# A Few Hidden Features in Q-CHEM for Geometry Optimization, Wavefunction Analysis, and QM/MM Calculations

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- Potential Energy Scan
- 2 Electrostatic-Potential-Derived (ESP) Charges
- 3 Decomposition of TDDFT excitation energies
- 4 Multiple-Environment Single-System QM/MM
- General Research Interest

## Relaxed Potential Energy Scan

A series of constrained geom opt



 Initial angle: -180° Final angle: 180° Increment: 15°

- constraints: bond lengths, bond angles, torsional angles
- Initial value can be larger than the final value
- 2-d scan is allowed

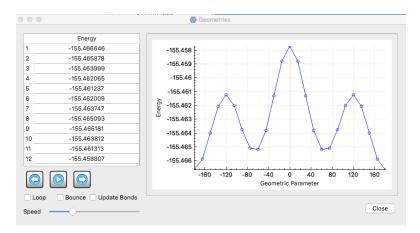
```
$molecule
0 1
                     -0.128781
        1.934574
                                   -0.000151
        0.556601
                      0.526657
                                    0.000200
       -0.556627
                     -0.526735
                                    0.000173
       -1.934557
                      0.128837
                                   -0.000138
Н
        2.720125
                      0.655980
                                   -0.000236
        2.061880
                     -0.759501
Н
                                   -0.905731
Н
        2 062283
                     -0.759765
                                    0.905211
        0.464285
                      1.168064
                                   -0.903444
Н
        0.464481
                      1.167909
                                    0.903924
Н
Н
       -0.464539
                     -1.167976
                                    0.903964
Н
       -0.464346
                     -1.168166
                                   -0.903402
       -2.062154
                      0.759848
                                    0.905185
                     -0.655832
       -2.720189
                                   -0.000229
       -2.061778
                      0.759577
                                   -0.905748
Send
```

```
$rem
jobtype pes_scan
method hf
basis sto-3g
$end
```

```
$scan
tors 1 2 3 4 -180 180 15
Send
```

## Relaxed Potential Energy Scan Results

IQmol visualization of 1-d scan results



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## Frozen Potential Energy Scan

 A series of single-point energy calculations



- Initial bond length: 1.0Å
- Final bond length: 2.0 Å
- Increment: 0.5 Å
- No geometry optimization
- Input geometry must be z-matrix

```
Smolecule
   C RCO
H1 C RCH1 O H1CO
  C 1.0
          O XCO
                      180.0
H2 C RCH2 X H2CX H1
                      90 0
H3 C RCH2 X H2CX H1
                      -90.0
                      180.0
H4 O ROH C HOC
     = 1.421
RCH1 = 1.094
RCH2 = 1.094
     = 0.963
H1CO = 107.2
XCO = 129.9
H2CX = 54 25
HOC = 108.0
Send
$rem
JOBTYPE
                  pes scan
FROZEN SCAN
                  TRUE
EXCHANGE
CORRELATION
                  WWW
BASTS
                  3 - 21G
$end
$scan
stre 1 2 1.0 2.0 0.5
$end
```

## Restrained Potential Energy Scan

A series of restrained geom opt



Geometry restraints

$$k\left(R_{12}-R_{16}-R\right)^2$$

- R<sub>min</sub>: -2.0Å • Rmax: 2.0 Å
- Increment: 0.2 Å
- Force constant (k): 1000.0
- 1-d scan only
- Can also use "r12pr34"

```
$molecule
            0.418808
                         -1.240869
                                        0.249048
           -0.775224
                         -1.495584
                                        1.586668
            1.408172
                         -1.490565
                                        0.631227
            0.147593
                         -1.907736
                                      -0.568952
            0.413296
                         -0.199000
                                      -0.092071
            1.947359
                          1.619163
                                      -1.747832
```

