

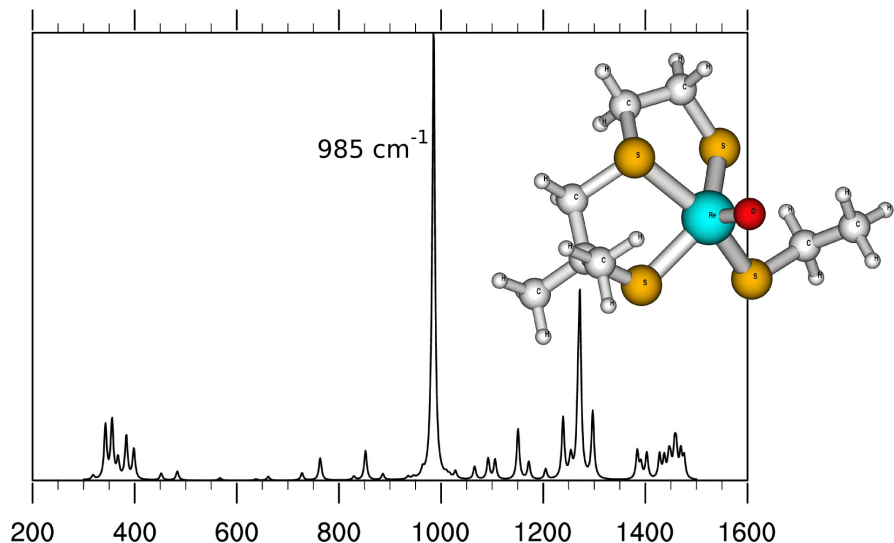
Parity violation in molecular systems: what we can learn from theory

Trond Saue

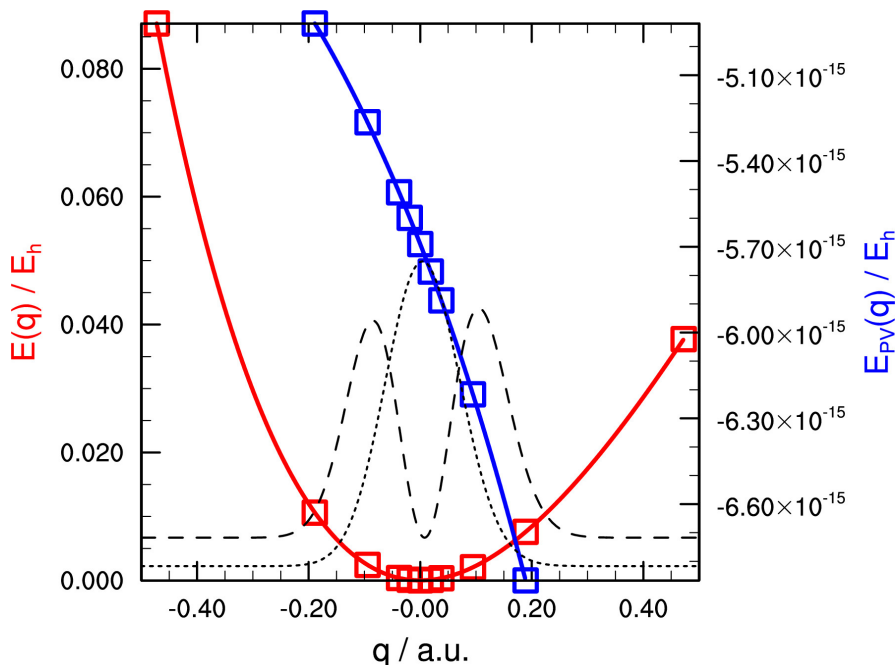
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Reminder: Computational procedure



1. **Geometry optimization + rovibrational analysis** (gaussian/B3LYP): the infrared spectrum is simulated using calculated frequencies and intensities



2. Calculation of total energy $E(q)$ and PV energy contribution $E_{PV}(q)$ along selected vibrational mode (DIRAC/X2C/B3LYP or HF)

