

pynta: An automated workflow code for reaction path exploration on surfaces

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Open-Source Software for Kinetics, Chemical Networks, & Reactor Modeling

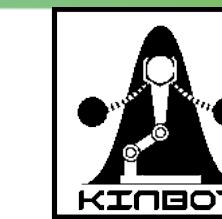
2022 Fall ACS Meeting

8/21 – 8/25, 2022

Funding: DOE BES CTC

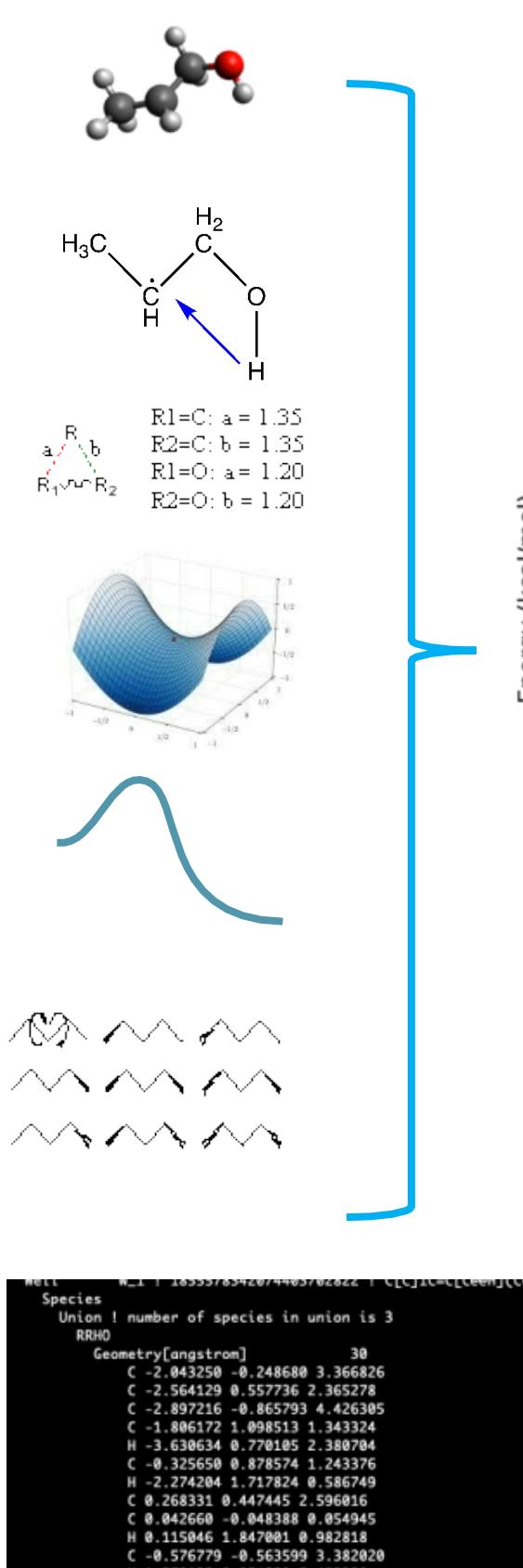
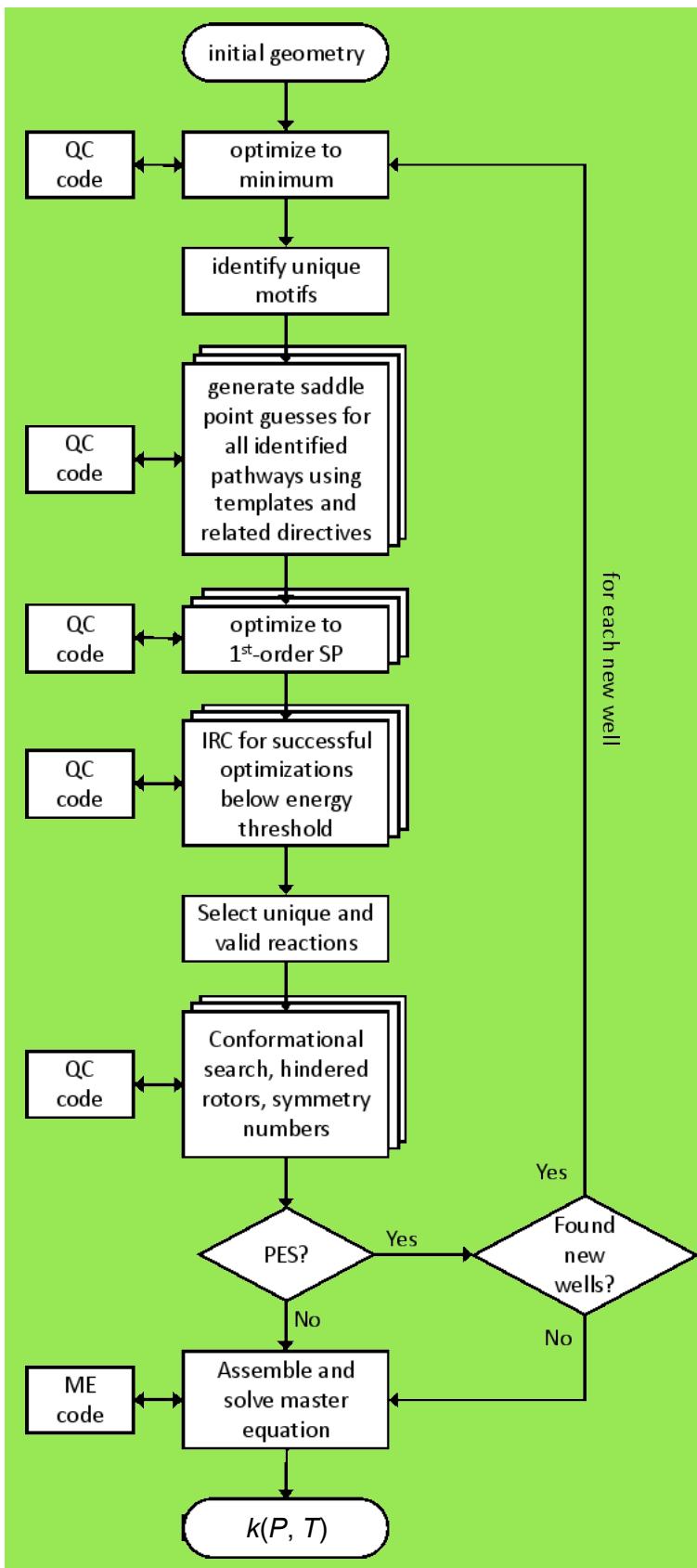


Automation for gas-phase kinetics

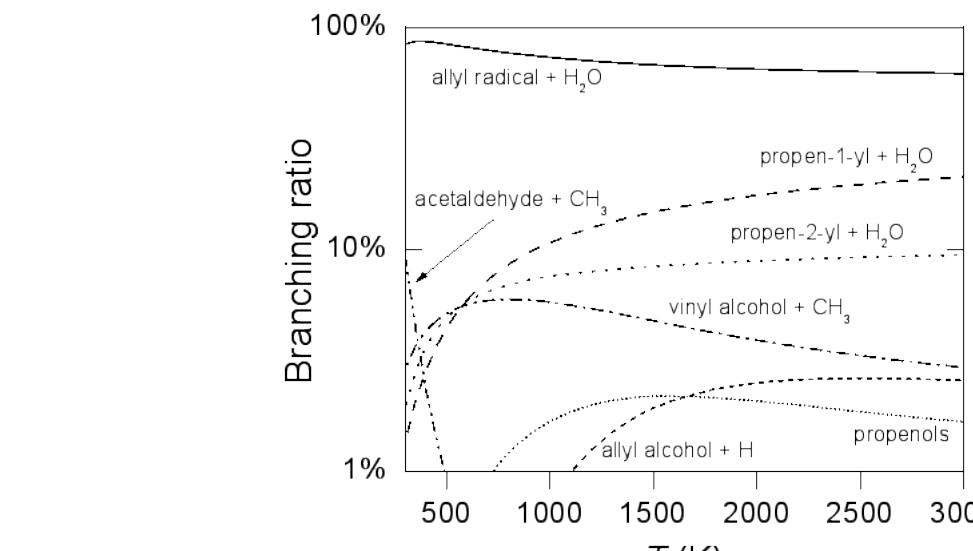


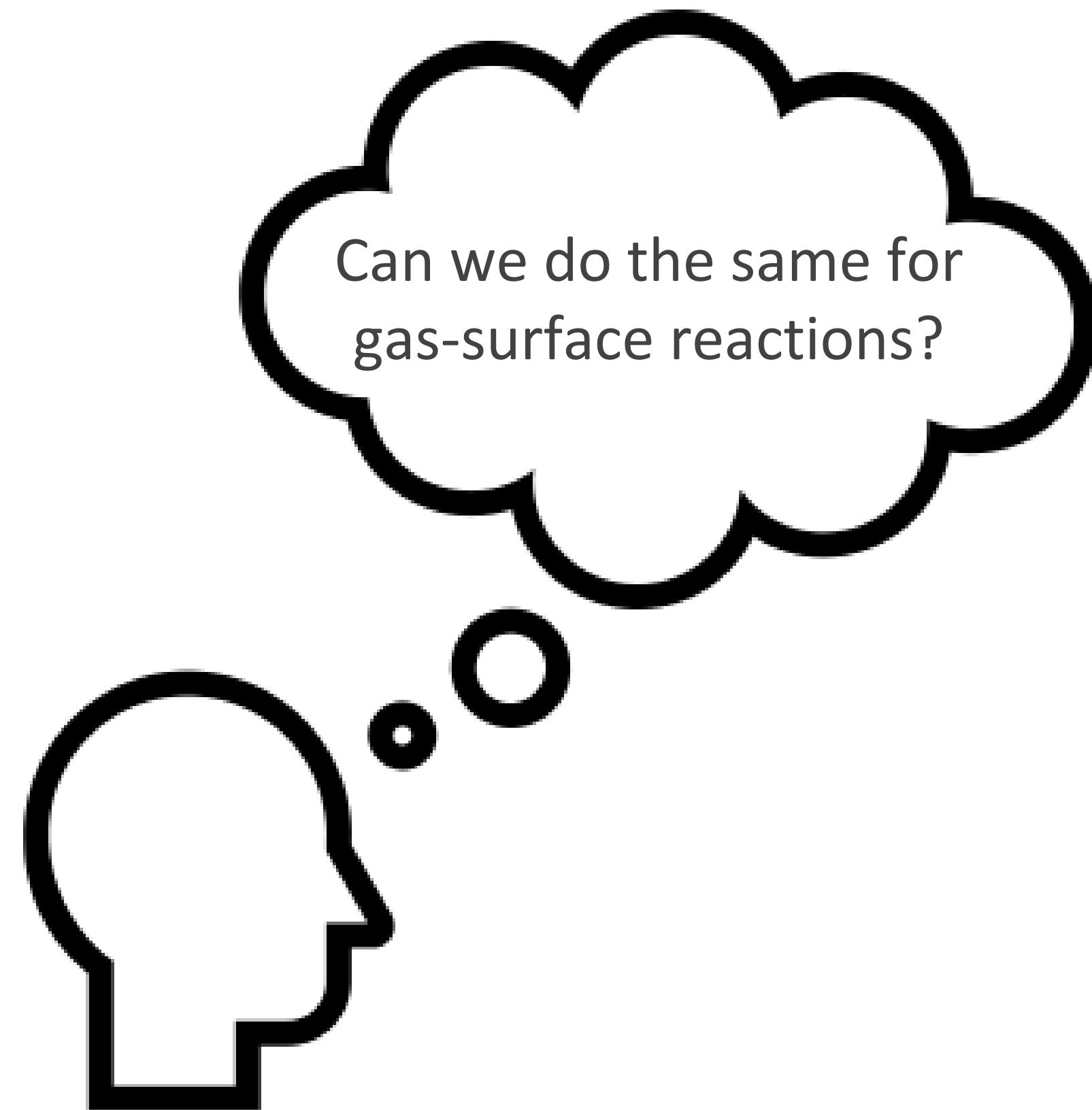
github.com/zadorlab/KinBot

Van de Vijver, JZ, Computer Physics Communication, 2020, 248, 106947



Well Species Union ! number of species in union is 3
RRHO
Geometry[angstrom] 30
C -2.043250 -0.248680 3.366826
C -2.564129 0.557736 2.365278
C -2.897216 -0.865793 4.426305
C -1.806172 1.098513 1.343324
H -3.630634 0.770105 2.380704
C -0.325650 0.878574 1.243376
H -2.274204 1.717824 0.586749
C 0.268331 0.447445 2.596016
C 0.042660 -0.048388 0.054945
H 0.115046 1.847001 0.982818
C -0.576779 -0.563599 3.382020
H -0.226223 -0.251220 3.452200





Can we do the same for
gas-surface reactions?