Send Śrem

jobtype pes\_scan exchange b3lyp basis 6-31G\*

Send

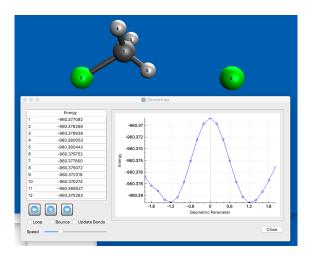
\$scan

r12mr34 1 2 1 6 -2 0 2 0 0 2 1000 0

\$end

## Restrained Potential Energy Scan Results

IQmol visualization of 1-d scan results



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## **Restrained Geometry Optimization**

```
$molecule
-1 1
   C
         0.4188082876 -1.2408695705 0.2490483370
   C1
       -0.7752244258 -1.4955845511 1.5866685479
          1.4081726568 -1.4905651530
                                         0.6312274608
   Н
   Н
          0.1475935065
                         -1.9077368904
                                        -0.5689527615
   Н
        0.4132963138
                         -0.1990006891
                                        -0.0920715962
          1.9473593327 1.6191631608
                                       -1.7478328574
Send
$rem
jobtype opt
exchange b3lyp
```

```
$opt2
r12mr34 1 2 1 6 0.0 1000.0
$end
```

basis 6-31G\*

\$end

- Potential Energy Scar
- Electrostatic-Potential-Derived (ESP) Charges
- 3 Decomposition of TDDFT excitation energies
- 4 Multiple-Environment Single-System QM/MM
- General Research Interest

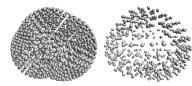
## Charge Populations in Q-CHEM

- Mulliken charges
  - Can be non-physical with larger basis sets
- NBO charges
- CHELPG charges (CHELPG = TRUE)
  - Well reproduce the dipole moment
- Hirshfeld charges (HIRSHFELD = TRUE)
  - Usually too small
- CM5 charges (CM5 = TRUE)
  - Improvement by Don Truhlar
- Iterative Hirshfeld charges (HIRSHITER = TRUE)
  - Much better than Hirshfeld charges

J. Phys. Chem. A, 119, 5865 (2015).

## ESP/RESP charge calculations

- ESP Grid Options:
  - 1: Spherical harmonics
  - 2: Lebedev



RESP CHARGES = TRUE

```
$molecule
0 1
            1.22227
                       -0.2466
                                   0.0178
            2.09272
                        0.3971
                                  -0.0423
                       -0.9548
            1.27594
                                  -0.8023
            1.26663
                       -0.7997
                                  0.9461
           -0.07041
                       0.5616
                                  -0.0459
                       1.1336
           -0.12063
                                  -0.9663
           -0.12012
                       1.2567
                                  0.7739
           -1.24418
                       -0.2540
                                  0.1065
           -1.35216
                       -0.8912
                                  -0.5924
$end
Śrem
   BASIS
                = 6-31G*
  METHOD
                = B3LYP
  JOBTYPE
                = SP
  ESP CHARGES
                       # or 2
```

\$end

## ESP charges with different grids

- Dipole moment: X 1.0606; Y -0.2250; Z -1.2090; Tot 1.6239
- ESP\_Charges = 1

Merz-Kollman ESP Net Atomic Charges

Ato	mc	Charge (a.u.)
2 3 4 5 6 7	С Н Н С С Н	-0.359692 0.075218 0.088502 0.105036 0.356774 -0.035130 0.032457
9	Н	-0.673003 0.409839

Sum of atomic charges = 0.000000

Related Dipole Moment = 1.6305

Related Dipole Moment = 1.63(X 1.0669 Y -0.2059 Z -1.2157)

#### ESP\_Charges = 2

Merz-Kollman ESP Net Atomic Charges

Atom	Charge (a.u.)
1 C	-0.362941
2 H	0.076361
3 H	0.090774
4 H	0.107454
5 C	0.338283
6 н	-0.028913
7 н	0.038096
8 0	-0.669108
9 н	0.409994
Sum of atomic charges =	0.000000

Related Dipole Moment = 1.6285 (X 1.0631 Y -0.2069 Z -1.2162)