3. Numerical solution of vibrational problem (Numerov-Cooley) $\rightarrow \psi_\nu^{\text{vib}}$

4. Integration of $E_{PV}(q)$ with vibrational solutions:
$$E_{PV}(\nu) = \langle \psi_\nu^{\text{vib}} | E_{PV}(q) | \psi_\nu^{\text{vib}} \rangle$$

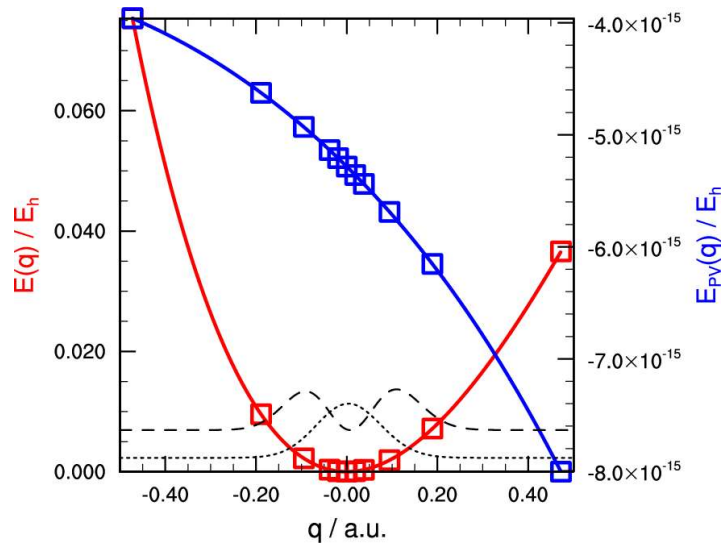
Reminder: Perturbational approach

- Series expansion:

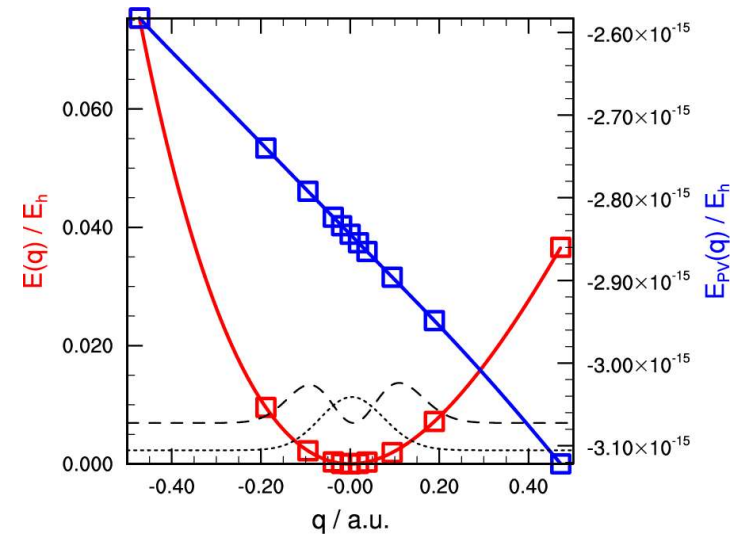
$$E(q) = V(q=0) + V^{[1]}q + \frac{1}{2}V^{[2]}q^2 + \dots \quad E_{PV}(q) = P(q=0) + P^{[1]}q + \frac{1}{2}P^{[2]}q^2 + \dots$$

- PV shift

$$\Delta\nu_{RS} = \frac{\hbar}{\sqrt{\mu V^{[2]}}} \left\{ P^{[2]} - \frac{V^{[3]}}{V^{[2]}} P^{[1]} \right\}$$



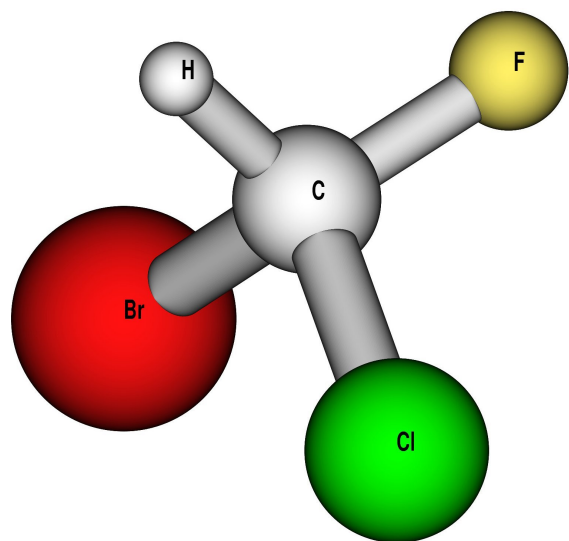
5b (HF): $\Delta_{RS} = -1020.7$ mHz
(harmonic: -390.4 mHz)



