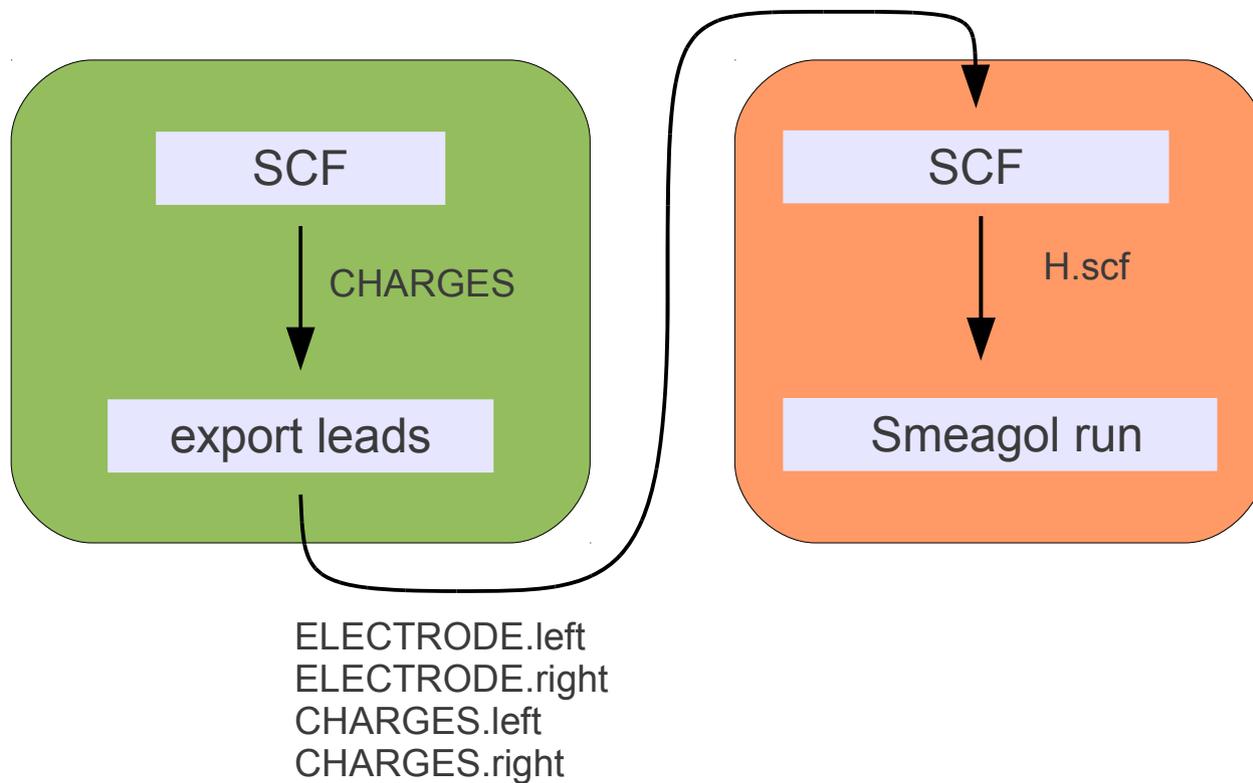


Basic scheme of Fireball2Smeagol Run

PRINCIPAL LAYER
directory

EXTENDED MOLECULE
directory



BULK calculation

```
answer.bas [----] 0 L:[ 1+ 0 1/ 7]
element 6 x y z
1 3.000000 3.000000 1.000000
1 3.000000 3.000000 2.000000
1 3.000000 3.000000 3.000000
1 3.000000 3.000000 4.000000
1 3.000000 3.000000 5.000000
1 3.000000 3.000000 6.000000
```

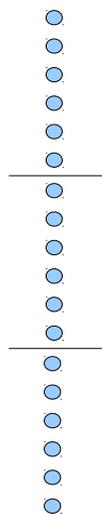
```
cel.lvs [----] 20 L:[ 1+ 2 3/ 4]
20.000000000 0.000000000 0.000000000
0.000000000 20.000000000 0.000000000
0.000000000 0.000000000 6.000000000
```

```
input.kpts [----] 0 L:[ 1+ 0 1/102] *(0 /6956b)= 32
100
0.000000000 0.000000000 -0.5183627873 0.0100000000
0.000000000 0.000000000 -0.5078908118 0.0100000000
0.000000000 0.000000000 -0.4974188363 0.0100000000
0.000000000 0.000000000 -0.4869468608 0.0100000000
0.000000000 0.000000000 -0.4764748853 0.0100000000
0.000000000 0.000000000 -0.4660029098 0.0100000000
```

k-points in x,y,z (for bulk)

```
MOLECULE.kpts [----] 0 L:[ 1+ 2 3/ 3] *(73 / 73b)= <E0F>
1
0.000000000 0.000000000 0.000000000 1.000000000
```

k-points in x,y (for exported leads)



```
fireball.in [----] 13 L
&OPTION
basisfile = answer.bas
lvsfile = cel.lvs
icluster = 0
nstepf = 1
sigmatol = 0.0000000001
max_scf_iterations = 100
dt = 0.5
iqout = 1
ismeargol = 0
ifixcharge = 0
&END

&OUTPUT
iwrthSrho = 0
iwrteigen = 0
iwrtdos = 0
&END

&QUENCH
&END

&MESH
&END
```

SCF

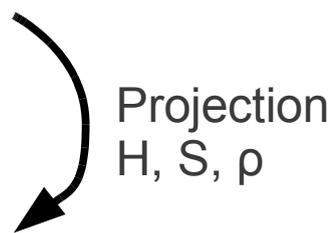
```
fireball.in [----] 1
&OPTION
basisfile = answer.bas
lvsfile = cel.lvs
icluster = 0
nstepf = 1
sigmatol = 0.0000000001
max_scf_iterations = 100
dt = 0.5
iqout = 1
ismeargol = 0
ifixcharge = 1
&END

&OUTPUT
iwrthSrho = 1
iwrteigen = 0
iwrtdos = 0
&END

&QUENCH
&END

&MESH
&END
```

export leads



output to "ELECTRODE" file

Extended molecule calculation

```
answer.bas [----] 39 L:[ 1+12 13/ 25] *(5
24
1 3.0000000000 3.0000000000 1.0000000000
1 3.0000000000 3.0000000000 2.0000000000
1 3.0000000000 3.0000000000 3.0000000000
1 3.0000000000 3.0000000000 4.0000000000
1 3.0000000000 3.0000000000 5.0000000000
1 3.0000000000 3.0000000000 6.0000000000
1 3.0000000000 3.0000000000 7.0000000000
1 3.0000000000 3.0000000000 8.0000000000
1 3.0000000000 3.0000000000 9.0000000000
1 3.0000000000 3.0000000000 10.0000000000
1 3.0000000000 3.0000000000 11.0000000000
1 3.0000000000 3.0000000000 13.5000000000
1 3.0000000000 3.7000000000 13.5000000000
1 3.0000000000 3.0000000000 16.0000000000
1 3.0000000000 3.0000000000 17.0000000000
1 3.0000000000 3.0000000000 18.0000000000
1 3.0000000000 3.0000000000 19.0000000000
1 3.0000000000 3.0000000000 20.0000000000
1 3.0000000000 3.0000000000 21.0000000000
1 3.0000000000 3.0000000000 22.0000000000
1 3.0000000000 3.0000000000 23.0000000000
1 3.0000000000 3.0000000000 24.0000000000
1 3.0000000000 3.0000000000 25.0000000000
1 3.0000000000 3.0000000000 26.0000000000
```

```
smeagol.optional [----] 3 L:
5 NEnergyR
90 NEnergyIC
20 NEnergyIL
10 NPolos
0.001 Delta
-45.0 EnergyLB
1 NSlices
T TrCoeff
1000 NeneT
-20.0 TEnergI
30.0 TEnergF
-4.299536 Fermi_level
-5.0 V_Bias
12.0 r_left
12.5 r_right
1 useLeads?
4.50 r_start_fithop
0.25 r_scale_fithop
```

