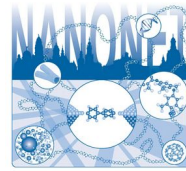




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# Evolution of Molecular Binding in the MCBJ Junctions

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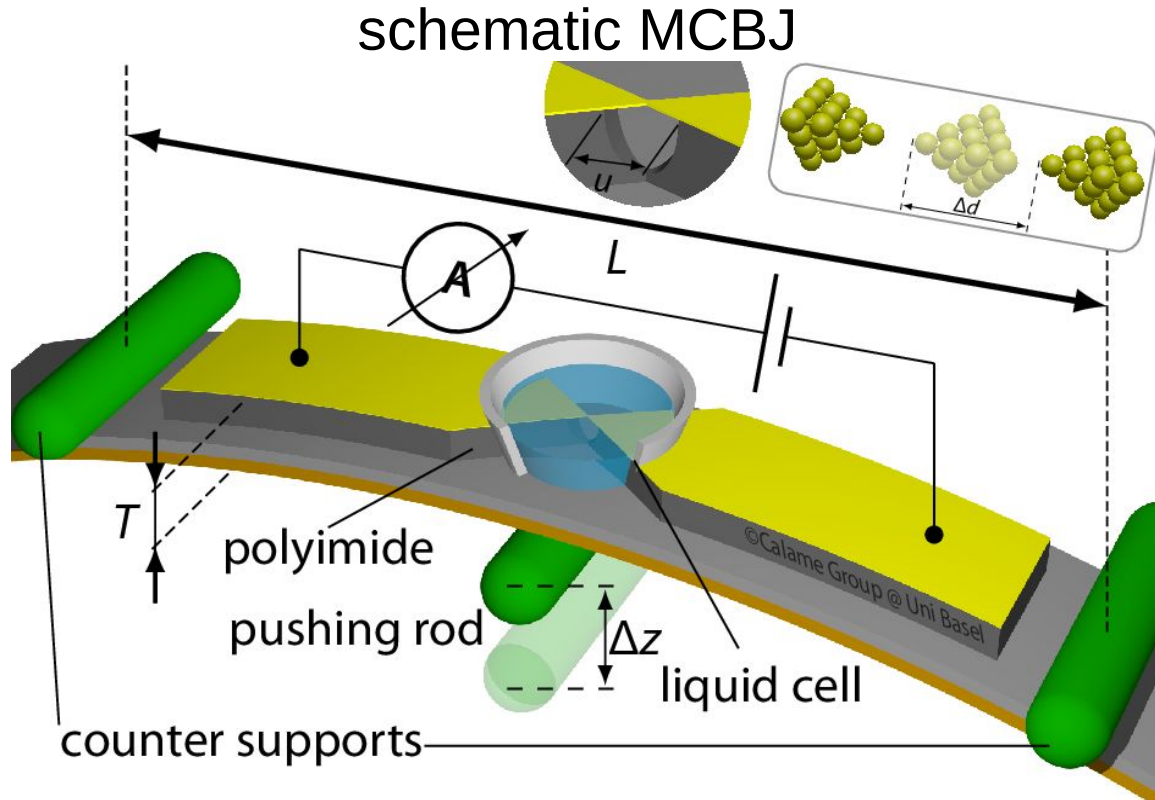


