

Model Reduction

RMG Study Group Presentation

Kehang Han

Apr. 28, 2015

Outline

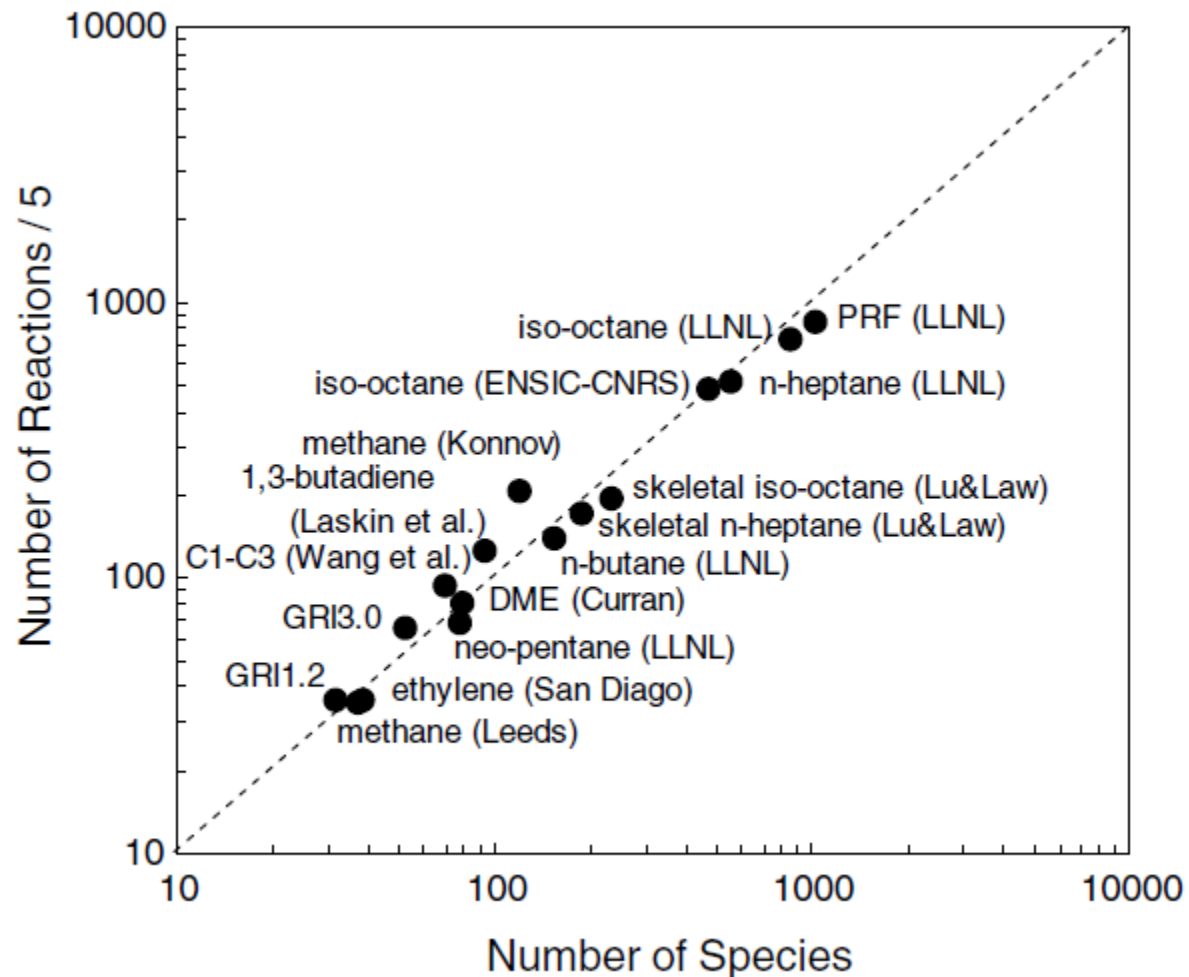
- Why we need model reduction
- Two levels of model reduction
- Most recent methods
- Personal opinion

High CPU time in predictive combustion

- Predictive combustion: detailed model + CFD
- For normal detailed kinetic mechanisms, over 90% of the CPU time of reactive flow simulations is spent solving the ODE systems.[1]
- Why high?
 - Many species and reactions in detailed kinetic models
 - Very different time scales in the models → computational stiffness

[1] Long Liang, etc. The use of dynamic adaptive chemistry in combustion simulation of gasoline surrogate fuels (2009)

Many species and rxns in detailed models



- Tens to hundreds of species
 - Lots of state variables in CFD
 - (x1, x2, x3, ..., T, P)
- Hundreds to thousands of rxns [1]
 - RMG gives even more rxns
- Butanol kinetics by Dr. Cai [2]
 - ~90% rxns are insignificant under conditions of his interest