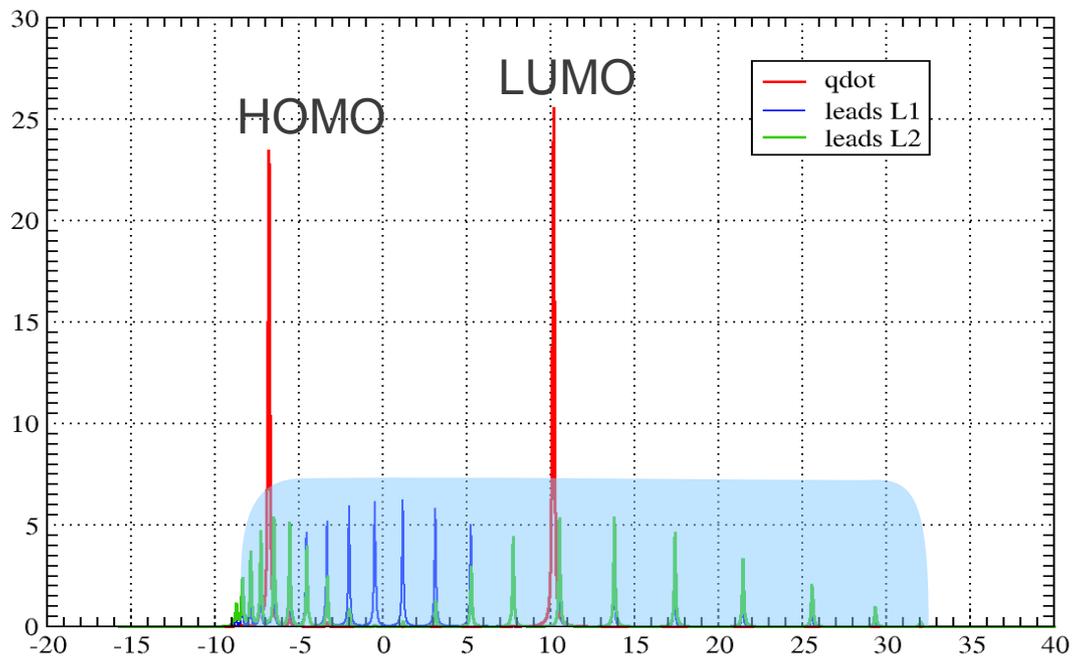
```
cel.lvs [----] 0 L:[ 1+ 2 3/ 4] *(9
20.0000000000 0.0000000000 0.0000000000
0.0000000000 20.0000000000 0.0000000000
0.0000000000 0.0000000000 26.0000000000
```

```
input.kpts [----] 0 L:[ 1+ 0 1/ 3] *(0 / 73b)= 32
1
0.0000000000 0.0000000000 0.0000000000 1.0000000000
```

Example 1

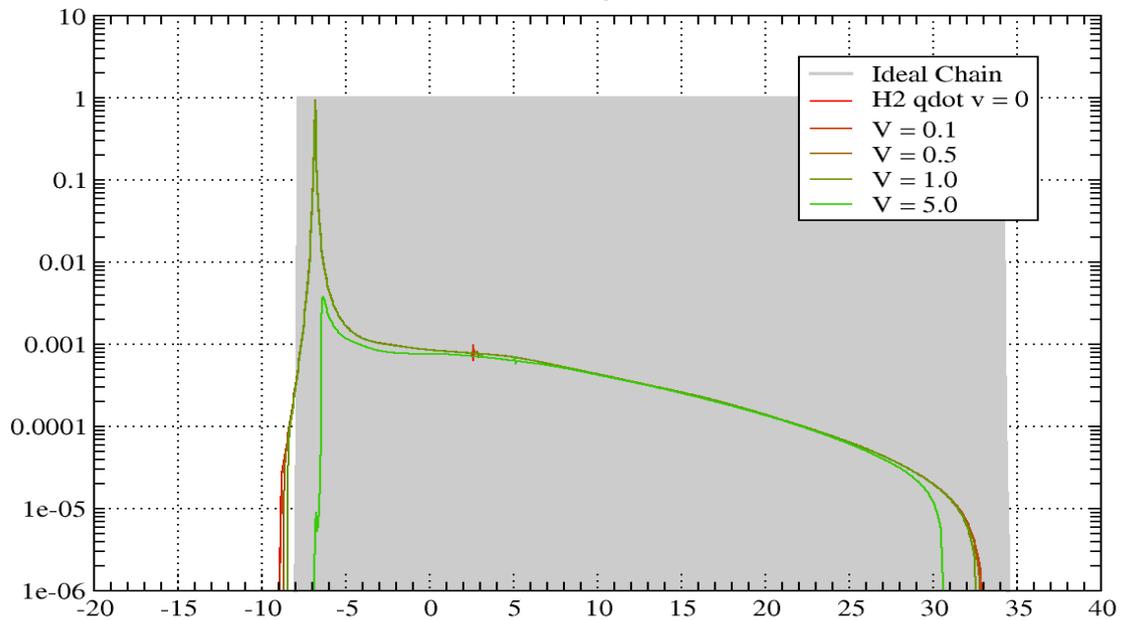
**H₂ quantum dot
in nonselfconsistent
potential**