# Introduction

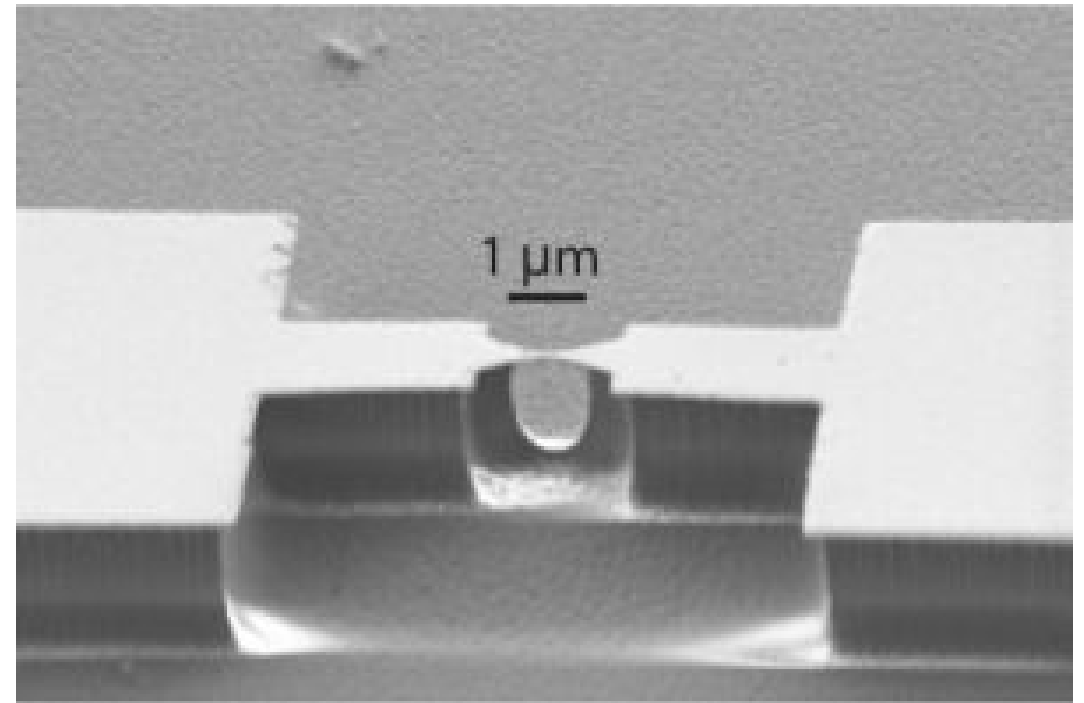
Mechanically controlled Break-Junctions



# Mechanically controlled Break-Junctions

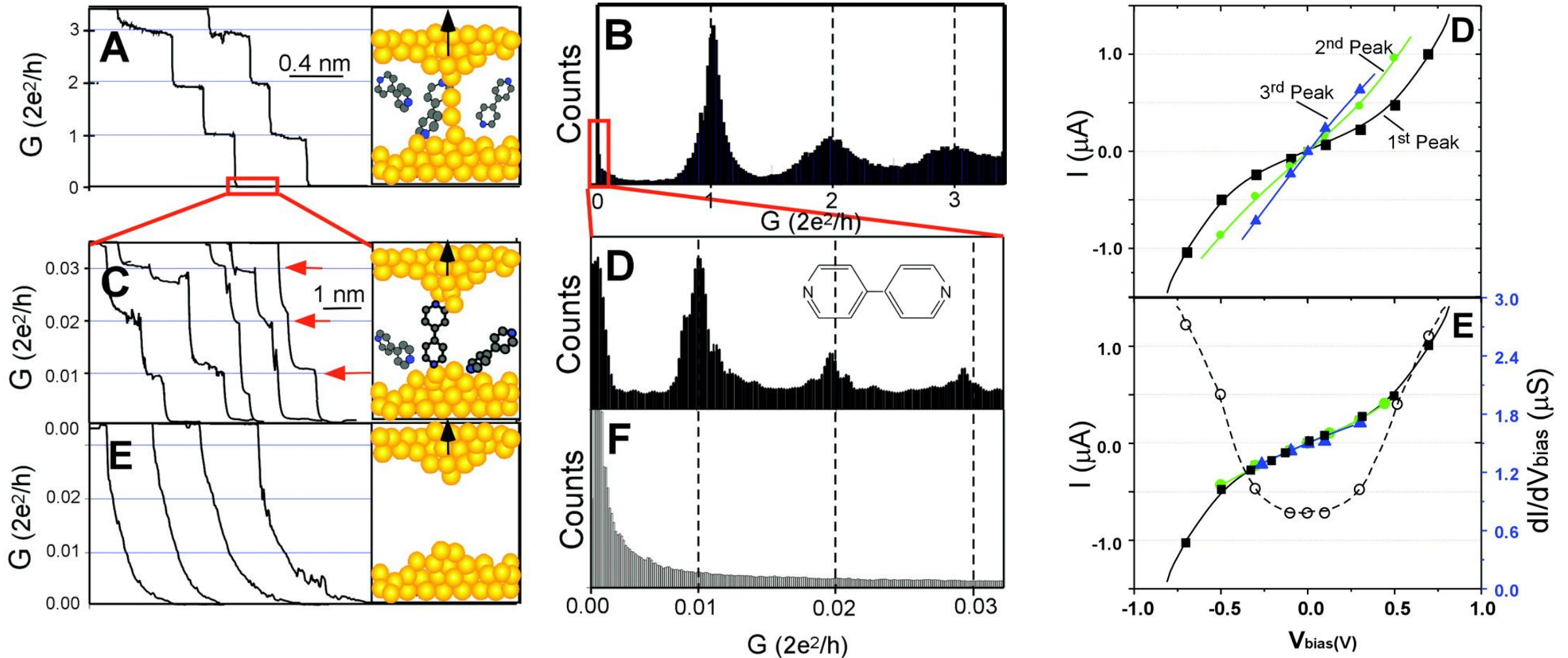


SEM Image



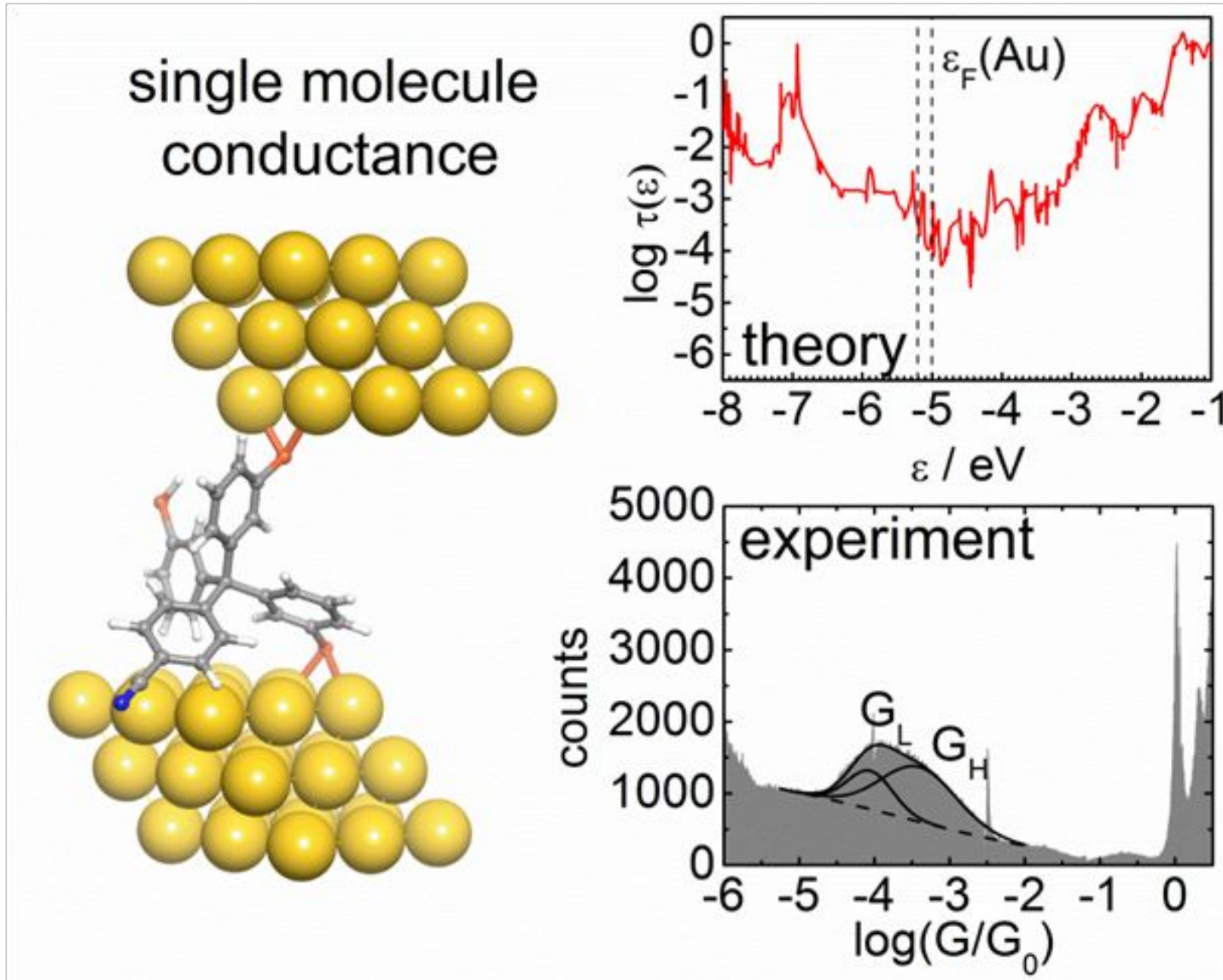
- Au-Au nanoconstriction fabricated using lithographic techniques
- Main principle: Controllably break a metallic wire to form atomistically shaped electrodes
- Probe molecules trapped btw. the sub-nanometer gap
- Attenuation factor?

# Electrical signature of single molecules



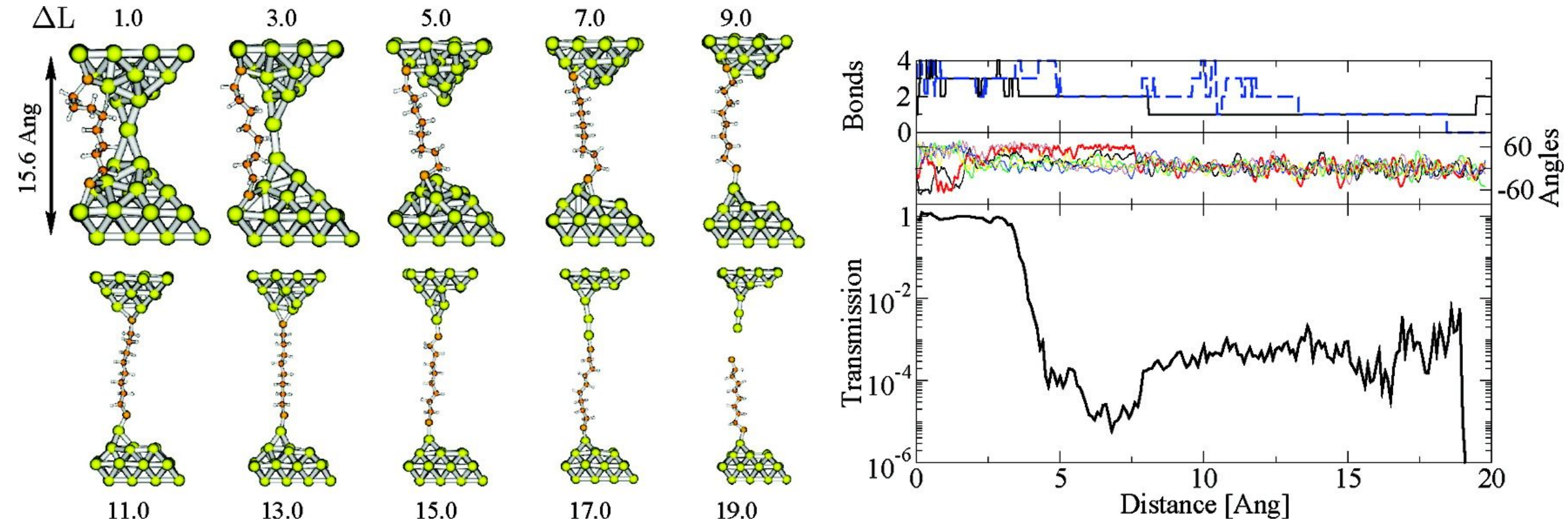
- Metallic bridges  $\triangleright$  flat conductance plateaus | molecules  $\triangleright$  reclining conductance plateaus
- Metallic bridges  $\triangleright$  linear IV | molecules  $\triangleright$  S-shaped IV  $\triangleright$  *Fit single level model* [ $\epsilon$  |  $\Gamma$ ]
- Typically multiple measurements are performed (statistical significance)

# Electrical signature of single molecules using NEGF + static configurations



- Representative | energetically most favorable configuration
  - Transmission using NEGF to evaluate HOMO/LUMO mediated transport
  - NEGF calculations computationally expensive ◀ Measurements are performed numerous times
- 
- ▶ strongly reduced coverage of the configurational space
  - ▶ Steady state assumption
    - ▶ Tip shape do not change considerably
    - ▶ several measurement filter out statistically insignificant configurations

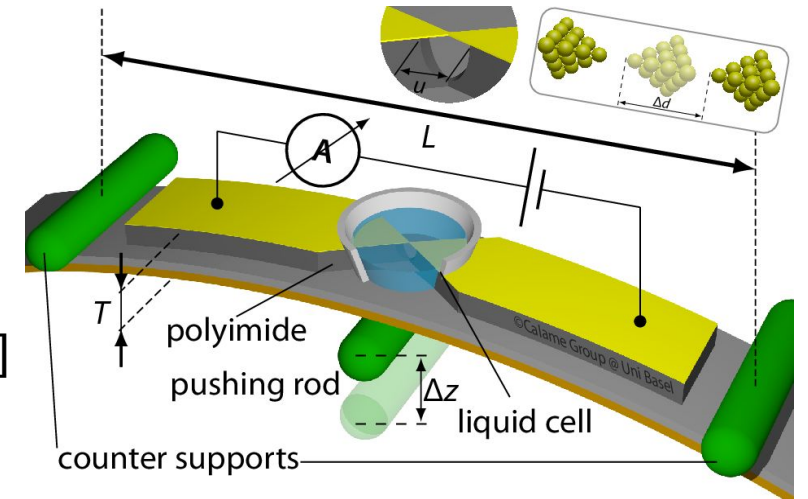
# Electrical signature of single molecules using NEGF + MD Simulations



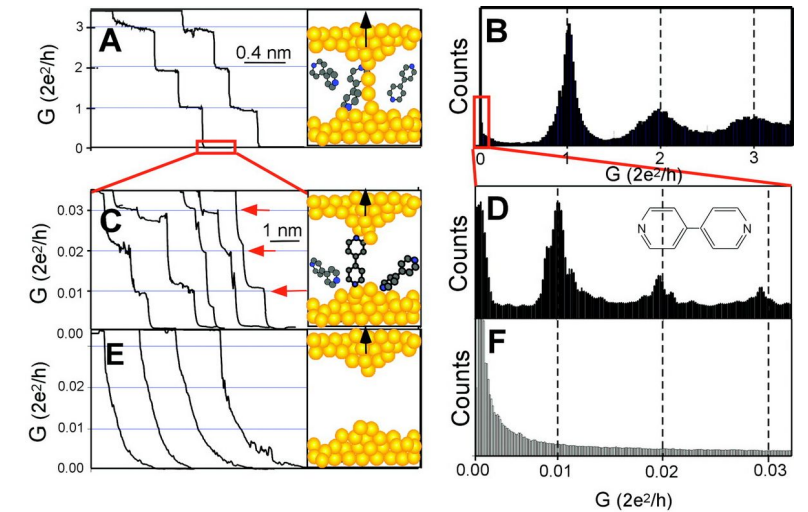
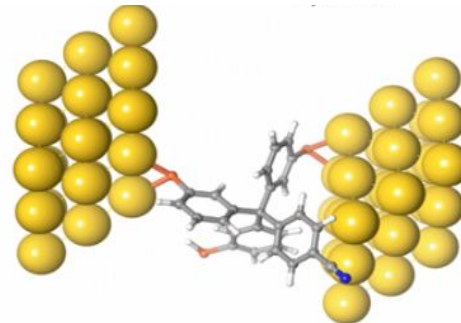
- Representative configuration for each tip-tip separation
- Energetically most favorable configuration | Snapshots from limited trajectories using MD-Simulations
- ▶ Are molecules in steady state during the IV-measurement?
- ▶ Adequate coverage of the configurational space?