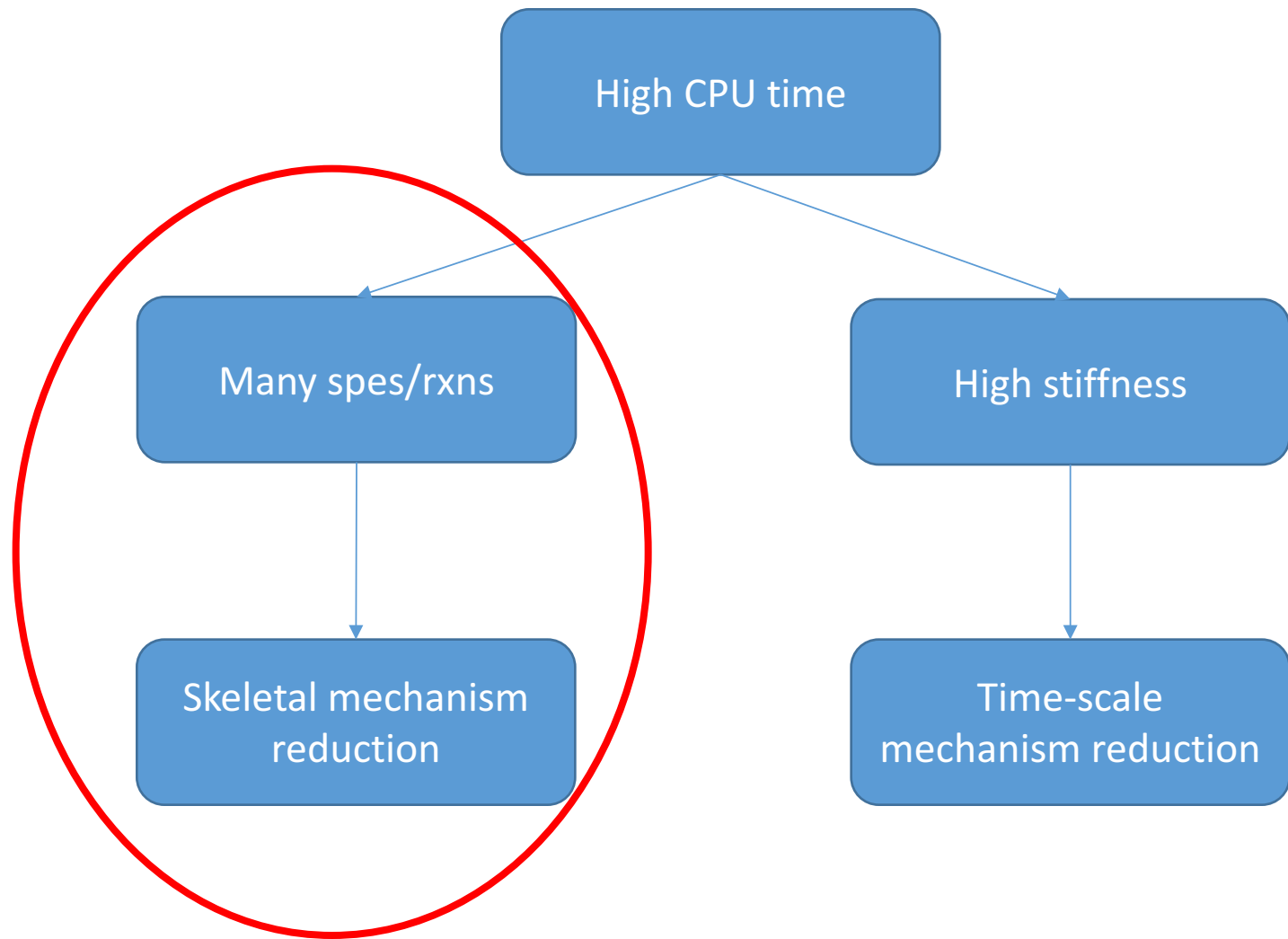
[1] C.K. Law, *Proc. Combust. Inst.* 31 (2007) 1–29.

[2] Personal conversation with Dr. Jianghua Cai on Apr. 22, 2015

Computational stiffness

- Highly reactive radicals and reactions → very short time scales (usually cause high stiffness ratio*)
- Short time scales → small integration steps → long CPU time

Note: *stiffness ratio* = $\frac{\text{max eigen value}}{\text{min eigen value}}$



High CPU time

Many spes/rxns

High stiffness

Skeletal mechanism
reduction

Time-scale
mechanism reduction

Skeletal mechanism reduction

- Identify and eliminate unimportant species and reactions
- Various methods along history
 - Sensitivity analysis
 - Principal component analysis
 - Jacobian analysis
- Most recent methods
 - Directed relation graph (DRG) by Tianfeng Lu (Princeton)
 - Optimization-based approach by Binita Bhattacharjee (MIT)