## ESP and RESP charges

- Dipole moment: X 1.0606; Y -0.2250; Z -1.2090; Tot 1.6239
- ESP\_Charges = 1

Merz-Kollman ESP Net Atomic Charges

Ato	m	Charge (a.u.)
1 2 3 4 5 6	Н Н Н	-0.359692 0.075218 0.088502 0.105036 0.356774 -0.035130 0.032457
8	0	-0.673003
-	Н	0.409839

Sum of atomic charges = 0.000000

Related Dipole Moment = 1.6305 (X 1.0669 Y -0.2059 Z -1.2157) RESP\_Charges = 1

Merz-Kollman RESP Net Atomic Charges

Atom	Charge (a.u.)
1 C	-0.272946
2 H	0.067338
3 H	0.067338
4 H	0.067338
5 C	0.346165
6 H	-0.015626
7 H	-0.015626
8 0	-0.608147
9 H	0.364167
Sum of atomic char	ges = 0.000000

Related Dipole Moment = 1.7694 (X 1.0670 Y -0.1781 Z -1.4002)

## ESP charges for TDDFT excited states

```
$molecule
        -0.00000 0.00000 0.7625
         0.00000 0.00000 -0.6375
   H 0.86602 -0.00000 -1.1375
        -0.86602 0.00000 -1.1375
$end
$rem
         = 6-31G*
  BASTS
  METHOD = B3LYP
  JOBTYPE = SP
  CIS N ROOTS = 3
  SET TRIP
             = FALSE
  RPA
             = 2
                      # No TDA
  POP MULLIKEN = -1
                      # Mulliken charges for all states
  ESP CHARGES = 1
                      # ESP charges for all states
  CIS RLX DNS
             = 1
                      # We should use the relaxed density
Send
```

## ESP and RESP charges

#### CIS\_RLX\_DNS = 0

Merz-Kollman ESP Net Atomic Charges

Atom	Charge (a.u.)		
1 O 2 C 3 H 4 H	-0.365648 0.185380 0.090134 0.090134		
Sum of atomic charges =	0.000000		
Related Dipole Moment = 2.8917 (X 0.0000 Y 0.0000 Z -2.8917)			
ESP charges for excited	states		

	1	2	3
1	0.2938661	0.2277436	-0.0598282
2	-0.8473374	-0.8761796	0.0035142
3	0.2767357	0.3242180	0.0281570
4	0 2767357	0 3242180	0.0281570

#### CIS\_RLX\_DNS = 1

Merz-Kollman ESP Net Atomic Charges

Atom	Charge (a.u.)
1 O 2 C 3 H 4 H	-0.365648 0.185380 0.090134 0.090134
Sum of atomic charges	= 0.000000
Related Dipole Moment	= 2.8917

(X 0.0000 Y 0.0000 Z -2.8917)

#### ESP charges for excited states

	1	2	3
1	-0.0914159	-0.0518477	-0.3150189
2	-0.3737757	-0.5369765	0.2755906
3	0.2325958	0.2944121	0.0197142
4	0.2325958	0.2944121	0.0197141

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## **TDDFT** excitation energies

TDDFT working equation is

$$\begin{pmatrix} \mathbf{A} & \mathbf{B} \\ \mathbf{B} & \mathbf{A} \end{pmatrix} \begin{pmatrix} \mathbf{X} \\ \mathbf{Y} \end{pmatrix} = \Delta E \begin{pmatrix} \mathbf{X} \\ -\mathbf{Y} \end{pmatrix}$$

Hirata and Head-Gordon, Chem. Phys. Lett. 302, 375 (1999);

## **TDDFT** excitation energies

TDDFT working equation is

$$\begin{pmatrix} \mathbf{A} & \mathbf{B} \\ \mathbf{B} & \mathbf{A} \end{pmatrix} \begin{pmatrix} \mathbf{X} \\ \mathbf{Y} \end{pmatrix} = \Delta E \begin{pmatrix} \mathbf{X} \\ -\mathbf{Y} \end{pmatrix}$$

We can write the vertical excitation energy (VEE) as

$$\Delta E = \Delta E_1 + \Delta E_2$$

$$\Delta E_1 = \sum_{ai} (X_{ai}^2 + Y_{ai}^2) (\varepsilon_a - \varepsilon_i)$$

$$\Delta E_2 = \Delta E_{2,J} + \Delta E_{2,K} + \Delta E_{2,XC}$$

Hirata and Head-Gordon, Chem. Phys. Lett. 302, 375 (1999); Mol. Phys. 108, 2791 (2010).