# Interpretation MCBJ experiments using NEGF

- Attenuation factor in experiments
- Real tip-tip separation cannot be determined precisely  
[ substrate bending non-uniform | different tip shapes | tip deformation ]
- Statistical measurements



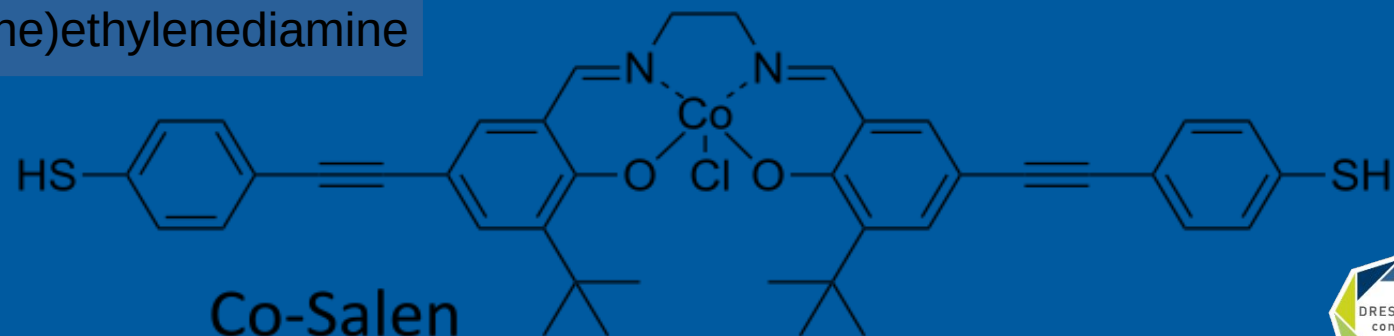
- NEGF calculations computationally expensive
- Strongly reduced configurational space
- Steady state assumption



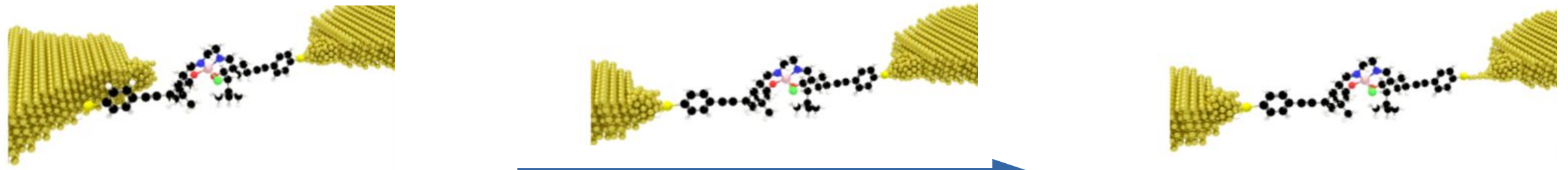
# „Time“ evolution of Molecular binding in MCBJs

Recent experimental results on SALEN

N,N'-Bis(5-ethynylbenzenthioisalicylidene)ethylenediamine







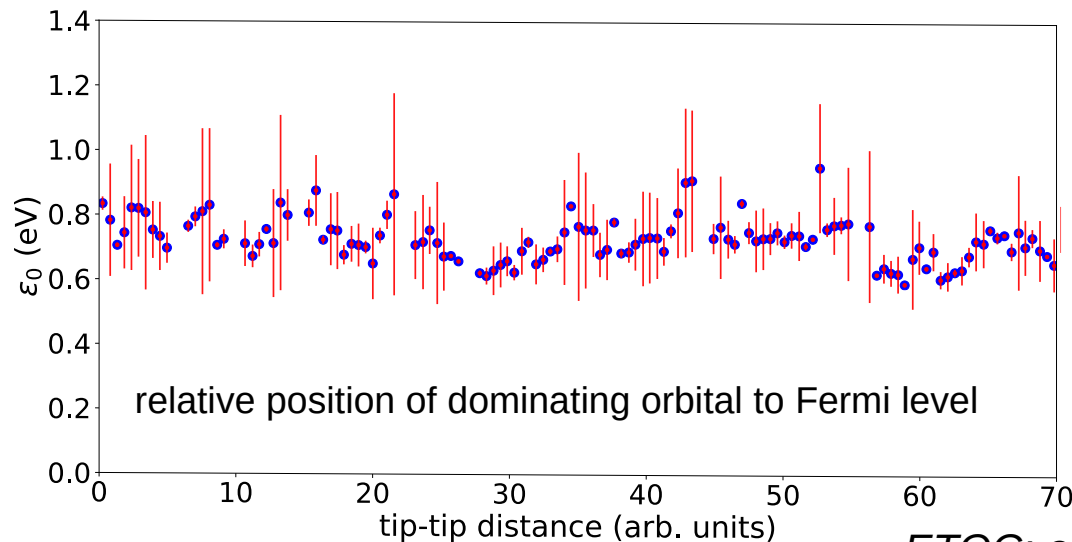
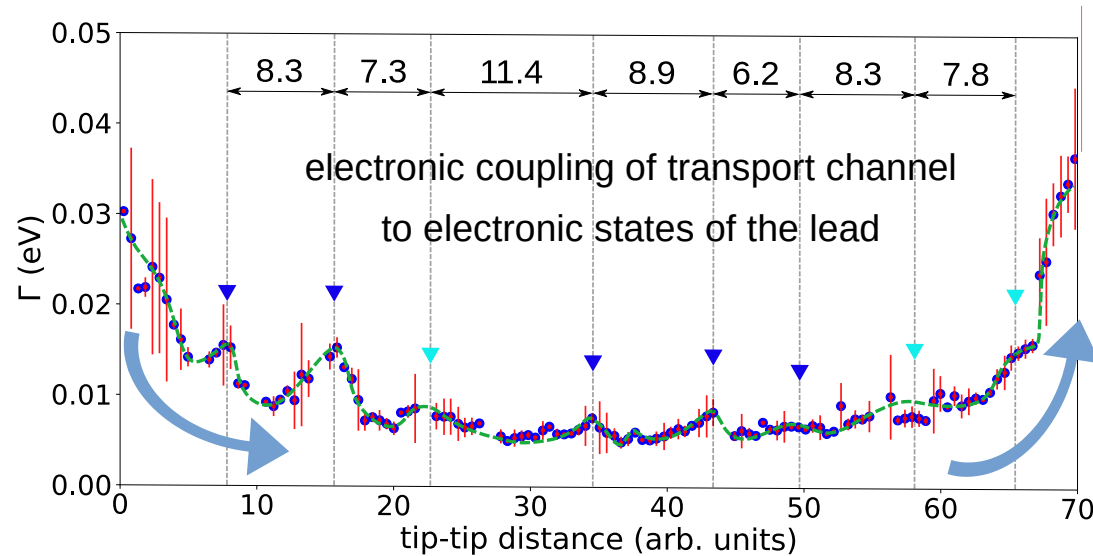
Opening curve measurement MCBJ

## Difference to previous MCBJ experiments

- Focus on single measurements | not statistics
- Follow „time“ evolution of ETCQs at the time scale of measurements

## „Time“ Evolution of ETCQs

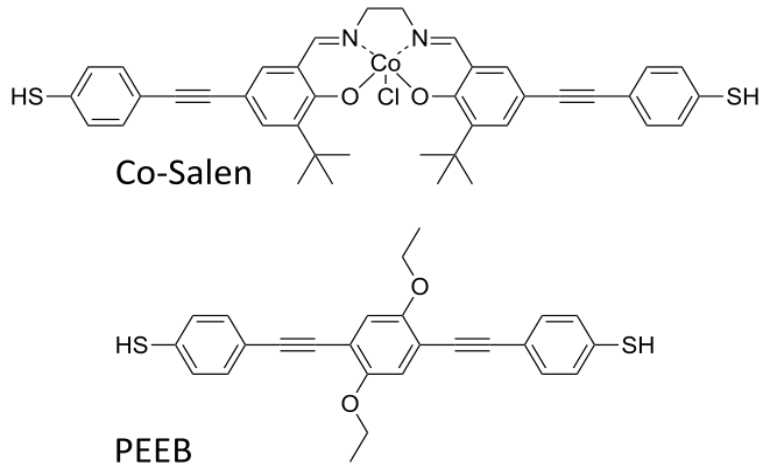
- $\Gamma$  „bathtub“-curve with recurrent peak-features
- $\varepsilon_0$  oscillates in the range [0.6eV , 1.0eV]
- ▶ steady state not necessarily presumable
- ▶ need to include geometrical distortions of the molecule and docking dynamics on gold facets
- ▶ „perform“ statistics using DFT+NEGF



ETQC: *electronic transport characteristic quantities*

**Scheme to follow  
„Time“ evolution ETCQs using DFTB+NEGF**

# „Time“ Evolution of ETCQs ▶ „perform“ statistics using DFT+NEGF



- reduce computational effort ▶ PEEB instead of SALEN
- both bear thiol-anchoring-groups ▶ similar binding to Au
- „Time“-evolution of  $\Gamma$  expected to be similar
- Slakos for Co-Salen unavailable

- local-coordination of gold facets [ fcc ] ▶ numerous docking possibilities
- PEEB flexible ▶ include geometric deformation [ bending | stretching ]