DRG method

- Being Unimportant: negligible contribution to production rate of every other species [1]

$$R_A = \sum_{i=1,I} v_{A,i} \omega_i, \quad (1)$$

$$\omega_i = k_{fi} \prod_{j=1}^K C_j^{v'_{ij}} - k_{bi} \prod_{j=1}^K C_j^{v''_{ij}}, \quad (2)$$

$$k_{fi} = [A_i T^{n_i} \exp(-E_i/RT)] F_i, \quad (3)$$

- Dependence of A on B

$$r_{AB} \equiv \frac{\sum_{i=1,I} |v_{A,i} \omega_i \delta_{Bi}|}{\sum_{i=1,I} |v_{A,i} \omega_i|}, \quad (4)$$

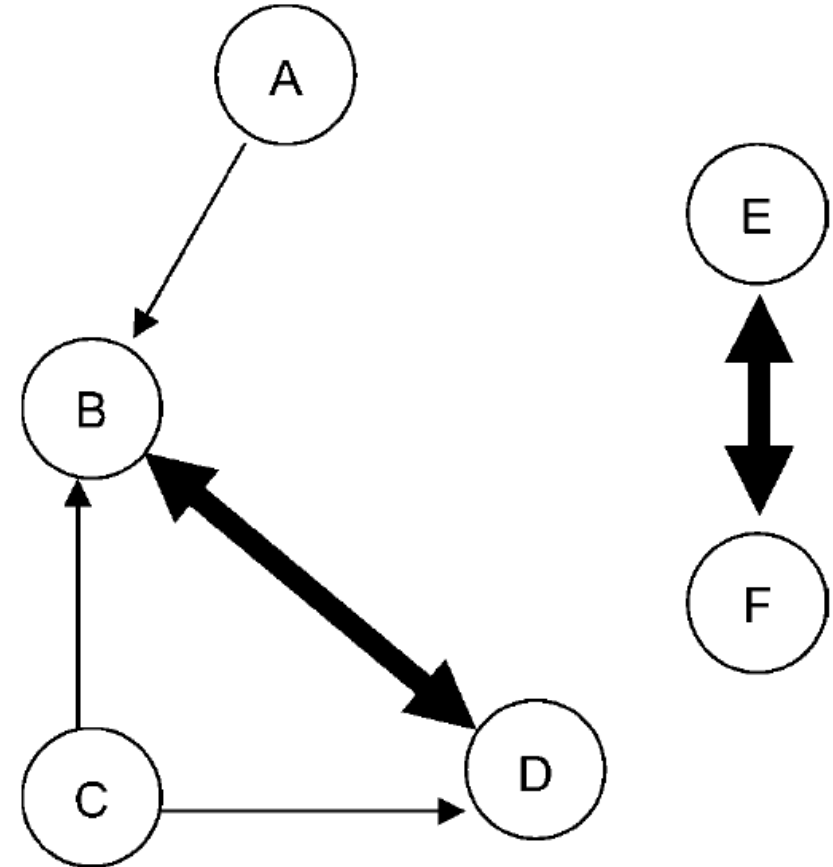
$$\delta_{Bi} = \begin{cases} 1 & \text{if the } i\text{th elementary reaction} \\ & \text{involves species } B, \\ 0 & \text{otherwise.} \end{cases} \quad (5)$$

[1] Tianfeng Lu, Chung K. Law, *A directed relation graph method for mechanism reduction* 2005

Eliminate negligible dependence

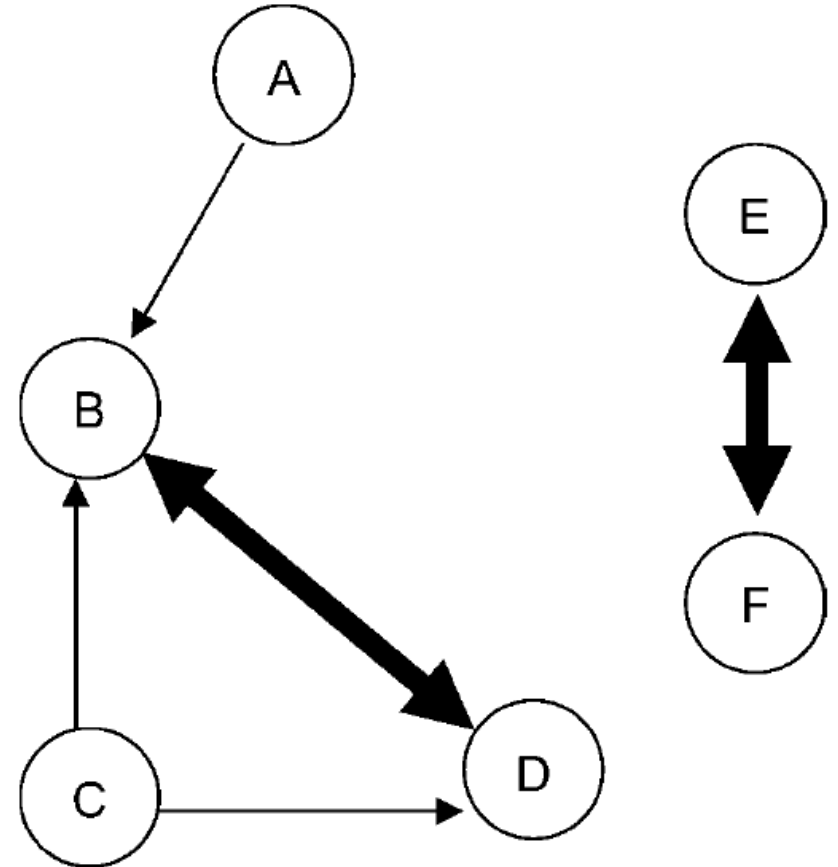
Any $r_{ij} < \varepsilon$ (say, 0.01) would be regarded as negligible dependence