From formal catalytic reactions to *ab initio* kinetics

RMG reaction representation

- index: 0

reaction: CH3X + X <=> CH2X + HX

reaction_family: Surface_Abstraction

reactant: |

1 *1 C u0 p0 c0 {2,S} {3,S} {4,S} {5,S}

2 *2 H u0 p0 c0 {1,S}

3 H u0 p0 c0 {1,S}

4 H u0 p0 c0 {1,S}

5 X u0 p0 c0 {1,S}

6 *3 X u0 p0 c0

product: |

1 *1 C u0 p0 c0 {3,S} {4,S} {5,D}

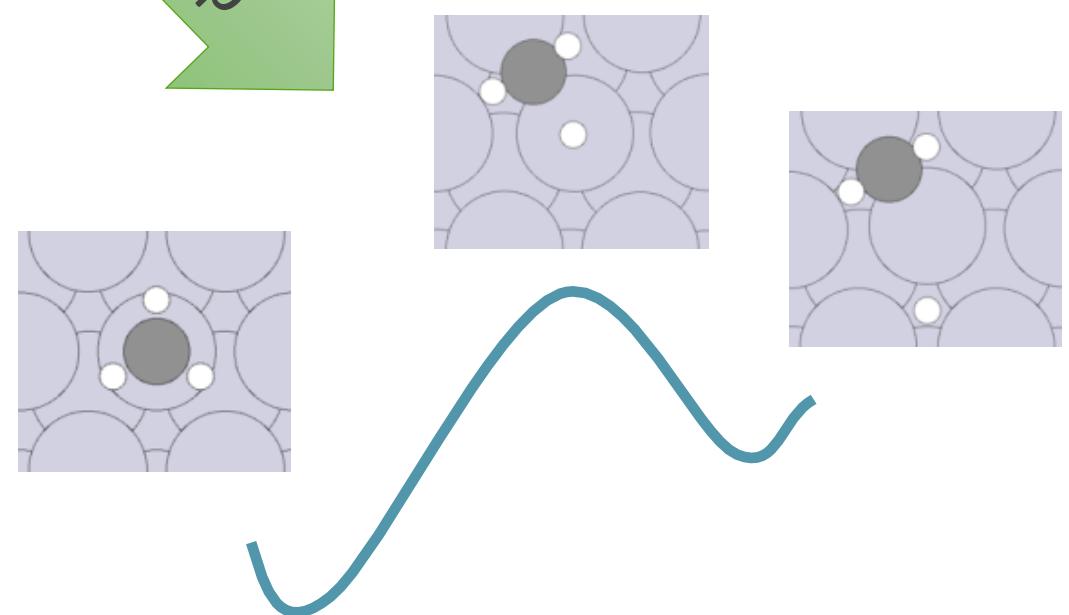
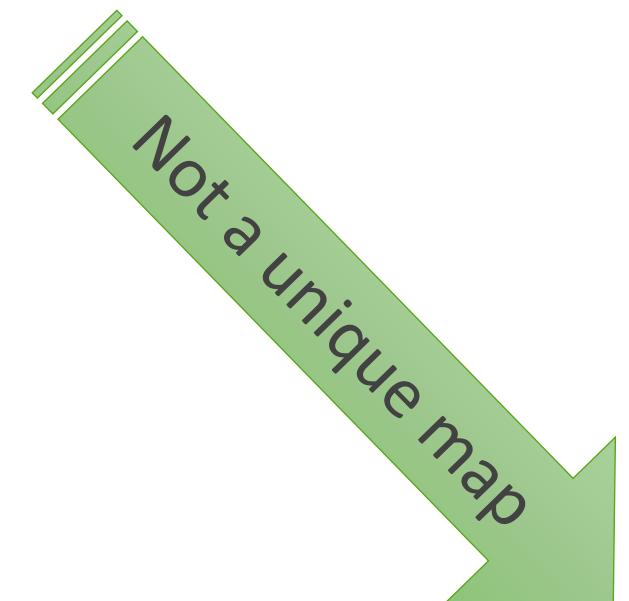
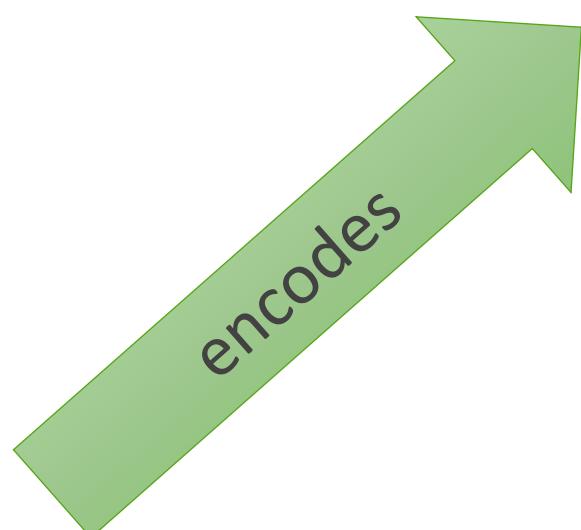
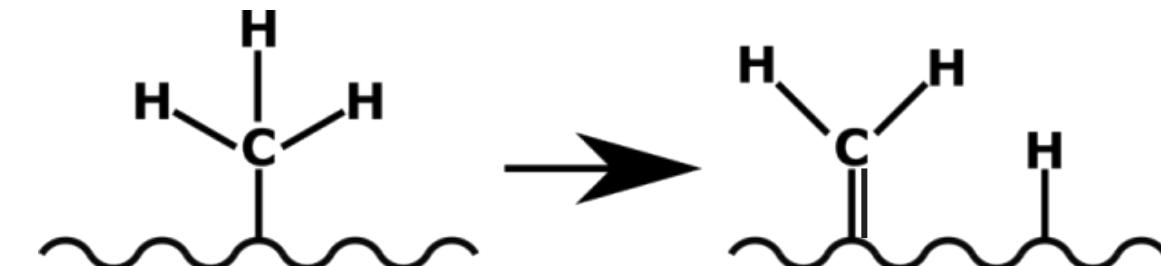
2 *2 H u0 p0 c0 {6,S}

3 H u0 p0 c0 {1,S}

4 H u0 p0 c0 {1,S}

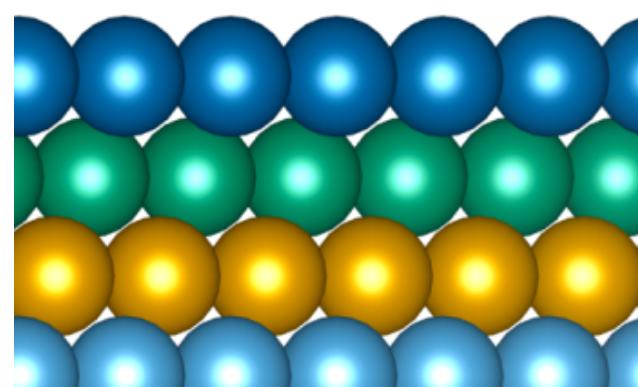
5 X u0 p0 c0 {1,D}

6 *3 X u0 p0 c0 {2,S}



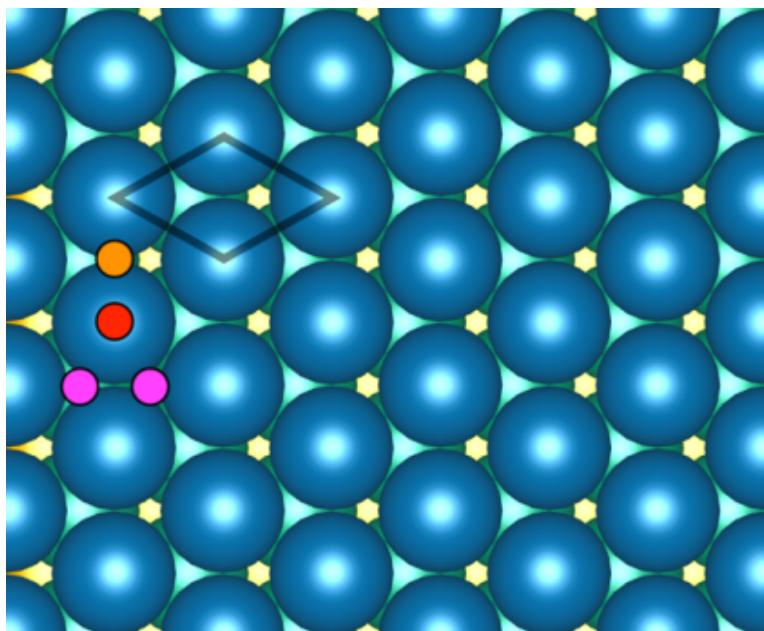
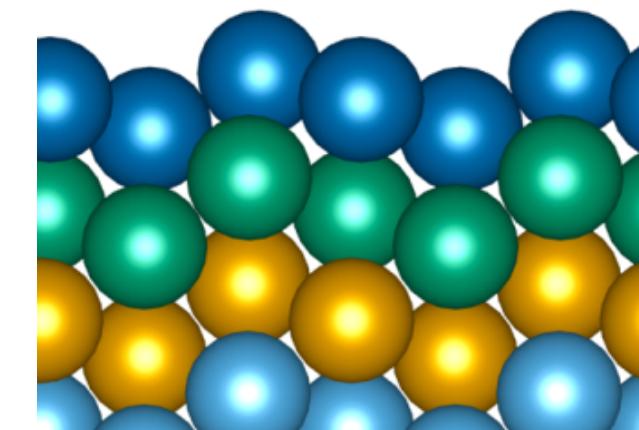
There are many potential binding sites even for simple surfaces

FCC(111)

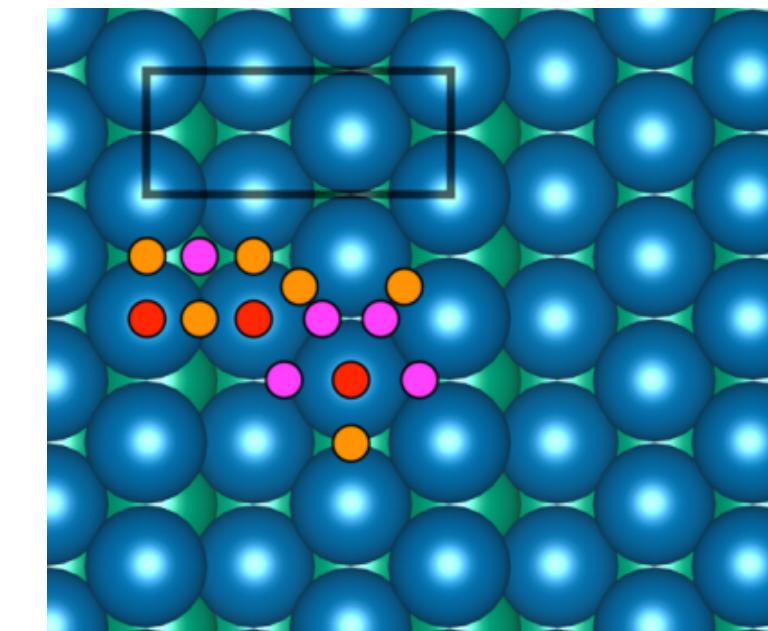


Side view

FCC(211)