H2 qdot DOS

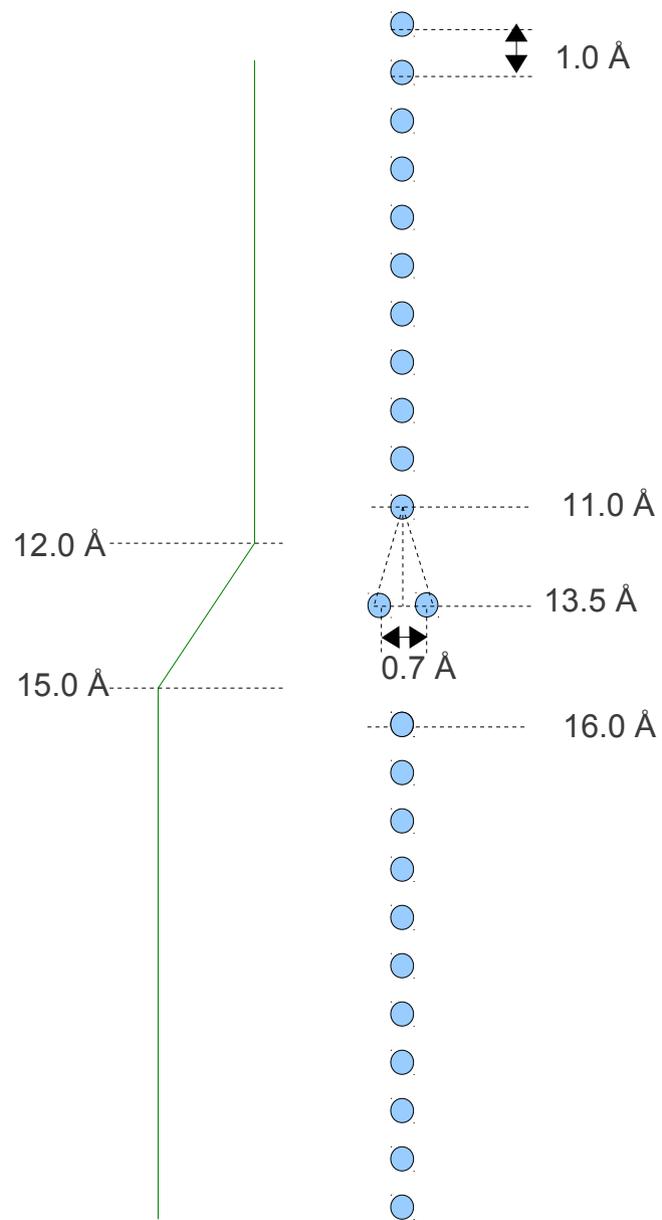


Transmission Coefficient H2 qdot

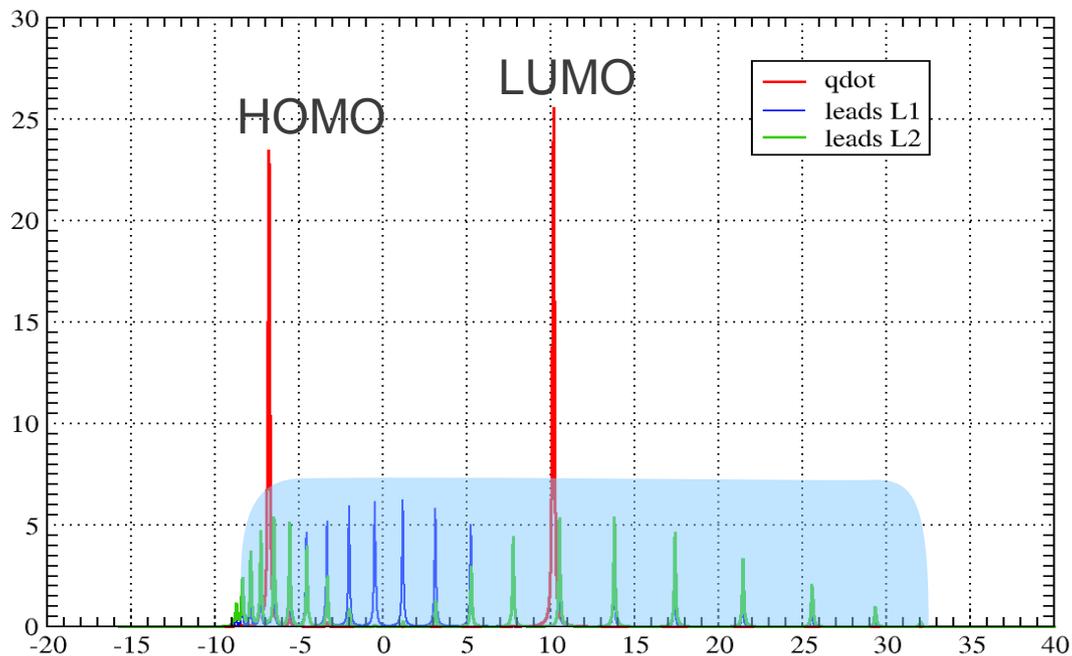
symetric potential



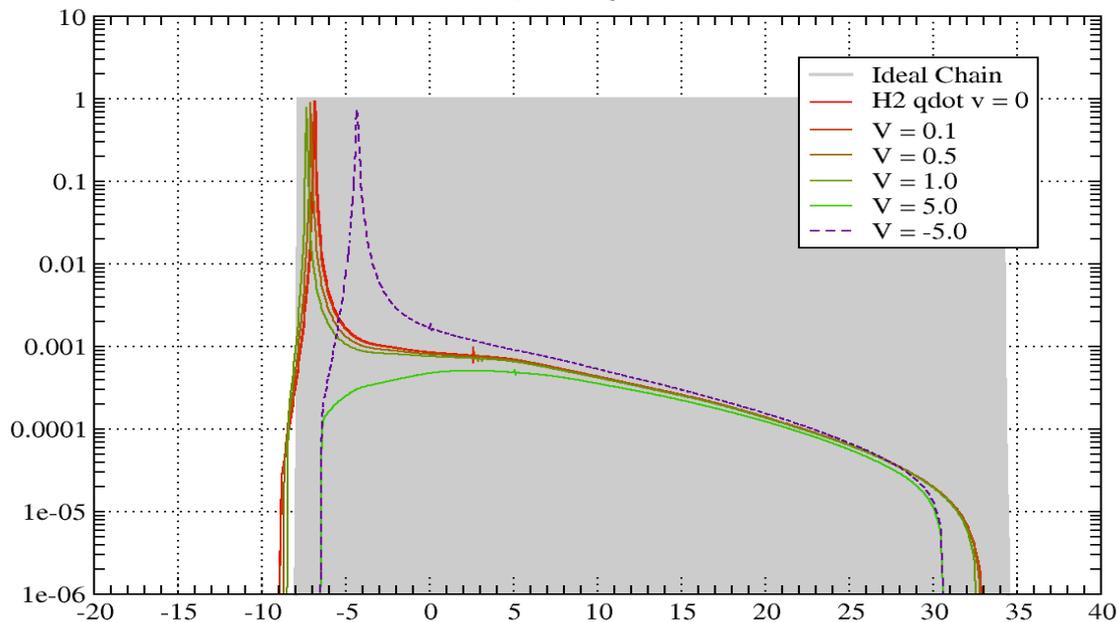
symetric potential symetric position



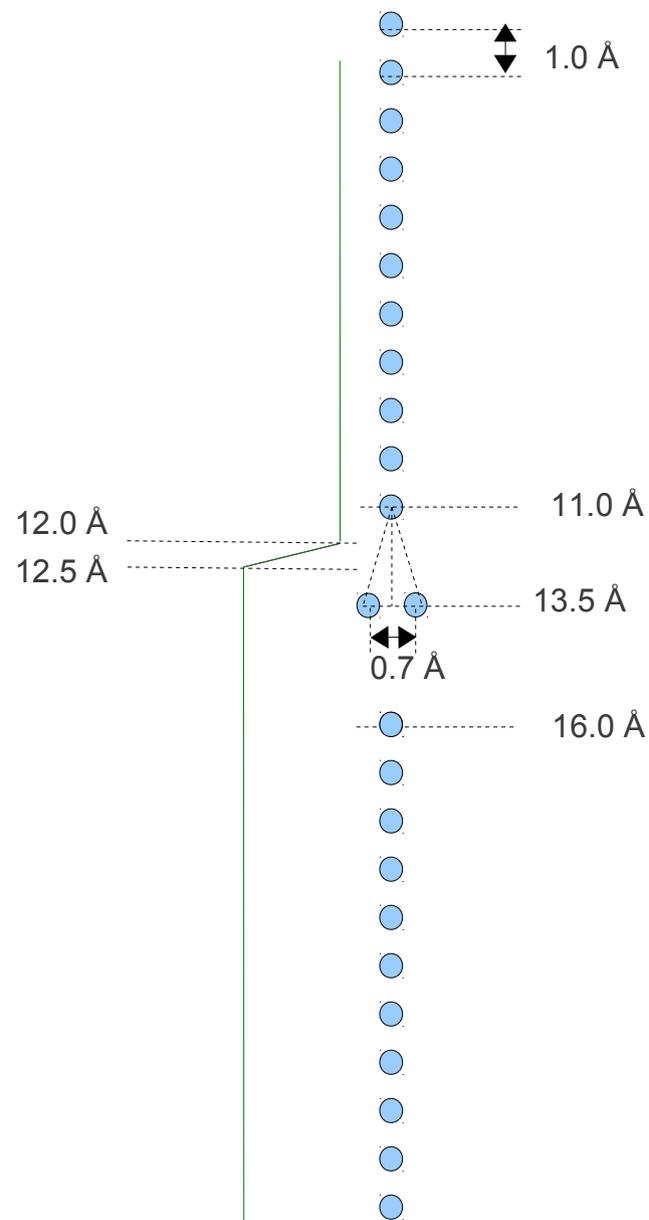
H2 qdot DOS



Transmission Coefficient H2 qdot asymmetric potential

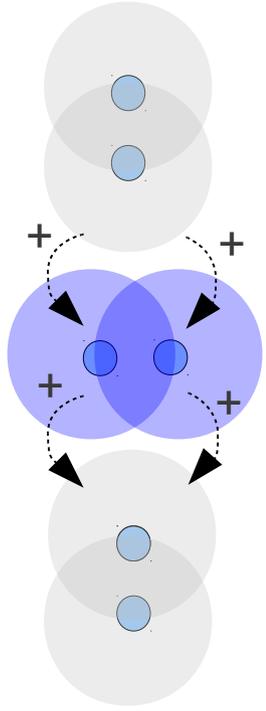


asymmetric potential symetric position



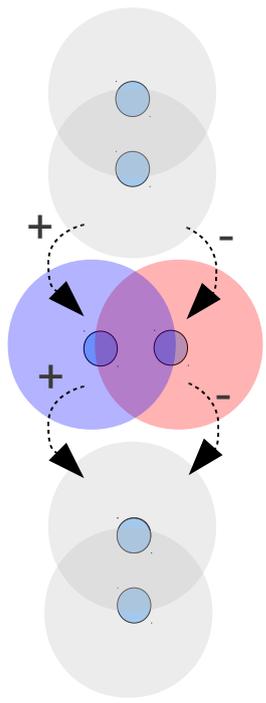
H2 quantum dot (explanation)