r_{ij}	A	B	C	D	E	F
A	1	0.2	0.001	0.001	0.002	0.001
B	0.001	1	0.001	0.8	0.002	0.002
C	0.002	0.2	1	0.15	0.002	0.001
D	0.001	0.9	0.002	1	0.001	0.002
E	0.001	0.002	0.001	0.002	1	0.9
F	0.001	0.002	0.002	0.001	0.9	1



Get dependence set of species A

- It seems that A only depend on B
 - To estimate A's flux, B should be included in skeletal model
- But B depends on D heavily
 - To estimate B's flux, D should be included in skeletal model
- So A's dependence set is {B, D}
- Dependence set of A contains all the vertices to which there's at least a path from A



DRG Algorithm

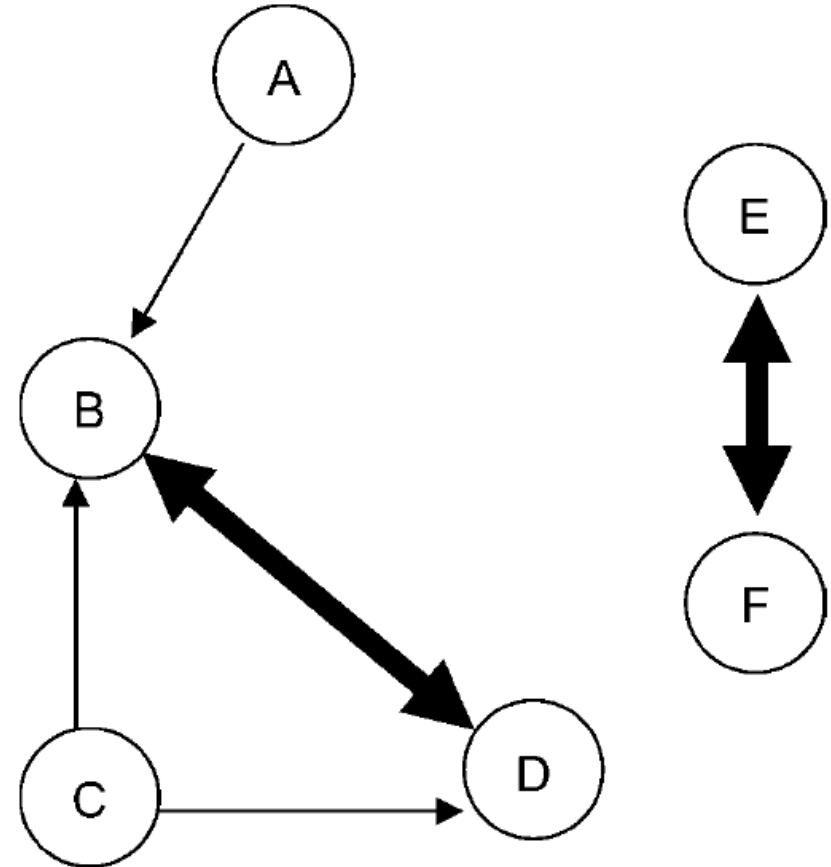
- Step1: select a set of starting species like A
- Step2: calculate r_{ij} (an $N \times N$ matrix where N is the number of species in full mechanism) under selected condition (T, P...)
- Step3: select tolerance ε (there's a paper studying how to choose it [1])
- Step4: draw the dependence graph
- Step5: determine dependence set for each starting species
- Step6: get the final skeletal mechanism by forming union of all the dependence sets f starting species

[1] Xiaolong Gou, Zheng Chen, Wenting Sun, Yiguang Ju. A dynamic adaptive chemistry scheme with error control for combustion modeling with a large detailed mechanism. 2012.

Improved DRG

- Example:
 - Condition: post-ignition
 - Starting species includes CO
 - Skeletal mechanism contains lots of heavy hydrocarbon species
- Reason: strong back propagating connections from CO to heavy HC
- Add search-depth control [1]

$$R_{V_0}(V) = \max_{\Omega} \{\Pi r_{ij}\},$$



Optimization-based reduction

- State variable vector: $x = (x_1, x_2, \dots, x_n, T, P)$
- The optimization formulation would be

$$\min \sum_{k=1}^{N_R} z_k$$

$$G[x(t, z), x_{ref}(t)] \leq 0$$

$$z_k \in \{0, 1\}$$

$$\left. \begin{aligned} \dot{x}_{j,ref} &= \Gamma_{j,ref}(\mathbf{x}_{ref}), & x_{j,ref}(t_0) &= x_{j,ref,t_0} \\ \dot{x}_j &= \Gamma_j(\mathbf{x}, \mathbf{z}), & x_j(t_0) &= x_{j,ref,t_0} \end{aligned} \right\}$$

$$\forall j = 1 \dots N_S + 1 \quad (3)$$

$$G = \sum_{j=1}^{N_S+1} \alpha_j \left(\int_0^{t_f} \frac{x_j(t, z) - x_{j,ref}(t)}{x_{j,ref}(t)} dt \right)^2 - \delta_1 \leq 0$$