- can PEEB slide on Au-Facet? ▶ ensemble of configuration

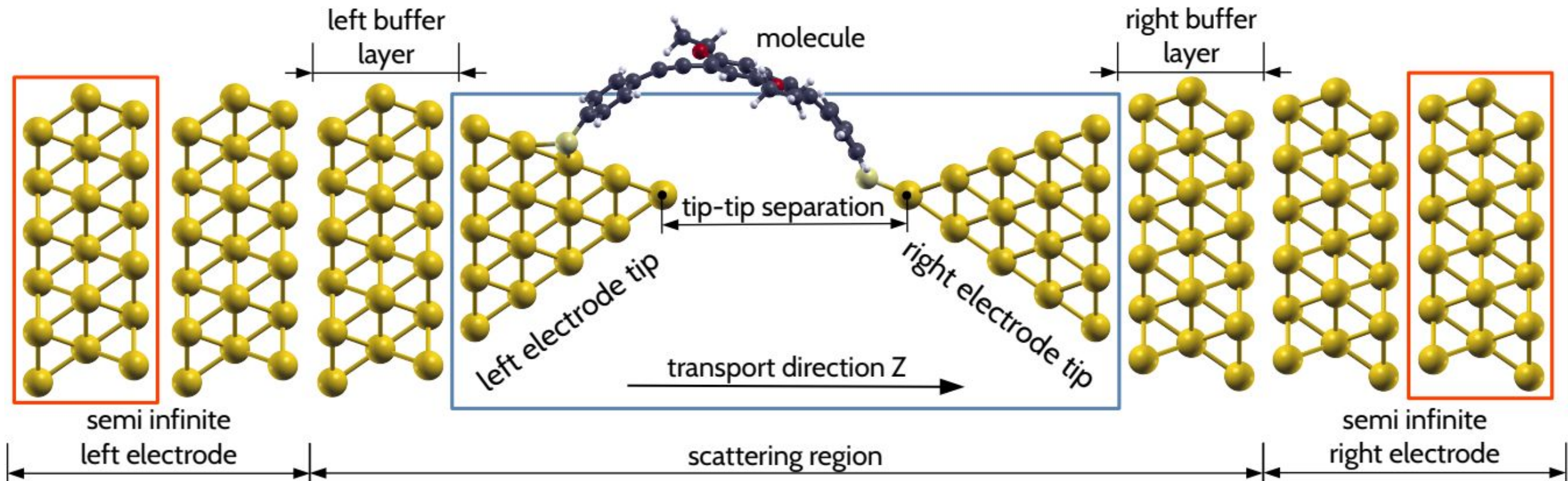


suitable characterization of the configuration ensemble

- ▶ Transport setup | Docking Grid
- ▶ Parse facet docking energy landscape
- ▶ include geometric deformations
- ▶ „Random Walk“ more suitable instead of „normal boltzmann“ distribution
- ▶ incorporates surface dynamics

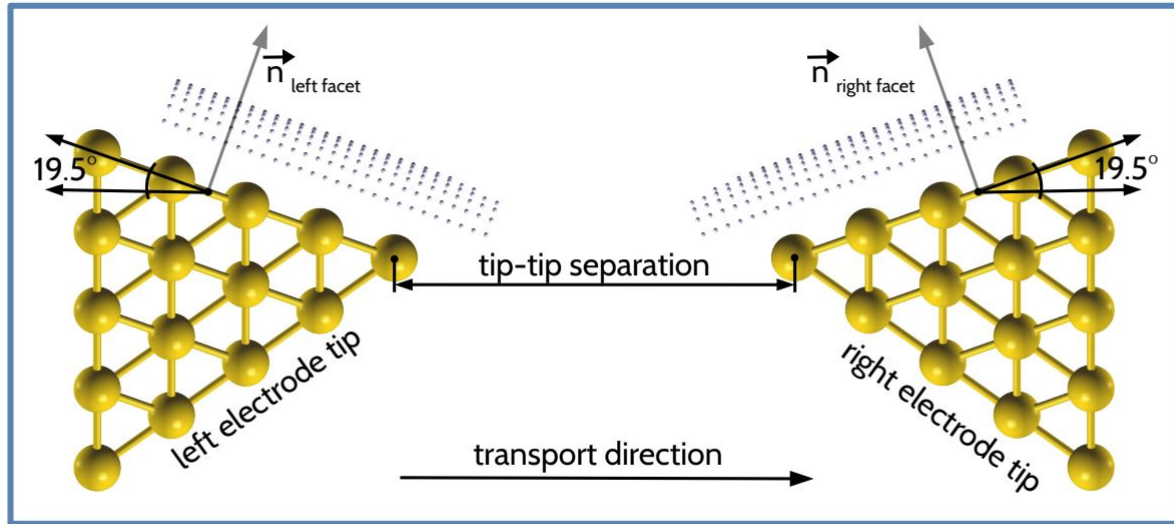
# Transport Setup | Docking Grid | Facet Energy Landscape



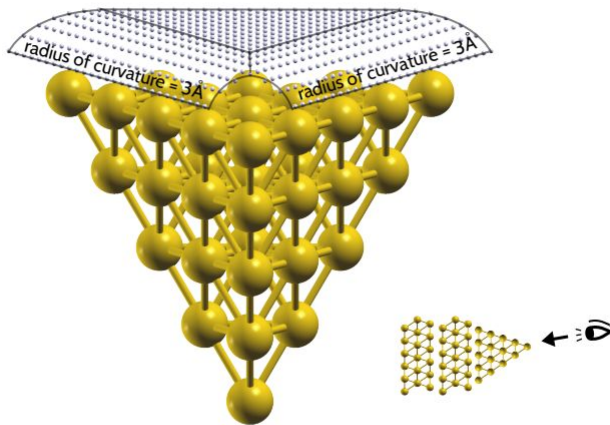


- electrode tips – equilateral triangular pyramids (baselength 5x Au-Au)
- buffer/semi-infinite layers (3 Au Layer 6x6)
- local-coordination-environment ▶ face-centered-cubic
- tip-tip separation [ 11.54Å, 27.24Å ]

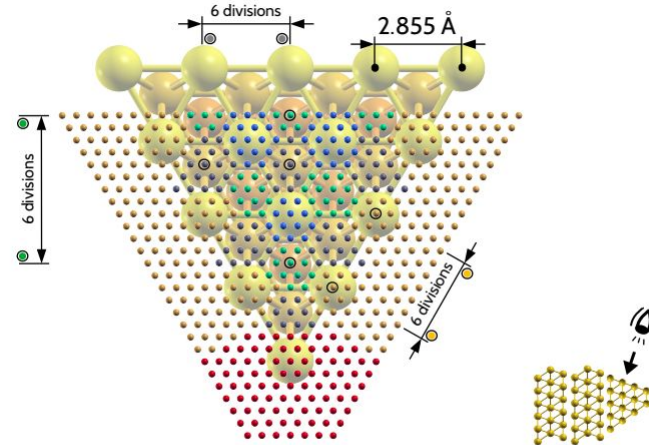
# Docking grid



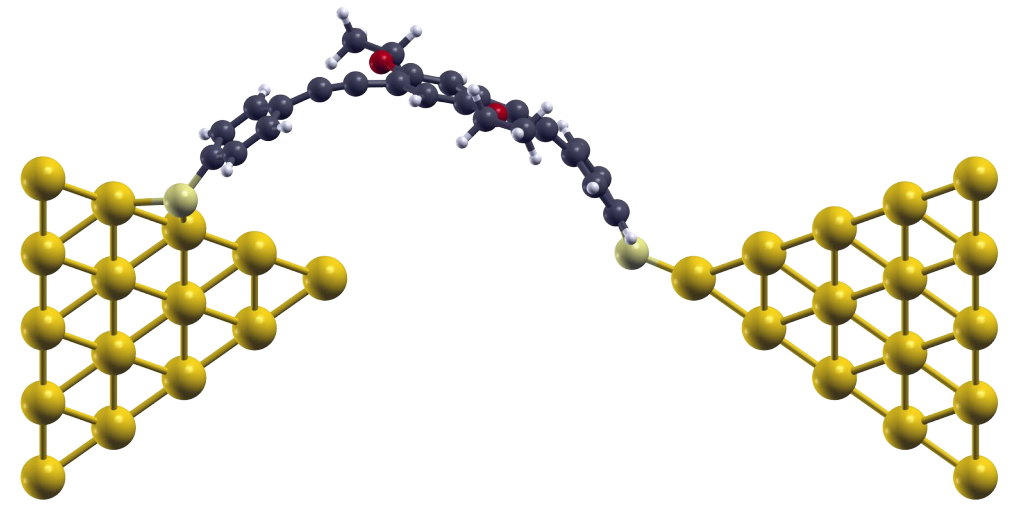
(a) docking grids on left-/right facets



(b) curvature of docking grid about the facet edges

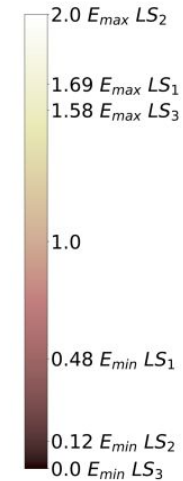
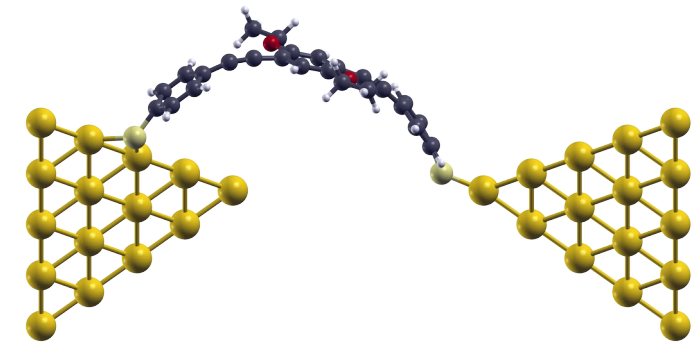
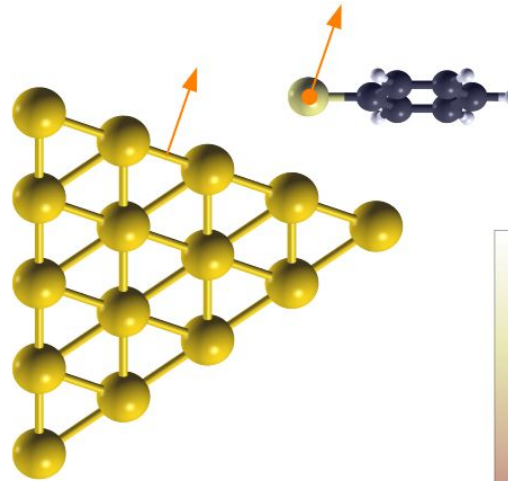
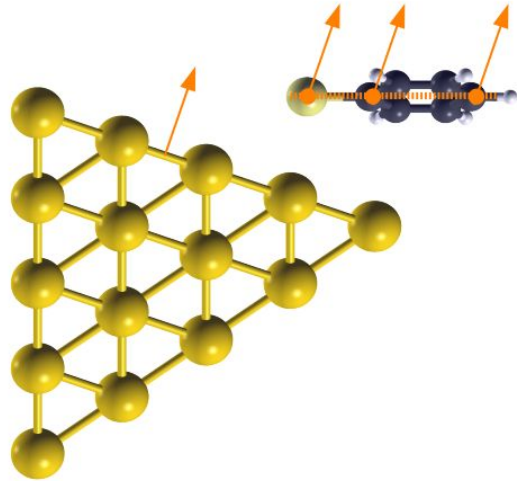
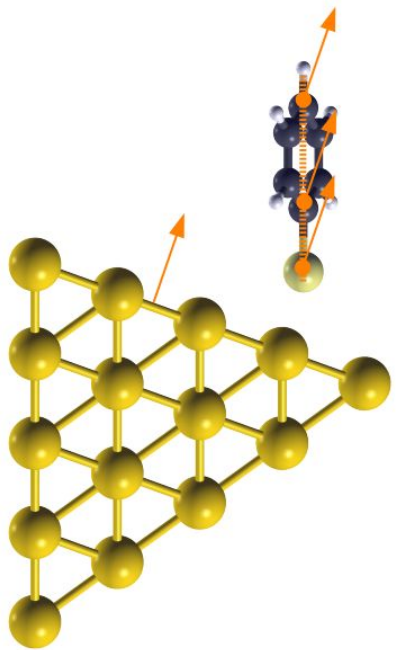


