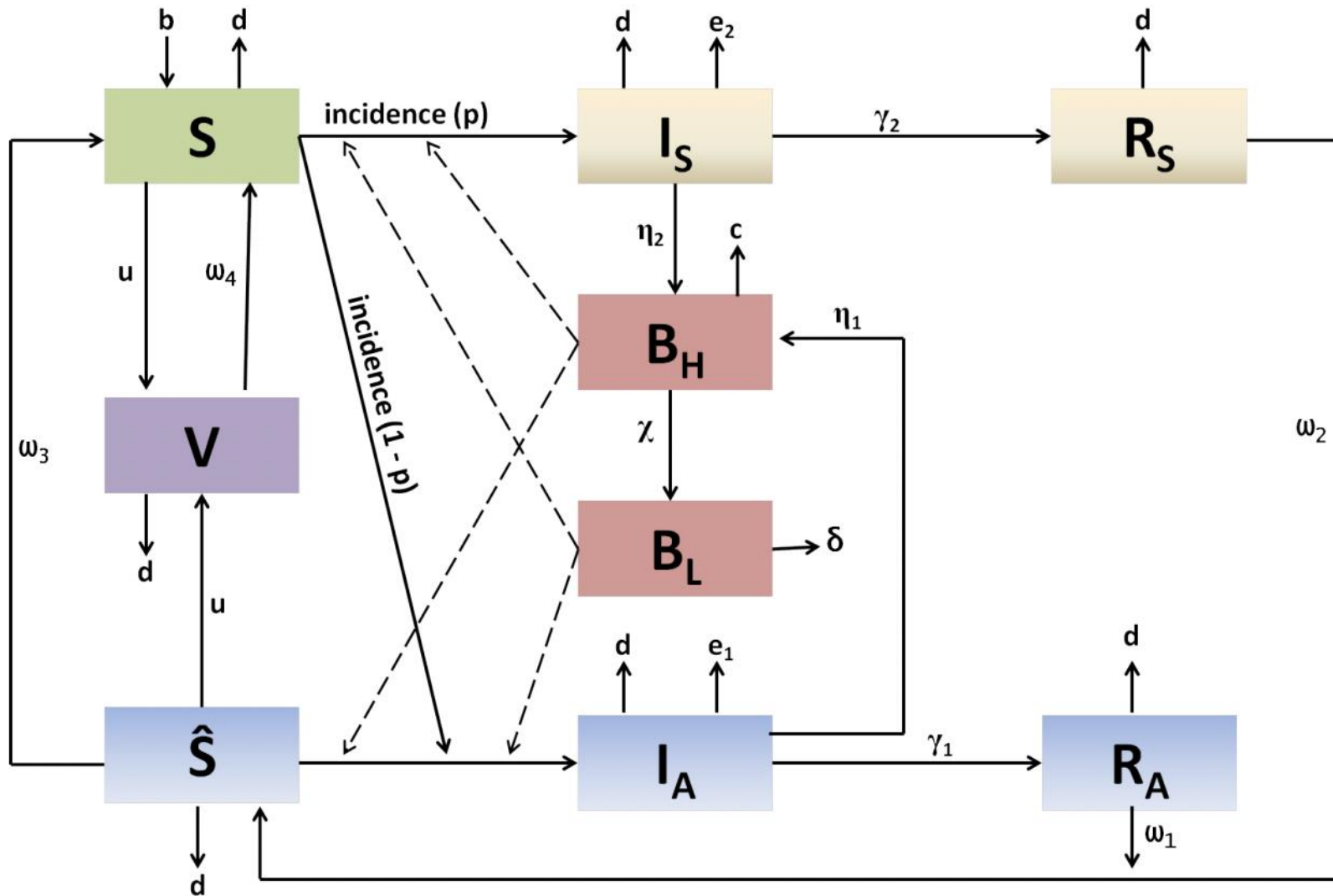
$e_1$	Cholera-related death rate for asymptomatic infecteds	$e_2/20 \text{ day}^{-1}$
$e_2$	Cholera-related death rate for symptomatic infected	$0.03 \text{ day}^{-1}$
$\gamma_1$	Cholera recovery rate (asymptomatic)	$0.75 \text{ day}^{-1}$
$\gamma_2$	Cholera recovery rate (symptomatic)	$0.1 \text{ day}^{-1}$
$\omega_1$	Rate of waning cholera immunity from asymptomatic infecteds to susceptibles with partial immunity	$1/180 \text{ day}^{-1}$
$\omega_2$	Rate of waning cholera immunity from symptomatic infecteds to susceptible humans with partial immunity	$1/(365 * 2) \text{ day}^{-1}$
$\omega_3$	Immunity waning rate: susceptibles without partial immunity $\rightarrow$ susceptibles with partial immunity	$1/(10 * 365) \text{ day}^{-1}$
$\omega_4$	Immunity waning rate: humans with vaccinated immunity $\rightarrow$ susceptibles without partial immunity	$0.001 \text{ day}^{-1}$
$s$	Scaling factor used to compute $\eta_2$ from $\eta_1$	100
$\eta_1$	Rate of contribution to HI vibrios in environment by asymptomatic infecteds	$0.008 \text{ cells/ml-day-human}$
$\eta_2$	Rate of contribution to HI vibrios in environment by symptomatic Infected.	$s * \eta_1 \text{ cells/ml-day-human}$
$\chi$	Transaction rate of vibrios from HI to non-HI state	$5 \text{ day}^{-1}$
$d$	Death rate of vibrios	$1/30 \text{ day}^{-1}$
$u$	Rate at which susceptible and asymptomatic infecteds are vaccinated daily	$0 \text{ day}^{-1}$
$b$	Natural birth rate of humans	$0.03/365 \text{ day}^{-1}$
$d$	Natural death rate of humans	$0.02/365 \text{ day}^{-1}$