## **TDDFT** excitation energies

TDDFT working equation is

$$\begin{pmatrix} \mathbf{A} & \mathbf{B} \\ \mathbf{B} & \mathbf{A} \end{pmatrix} \begin{pmatrix} \mathbf{X} \\ \mathbf{Y} \end{pmatrix} = \Delta E \begin{pmatrix} \mathbf{X} \\ -\mathbf{Y} \end{pmatrix}$$

We can write the vertical excitation energy (VEE) as

$$\Delta E = \Delta E_1 + \Delta E_2$$

$$\Delta E_1 = \sum_{ai} (X_{ai}^2 + Y_{ai}^2) (\varepsilon_a - \varepsilon_i)$$

$$\Delta E_2 = \Delta E_{2,J} + \Delta E_{2,K} + \Delta E_{2,XC}$$

- VEE can be decomposed for
  - pure and hybrid functionals
  - two range-separated functionals:  $\omega$ B97X-D and LRC- $\omega$ PBEh
  - Results are slightly off for pure basis sets

Hirata and Head-Gordon, Chem. Phys. Lett. 302, 375 (1999); Mol. Phys. 108, 2791 (2010).

## Sample Input: H<sub>2</sub>CO

```
$molecule
0 1
C, 1, R1
H, 2, R2, 1, A
H, 2, R2, 1, A, 3, 180.
R1 = 1.4
R2=1.0
A=120.
$end
$rem
met hod
                                    wR97X-D
basis
                                     6-31++G**
                                     5
cis_n_roots
rpa
cis dynamic mem
                                    false
                                                #use old TDDFT implementation
excit_energy_components
                                    true
$end
```

#### HOMO: -0.369; LUMO: -0.028; Gap: 9.279 eV

#### Output file:

```
Excited state 1: excitation energy (eV) = 2.1475
Total energy for state 1:
                                   -114.33552540 au
   Multiplicity: Triplet
   Trans. Mom.: 0.0000 X 0.0000 Y 0.0000 Z
   Strength : 0.0000000000
   X: D( 8) --> V( 1) amplitude = 0.9968
   Details -- H: 26.7523 J1: -28.3251 K1: 8.1581 XC1: 3.0430
             J2: 0.0000 K2: -7.1299 XC2: -0.3510 Total: 2.1475
   dE1 = 9.6283 \text{ eV}; dE2 = -7.4809 \text{ eV} #not shown in output
Excited state 3: excitation energy (eV) = 2.8774
Total energy for state 3:
                                         -114.30870220 au
   Multiplicity: Singlet
   Trans. Mom.: -0.0000 X -0.0000 Y -0.0000 Z
   Strength : 0.0000000000
   X: D( 8) --> V( 1) amplitude = 0.9963
   Details -- H: 28.4549 J1: -30.3332 K1: 8.1053 XC1: 3.2048
             J2: 0.6325 K2: -6.9064 XC2: -0.2806 Total: 2.8774
   dE1 = 9.4318 \text{ eV}; dE2 = -6.5545 \text{ eV} #not shown in output
```