(c) grid division of docking positions on the facet edges



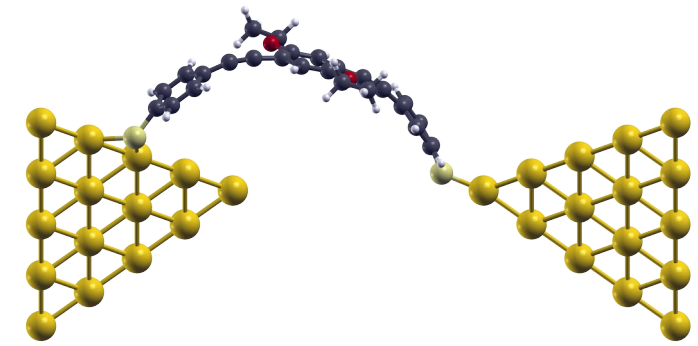
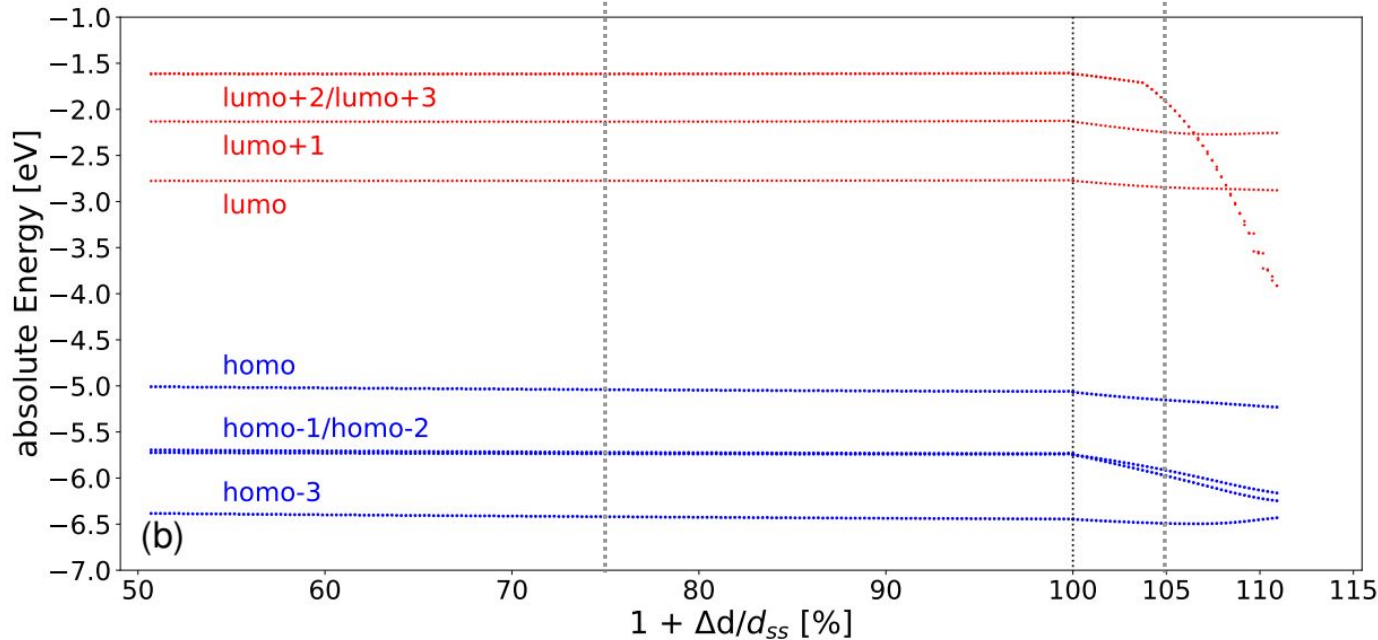
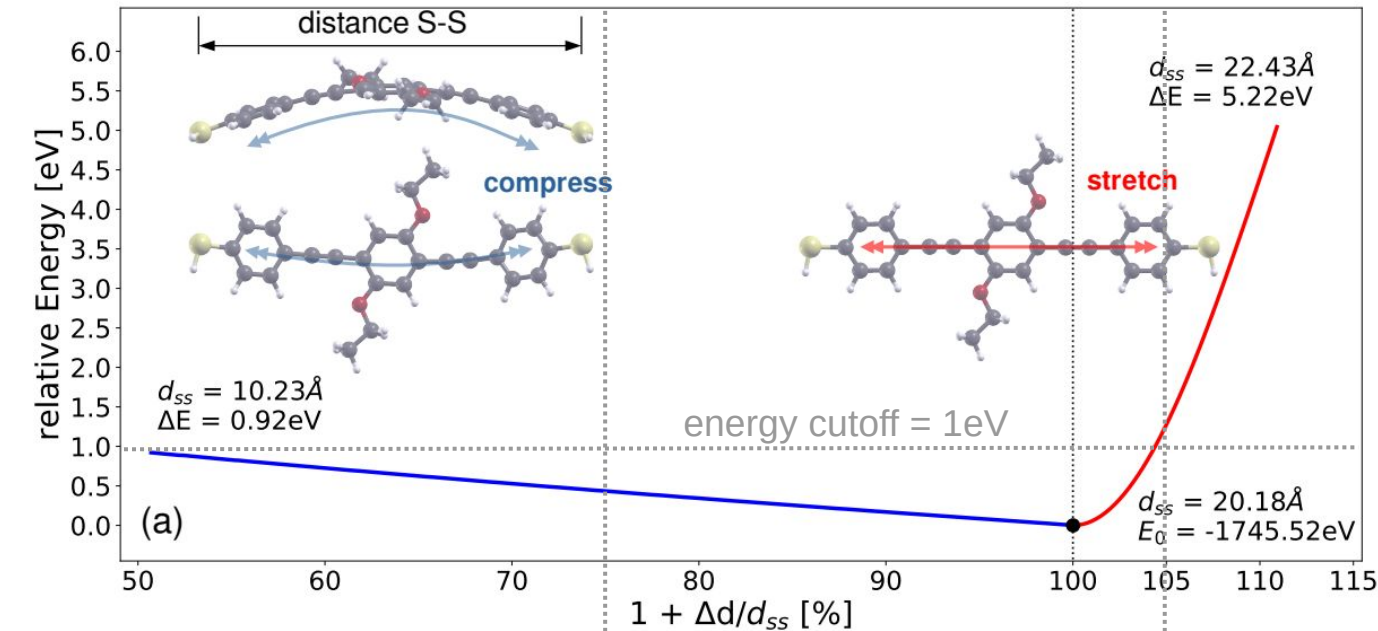
- Is the molecule with thiol anchoring groups at equilibrium for all tip-tip separations?
- Can the anchoring S-atoms slide freely on the gold facets?
- How much can the molecule be bend or stretched? Alteration of electronic states?
- Grid spacing accomodates midpoints between high symmetry sites
- Additional grid points chosen protruding over the facet-edges

# Facet Energy Landscape



- Thiophenyl
  - vertical, horizontal orientation
  - unrestricted
- Anchoring S-Atom moves along the facet normal
- Energy landscape
  - recurrent features and energy minimum at edges
  - tip region energetically unfavorable

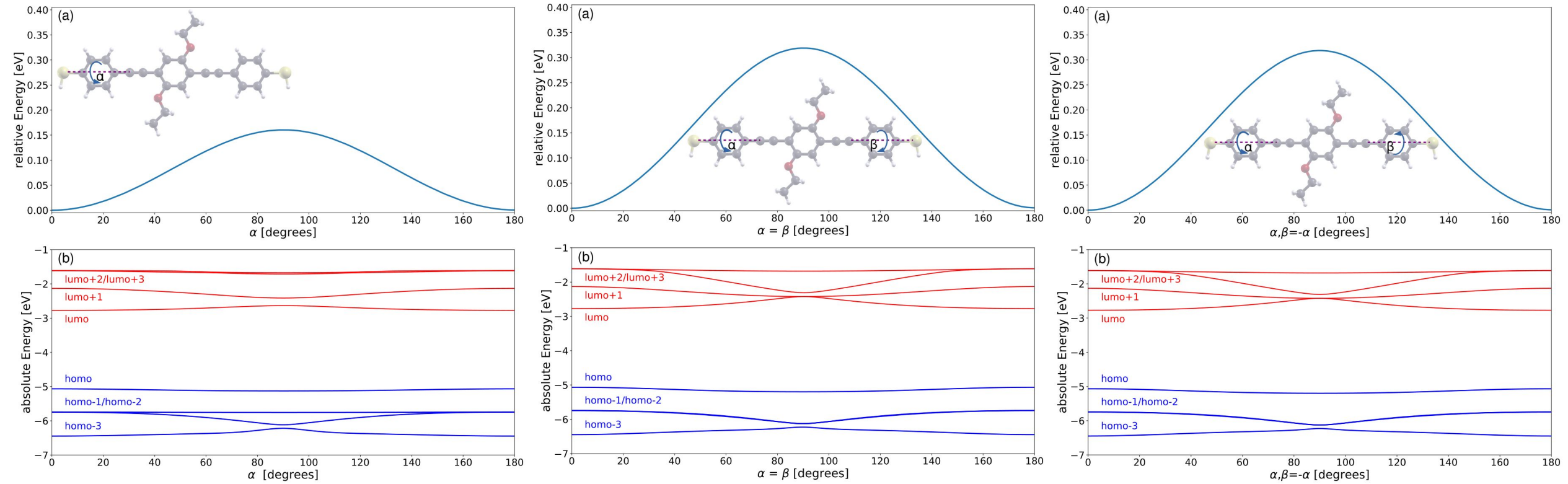
# Geometric Deformation



- Stretching follows Hooke's Law (parabola)
- Compression linear ▶ in-plane and out-of-plane bulging
- Initial energy cutoff 1eV
- Distortions included  $[0.75, 1.05] * d_{SS}$
- Energy crossover of unoccupied levels  $[1.05, 1.10] * d_{SS}$
- Splitting on HOMO-1/-2 negligible