# Outcomes of interest

1. Total number of infected individuals
2. Total number of symptomatic infected individuals

Make a guess: which parameters will be most influential on the outcomes of interest, and by how much?



### Outcomes of interest:

1. Total number of infected individuals
2. Total number of symptomatic infected individuals

### Comments on question above

- Difficult to tell by observation!
- PRCC allows us to get answers using a principled procedure.

# Testing monotonicity assumptions

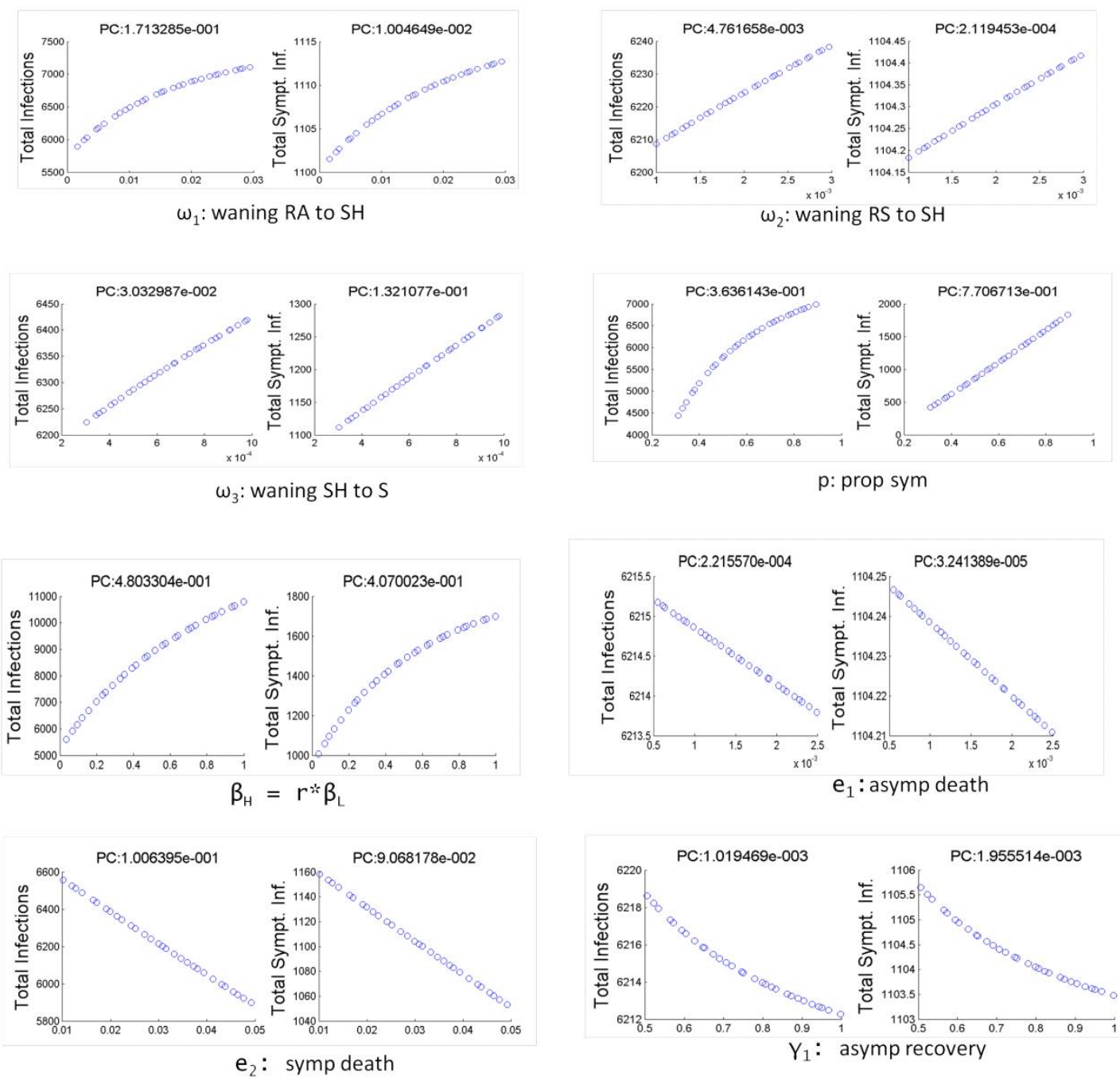
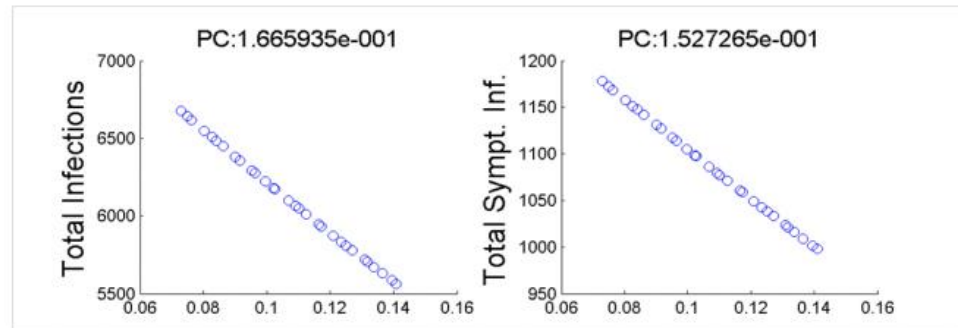
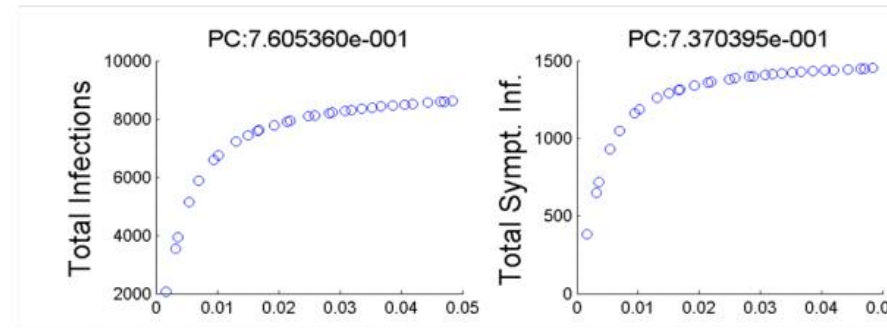