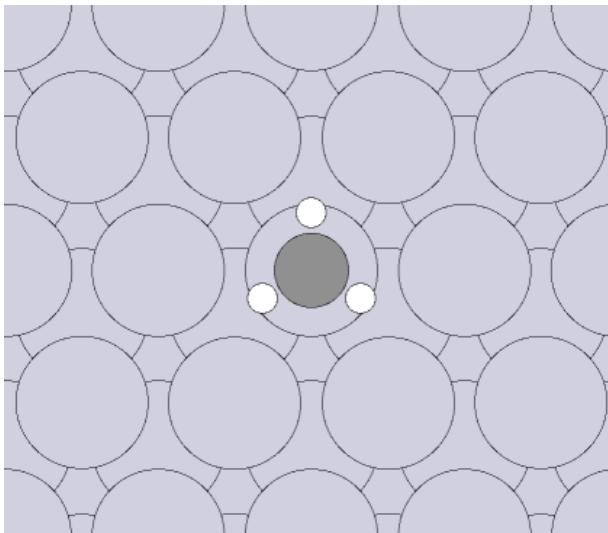
Top view



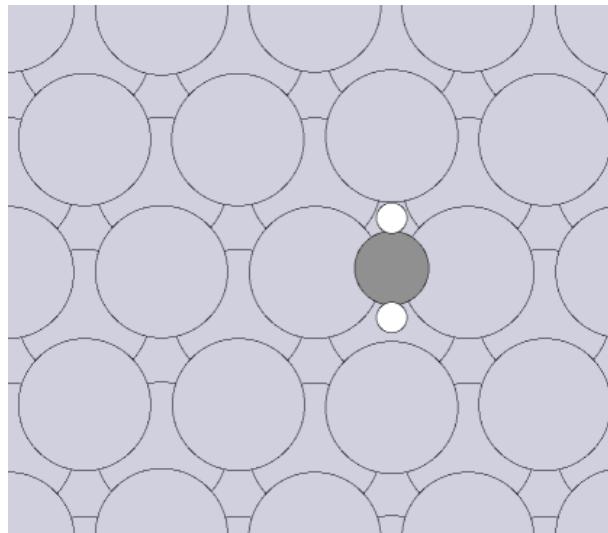
...and there are even more for high-index metal facets or nanoparticles.

There are only a few sites that are minima on Pt(111) for these species

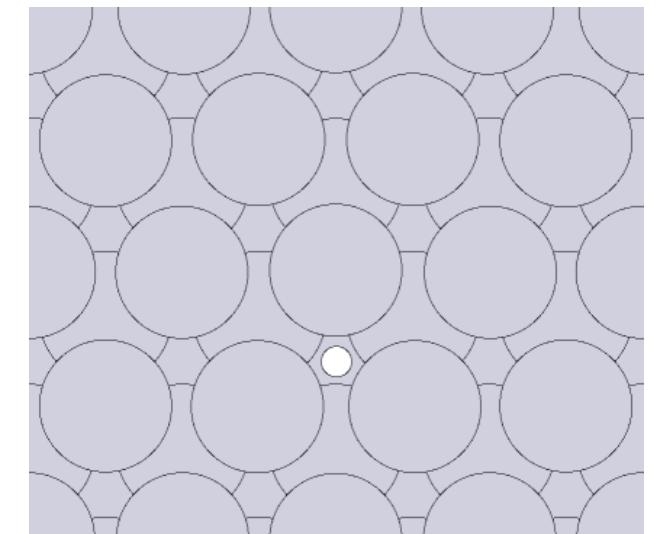
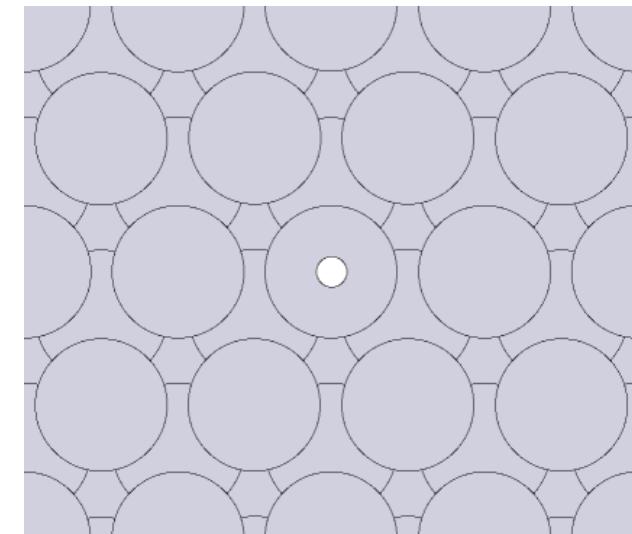
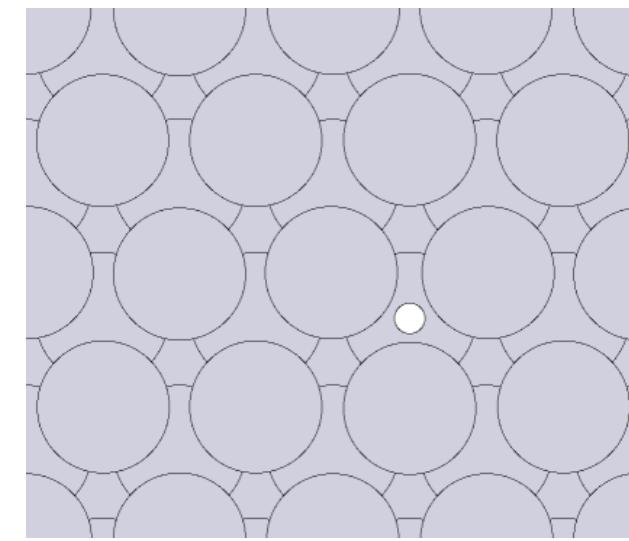
CH_3



CH_2

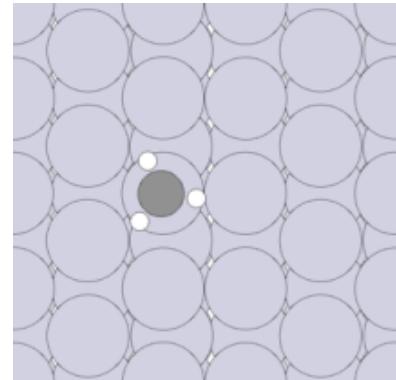


H

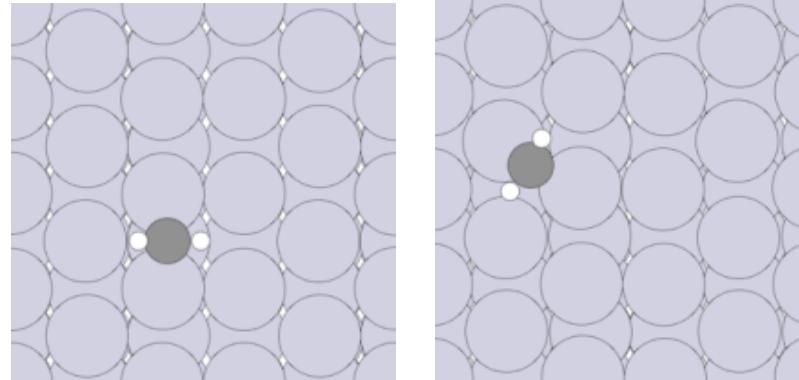


But there are a lot more already on Pt(211)