HOMO bonding

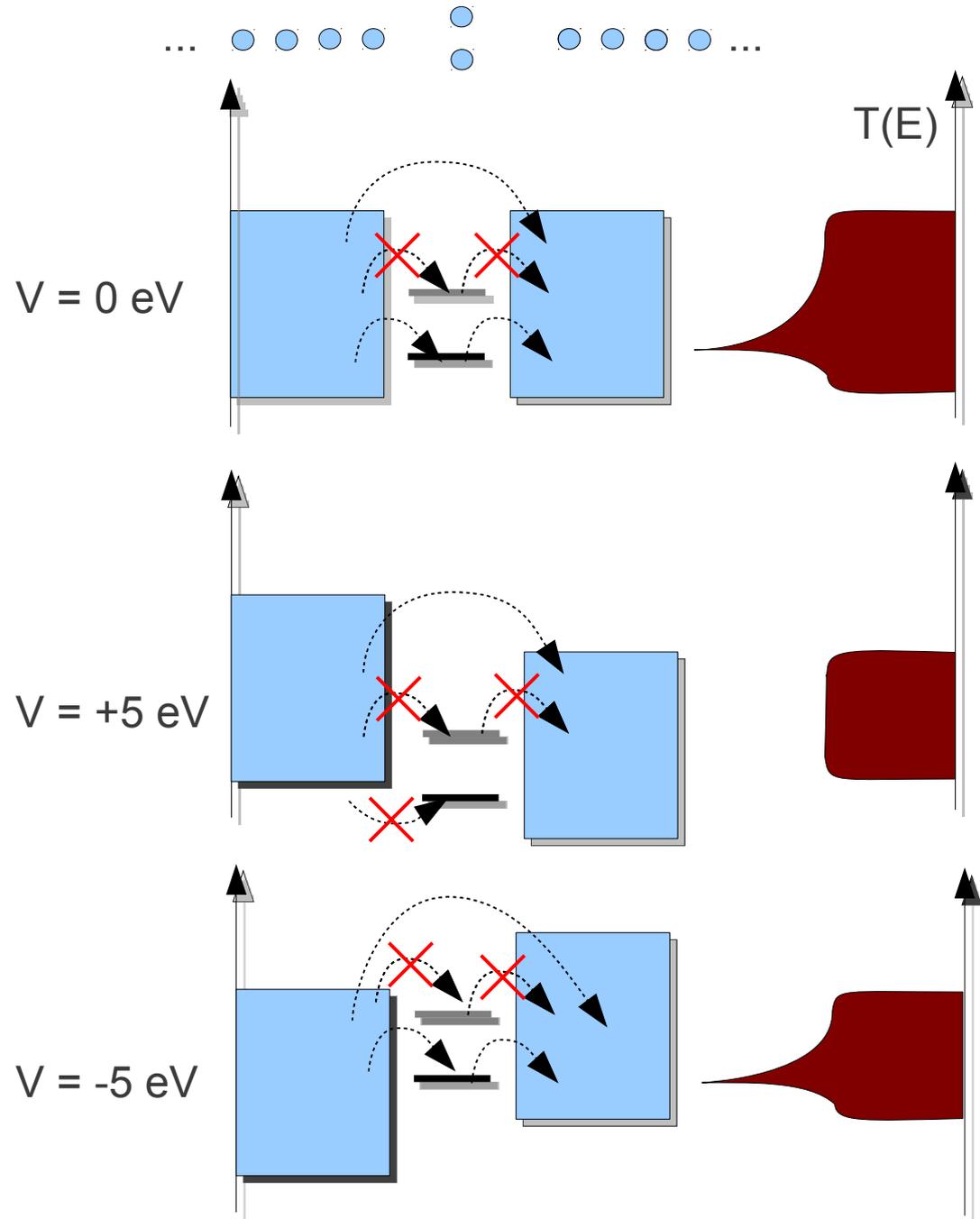


$$t_{12} \sim \int s \cdot (s + s) = 0$$

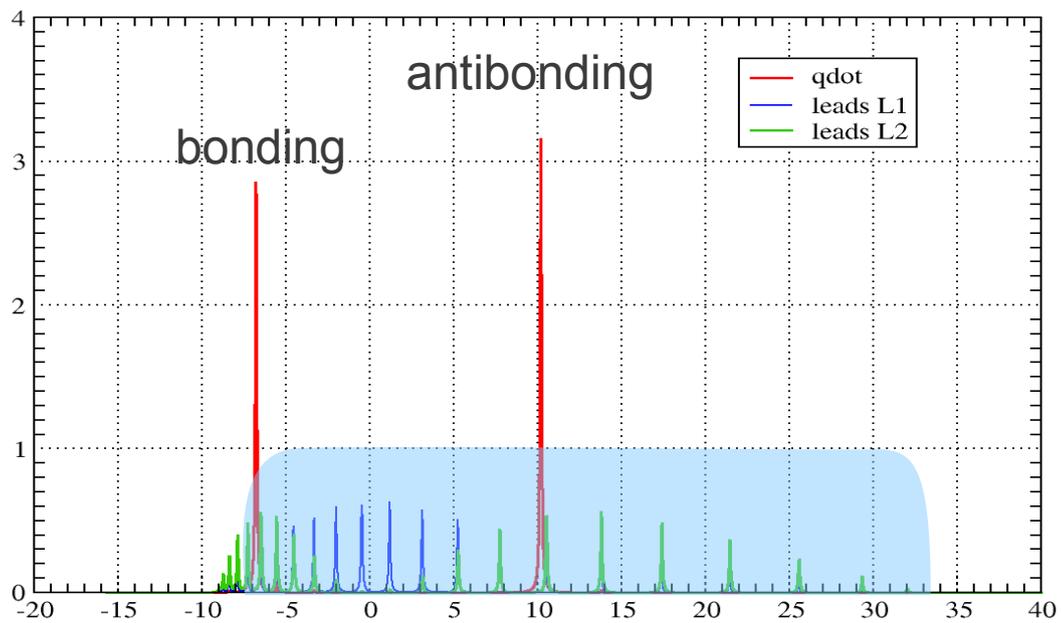
LUMO antibonding



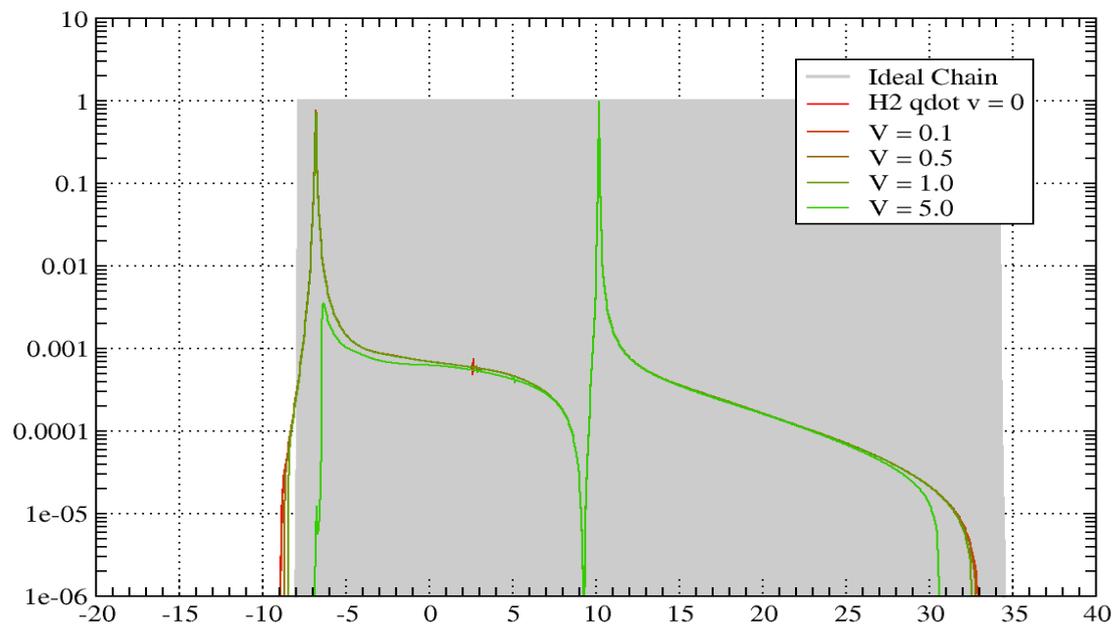
$$t_{12} \sim \int s \cdot (s - s) = 0$$