5b (B3LYP): $\Delta_{RS} = -84$ mHz
(harmonic: +0.02 mHz)

PV shifts for all normal modes of CHFCIBr

B3LYP	$\bar{\nu}(\text{cm}^{-1})$	μ (amu)	Displacement (mHz)		Attribution
			Total	Harmonic	
1	217.7	26.6781	-1.186		Bending Cl - C - Br
2	307.2	21.0297	9.85	2.184	Bending F - C - Br
3	418.9	17.6804	-5.774	-2.126	Bending Cl - C - F
4	642.2	5.6584	-5.807	-3.65	Stretching C - Br
5	744.2	8.1195	12.489	9.715	Stretching C - Cl
6	1092.3	11.3289	3.432	-1.876	Stretching C - F
7	1215.3	1.0943	1.261	0.942	Bending H - C - Br
8	1328.0	1.1154	-3.436	-1.638	Bending F - C - H
9	3150.4	1.0885	-1.649	-0.097	Stretching H - C

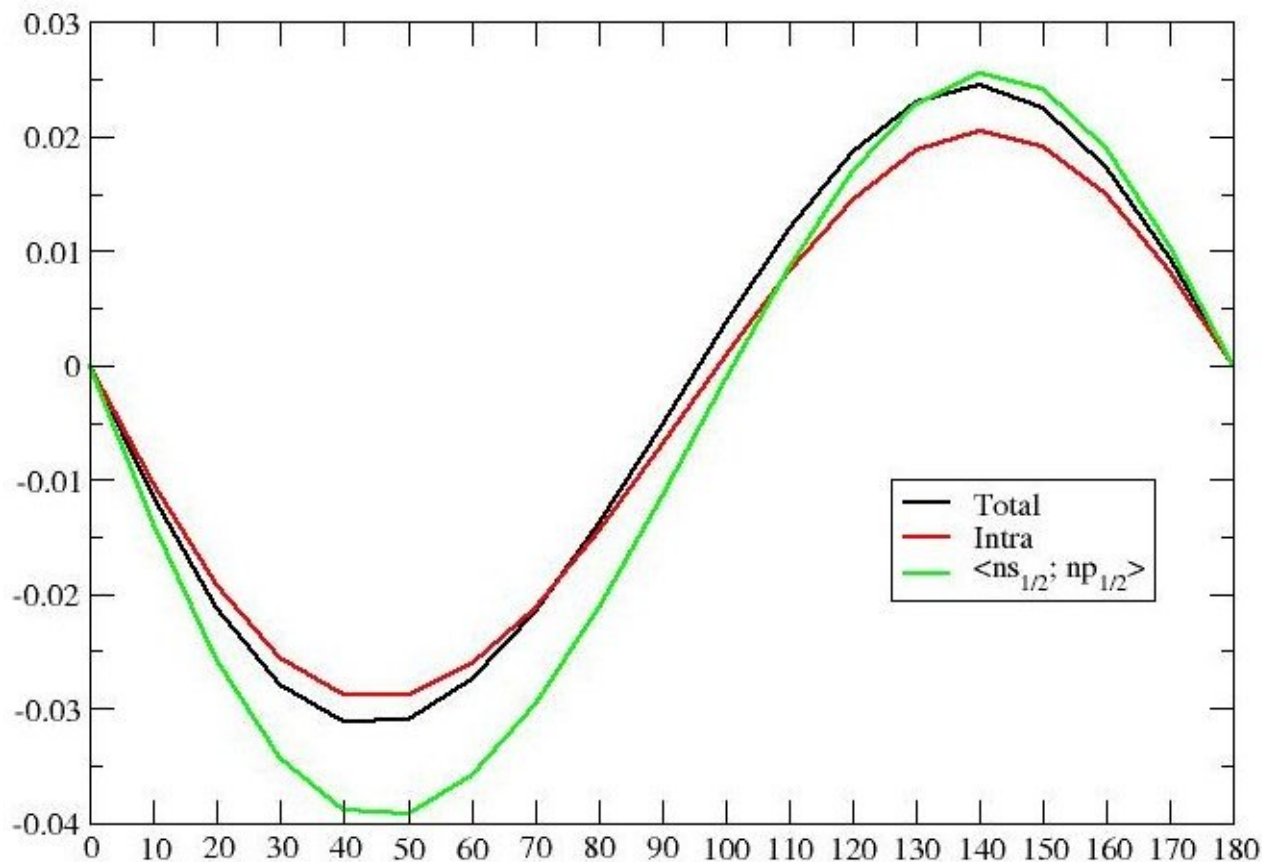