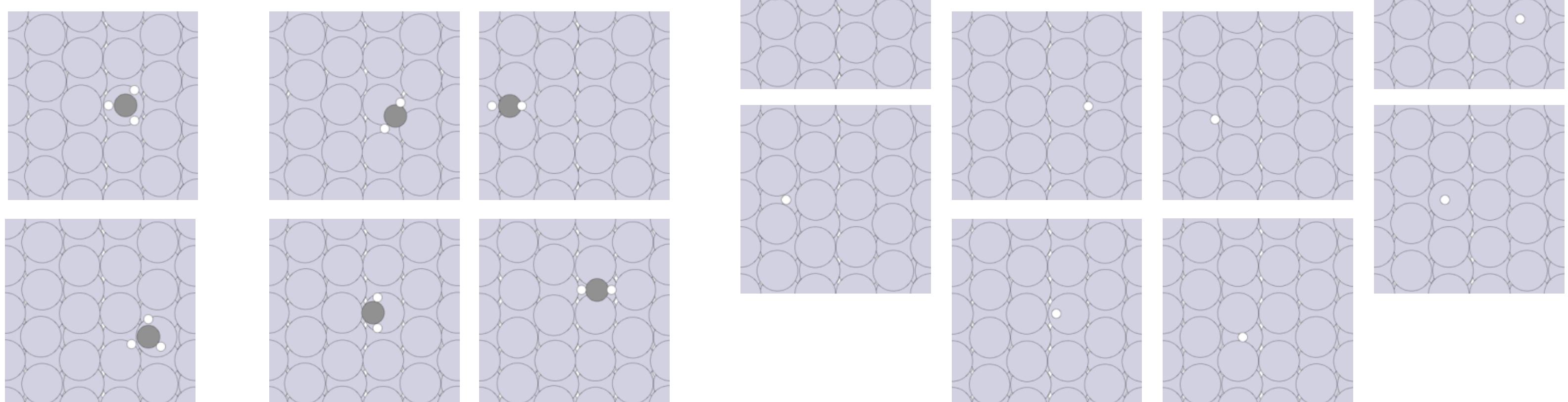
CH₃



CH₂



H

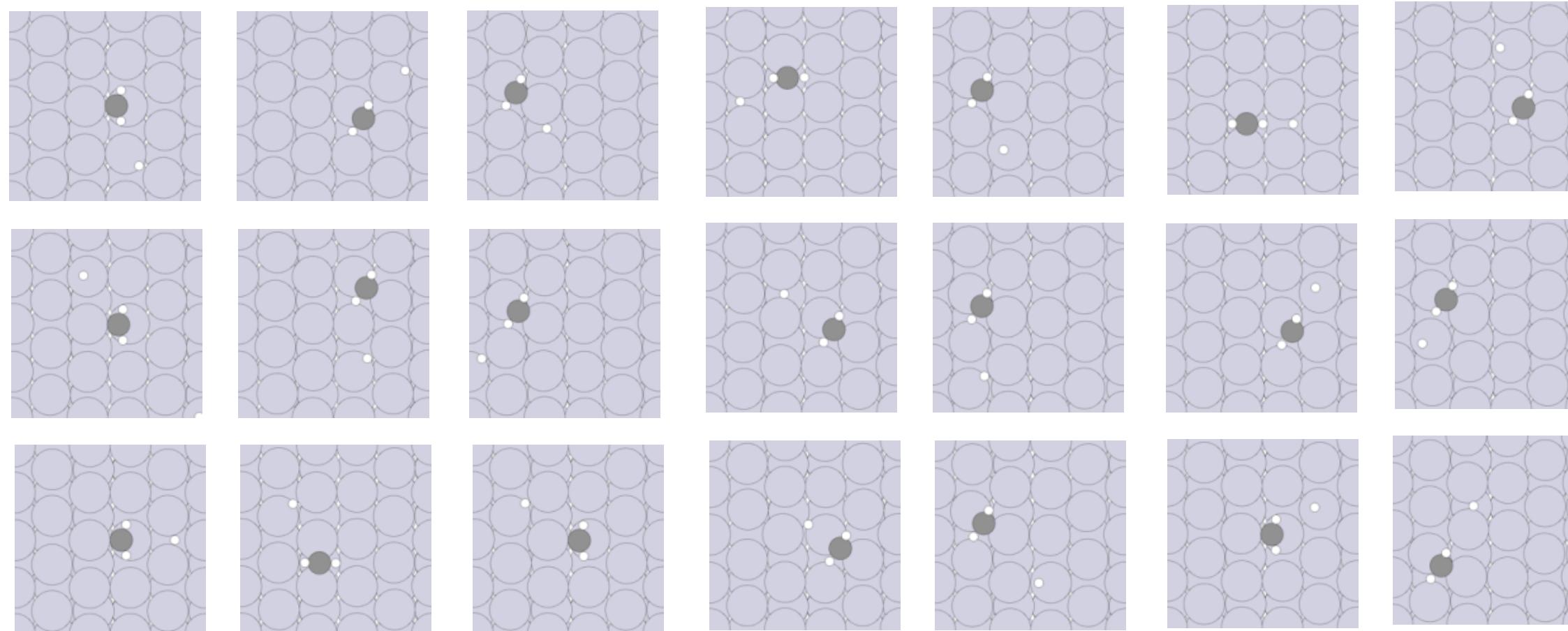


Thermochemistry can be strongly dependent on the choice of site

Pt(111): $\Delta_r E: 0.267 +/- 0.053$ eV

Pt(211): $\Delta_r E: 0.064 +/- 0.415$ eV

$\text{CH}_2 + \text{H}$ co-adsorbed minima on Pt(211)



...There are **73** total minima!

It is difficult or impossible to know a priori which minima are directly connected to the reaction of interest, and which involve diffusion of adsorbates in addition to reaction.

→ Can we still do a systematic search at a cost lower than a full NEB search?

We started developing *pynta* <https://github.com/zadorlab/pynta>
("pinta" means surface in Finnish)

Initial workflow for dissociation/association reactions

1. Prepare surface → Saved and reused

- Size
- Symbol
- Lattice constant
- Vacuum
- Type
- Level of theory

2. Place reactant and product adsorbate(s) extracted from RMG notation on all unique sites

→ Optimize, filter

→ Calculate vibrational frequencies

→ This step provides thermo information

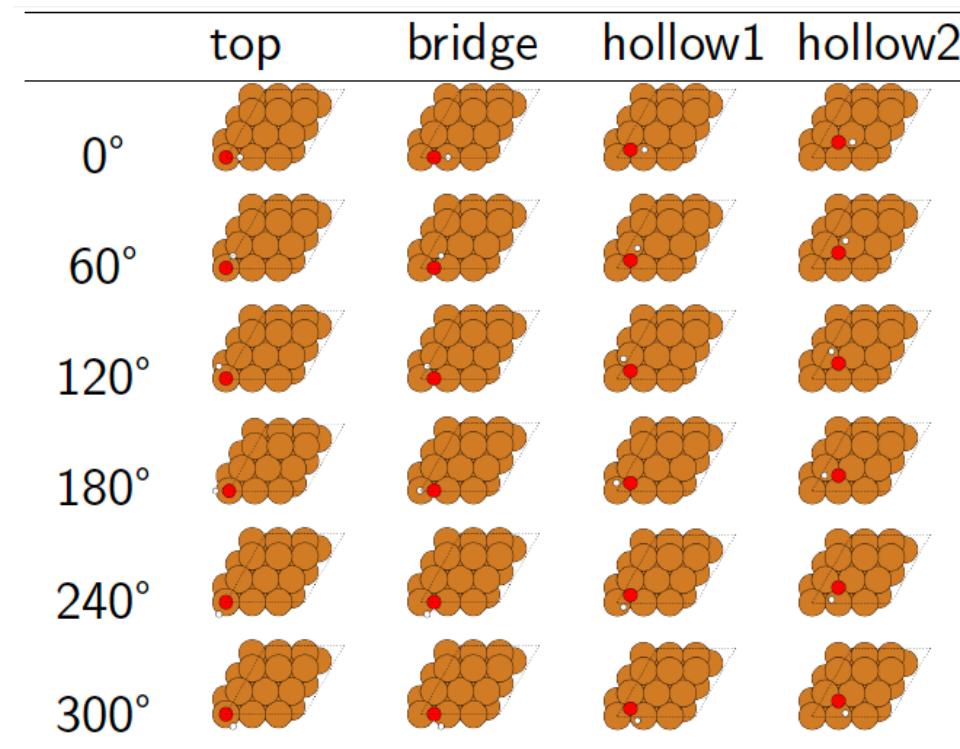
- Lowest energy structures
- Aids more accurate partition function calculations



Initial workflow for dissociation/association reactions

3. Create saddle point guesses

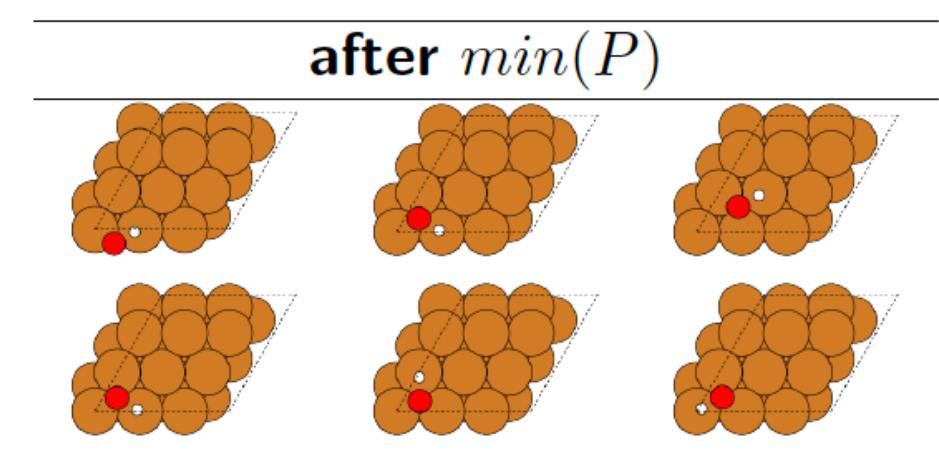
- Place unimolecular species (AB) on all sites and create rotamers
- Filter



- Stretch bond to 1.4x and apply penalty function, surface is not relaxed

$$P = (r_{A-X} - \overline{r_{A-X}})^2 + (r_{B-X} - \overline{r_{B-X}})^2 + E_{xTB}$$

- Filter



Initial workflow for dissociation/association reactions

4. Optimize to saddle points with all constraints lifted at DFT level

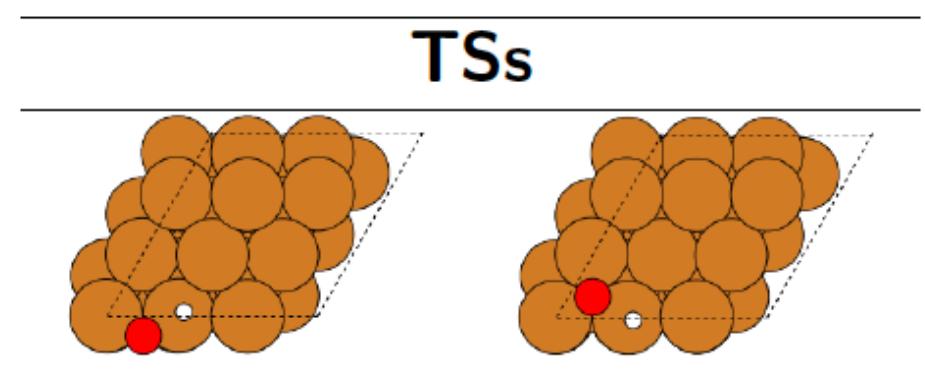
- Geometry optimization is done with **Sella**, our own optimizer (sella means saddle in Latin)

<https://github.com/zadorlab/sella>

E. D. Hermes, K. Sargsyan, H. N. Najm, JZ: JCTC. 2019, 15 6536-6549.

E. D. Hermes, K. Sargsyan, H. N. Najm, JZ: JCP, 2021, 155 094105.

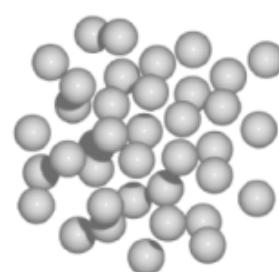
E. D. Hermes, K. Sargsyan, H. N. Najm, JZ: JCTC, 2022, submitted.



Sella has superior performance for saddle points optimizations for solids and molecular systems, and has featureful optimizations with constraints, and can carry out IRC as well.

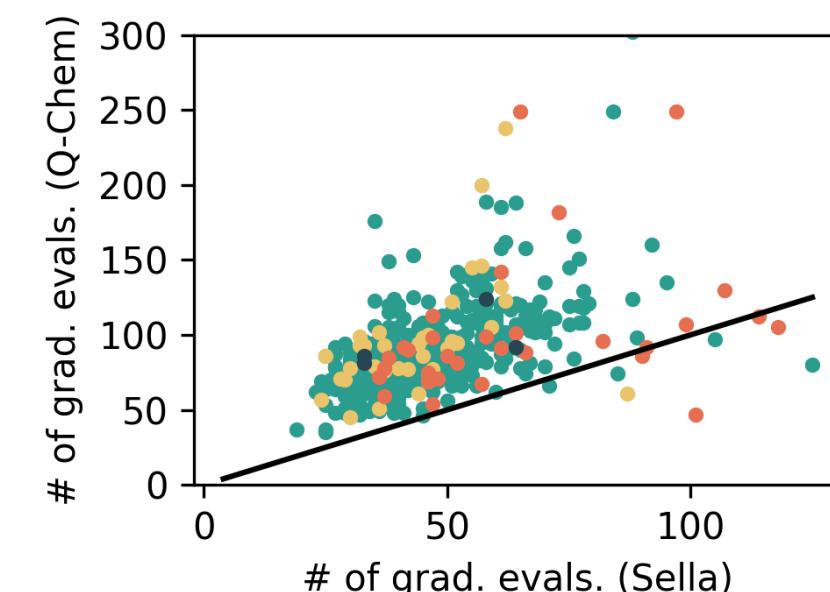
Example comparison from our own molecular saddle benchmark

<https://www.optbench.org>



Algorithm	<FCs>	min(FCs)	median(FCs)	max(FCs)	nfailed
Sella	81	37	75	166	0
optim_ffd	145	57	119	565	0
pele_ffd	192	59	150	1488	0
eon-lanczos-improved ¹	237	65	181	1898	0
eon-lanczos ¹	321	83	220	8857	0
eon-dimer ¹	528	92	388	3581	0
Show/Hide Additional Entries...					

Algorithm	<FCs>	min(FCs)	median(FCs)	max(FCs)	nfailed
Sella	49	32	45	98	0
optim_ffd	71	43	70	143	0
pele	88	52	78	198	0
optim_cd	90	52	86	199	0
eon-lanczos-improved ¹	106	71	100	163	0
eon-dimer ¹	116	83	114	160	0
eon-lanczos ¹	143	106	139	292	0
Show/Hide Additional Entries...					