# Inducing torsion – perturbation of electronic states

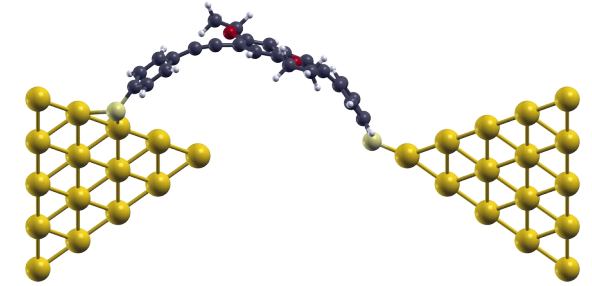
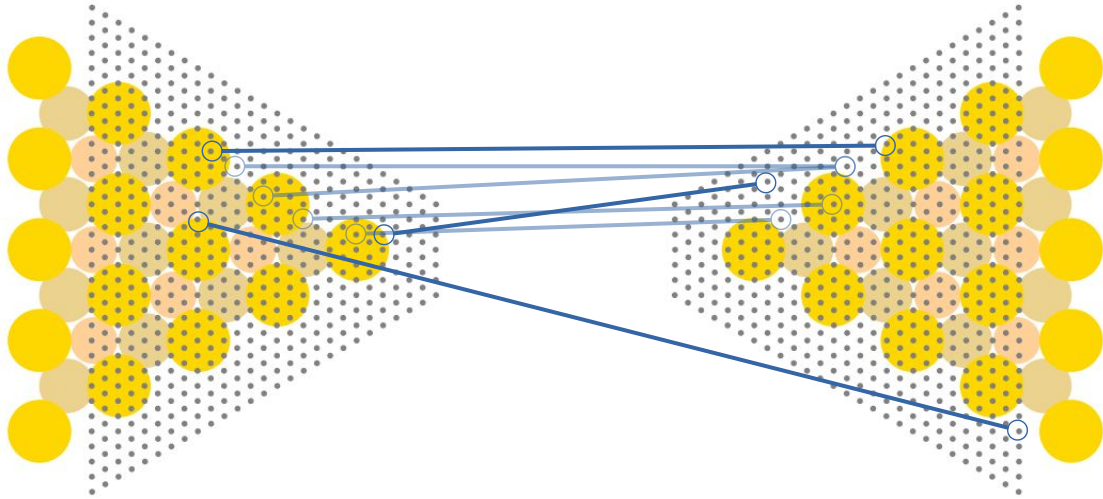


- In all three cases, crossover of unoccupied/occupied level occurs at torsional distortion [  $60^\circ$ ,  $120^\circ$  ]

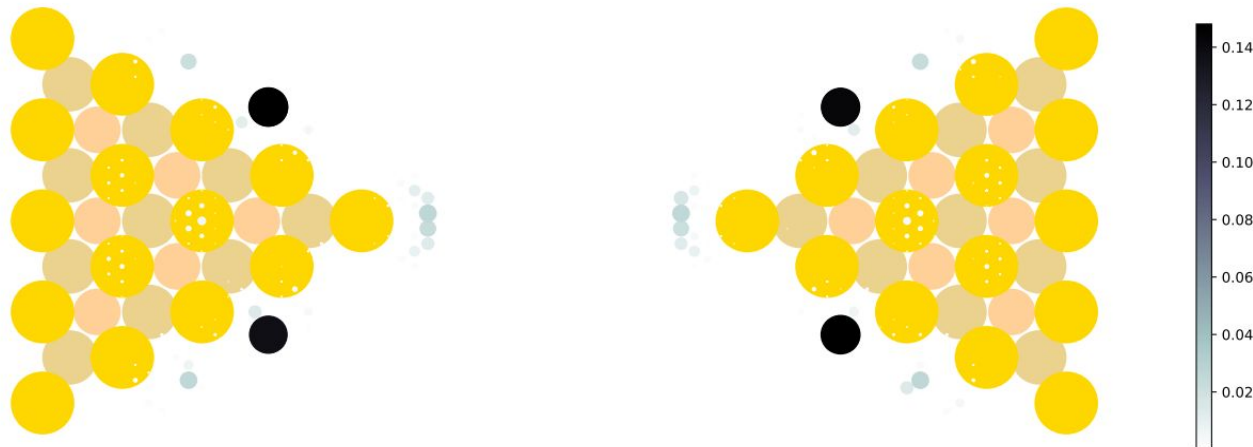
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# Random Walk | Surface Dynamics

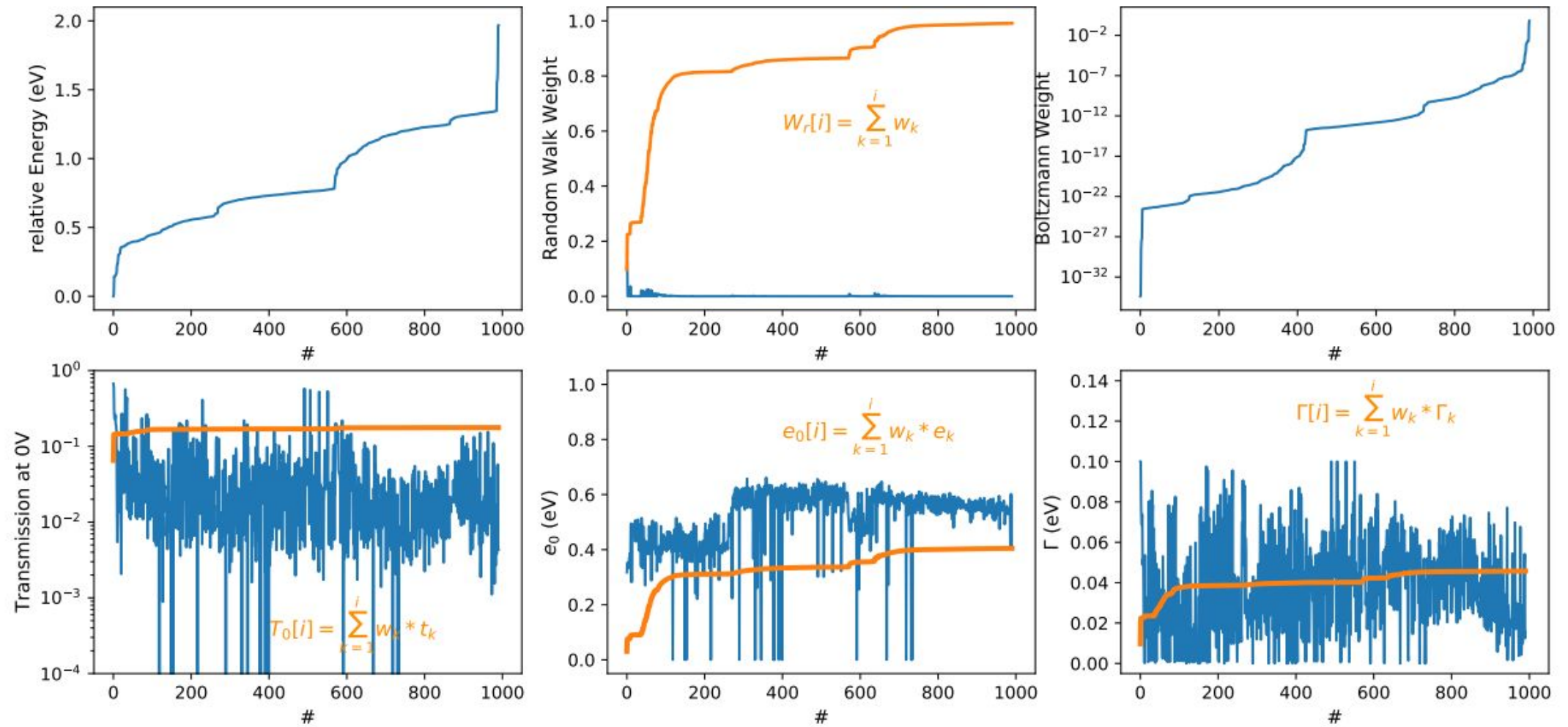
# Random Walk | Surface Dynamics



- Driving Force for surface dynamics
- Random walk – transitions to nearest neighbours allowed