Sampling point must be adapted to curvature of potential curve; use classical turning points

$$x_{max} = \sqrt{\frac{2\hbar}{\mu\omega} \left(\nu + \frac{1}{2} \right)}$$

Make robust scheme for numerical differentiation.

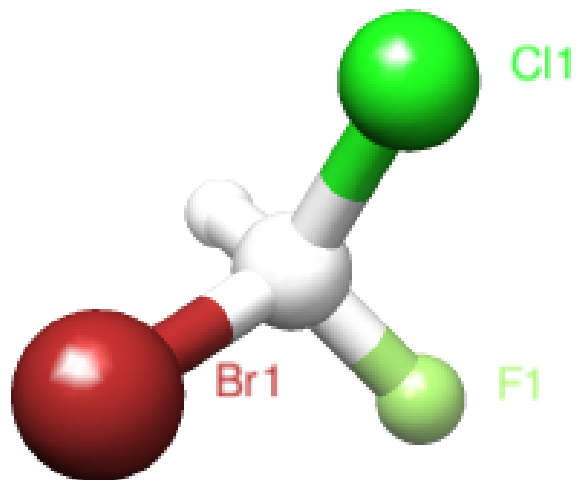
Reminder: Towards deeper understanding



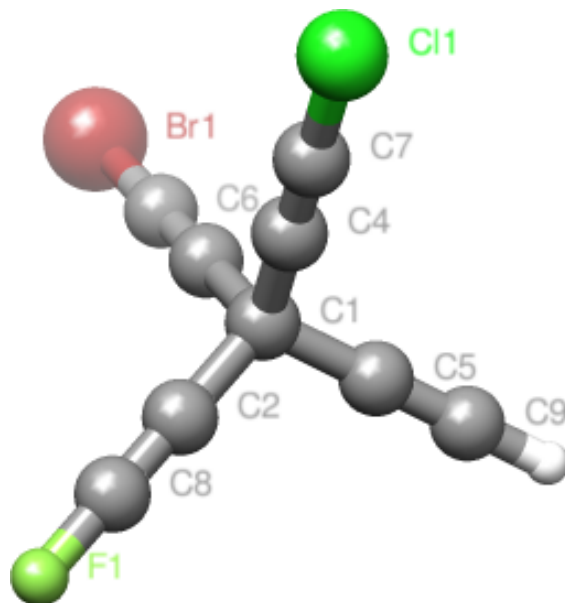
HF	H_2O_2	H_2S_2	H_2Se_2	H_2Te_2	H_2Po_2
Full	-6.729E-06	-7.435E-05	-3.163E-03	-2.787E-02	-7.955E-01
Intra	-5.879E-06	-6.876E-05	-2.717E-03	-2.459E-02	-6.334E-01
$\langle ns_{1/2} \hat{H}_{pv}^X np_{1/2} \rangle$	-8.819E-06	-8.548E-05	-3.773E-03	-3.216E-02	-7.728E-01

A non-zero contribution requires an imaginary phase in the mixing coefficient between $s_{1/2}$ and $p_{1/2}$.