Optimization-based reduction

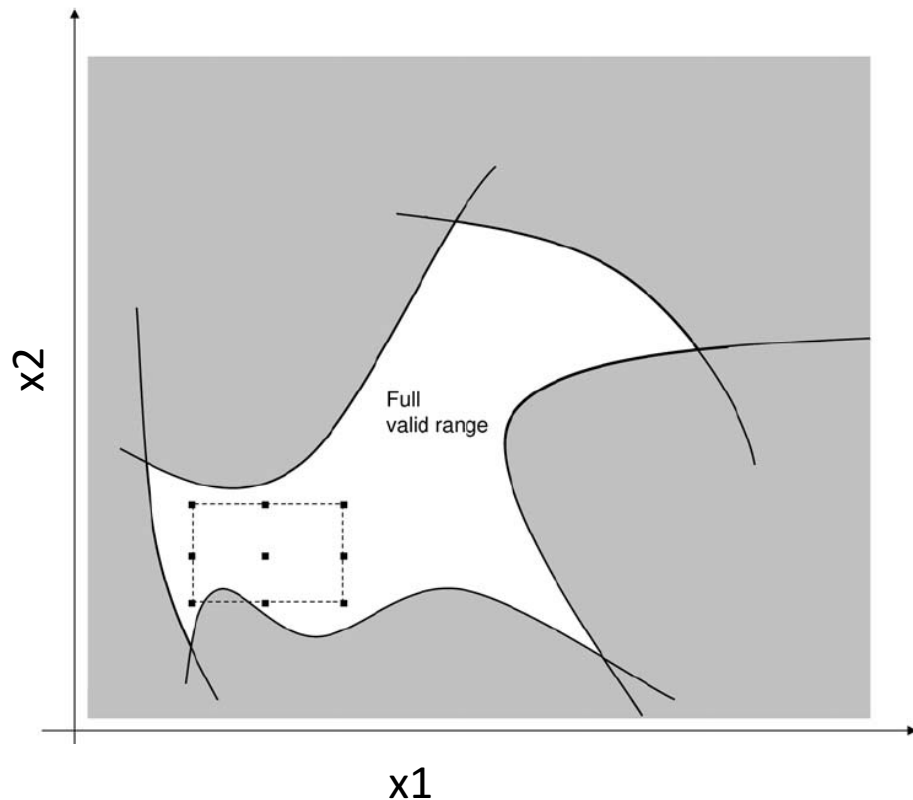
- State variable vector: $x = (x_1, x_2, \dots, x_n, T, P)$
- The optimization formulation would be

$$\begin{aligned}
 & \min \sum_{k=1}^{N_R} z_k & \left. \begin{aligned} \dot{x}_{j,ref} &= \Gamma_{j,ref}(\mathbf{x}_{ref}), & x_{j,ref}(t_0) &= x_{j,ref,t_0} \\ \dot{x}_j &= \Gamma_j(\mathbf{x}, \mathbf{z}), & x_j(t_0) &= x_{j,ref,t_0} \end{aligned} \right\} \\
 & G[x(t, \mathbf{z}), x_{ref}(t)] \leq 0 & \forall j = 1 \dots N_S + 1 & \quad (3) \\
 & z_k \in \{0, 1\} & G_{j,i} = |\Gamma_j(\mathbf{x}_{ref}(t_i), \mathbf{z}) - \Gamma_{ref,j}(\mathbf{x}_{ref}(t_i))| - (atol_j \\
 & & + rtol_j |\Gamma_{ref,j}|) \leq 0 \quad \forall j = 1 \dots N_S + 1, i = 1 \dots N_t
 \end{aligned}$$

Global minimum is guaranteed!

Identify valid range for reduced model

- We want to use the reduced model over a range of conditions
- But the reduced model is produced under a single condition



$$\left. \begin{aligned} \dot{x}_{j,ref} &= \Gamma_{j,ref}(\mathbf{x}_{ref}), & x_{j,ref}(t_0) &= x_{j,ref,t_0} \\ \dot{x}_j &= \Gamma_j(\mathbf{x}, \mathbf{z}), & x_j(t_0) &= x_{j,ref,t_0} \end{aligned} \right\}$$