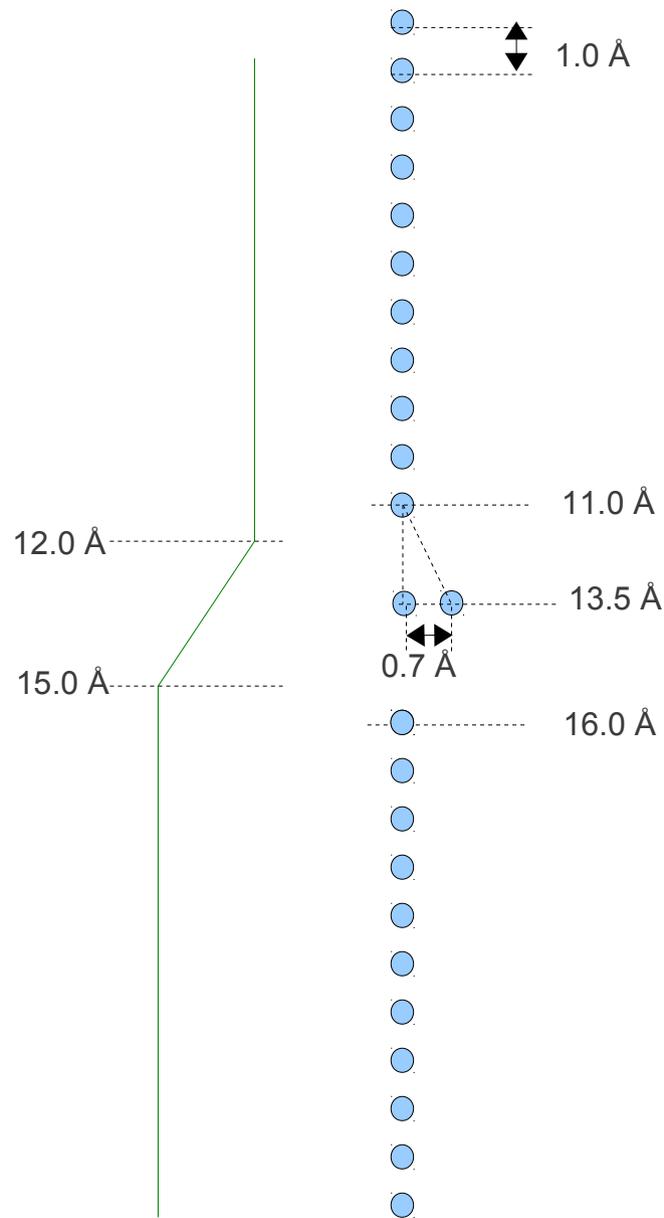
H2 qdot DOS



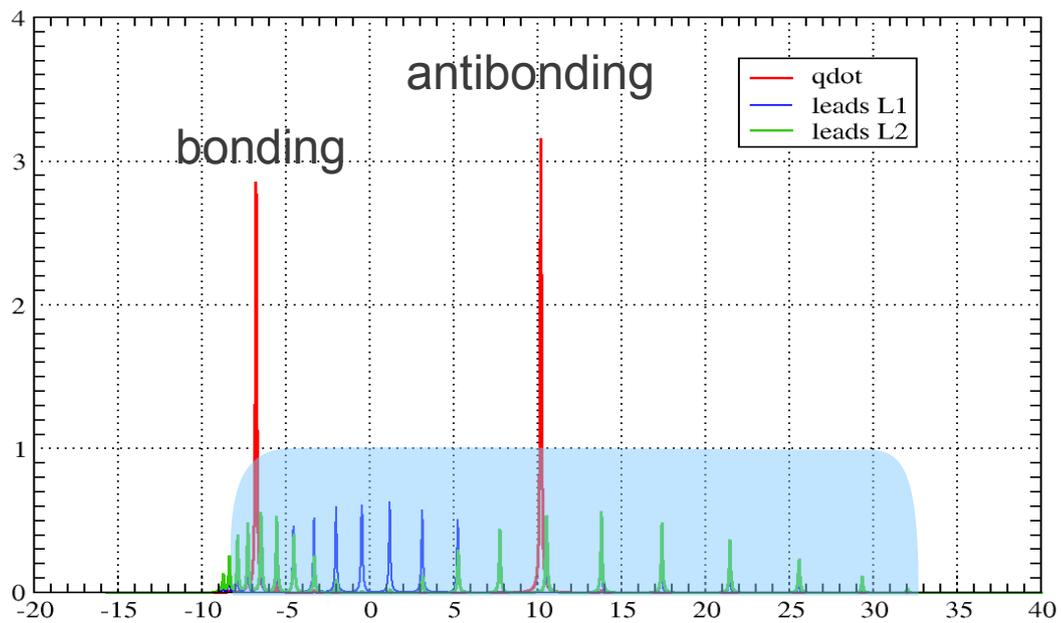
Transmission Coefficient H2 qdot



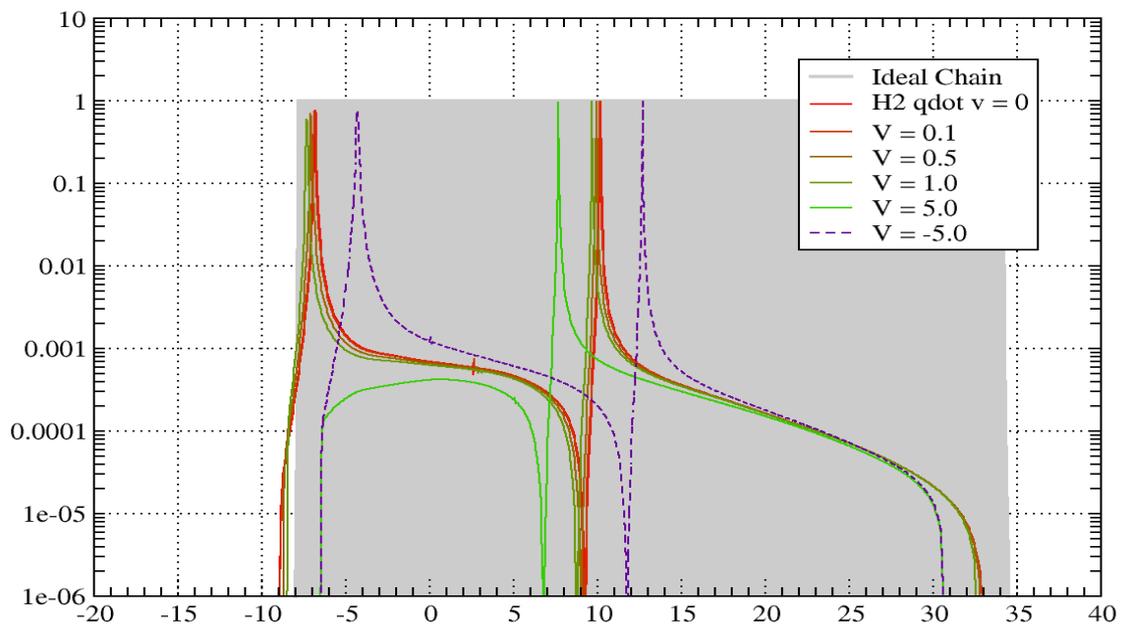
symetric potential asymmetric position



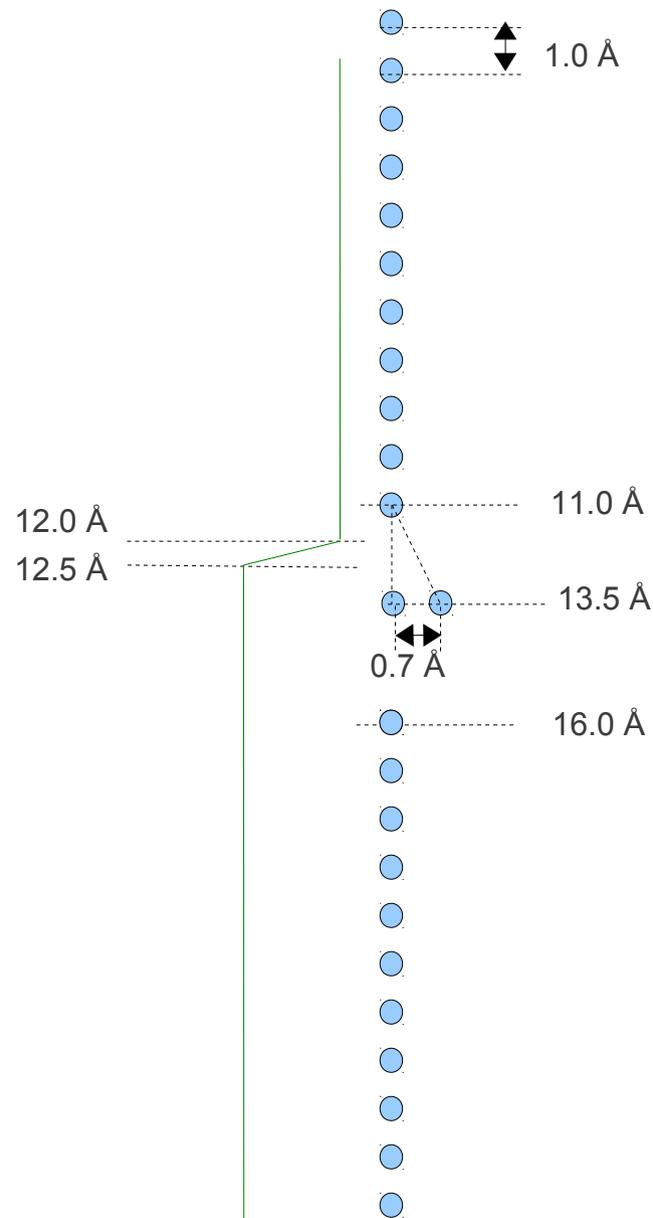
H2 qdot DOS



Transmission Coefficient H2 qdot



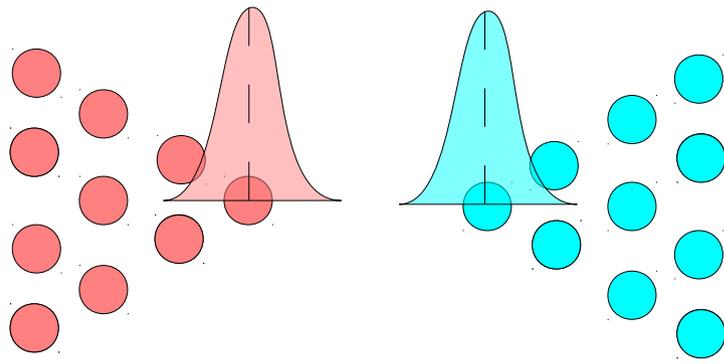
asymmetric potential asymmetric position