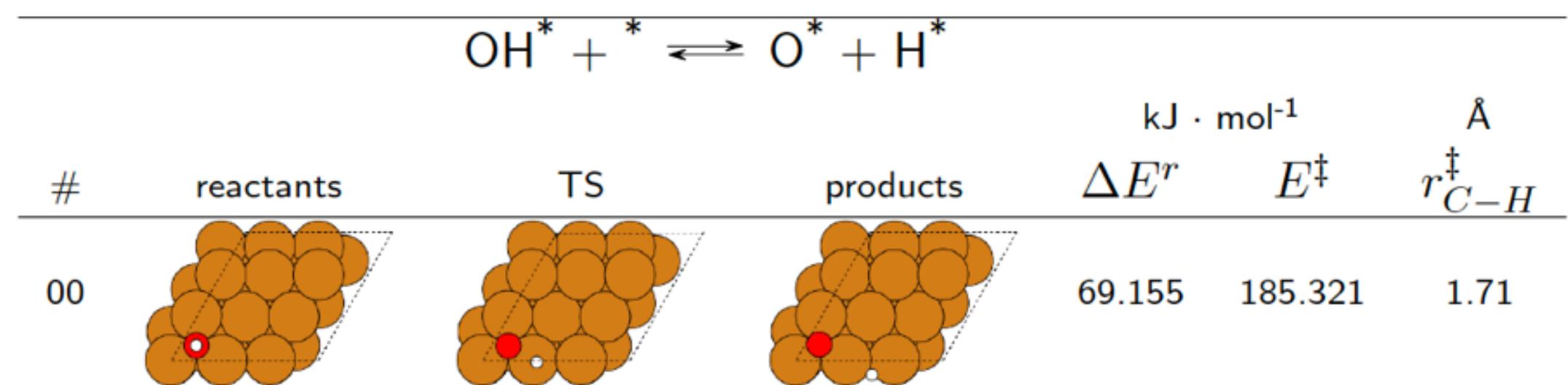
Initial workflow for dissociation/association reactions