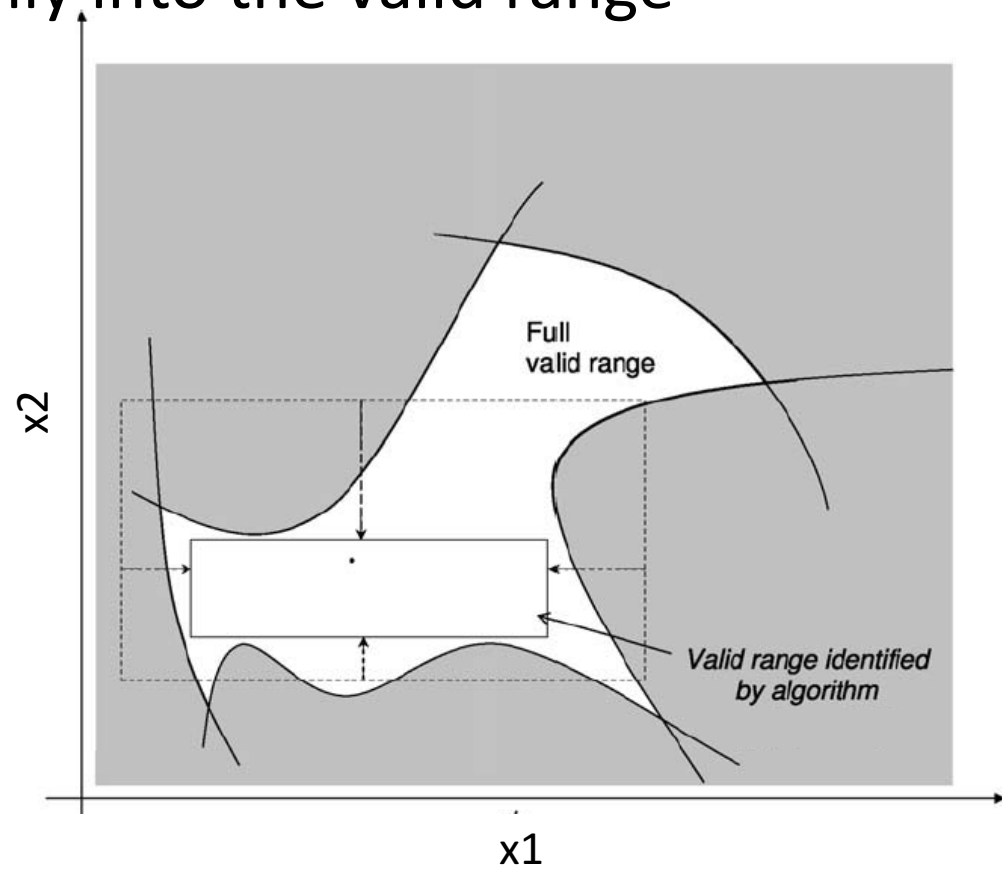
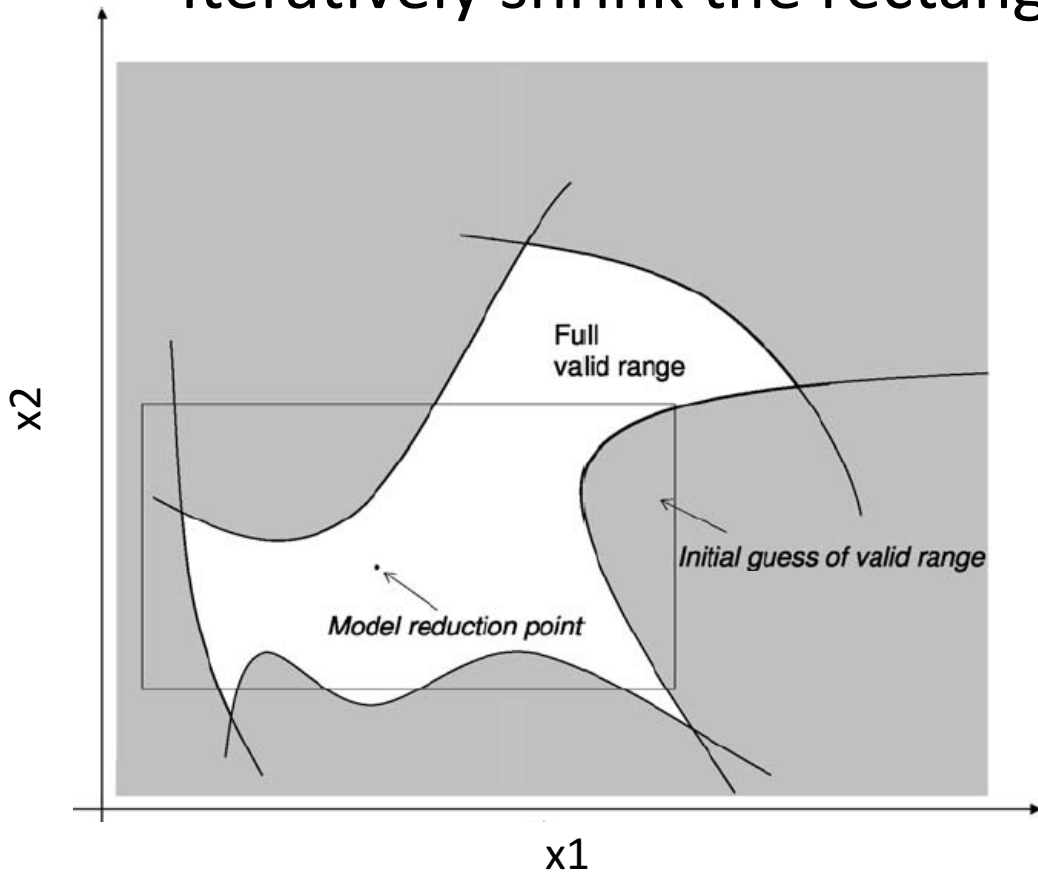
$$\forall j = 1 \dots N_S + 1$$