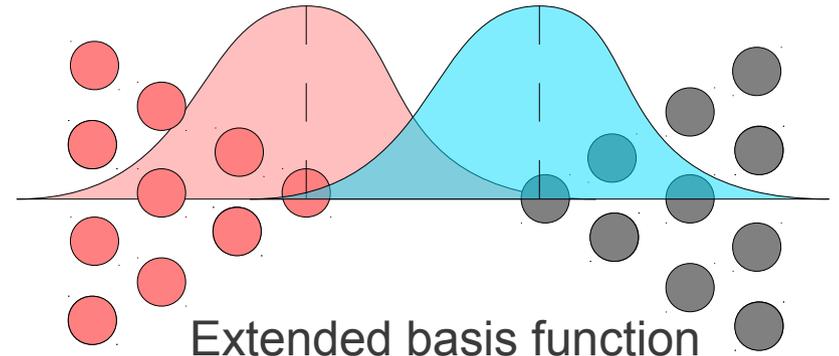
Example 2

**Braking Gold Chain
width extendet
hopping**

extended hopping



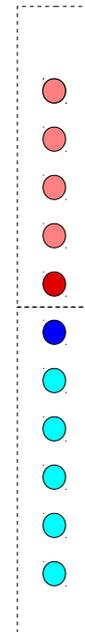
Original basis function



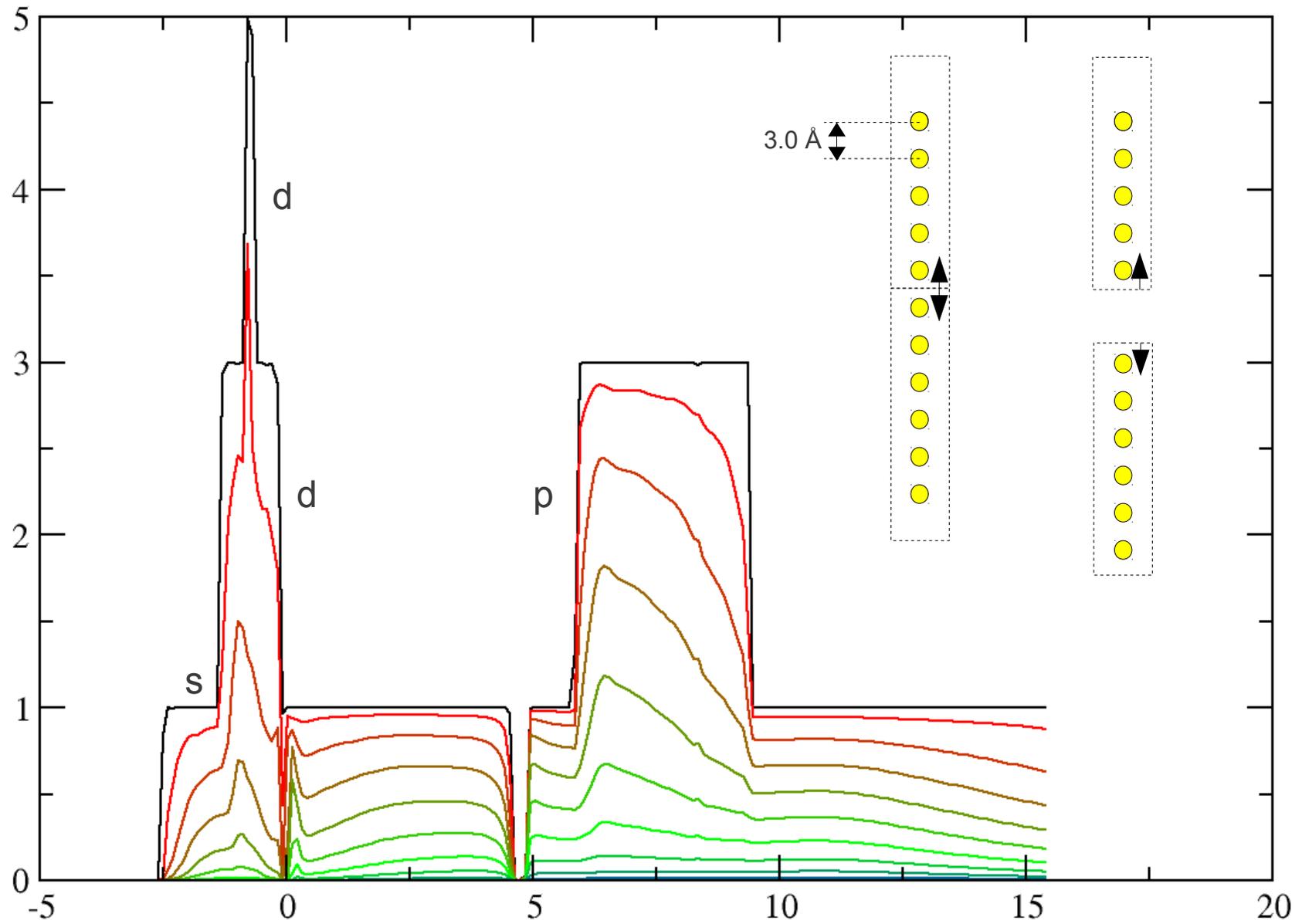
Extended basis function
in apex region

This is replaced to H and S matrix
in apex region before
before construction of Green's function

```
interaction.optional [----] 1 L:[ 1+11 12/ 13] *(650 / 686b)= 32 0x20
0 ! ncell sample1
12 ! total number of atoms in the sample1
1 ! number of intervals defining the sample1
1 12 ! atom1 atom2 of the interval
1 ! number of atoms in tip1
6 ! list of atoms in the tip1
0 ! ncell sample1
12 ! total number of atoms in the sample2
1 ! number of intervals defining the sample2
1 12 ! atom1 atom2 of the interval
1 ! number of atoms in tip2
7 ! list of atoms in the tip2
```