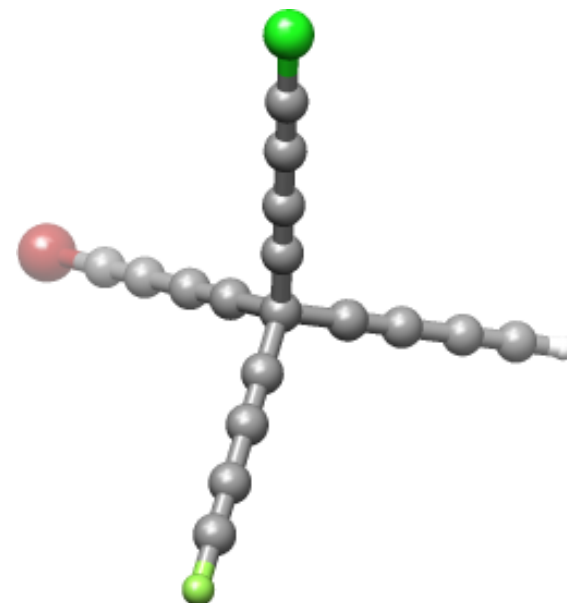
Exploring the chiral field: carbomers



$r(\text{C1-Br})$: 1.945 Å
 $r(\text{C1-Cl})$: 1.766 Å
 $r(\text{C1-F})$: 1.346 Å
 $r(\text{C1-H})$: 1.087 Å
Total energy: -3206.489586 E_h
HOMO-LUMO gap:
0.248490 E_h
Total EPV = 1.001E-18 E_h
EPV (central carbon C1):
5.523E-20 E_h