- HOMO: -0.369; LUMO: -0.028; Gap: 9.279 eV
- Output file:

```
Excited state 1: excitation energy (eV) = 2.1475
                             -114.33552540 au
Total energy for state 1:
   Multiplicity: Triplet
   Trans. Mom.: 0.0000 X 0.0000 Y 0.0000 Z
   Strength : 0.0000000000
   X: D( 8) --> V( 1) amplitude = 0.9968
   Details -- H: 26.7523 J1: -28.3251 K1: 8.1581 XC1: 3.0430
             J2: 0.0000 K2: -7.1299 XC2: -0.3510 Total: 2.1475
   dE1 = 9.6283 \text{ eV}; dE2 = -7.4809 \text{ eV} #not shown in output
Excited state 3: excitation energy (eV) = 2.8774
Total energy for state 3:
                             -114.30870220 au
   Multiplicity: Singlet
   Trans. Mom.: -0.0000 X -0.0000 Y -0.0000 Z
   Strength : 0.0000000000
   X: D( 8) --> V( 1) amplitude = 0.9963
   Details -- H: 28.4549 J1: -30.3332 K1: 8.1053 XC1: 3.2048
             J2: 0.6325 K2: -6.9064 XC2: -0.2806 Total: 2.8774
   dE1 = 9.4318 \text{ eV}; dE2 = -6.5545 \text{ eV} #not shown in output
```

HOMO-LUMO gap >> lowest singlet/triplet excitation energies

#### Why is the triplet state lower in energy?

```
Excited state 1: excitation energy (eV) = 2.1475
Total energy for state 1:
                                -114.33552540 au
   Multiplicity: Triplet
   Trans. Mom.: 0.0000 X 0.0000 Y 0.0000 Z
   Strength: 0.0000000000
   X: D( 8) --> V( 1) amplitude = 0.9968
   Details -- H: 26.7523 J1: -28.3251 K1: 8.1581 XC1: 3.0430
             J2: 0.0000 K2: -7.1299 XC2: -0.3510 Total: 2.1475
   dE1 = 9.6283 \text{ eV}; dE2 = -7.4809 \text{ eV} #not shown in output
Excited state 3: excitation energy (eV) = 2.8774
Total energy for state 3:
                                -114.30870220 au
   Multiplicity: Singlet
   Trans. Mom.: -0.0000 X -0.0000 Y -0.0000 Z
   Strength: 0.0000000000
   X: D( 8) --> V( 1) amplitude = 0.9963
   Details -- H: 28.4549 J1: -30.3332 K1: 8.1053 XC1: 3.2048
             J2: 0.6325 K2: -6.9064 XC2: -0.2806 Total: 2.8774
   dE1 = 9.4318 \text{ eV}; dE2 = -6.5545 \text{ eV} #not shown in output
```

#### Why is the triplet state lower in energy?

```
Excited state 1: excitation energy (eV) = 2.1475
Total energy for state 1:
                                          -114.33552540 au
   Multiplicity: Triplet
   Trans. Mom.: 0.0000 X 0.0000 Y 0.0000 Z
   Strength: 0.0000000000
   X: D( 8) --> V( 1) amplitude = 0.9968
   Details -- H: 26.7523 J1: -28.3251 K1: 8.1581 XC1: 3.0430
              J2: 0.0000 K2: -7.1299 XC2: -0.3510 Total: 2.1475
   dE1 = 9.6283 \text{ eV}; dE2 = -7.4809 \text{ eV} #not shown in output
Excited state 3: excitation energy (eV) = 2.8774
Total energy for state 3:
                                  -114.30870220 au
   Multiplicity: Singlet
   Trans. Mom.: -0.0000 X -0.0000 Y -0.0000 Z
   Strength: 0.0000000000
   X: D(8) \longrightarrow V(1) \text{ amplitude} = 0.9963
   Details -- H: 28.4549 J1: -30.3332 K1: 8.1053 XC1: 3.2048
              J2: 0.6325 K2: -6.9064 XC2: -0.2806 Total: 2.8774
   dE1 = 9.4318 \text{ eV}; dE2 = -6.5545 \text{ eV} #not shown in output
```

#### • The J2 tem is

$$\Delta E_{2,J} = \int \int \rho_{\rm trans}(\mathbf{r}_1) \frac{1}{r_{12}} \rho_{\rm trans}(\mathbf{r}_2) d\mathbf{r}_1 d\mathbf{r}_2$$