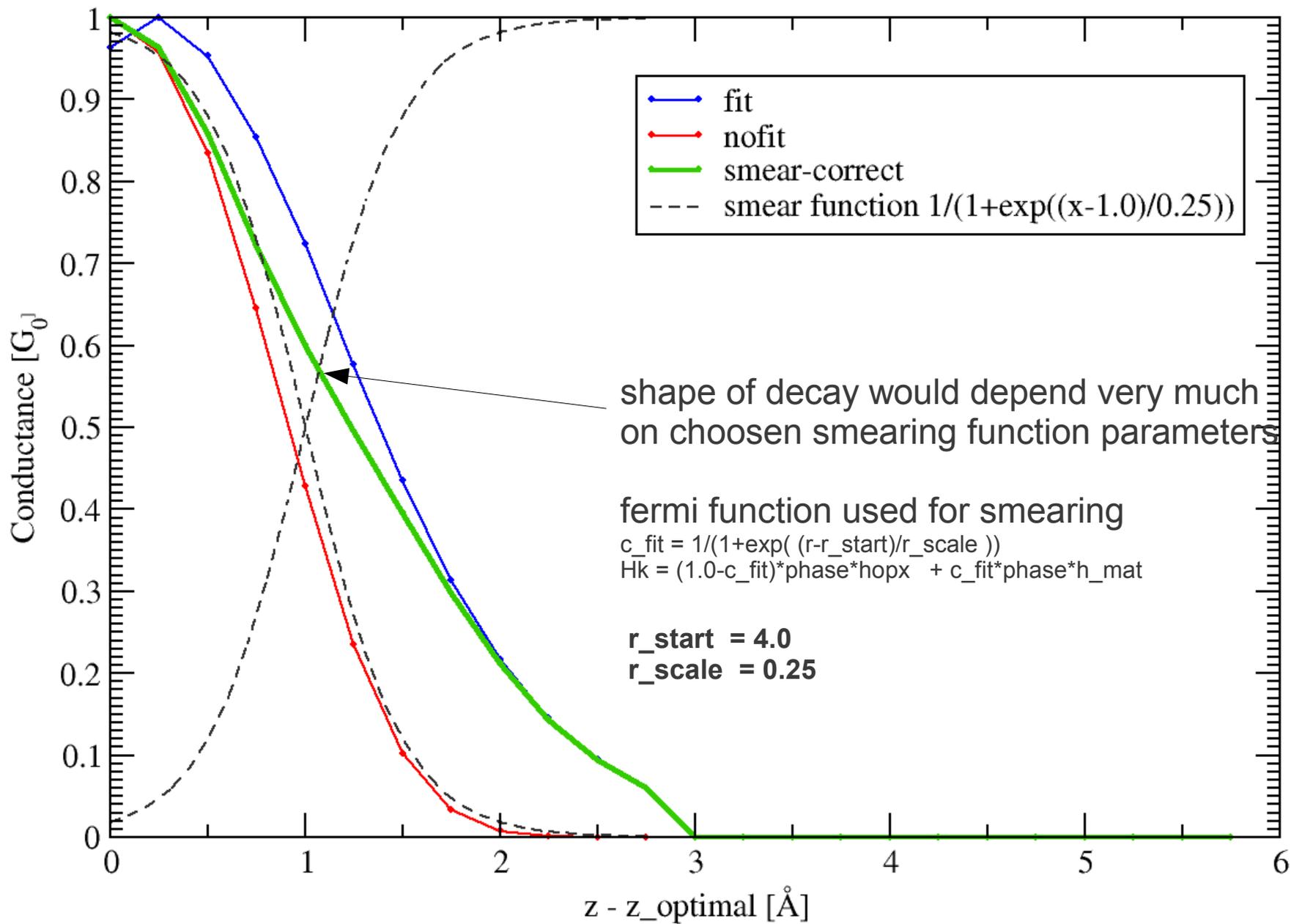


T(E) for 1D Gold chain by braking



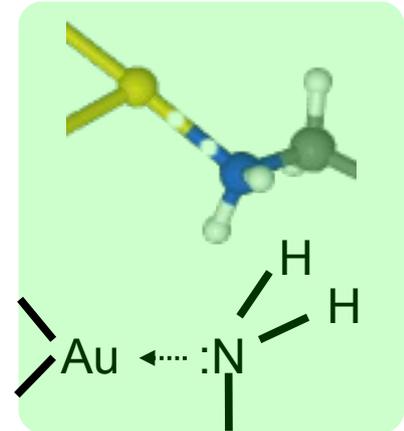
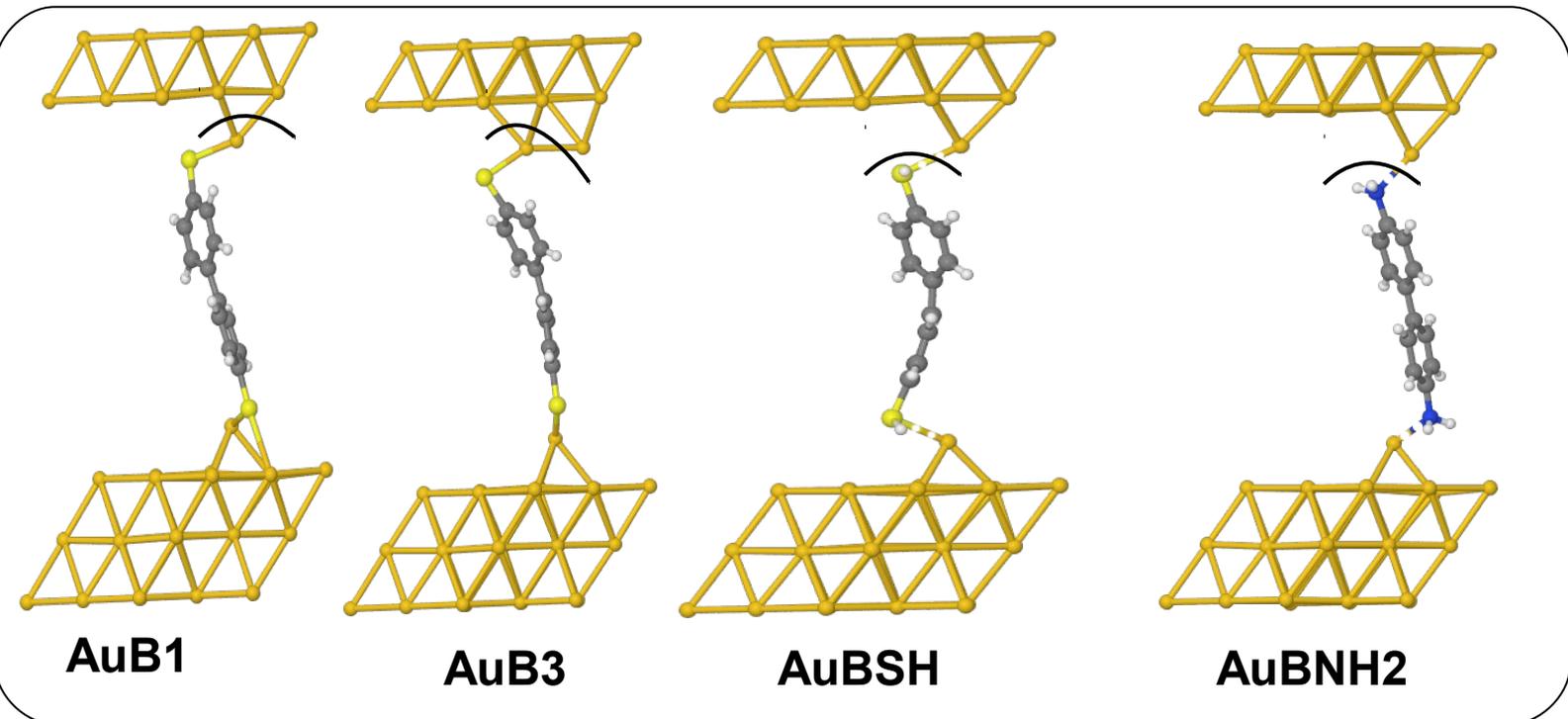
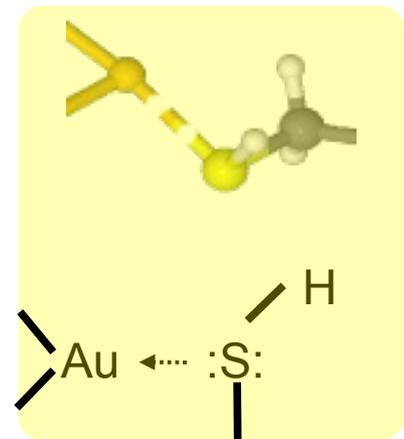
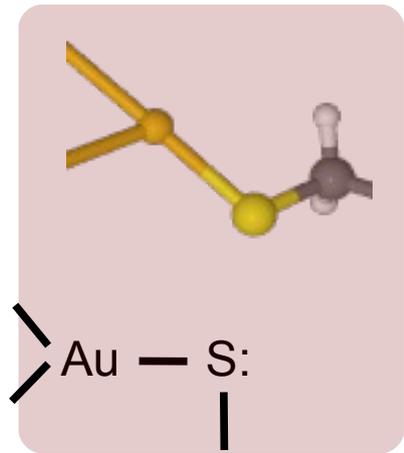
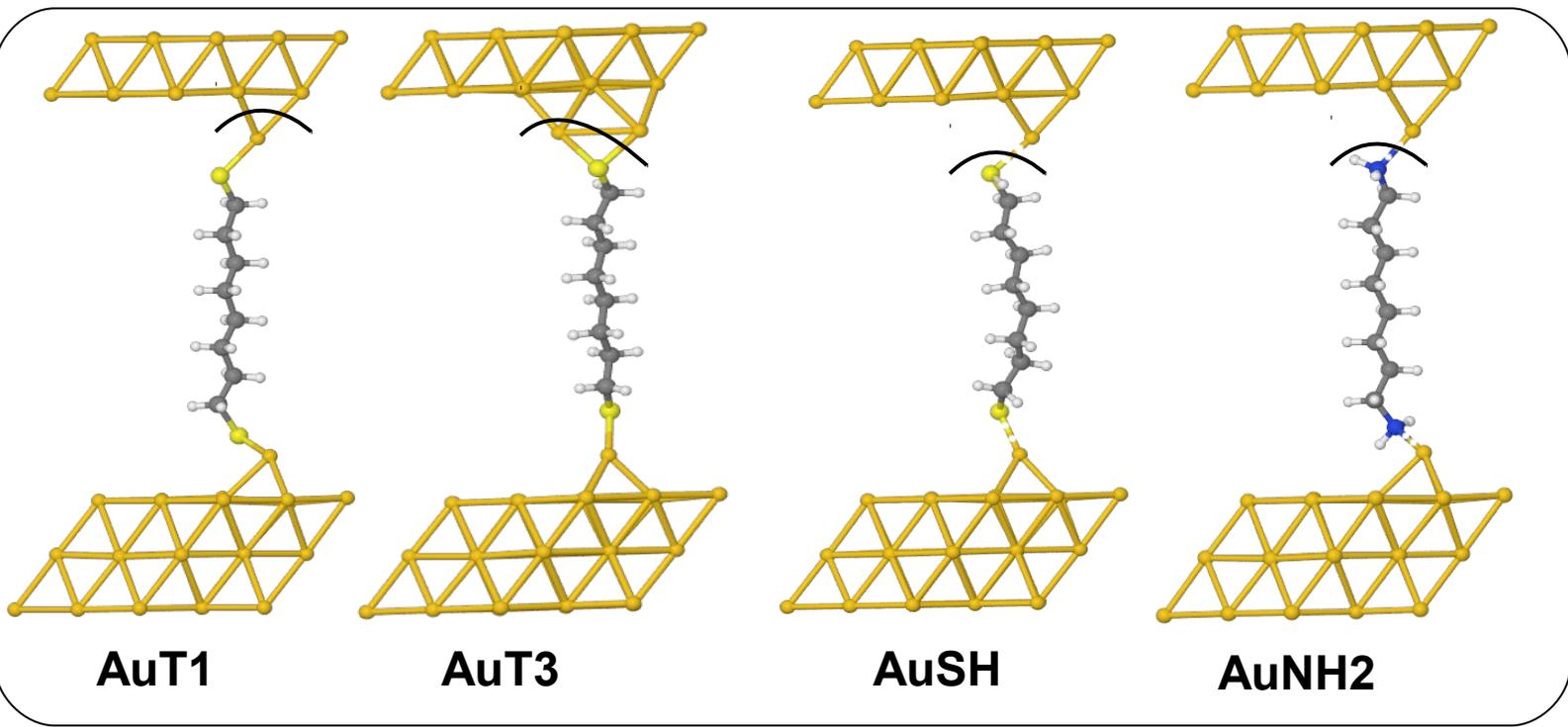
hopping fit smear Au-Chain 3.0Å