Figure 3.2: Monotonicity plots for  $\omega_1$ ,  $\omega_2$ ,  $\omega_3$ ,  $p$ ,  $\beta_H$ ,  $e_1$ ,  $e_2$ , and  $\gamma_1$

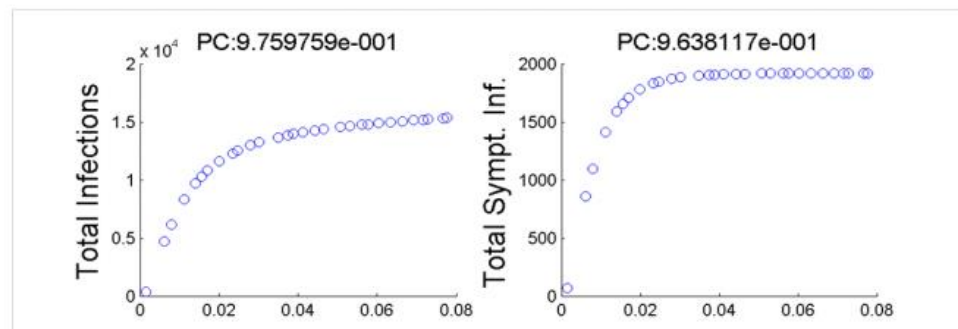
# Testing monotonicity assumptions (continued)