$$G_{j,i} = |\Gamma_j(\mathbf{x}_{ref}(t_i), \mathbf{z}) - \Gamma_{ref,j}(\mathbf{x}_{ref}(t_i))| - (atol_j + rtol_j |\Gamma_{ref,j}|) \leq 0 \quad \forall j = 1 \dots N_S + 1, i = 1 \dots N_t$$

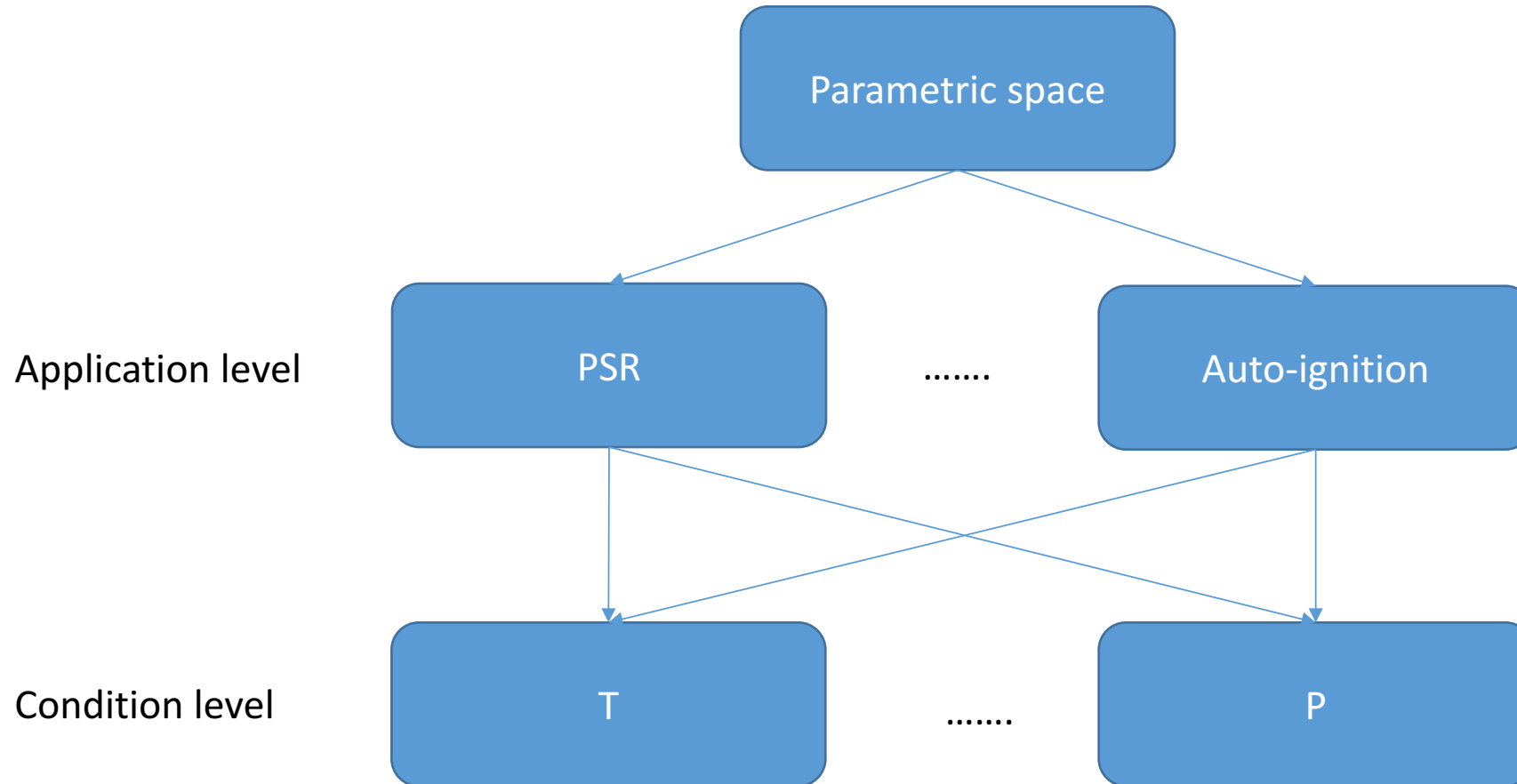
- Fix \mathbf{z}^* , solve for the range of condition