5. Verification

- IRC
- Frequency calculations

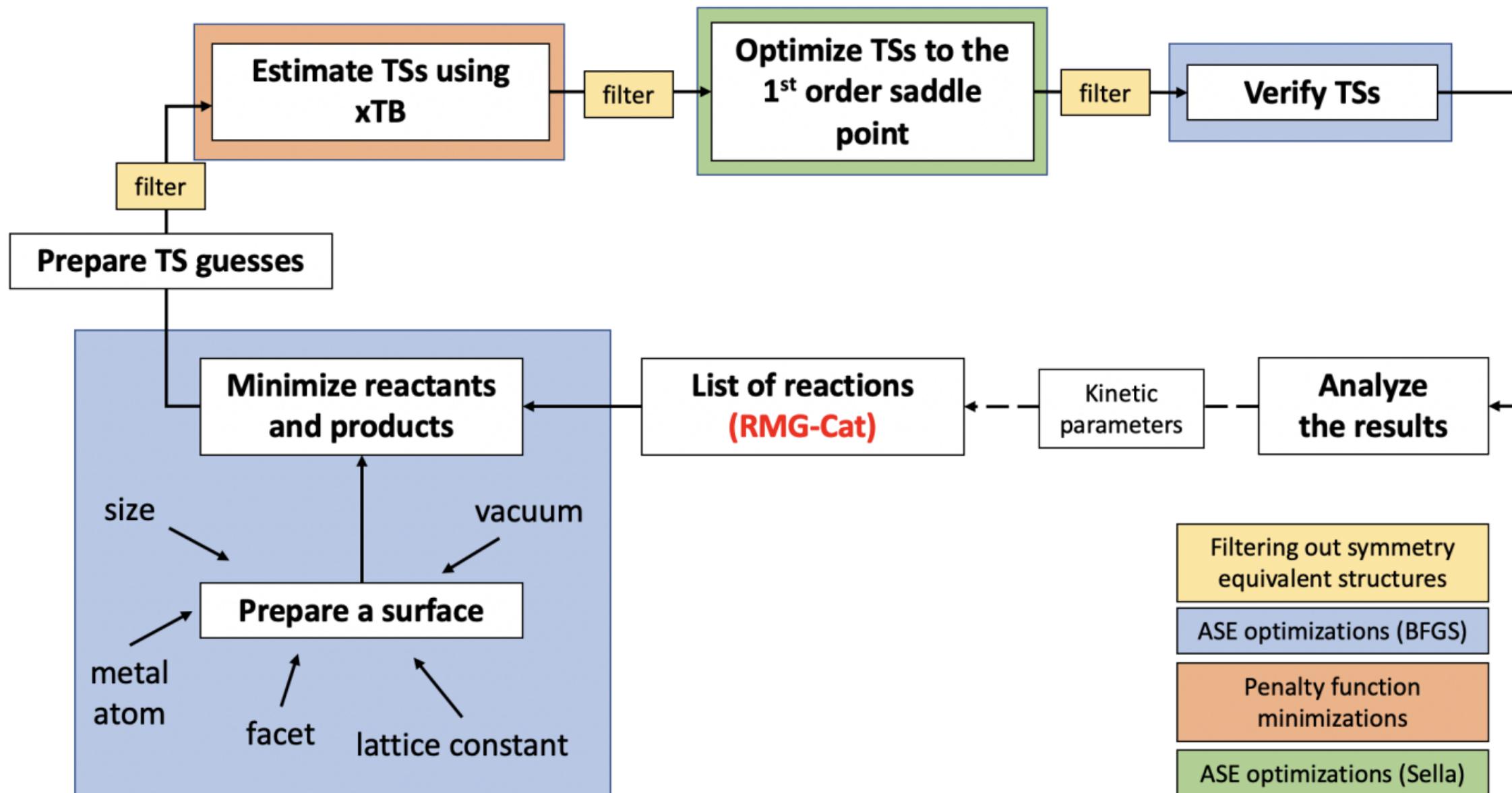
6. Analysis

- Reaction barrier
- Reaction enthalpy
- Geometry



7. Passing data to rate coefficient calculator

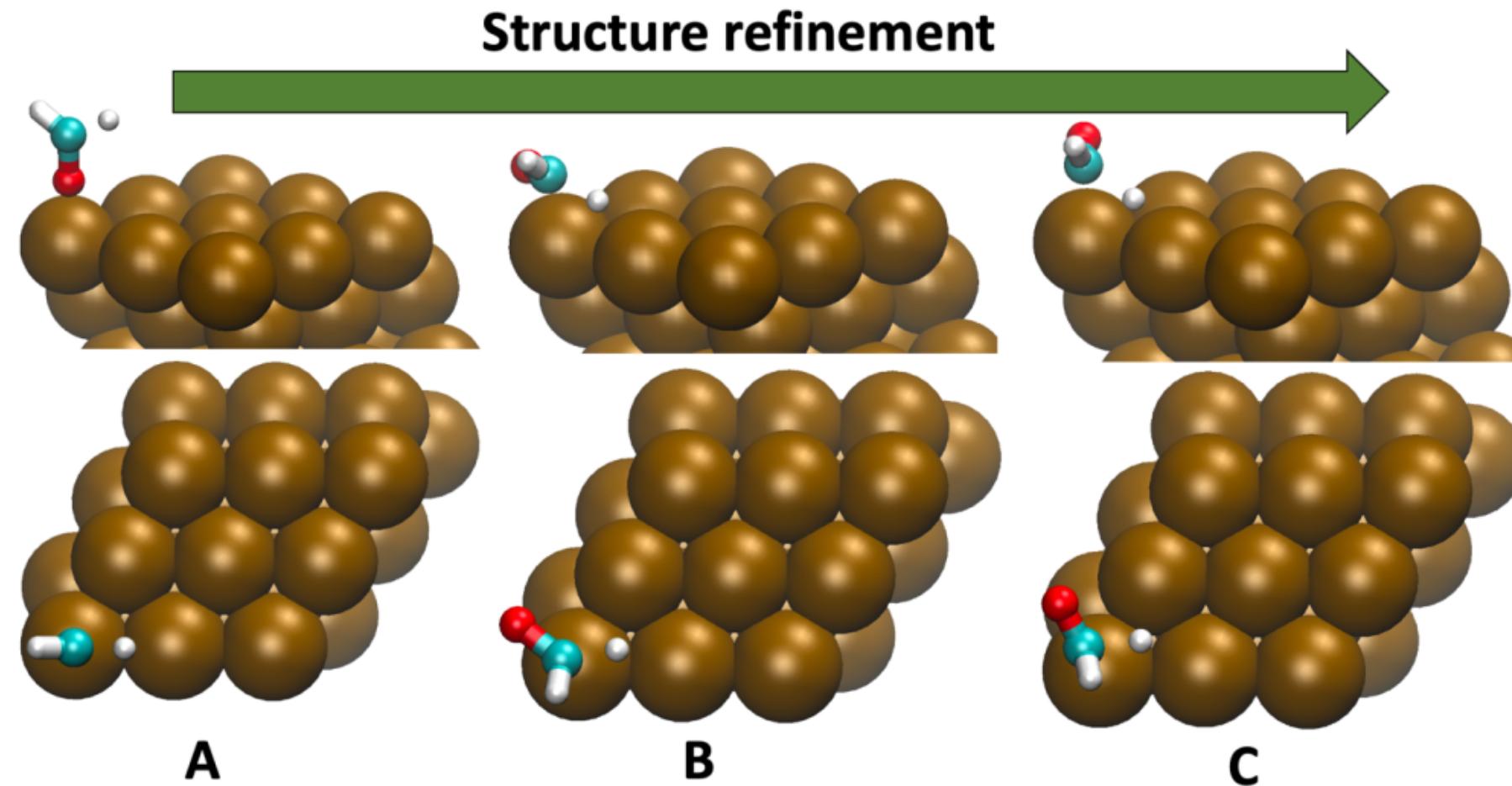
Summary and computational aspects



Important dependencies

- Balsam
- ASE
- Sella
- xTB
- Catkit

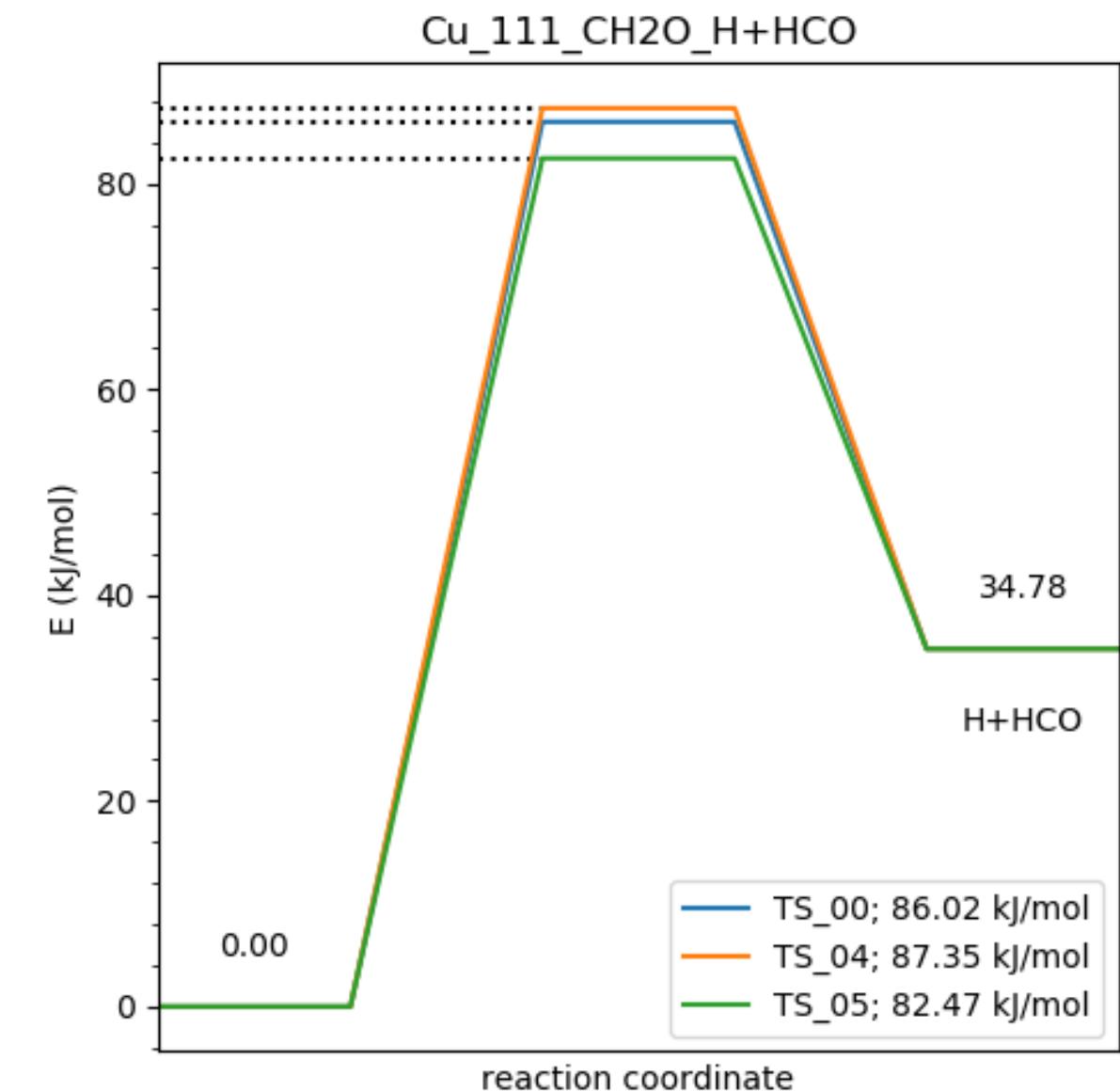
$\text{CH}_2\text{O} \rightarrow \text{CHO} + \text{H}$ on Cu(111)



“raw” guess
4 sites
292 initial structures

penalty fn optimization
42 unique starting geoms
10 final ones

final saddle point
3 unique saddles

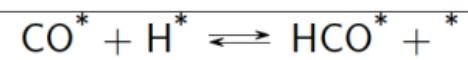


PBE functional results
Literature lists only one saddle

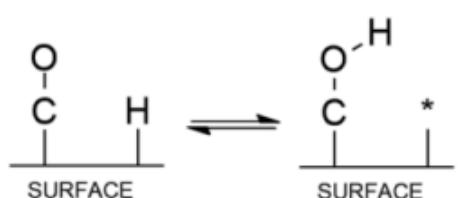
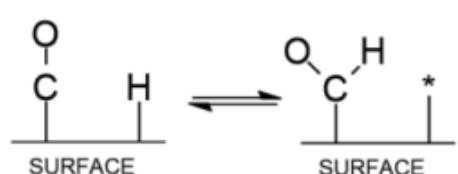
Further examples



#	reactants	TS	products	ΔE^r kJ · mol ⁻¹	E^\ddagger	$r_{\text{C}-\text{H}}^\ddagger$ Å
00				132.836	183.297	1.82
01				117.379	197.932	2.00



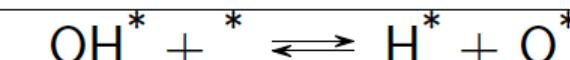
#	reactants	TS	products	ΔE^r kJ · mol ⁻¹	E^\ddagger	$r_{\text{X}-\text{H}}^\ddagger$ Å
00				93.809	103.403	1.52
01				123.530	223.419	1.28
02				86.842	96.120	1.41



SURFACE



#	reactants	TS	products	ΔE^r	E^\ddagger	$r_{\text{C}-\text{O}}^\ddagger$
00				-111.682	50.449	1.77
01				-118.958	46.756	1.82



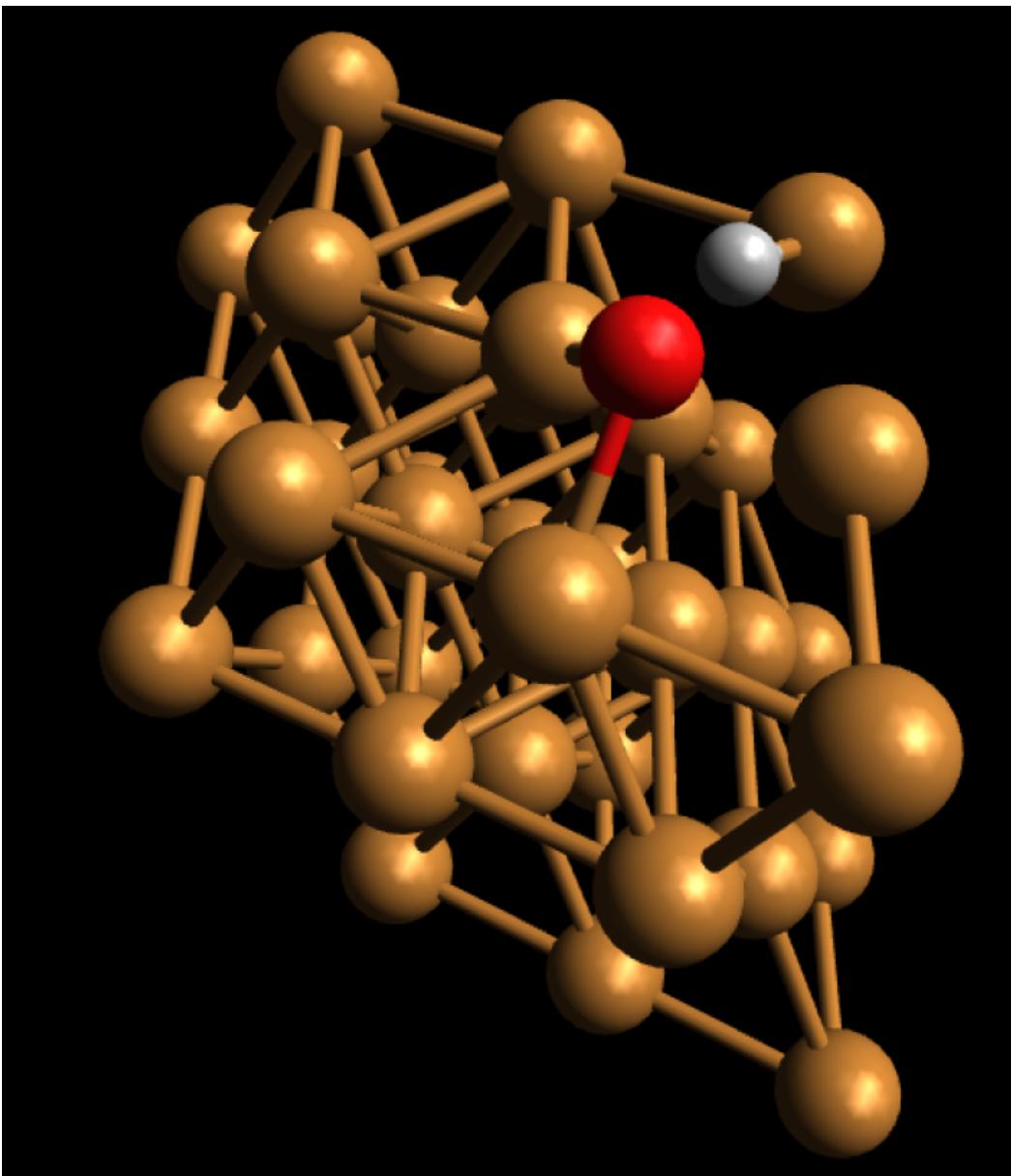
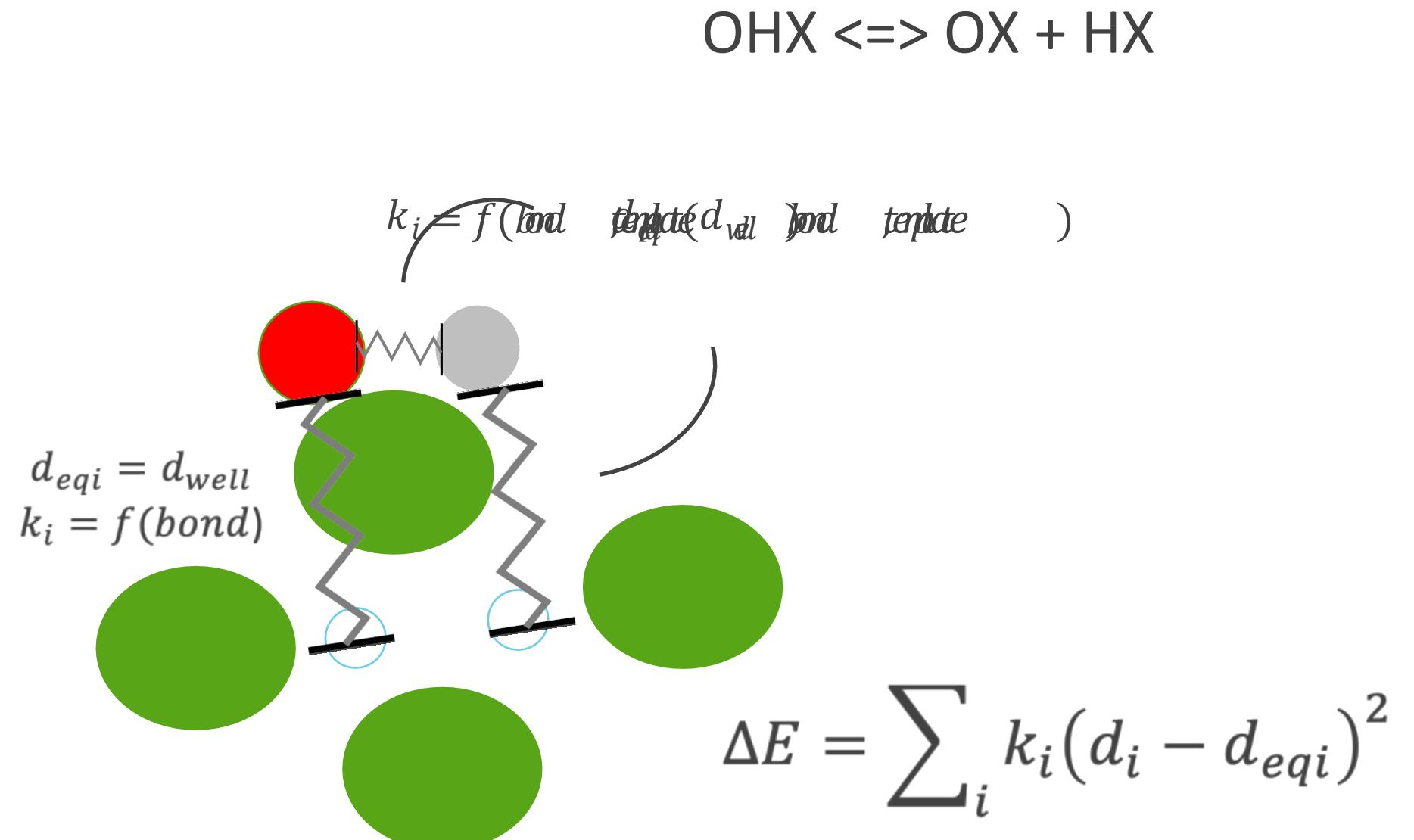
#	reactants	TS	products	ΔE^r kJ · mol ⁻¹	E^\ddagger	$r_{\text{X}-\text{H}}^\ddagger$ Å
03				69.598	157.963	1.57
05				95.090	148.133	1.69
07				63.890	150.247	1.55
08				114.593	200.130	1.64
11				23.954	107.021	1.53