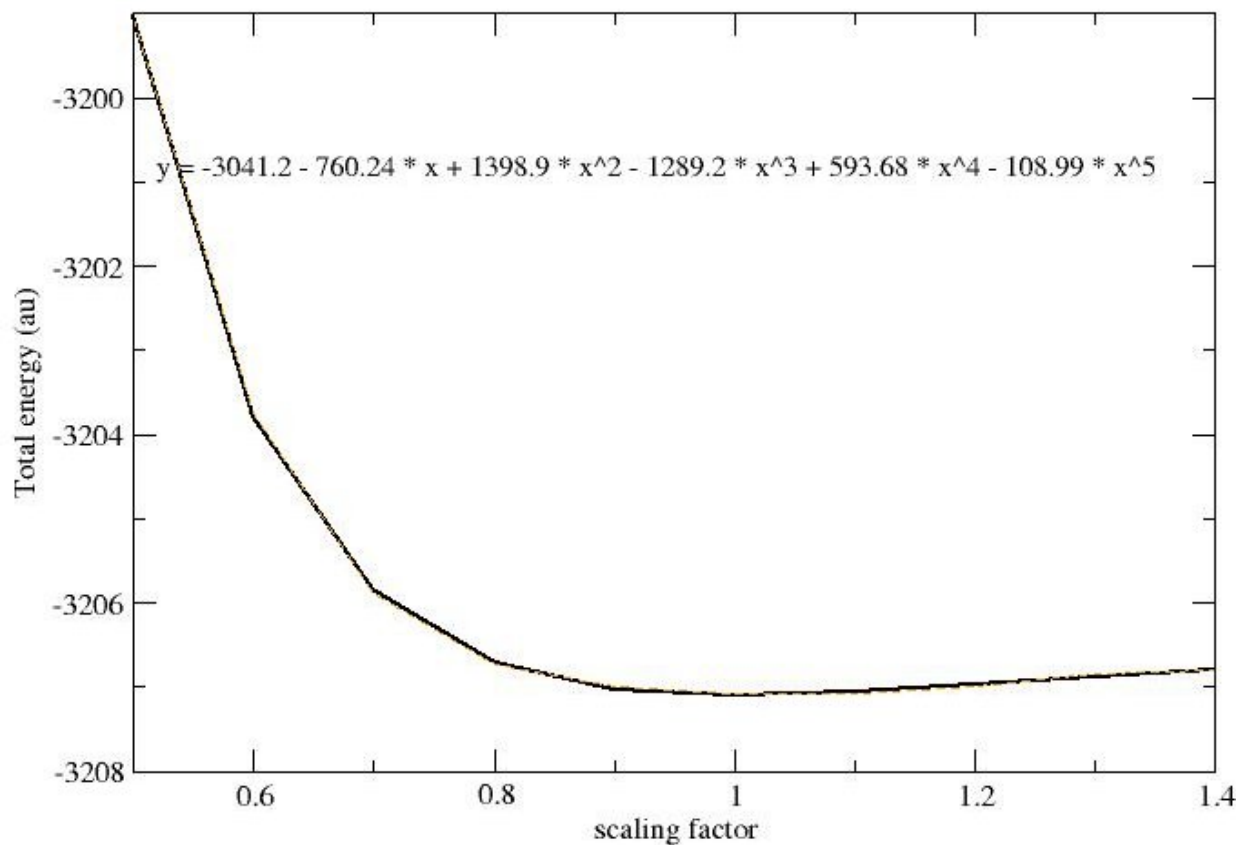
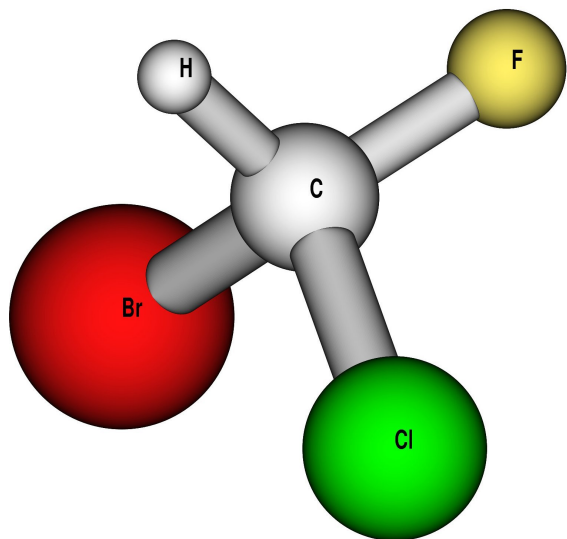
$r(\text{C1-Br})$: 4.469 Å
 $r(\text{C1-Cl})$: 4.316 Å
 $r(\text{C1-F})$: 3.947 Å
 $r(\text{C1-H})$: 3.744 Å
Total energy: -3510.990470 E_h
HOMO-LUMO gap:
0.242571 E_h
Total EPV = 1.629E-20 E_h
EPV (central carbon C1):
2.905E-23 E_h



$r(\text{C1-Br})$: 7.035 Å
 $r(\text{C1-Cl})$: 6.882 Å
 $r(\text{C1-F})$: 6.517 Å
 $r(\text{C1-H})$: 6.319 Å
Total energy: -3815.623118 E_h
HOMO-LUMO gap:
0.198085 E_h
Total EPV = -9.436E-23 E_h
EPV (central carbon C1): -
3.737E-25 E_h

