### Capstone Project Final Presentation Automated Model Reduction for Atmospheric Chemical Mechanisms (AMORE Project)

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### Atmospheric Chemistry Modeling: Bridging Scales





### AMORE (<u>A</u>tmospheric Chemistry <u>MO</u>del <u>RE</u>duction)





An automated tool for flexibly generating accurate reduced chemical mechanisms for use in atmospheric chemistry and air quality models.

Two-step model reduction: graph theorybased strongly-connected component analysis followed by simplified quasisteady state analysis



### Introduction Objective

- Develop and implement a graph-based algorithm to automatically generate reduced gas-phase chemical mechanisms.
- Improve accuracy and efficiency of large-scaled atmospheric chemistry models.

### Why it is meaningful?

- Enable the improved, sustained and consistent development of chemical mechanisms for air quality forecasting, research, and policy analysis.
- Understand the air quality, especially how air pollutants move in the atmosphere.
- Limitation of the computational complexity
- The chemical stiffness in the detailed chemical mechanisms



# **Dataset Overview**



Isoprene Mechanisms	Number of species	Number of reactions
MCM v.3.3.1	602	1926
CB05	18	9
CB6r2	38	22
GEOS-Chem v11.02c	106	335
GFDL-AM3	23	12
GISS ModelE	15	3
Paulot 2009 (high and low NOx, reduced)	45	28
Wennberg 2018 (full)	385	810

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#### Dataset Format: Kinetic Preprocessor (KPP) format

- A file of chemical reactions in the full mechanism.
- A file of species involved in the full mechanism.
- A file contains editable input parameters and initial values of included species, temperature, etc.

### **Procedure** Data Preprocessing

An Example of Chemical Reaction



• A list of tuple with reaction and reaction rate formula

('ISOP + OH = ISOP1OHc', 2.7E - 11 × EXP(390 / TEMP) × 0.63 × 0.5)



### **Procedure** Data Preprocessing

#### ISOP + OH = ISOP1OHc: 2.7E - $11 \times EXP(390 / TEMP) \times 0.63 \times 0.5$

The weight of the reaction for the product

$$R_A = \sum_{i=1,I} v_{A,i} w_i$$

 $v_{A,i}$  : the stoichiometric of species A

 $w_i$ : the production rate

 $\begin{aligned} r_{ISOP1OHc} &= v_{ISOP1OHc,i} \times w_i \\ &= v_{ISOP1OHc,i} \times k \times c[ISOP] \times c[OH] \\ &= 1 \times 2.7E - 11 \times EXP(390/TEMP) \times 0.63 \times 0.5 \times c[ISOP] \times c[OH] \end{aligned}$ 

- k : the reaction rate for this particular reaction (equation)
- c[ISOP] and c[OH] are the initial concentration values of ISOP and OH respectively

### **Procedure** Data Preprocessing

How to measure the influence of one species on another?

$$r_{AB} = \frac{\sum |v_{A,i}w_i\delta_{Bi}|}{\sum |v_{A,i}w_i|}$$

Normalized contribution of species B to the production rate of species A

$$\delta_{Bi} = \begin{cases} 1, \text{ if the ith elementary reaction involves species E} \\ 0, \text{ otherwise} \end{cases}$$



### Procedure

### Graph Construction

Directed Relation Graph (DRG)

- Node: a species in the detailed mechanism
- Edge from A to B: A is dependent on B (the removal of B would directly induce significant error to the production rate of A)

#### $ISOP + OH = ISOP1OHc: 2.7E - 11 \times EXP(390 / TEMP) \times 0.63 \times 0.5$





# Procedure

### Graph Construction

### Directed Relation Graph (DRG)

- Perform a depth first search starting from the 'starting sets' and obtain a skeleton graph
- We used organic species as starting sets while treating others as background species.





### **Procedure** Graph Reduction

Method:

- Define a threshold value  $\epsilon$
- Only keep the paths that all the edges has a weight greater than  $\epsilon$  such that  $r_{AB} \ge \epsilon$





# Results

### The number of Species for various threshold values



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# Product

#### Python Package Chem\_graph

We created a python package called chem\_graph accessible on Github. Our product contains following functionalities:

- Read and parse the raw file, and construct the graph accordingly
- Given threshold value and starting set, compute the reduced graph and dependent set
- Visualize the graph



## Product

Algorithm 1 Original DFS Approach	Algorithm 2 Modified DFS A	
Given $r_{AB}$ dictionary, list of $\epsilon E$ . starting set S	Given $r_{AB}$ dictionary, list of $\epsilon I$	
Create graph $q$ based on $r_{AB}$	Sort $r_{AB}$ dictionary based on the	
for each $\epsilon$ in $E$ do	Sort $E$ from high to low;	
create visited set $v$	create visited set $v$	
create stack $s$ , push $S$ into $s$	create set $c$ to keep track of gray	
<b>while</b> $s$ not empty <b>do</b>   pop one element $e$ from $s$	for each $\epsilon$ in $E$ do   include new edges with value	
for each neighbor $n$ of $e$ do   if the edge weight smaller than $\epsilon$ , continue	for each new edge do if the destination node of	
if $n$ visited, continue	DFS from the destination	
push $n$ to $v$ and $s$	$\mathbf{end}$	
end	output current $v$ for current	
end	$\mathbf{end}$	
output $v$ for $\epsilon$		
end		

Algorithm 2 Modified DFS Approach Given  $r_{AB}$  dictionary, list of  $\epsilon$  E. starting set Sort  $r_{AB}$  dictionary based on the edge weight from high to low; ort E from high to low; reate visited set vreate visited set vreate set c to keep track of graph or each  $\epsilon$  in E do include new edges with value smaller than last  $\epsilon_0$  but larger than current  $\epsilon$ for each new edge do if the destination node of the new edge is not visited, continue DFS from the destination node, push the newly visited to vend output current v for current  $\epsilon$ nd

Running time: O(n(V+E))

Running time: O(nlog(n) + V+E)



# Product

#### Sample Usage

from chem_graph import *		
demo_graph = ChemGraph()		
priority_species = ['ISOP1CO400Hc']		
<pre>demo_graph.construct(path='isoprene_full_v5',</pre>	starting_set=priority_species,	<pre>must_contain=['ISOP'])</pre>

File pathStarting setMust contain



# **Product** Sample Usage



Skeleton Graph

#### Whole Graph



## **Future Direction**

- Evaluation on the reduce mechanism
- Evaluation on the generated graph



### References

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# **Thank You!**