in +2,0 eV from fermi level

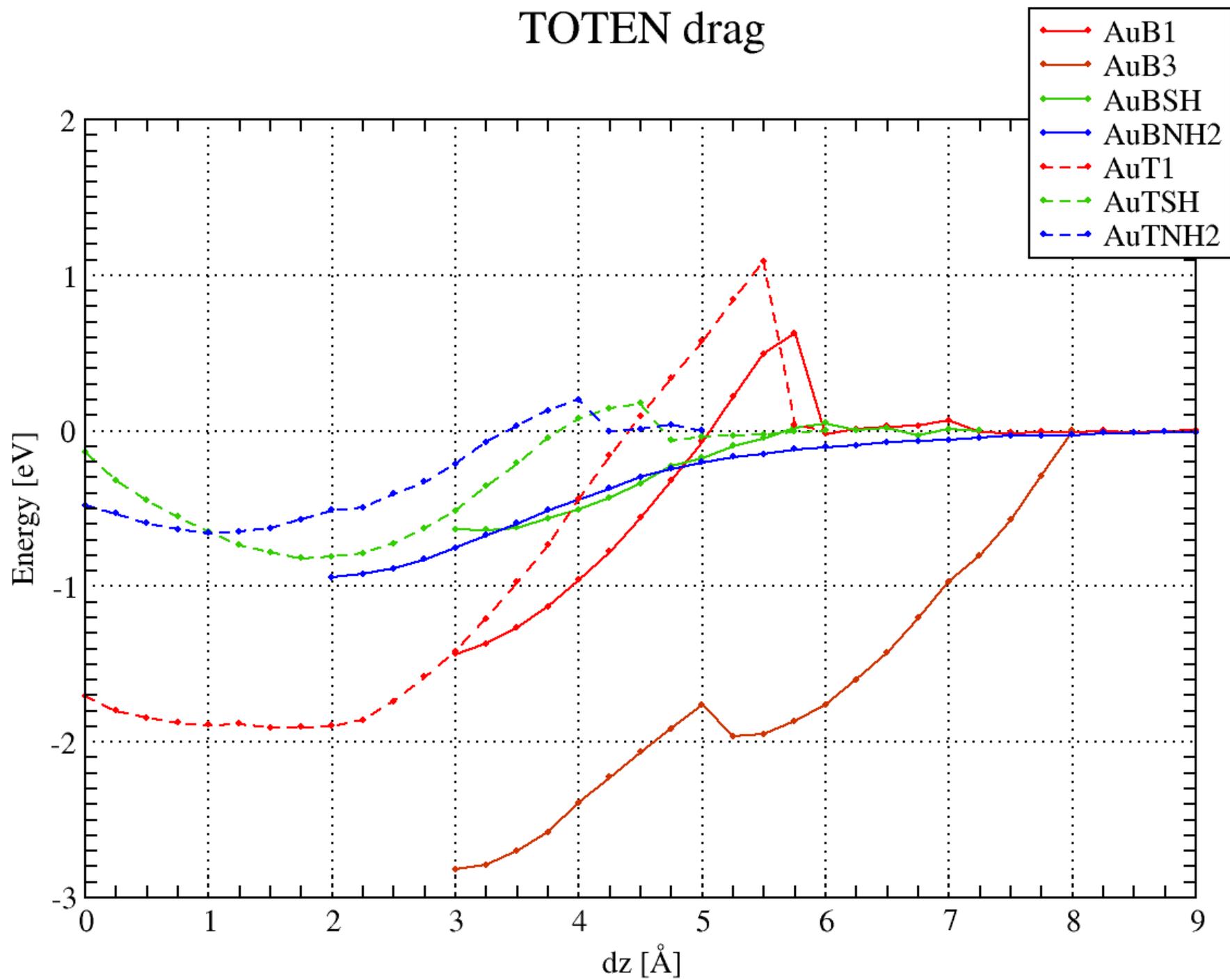


Example 3

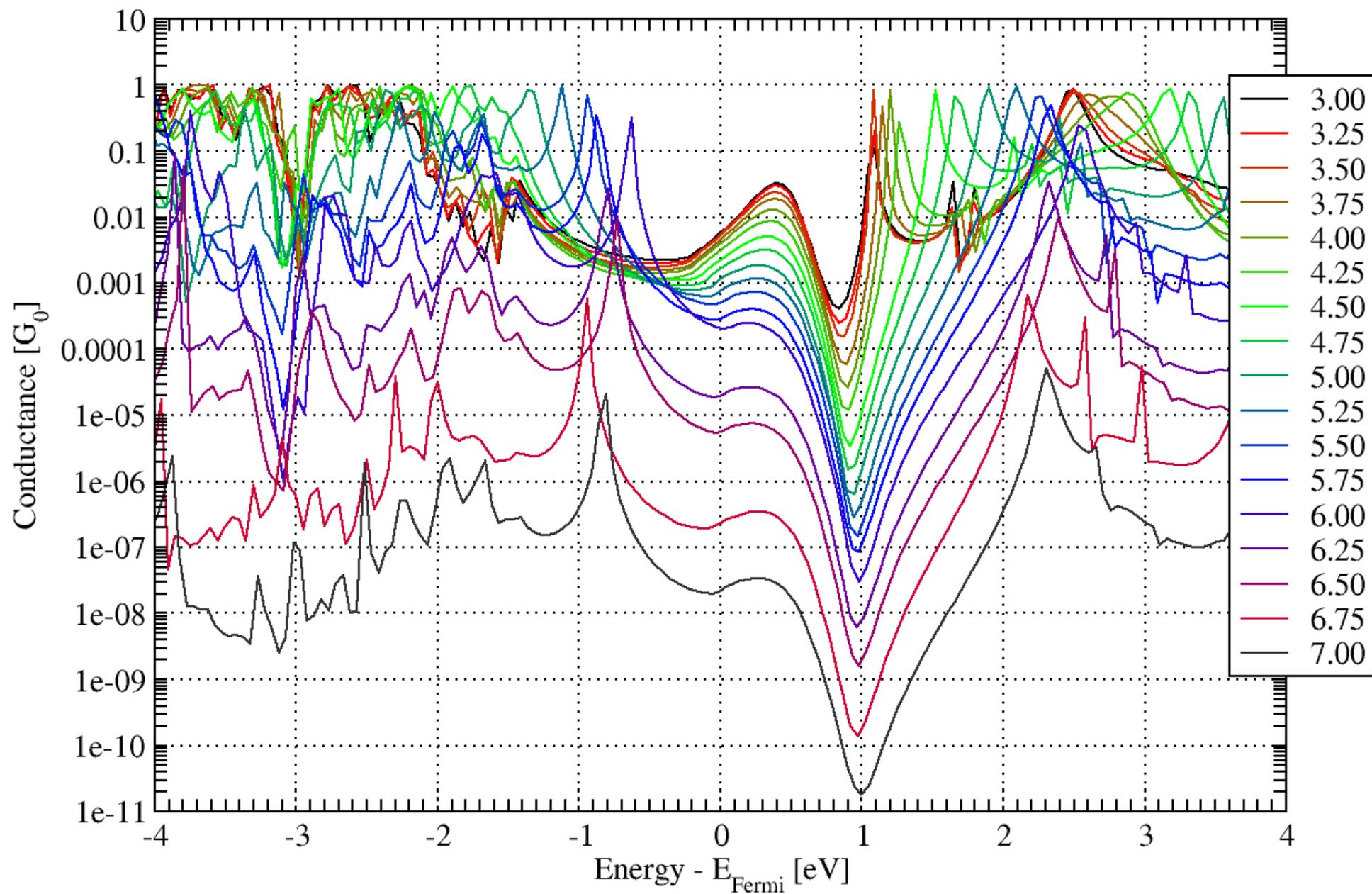
Conductance of molecules



TOTEN drag



Conductance drag AuBSH



AuBSH-drag I-z curve