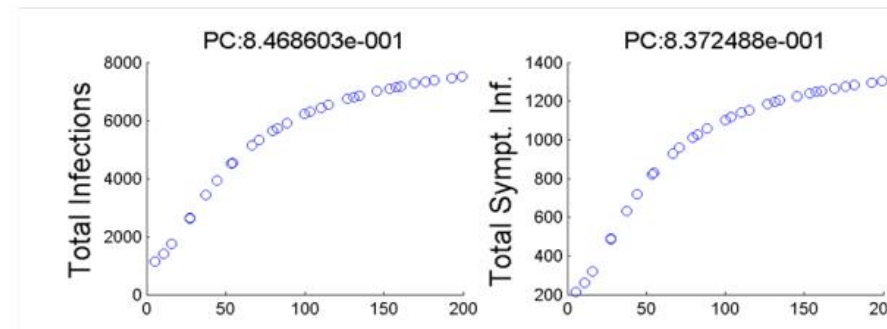
$\gamma_2$ : symp recovery



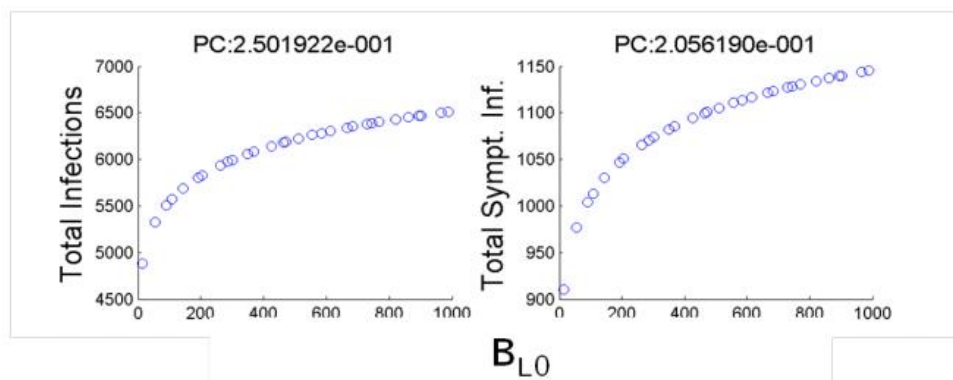
$\eta_1$



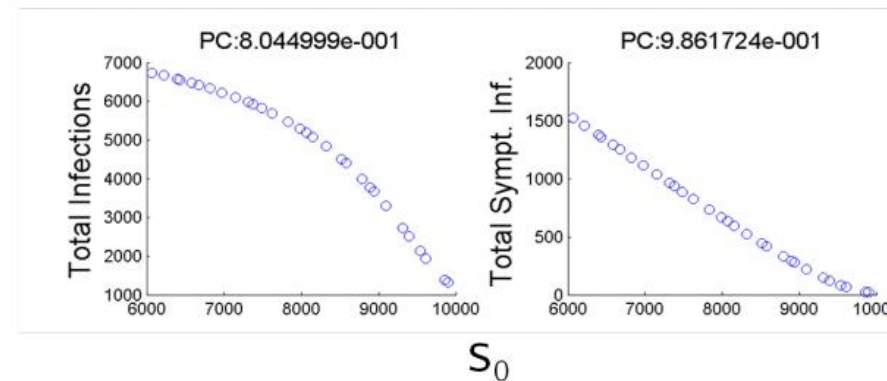
$\beta_L$  = low infectious



$\eta_2 = s \cdot \eta_1$



$B_{L0}$



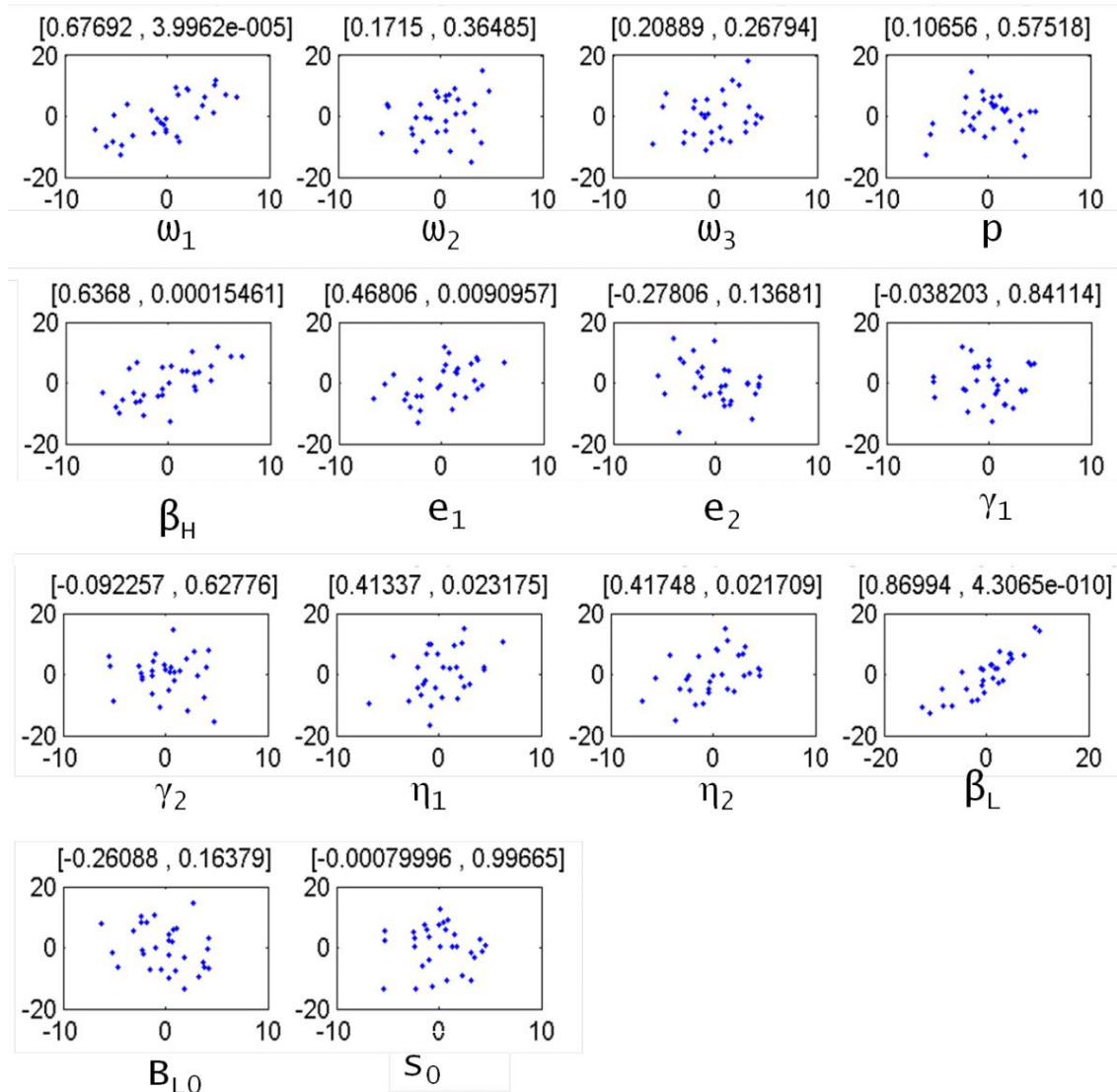
$S_0$

Figure 3.3: Monotonicity plots for  $\gamma_2$ ,  $\eta_1$ ,  $\eta_1$ ,  $\beta_L$ ,  $\eta_2$ ,  $B_{L0}$ , and  $S_0$

# LHS parameter ranges

Parameter	Min	Baseline	Max
$\omega_1$	0.001	1/180	0.03
$\omega_2$	0.001	1/(2*365)	0.003
$\omega_3$	0.0003	1/(10*365)	0.001
$p$	0.3	0.6	0.9
$r$	0.01	0.1	1
$e_2$	0.01	0.03	0.05
$e_1 = e_2/20$	0.0005	0.0015	0.0025
$\gamma_1$	1/2	0.75	1
$\gamma_2$	1/14	0.1	1/7
$\eta_1$	0.0001	0.008	0.05
$\eta_2$	1	100	200
$s$	0.001	0.008	0.08
$B_{L0}$	$\kappa_L/500$	$\kappa_L/2$	$\kappa_L$
$S_0$	6000	10000 - $\hat{S}_0$	10000