In Progress and Future Plans

- Integrate Pynta with the Fireworks workflow management software to
 - Operate efficiently on larger clusters
 - Improve restart functionality
 - Enable automatic workflow debugging
 - Make it easy to adapt the workflow as needed
- Integrate IRC calculations using Sella for TS verification
- Generalize TS estimation to most reaction templates of interest
 - Analyze reaction templates using RMG and the optimized reactant and product adsorbates using ASE
 - Use this information to generate a modified xTB potential that will optimize starting structures to good transition state guesses for the reaction of interest
- Integrate database for storing useful generated information from Pynta
 - Enable reuse of prior calculations
 - Make results accessible to scientific community

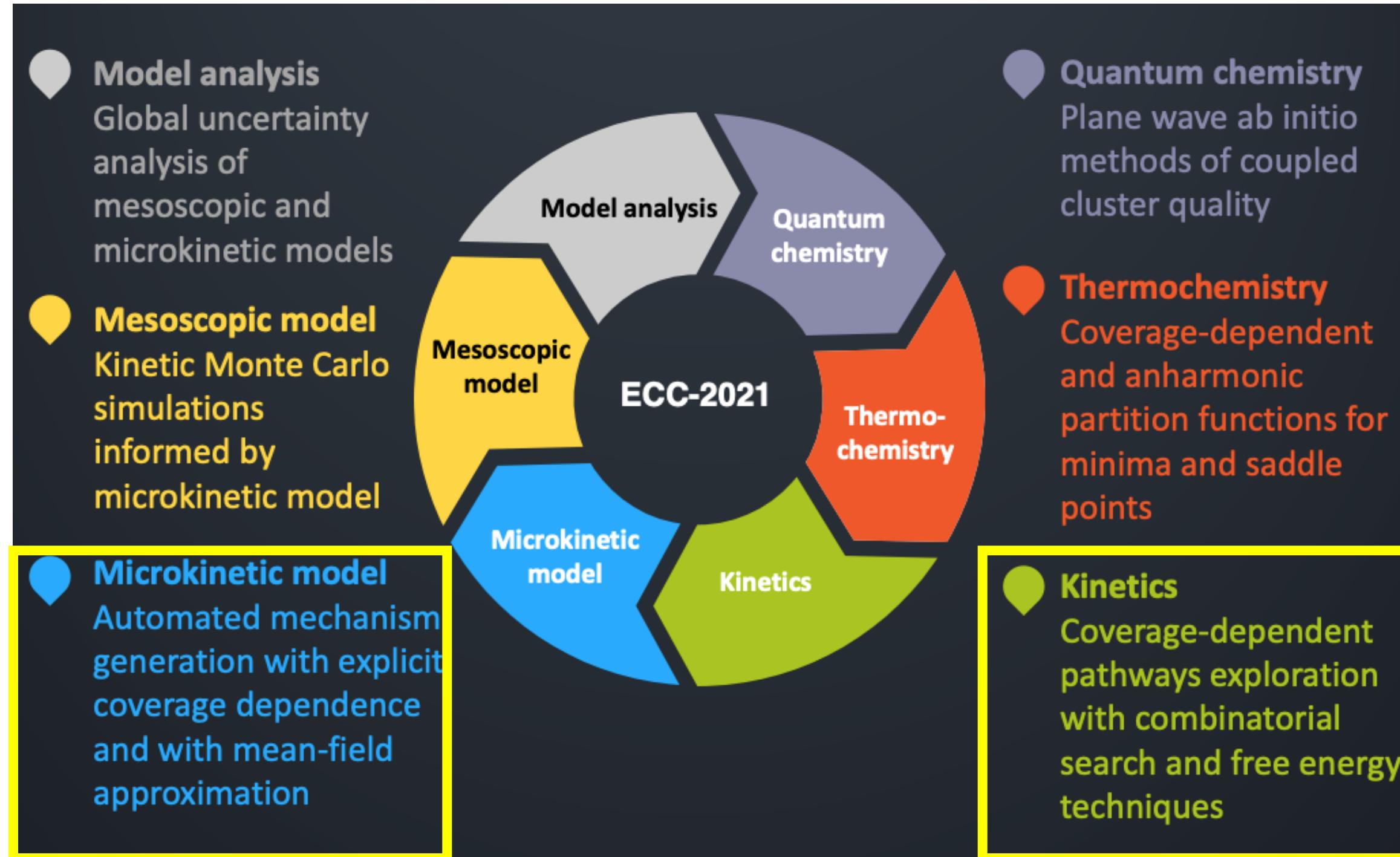


Harmonically Forced TS Searching



Where d_{eq} is the equilibrium bond length, k_i is the spring constant of the bond, d_{well} is the bond length in the associated adsorbed structure and d_i is the bond length.

pynta is part of the Exascale Catalytic Chemistry Project

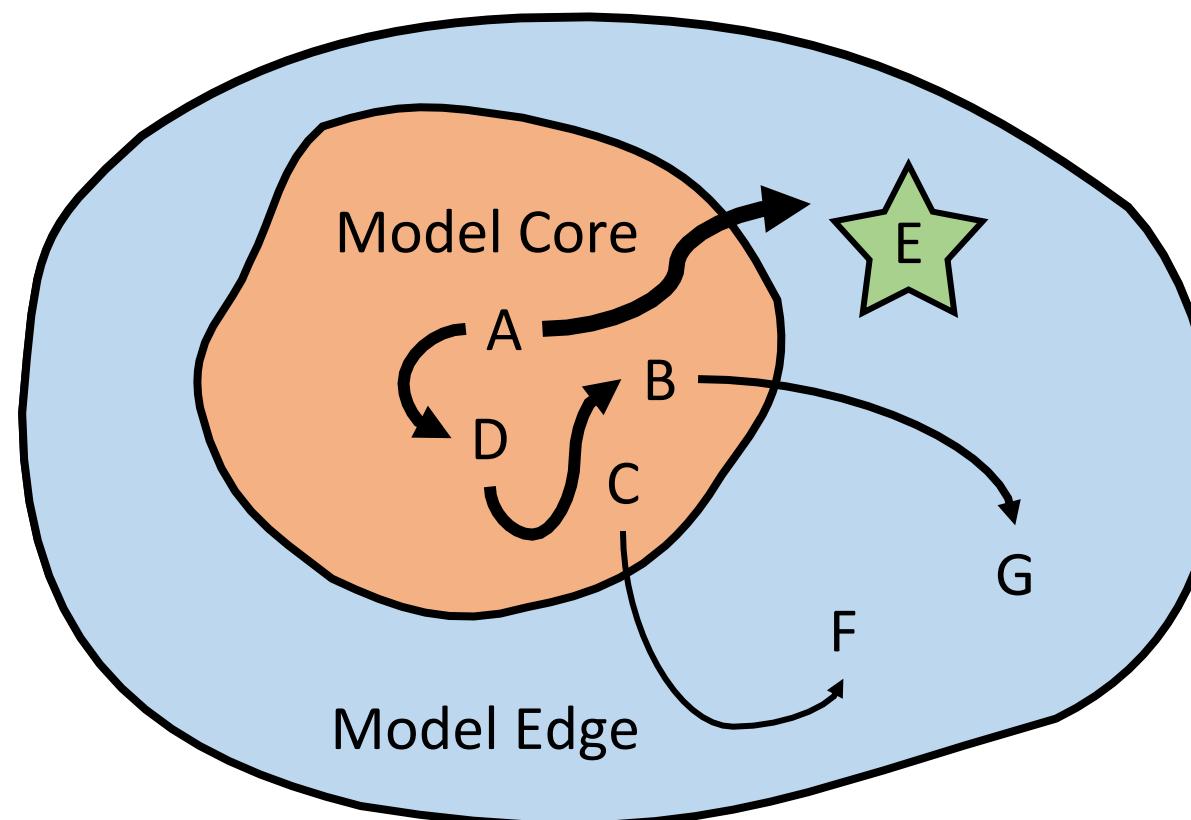


<https://www.ecc-project.org>

RMG can generate microkinetic models for catalytic processes

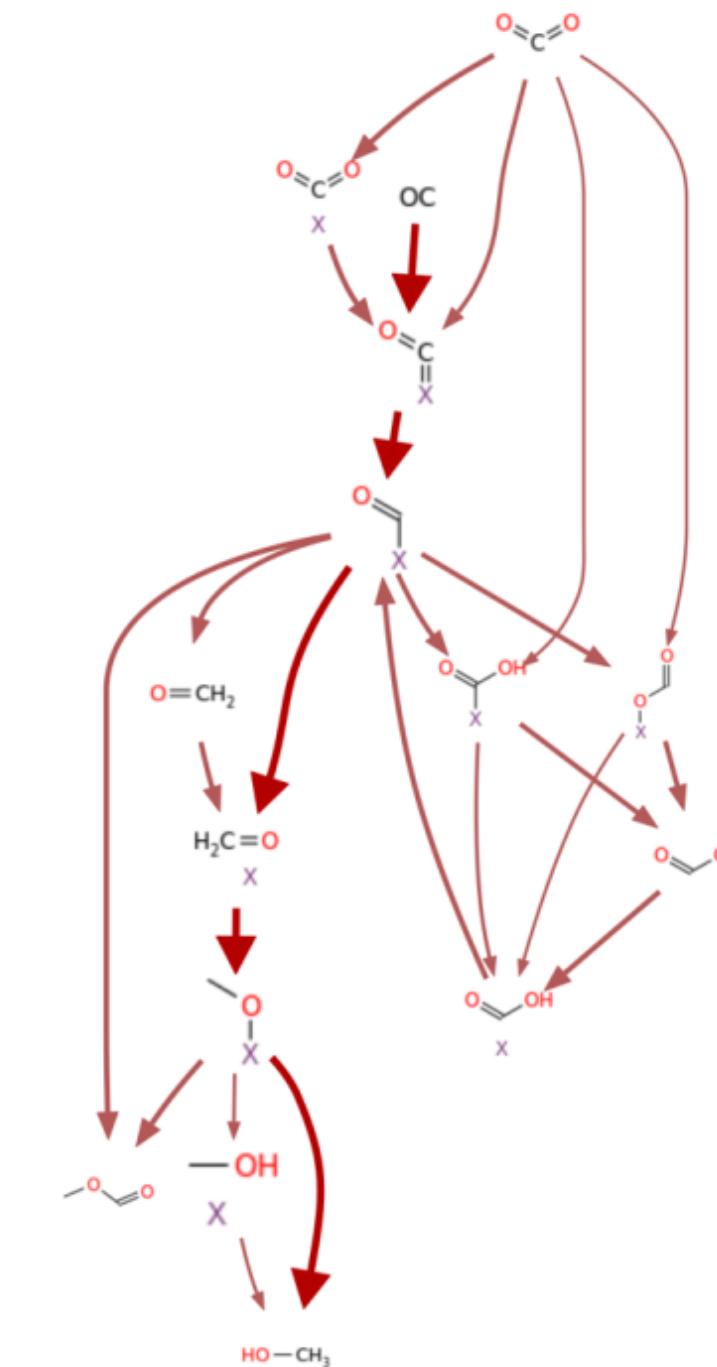
rmq

<https://github.com/ReactionMechanismGenerator/RMG-Py>



Methanol synthesis on Cu(111)

- Well studied mechanism
- Data exists for both experimental (Graaf) and theoretical work (Grabow)
- Good proof of concept mechanism for RMG

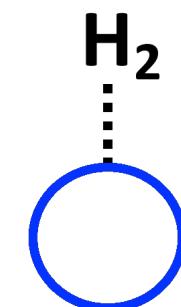


L. C. Grabow, *ACS Catal.* 2011, 1, 4, 365–384.

G. H. Graaf, *Chem. Eng. Sci.* 1988, 43, 3185.

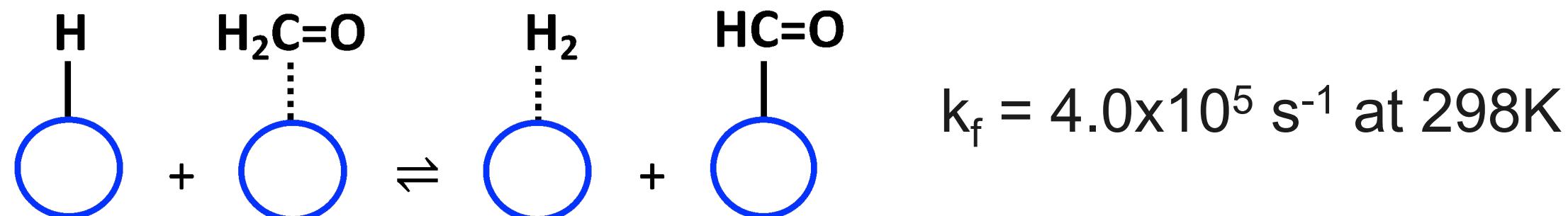
van der Waals (vdW) bonded H₂ was choking the surface

Literature data (Álvarez-Falcón) was incorporated for the binding of H₂ on Cu(111). RMG was estimating the thermo from Pt(111). Unexpectedly, we were underestimating the binding energy.