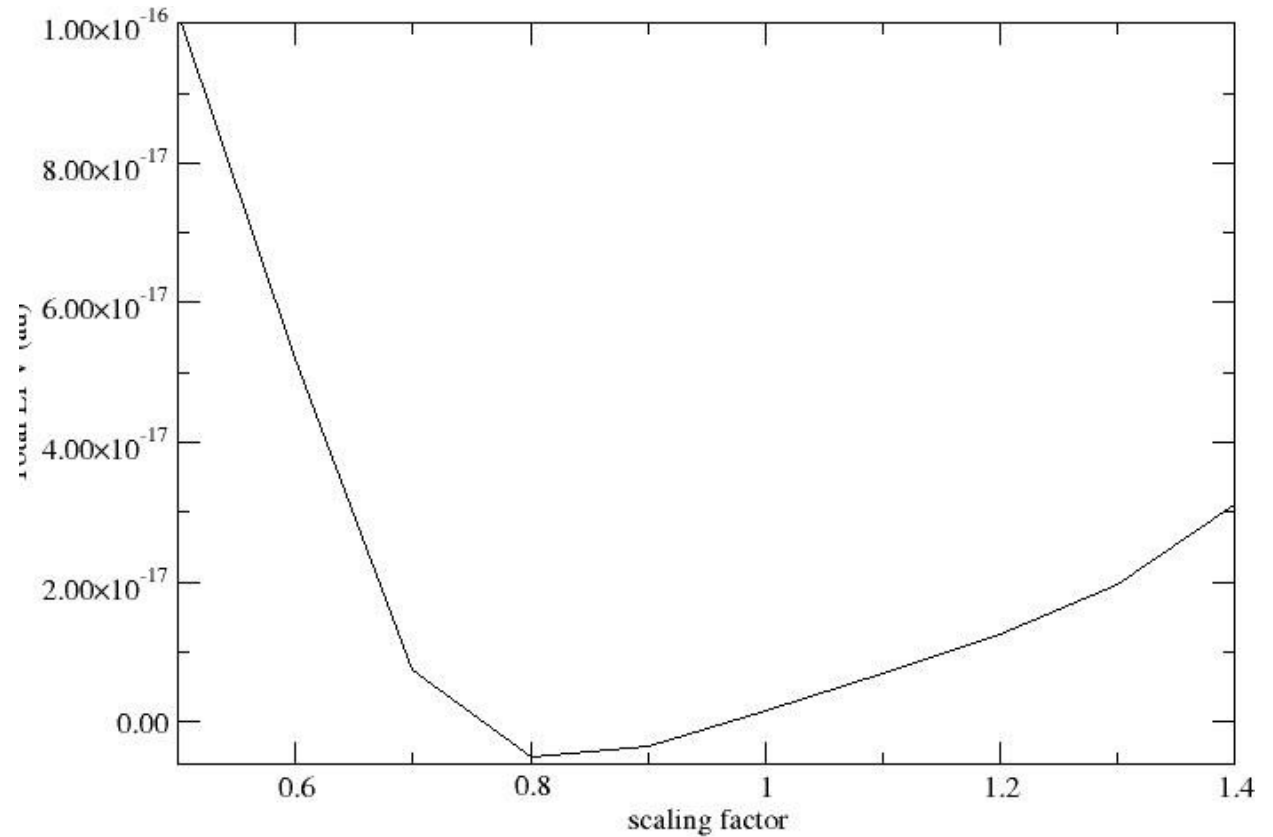
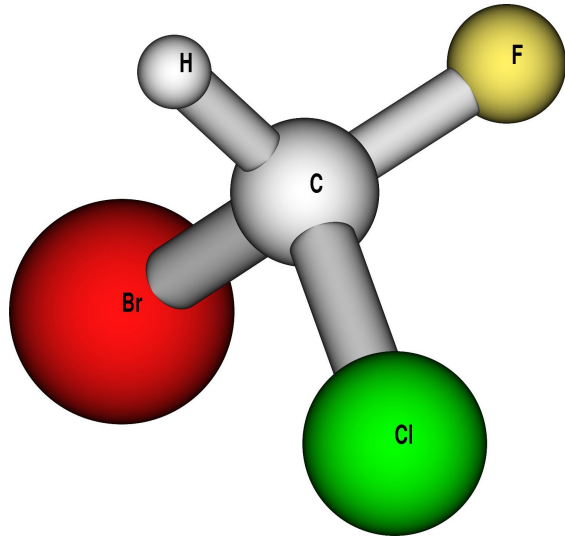
Exploring the chiral field: CHFCIBr