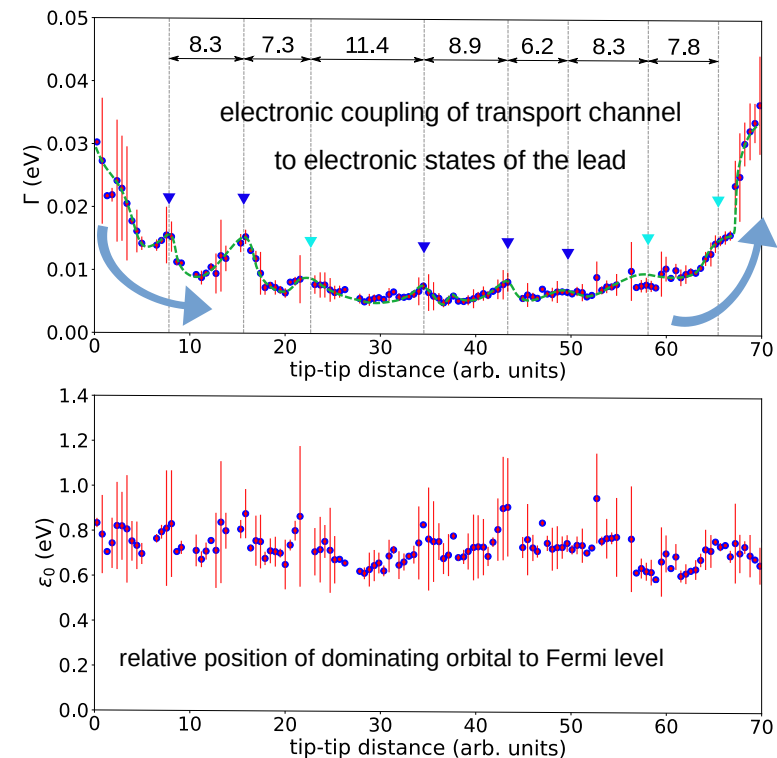
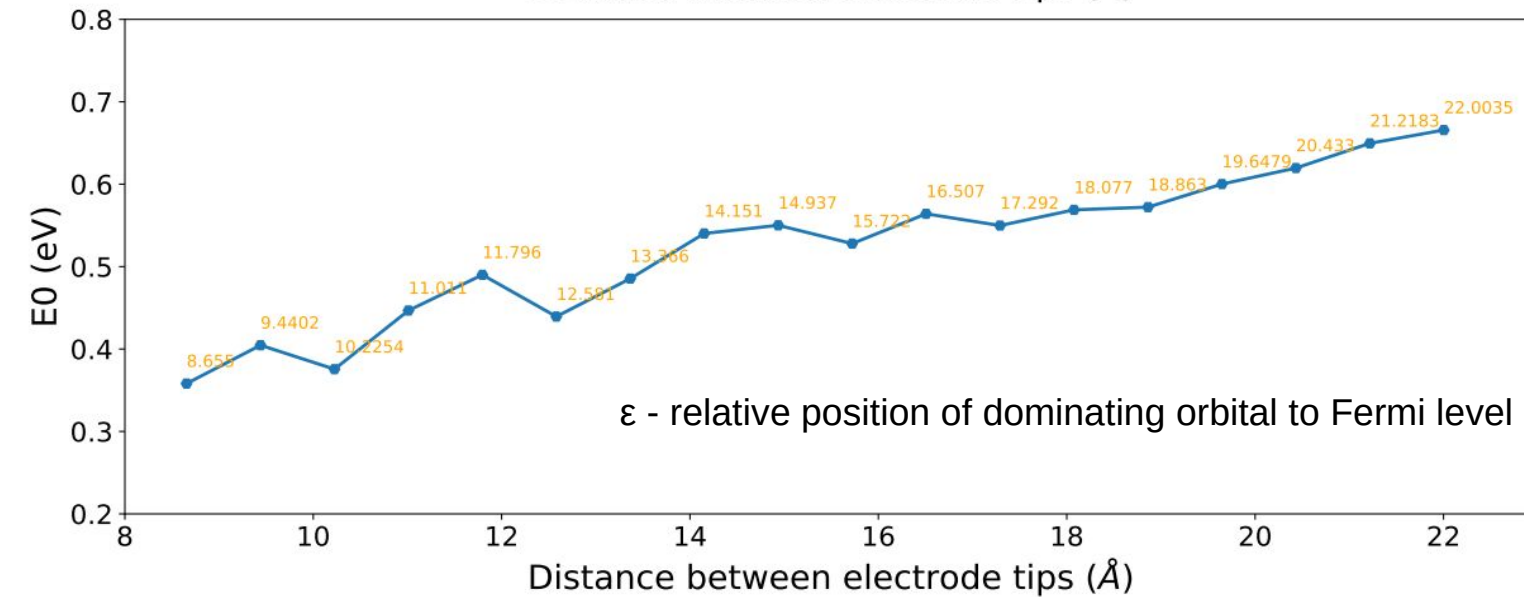
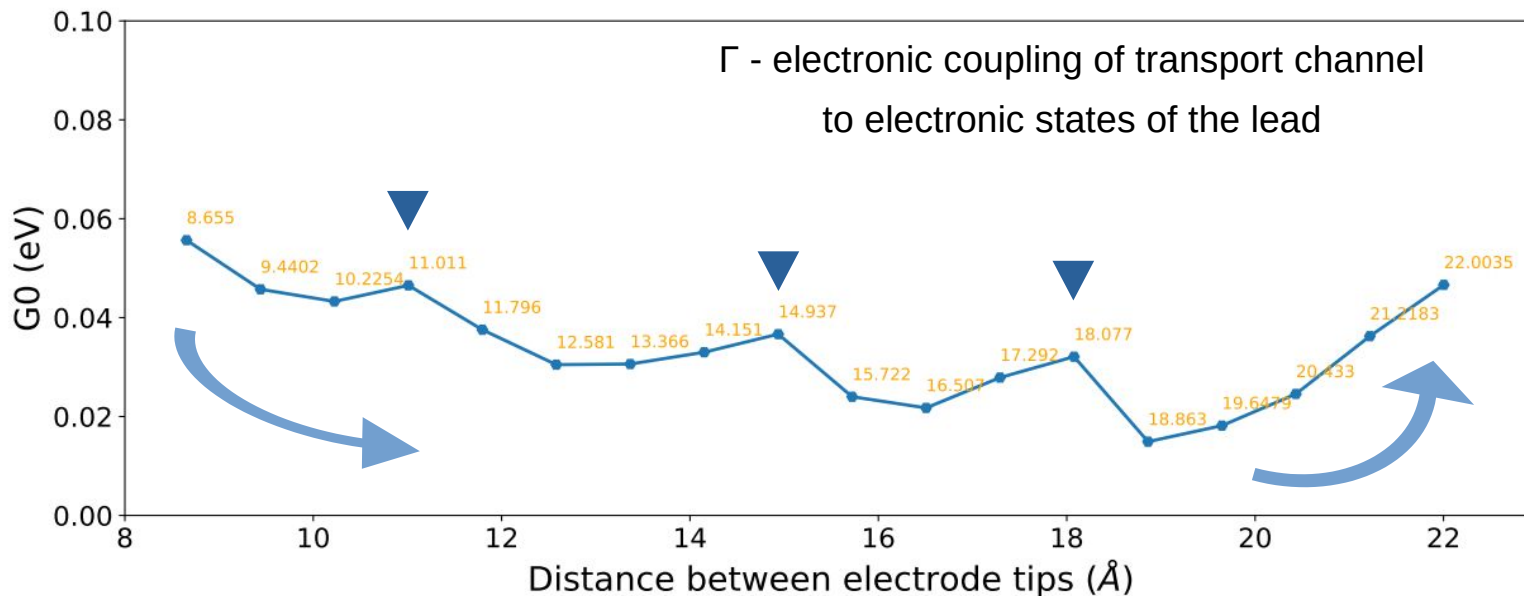


# Random Walk | Surface Dynamics



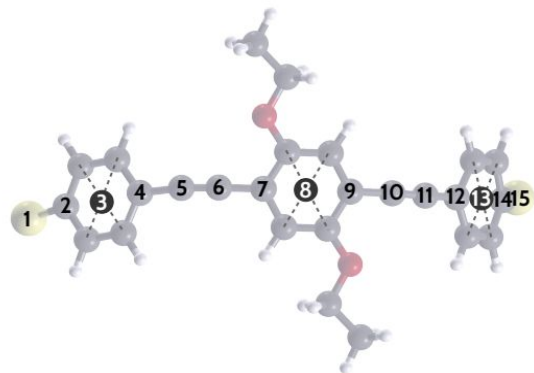
# „Time“ evolution ETCQs using DFTB+NEGF