#### Why is the triplet state lower in energy?

```
Excited state 1: excitation energy (eV) = 2.1475
                             -114.33552540 au
Total energy for state 1:
   Multiplicity: Triplet
   Trans. Mom.: 0.0000 X 0.0000 Y 0.0000 Z
   Strength: 0.0000000000
   X: D( 8) --> V( 1) amplitude = 0.9968
   Details -- H: 26.7523 J1: -28.3251 K1: 8.1581 XC1: 3.0430
             J2: 0.0000 K2: -7.1299 XC2: -0.3510 Total: 2.1475
   dE1 = 9.6283 \text{ eV}; dE2 = -7.4809 \text{ eV} #not shown in output
Excited state 3: excitation energy (eV) = 2.8774
Total energy for state 3:
                                 -114.30870220 au
   Multiplicity: Singlet
   Trans. Mom.: -0.0000 X -0.0000 Y -0.0000 Z
   Strength: 0.0000000000
   X: D(8) \longrightarrow V(1) \text{ amplitude} = 0.9963
   Details -- H: 28.4549 J1: -30.3332 K1: 8.1053 XC1: 3.2048
             J2: 0.6325 K2: -6.9064 XC2: -0.2806 Total: 2.8774
   dE1 = 9.4318 \text{ eV}; dE2 = -6.5545 \text{ eV} #not shown in output
```

• The J2 tem is

$$\Delta E_{2,J} = \int \int \rho_{\rm trans}(\mathbf{r}_1) \frac{1}{r_{12}} \rho_{\rm trans}(\mathbf{r}_2) d\mathbf{r}_1 d\mathbf{r}_2$$

• The transition density,  $\rho_{\rm trans}({\bf r})$ , is zero for triplet excitations.

#### TDDFT with B3LYP functional:

#### • TDDFT with $\omega$ B97X-D functional:

```
Excited state 3: excitation energy (eV) = 2.8774

Total energy for state 3: -114.30870220 au

Multiplicity: Singlet

Trans. Mom.: -0.0000 X -0.0000 Y -0.0000 Z

Strength : 0.00000000000

X: D(8) --> V(1) amplitude = 0.9963

Details -- H: 28.4549 JI: -30.3332 K1: 8.1053 XC1: 3.2048

J2: 0.6325 K2: -6.9064 XC2: -0.2806 Total: 2.8774

dE1 = 9.4318 eV; dE2 = -6.5545 eV #not shown in output

HOMO: -0.369; LUMO: -0.028; Gap: 9.279 eV #not shown in output
```

#### TDDFT with B3LYP functional:

```
Excited state 3: excitation energy (eV) = 2.8422

Total energy for state 3: -114.21958212 au

Multiplicity: Singlet

Trans. Mom.: 0.0000 X -0.0000 Y -0.0000 Z

Strength : 0.0000000000

X: D( 8) --> V( 1) amplitude = 0.9998

Details -- H: 28.8102 J1: -30.6356 K1: 4.2395 XC1: 3.0889

J2: 0.6445 K2: -3.0329 XC2: -0.2723 Total: 2.8422

dE1 = 5.5030 eV; dE2 = -2.6607 eV #not shown in output

HOMO: -0.297; LUMO: -0.098; Gap: 5.415 eV #not shown in output
```

#### • TDDFT with $\omega$ B97X-D functional:

```
Excited state 3: excitation energy (eV) = 2.8774

Total energy for state 3: -114.30870220 au

Multiplicity: Singlet

Trans. Mom.: -0.0000 X -0.0000 Y -0.0000 Z

Strength : 0.00000000000

X: D( 8) --> V( 1) amplitude = 0.9963

Details -- H: 28.4549 Jl: -30.3332 Kl: 8.1053 XCl: 3.2048

J2: 0.6325 K2: -6.9064 XC2: -0.2806 Total: 2.8774

dEl = 9.4318 eV; dE2 = -6.5545 eV #not shown in output

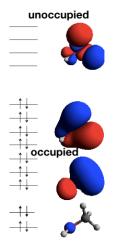
HOMO: -0.369; LUMO: -0.028; Gap: 9.279 eV #not shown in output
```