# LHS/PRCC results

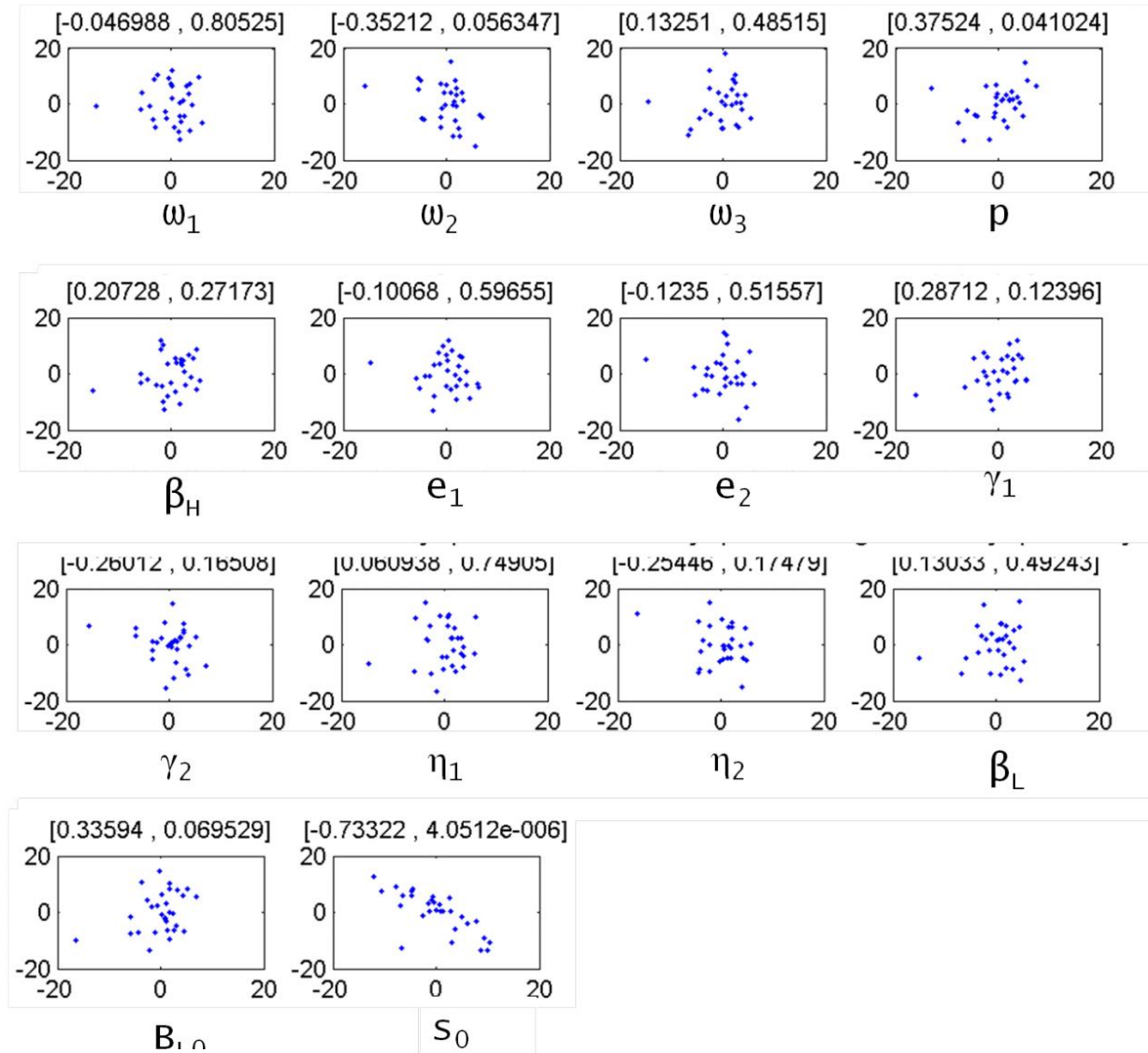


Annotations on top of plots: [PRCC, p-value]

PRCC Plots for Total Infecteds.



# LHS/PRCC results



Annotations on top of plots: [PRCC, p-value]

PRCC Plots for Total Symptomatic Infecteds.