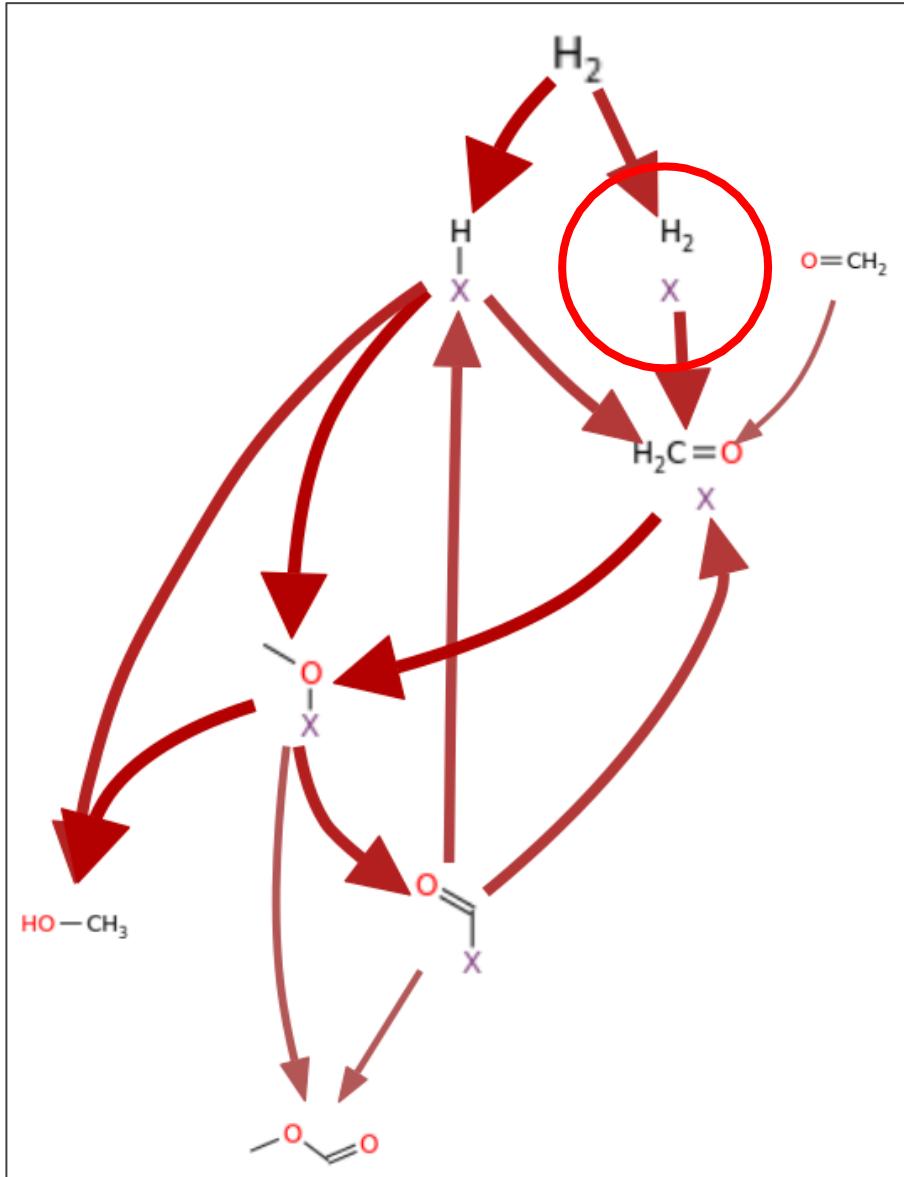


DFT BE on Pt(111): -0.054 eV.
DFT BE on Cu(111): -0.08 eV.

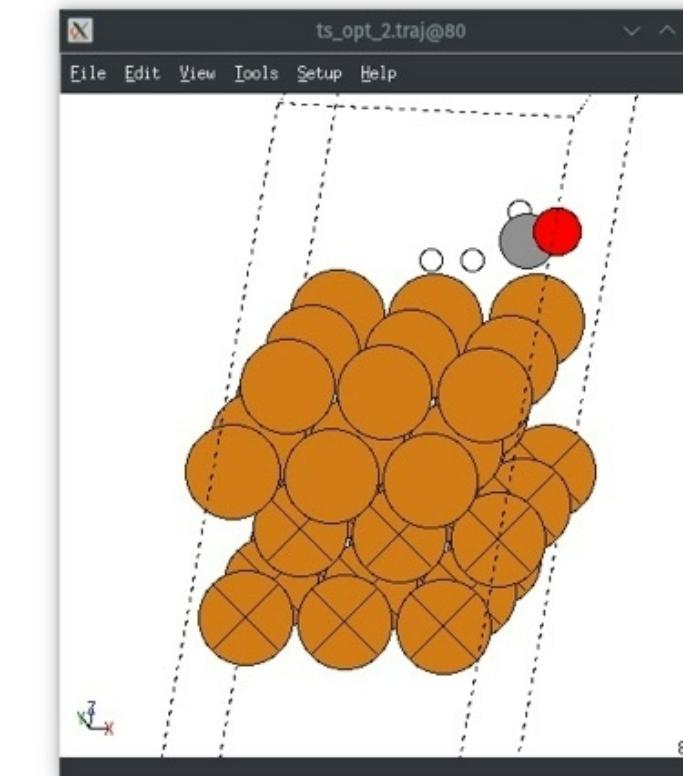
Sensitivity analysis revealed that the most sensitive reaction in the mechanism involved this H₂ species:



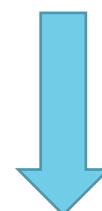
A saddle point search in pynta revealed the reaction was possible but very slow.



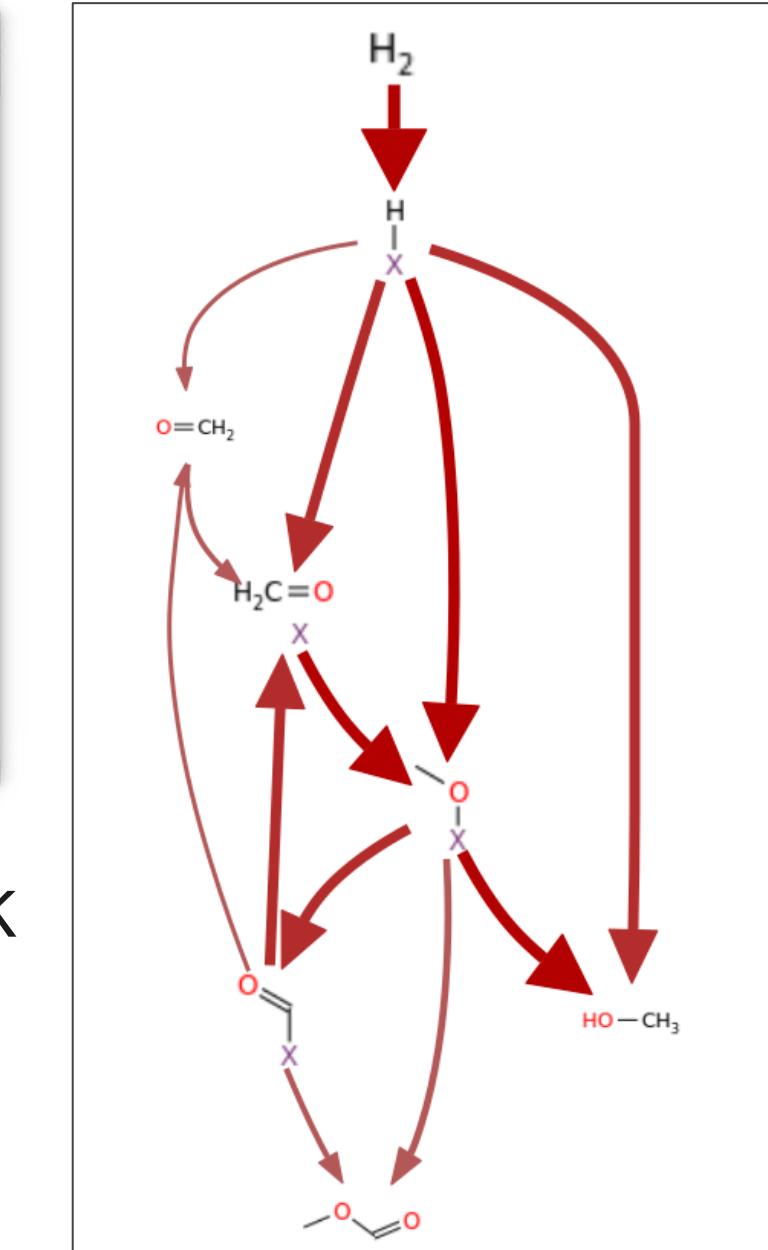
Model With Estimated Kinetics



$$k_f = 4.0 \times 10^5 \text{ s}^{-1} \text{ at 298K}$$



$$k_f = 6.5 \times 10^{-14} \text{ s}^{-1} \text{ at 298K}$$



Model With DFT calculated Kinetics

Further updates of the model:
Reaction enthalpies (kJ/mol)

Cu_111_O+HCO_OH+CO	-143.34
Cu_111_HCOOH_OH+HCO	76.33
Cu_111_H+HCO_CH2O	-9.88
Cu_111_CH3OH_H+CH3O	-12.01
Cu_111_CO2_O+CO	38.83
Cu_111_H+CH2O_CH3O	-90.99
Cu_111_H+CO2_COOH	33.79
Cu_111_HCO+CH2O_CO+CH3O	-184.8
Cu_111_CO2+HCO_CO+COOH	-60.03
Cu_111_H+CO_HCO	93.81
Cu_111_CH2O_H+HCO	9.88

Conclusions

- We are developing an open-source code, `pynta`, to explore surface-reactions automatically.
- The tool is designed to work with RMG, an automated reaction mechanism generator, which provides the reactions in the form of adjacency matrices.
- We demonstrated the viability of our approach in `pynta` through several examples, and outlined some of the immediate improvements that are currently under development.
- We provided an example in which `pynta` has improved a microkinetic model generated by RMG.