Identify valid range for reduced model

- Initially guess a range → a rectangle
- Iteratively shrink the rectangle fully into the valid range



Make skeletal mechanism valid in a range of parameters



Run model reduction many times under different points in the parametric space and form union of sub-skeletal mechanism

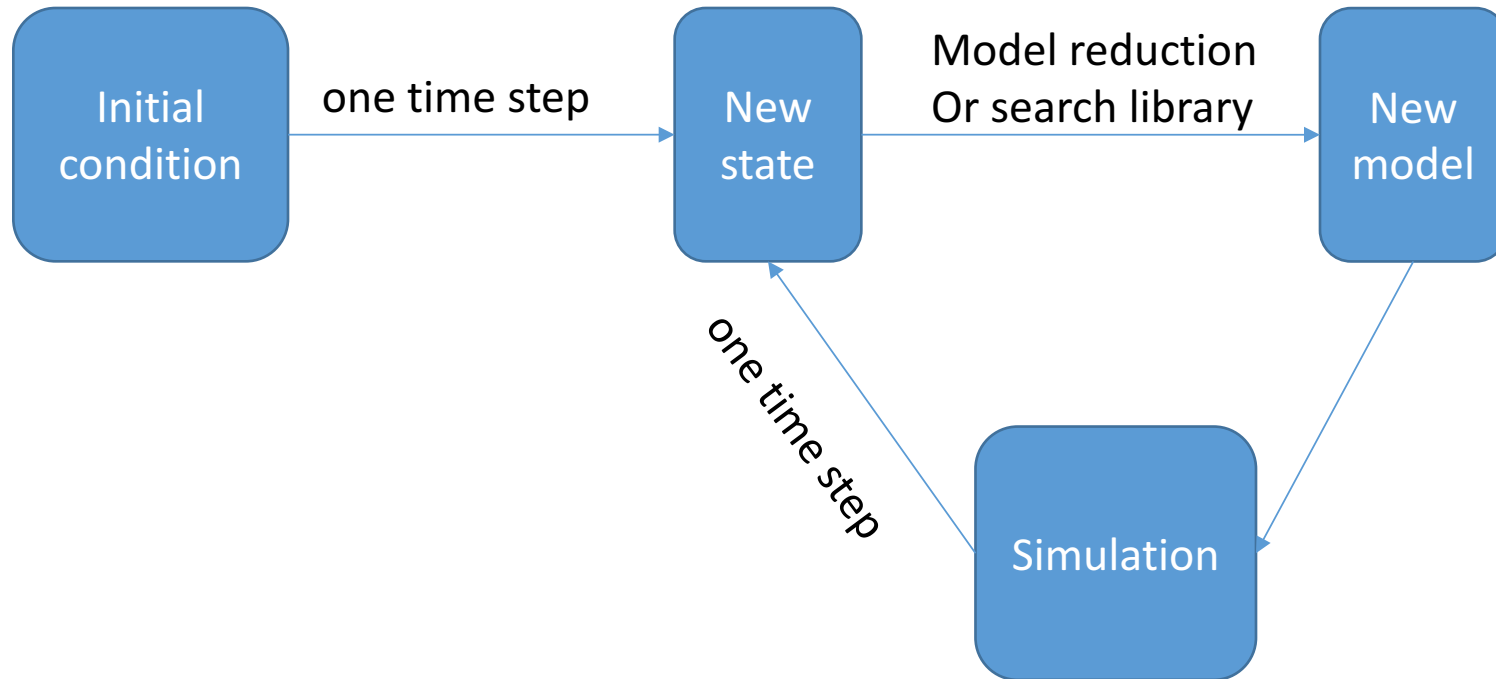
Adaptive chemistry

- Previous union of skeletal mechanisms
 - make big mechanism as well
- An alternative strategy by making use of the nature of local accuracy
- Two approaches to do adaptive chemistry
 - Create a library of reduced models before CFD and choose the most appropriate one during CFD [1]
 - Reduce model on-the-fly [2]

[1] Oluwayemisi O. Oluwole, Binita Bhattacharjee, John E. Tolsma, Paul I. Barton, William H. Green. Rigorous valid ranges for optimally reduced kinetic models. 2006

[2] Long Liang, John G. Stevens, John T. Farrell. A dynamic adaptive chemistry scheme for reactive flow computations. 2009

Adaptive chemistry scheme in CFD



Personal opinion about RMG and model reduction

- RMG is good at generating detailed mechanism over a certain valid condition range [1]
- If we want to add model reduction module,
 - Could use DRG or the optimization-based reduction
- How to pick condition for reduction
 - Single-condition RMG job, directly use that condition and figure out the valid condition range of the reduced mechanism
 - Multi-condition RMG job, run each condition and do union of sub-mechanism
- The reduced mechanism produced by RMG can still use Adaptive Chemistry fashion to conduct CFD

Backup: Homogeneous reduction \rightarrow diffusive applications