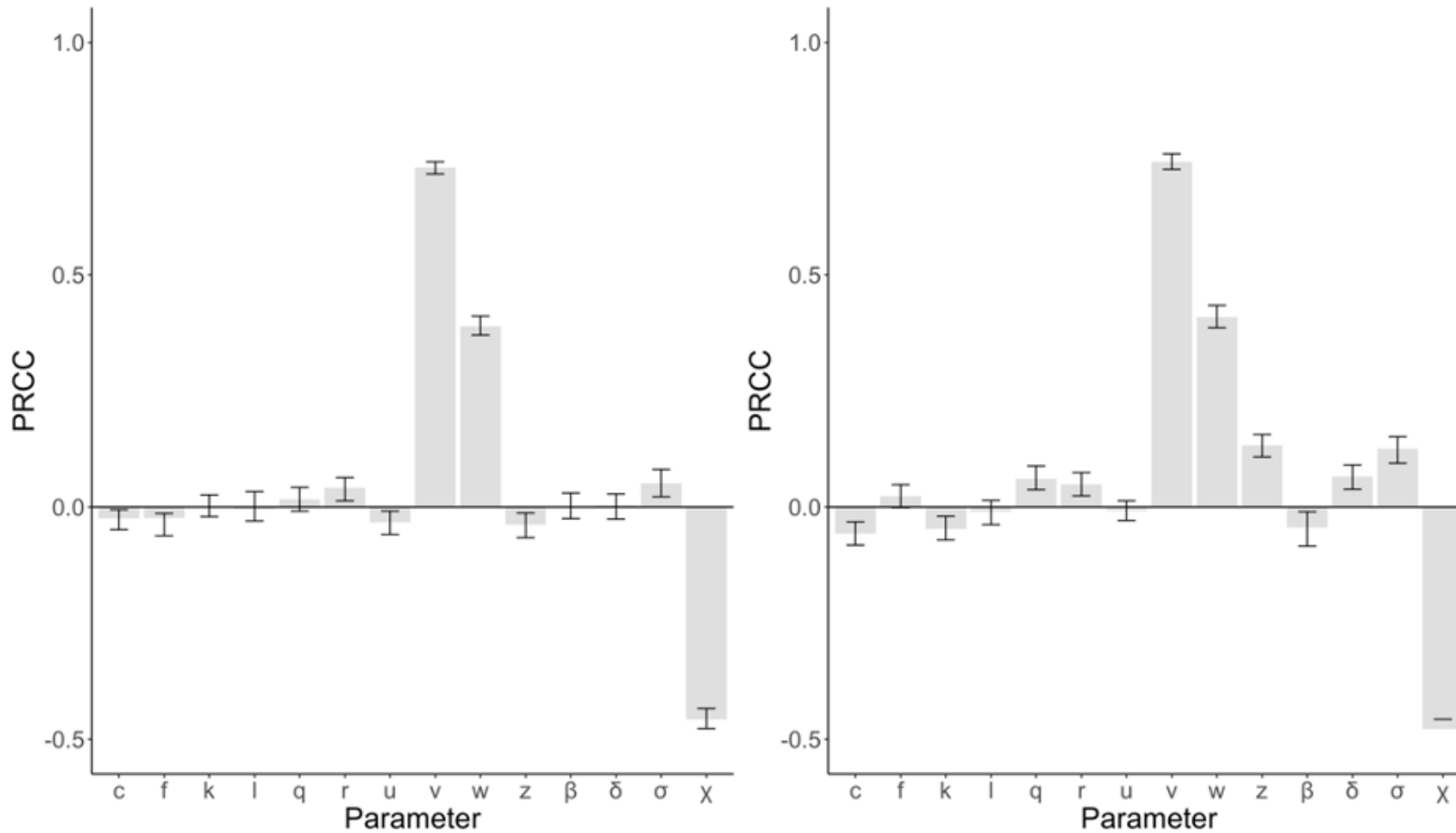
# LHS/PRCC results

Parameters	Total Infectious		Total Symptomatic Infecteds	
	<i>PRCC</i>	<i>p-value</i>	<i>PRCC</i>	<i>p-value</i>
$\omega_1$ : Waning $R_A$ to $\hat{S}$	*0.67692	4.00E-05	-0.04699	0.80525
$\omega_2$ : Waning $R_S$ to $\hat{S}$	0.1715	0.36485	-0.35212	0.056347
$\omega_3$ : Waning $\hat{S}$ to $S$	0.20889	0.26794	0.13251	0.48515
$p$ : Prop. Sympt.	0.10656	0.57518	0.37524	0.041024
$\beta_H = r * \beta_L$	*0.6368	0.000155	0.20728	0.27173
$e_1$ : Asympt. death rate	0.46806	0.009096	-0.10068	0.59655
$e_2$ : Sympt. death rate	-0.27806	0.13681	-0.1235	0.51557
$\gamma_1$	-0.0382	0.84114	0.28712	0.12396
$\gamma_2$	-0.09226	0.62776	-0.26012	0.16508
$\eta_1$	0.41337	0.023175	0.060938	0.74905
$\eta_2 = s * \eta_1$	0.41748	0.021709	-0.25446	0.17479
$\beta_L$ : Low infectious	***0.86994	4.31E-10	0.13033	0.49243
$B_{L0}$	-0.26088	0.16379	0.33594	0.069529
$S_0$	-0.0008	0.99665	** -0.73322	4.05E-06

(\*) is used to indicate possible contributors (PRCC values:  $\sim 0.5$  to  $0.69$  or  $-0.5$  to  $-0.69$ ), (\*\*) is used to indicate very likely contributors to uncertainty (PRCC values:  $\sim 0.7$  to  $0.79$  or  $-0.7$  to  $-0.79$ ) and (\*\*\*) is used to indicate highly likely contributors to uncertainty (PRCC values:  $\sim 0.8$  to  $0.99$  or  $-0.8$  to  $-0.99$ ). Grey shaded boxes are s



# Alternative means of presenting PRCC results



**Figure:** Global sensitivity analysis of the impact of parameters on **cumulative infections** (left panel) and **deaths averted** (right panel) over the range of daily vaccination rate using Partial rank correlation coefficient (PRCC).