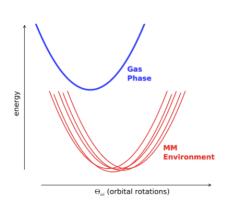
#### The K2 term is

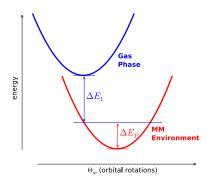
$$\Delta E_{2,K}(\text{PBE0}) = -\int\int\rho_{\text{detach}}(\mathbf{r}_1)\frac{0.25}{r_{12}}\rho_{\text{attach}}(\mathbf{r}_2)d\mathbf{r}_1d\mathbf{r}_2$$
 
$$\Delta E_{2,K}(\omega \text{B97X} - \text{D}) = -\int\int\rho_{\text{detach}}(\mathbf{r}_1)\frac{0.22\,\text{erfc}(\omega r_{12}) + \text{erf}(\omega r_{12})}{r_{12}}\rho_{\text{attach}}(\mathbf{r}_2)d\mathbf{r}_1d\mathbf{r}_2$$

- Potential Energy Scar
- 2 Electrostatic-Potential-Derived (ESP) Charges
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- Multiple-Environment Single-System QM/MM
- General Research Interest

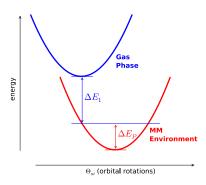
## Multiple-Environment Single-System QM/MM (MESS-QM/MM): QM/MM Calculations with a Fixed-Geometry QM Region



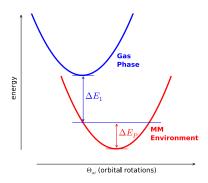




Assume a quadratic surface

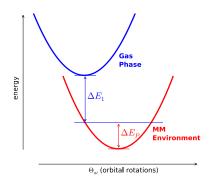


- Assume a quadratic surface
- Compute the Hessian for a reference environment



- Assume a quadratic surface
- Compute the Hessian for a reference environment
- For each MM environment
  - Compute the gradient (*g*)
  - Take a Newton step

$$\Delta\Theta = \mathbf{H}^{-1} \cdot g$$

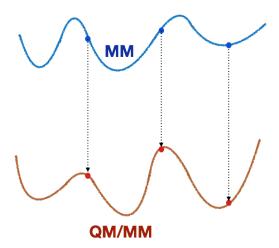


- Potential applications:
  - solvation free energies
  - enzymatic reaction free energy profiles
  - polarizable force-field development

## **Hydration Free Energy Calculations**

Yihan Shao (OU)

## Computational Strategy



Step 1: Compute hydration free energy using MM energy functions. Solvent: >1500 TIP3P water

Step 2: Compute QM/MM energy on some MM configurations. Obtain QM/MM correction to the hydration free energy. QM region: solute

## QM/MM Hydration Free Energy Calculations

- Computational Efficiency
  - MM: 100 CPU Hours
  - QM/MM correction: 400-2700 CPU Hours

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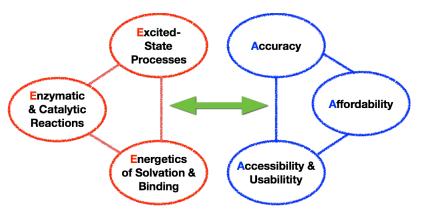
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- More sophisticated functionals yielded worse results.
- We also need to improve QM/MM vdW interactions.

- Potential Energy Scan
- Electrostatic-Potential-Derived (ESP) Charges
- 3 Decomposition of TDDFT excitation energies
- 4 Multiple-Environment Single-System QM/MM
- General Research Interest

### Computational Chemistry: Applications, Theory and Software



### Main Topics

- Fluorescence/bioluminescence, reaction rates, binding affinity
- Chemical, solvent, macromolecular, methodological perturbations

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- DOE, SBIR Phase I & II
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- University of Oklahoma