# „Time“ Evolution of ETCQs

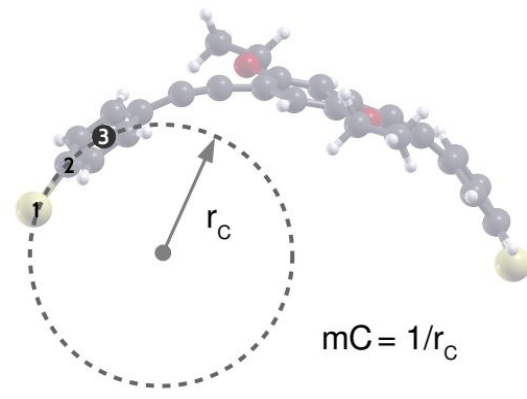


# Characterizing configuration Ensemble

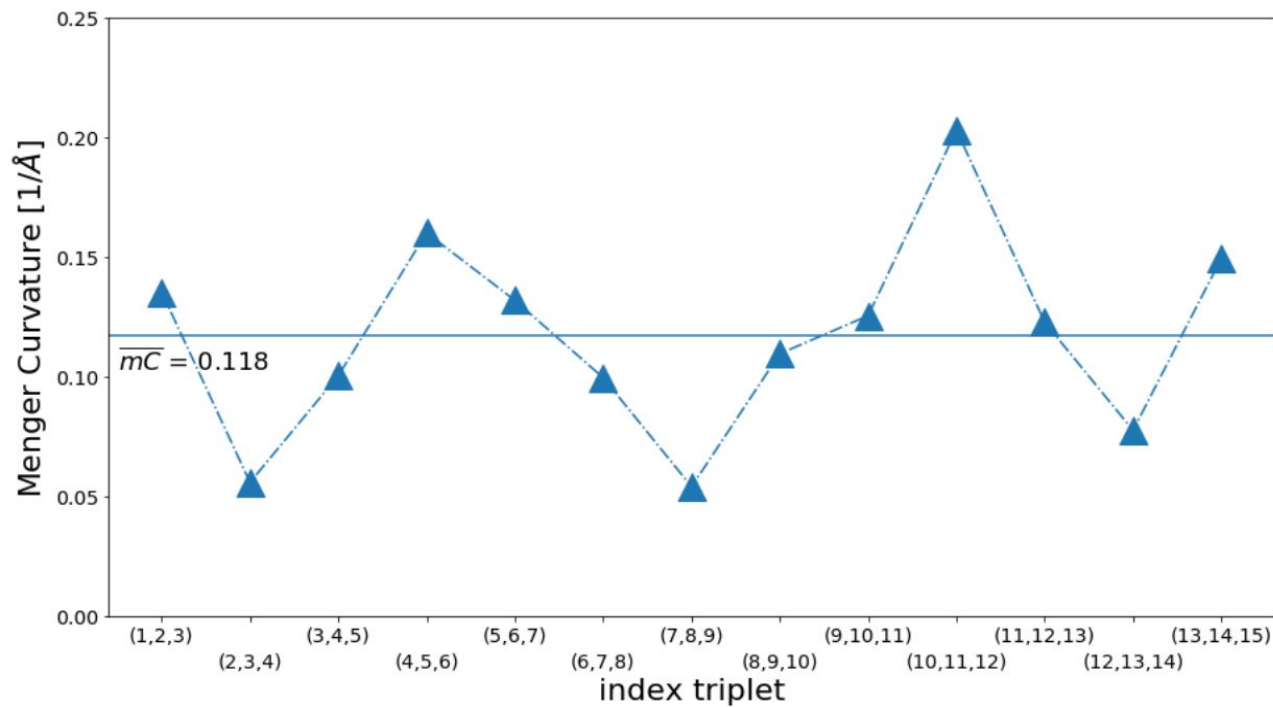
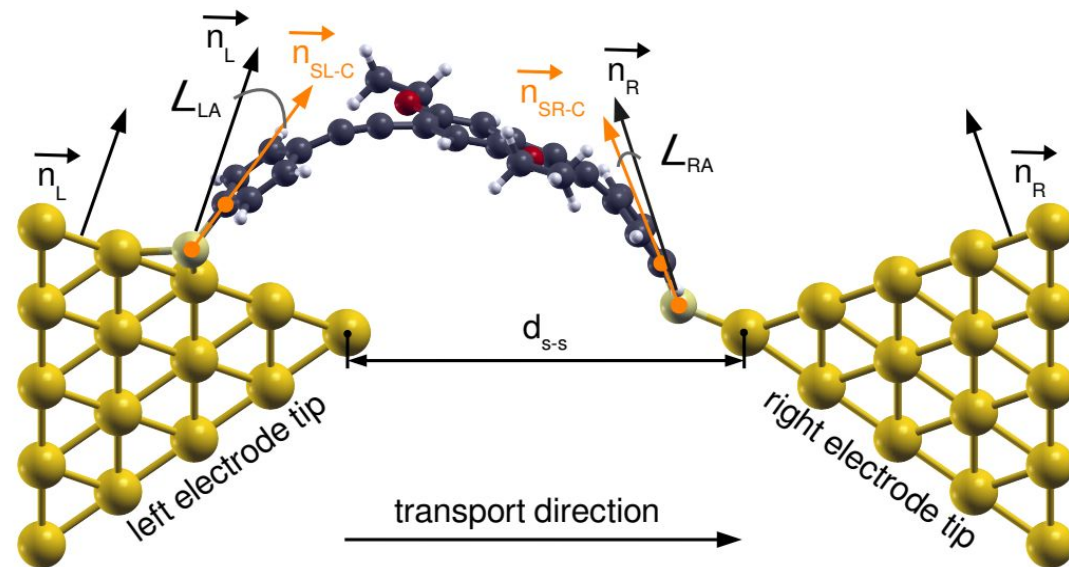
# Characterizing configuration ensemble



(a) enumeration of atoms and ring centers of the molecular backbone



(b) radius of curvature of a triplet



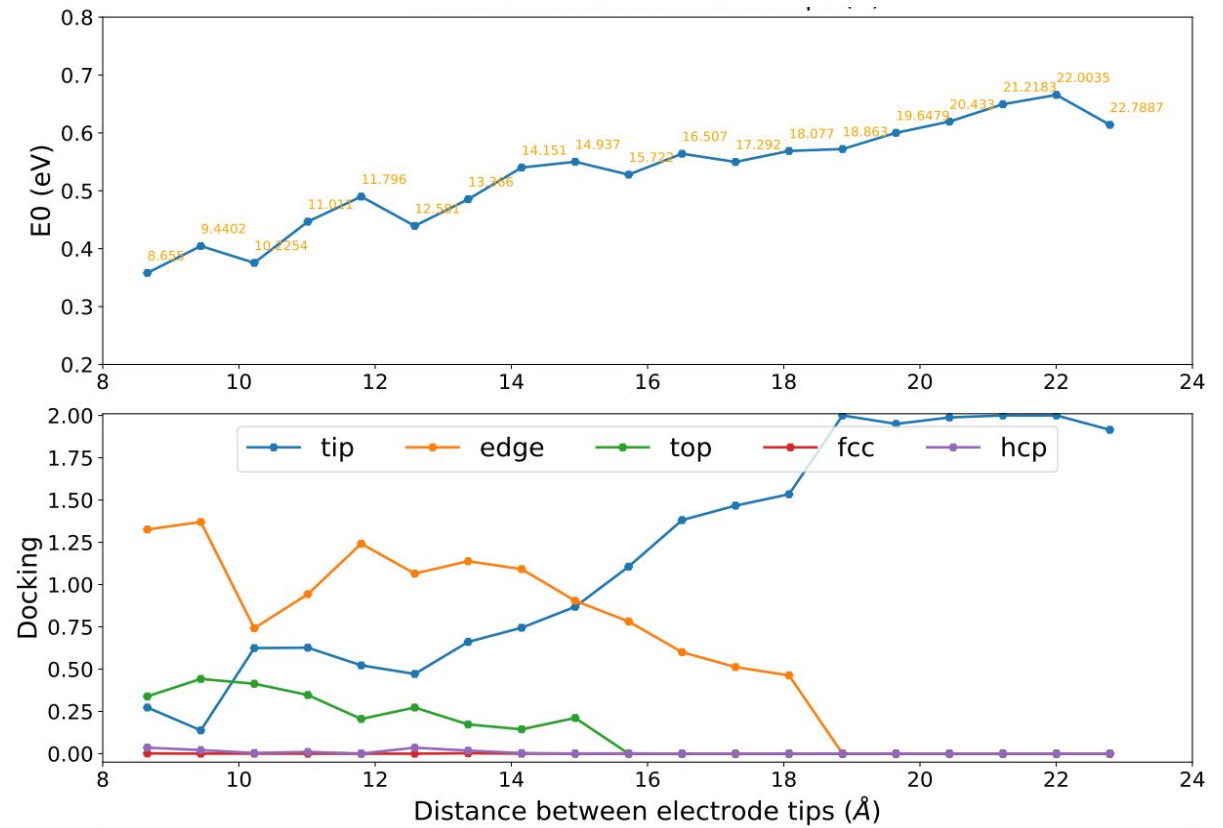
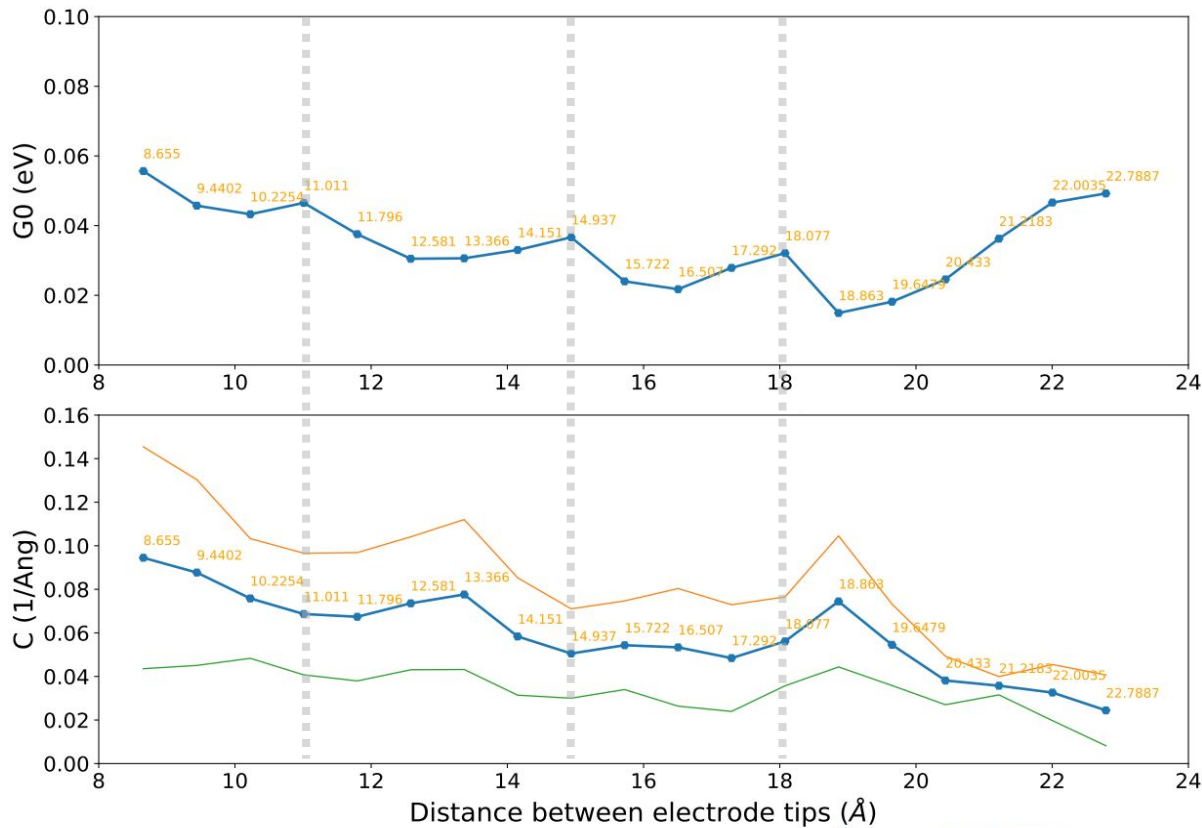
24  
(c) Menger Curvature

- Tip-tip separation
  - Average Menger Curvature of successive triplets along the molecular backbone
  - Local-coordination of docking position
- 
- S-S separation
  - Angle btw. Facet normals and S-C bond vector
  - Torsion angle

Source: J. Leger, *Annals of Mathematics*. 149 (3): 831–869 (1999)



# Characterizing configuration ensemble



- Recurrent feature in „time“ evolution of  $\Gamma$  linked to overall curvature of PEEB molecular backbone
- Binding predominantly on edges for shorter tip separations ▶ Coupling to multiple facet states
- At larger distances, binding to tip states ▶ Au-S-C bonding angle optimal
- Evolution of  $\varepsilon$  ▶ reduction of image charge ▶ reduction of renormalization

# Conclusion and Outlook

- JO

# Acknowledgements



- Florian Günther, Jeffery Kelling, Sibylle Gemming
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