We start from CCSD(T) optimized geometry and scale all bond lengths



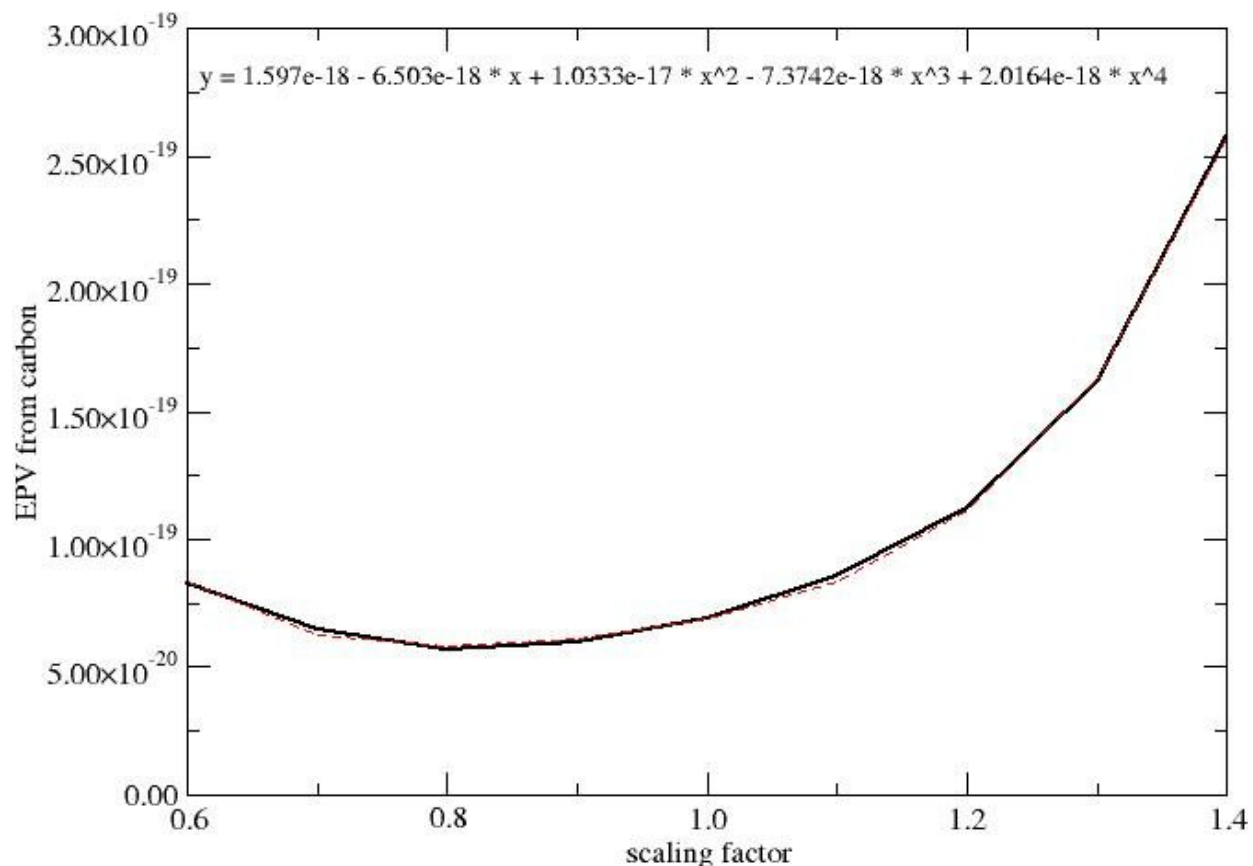
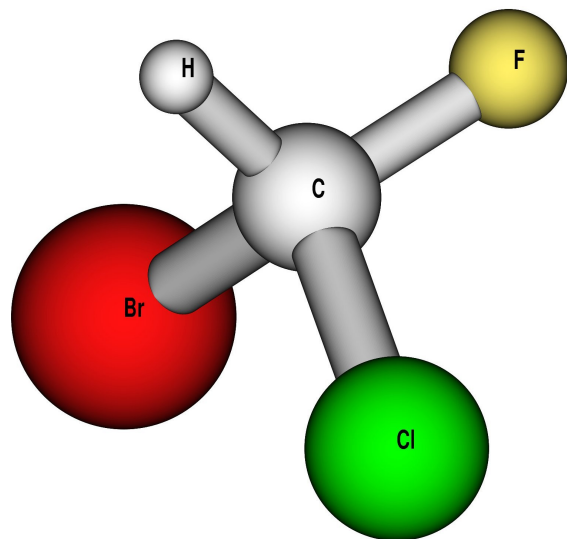
The total electronic energy shows a minimum near scaling factor 1.0.

Exploring the chiral field: CHFCIBr



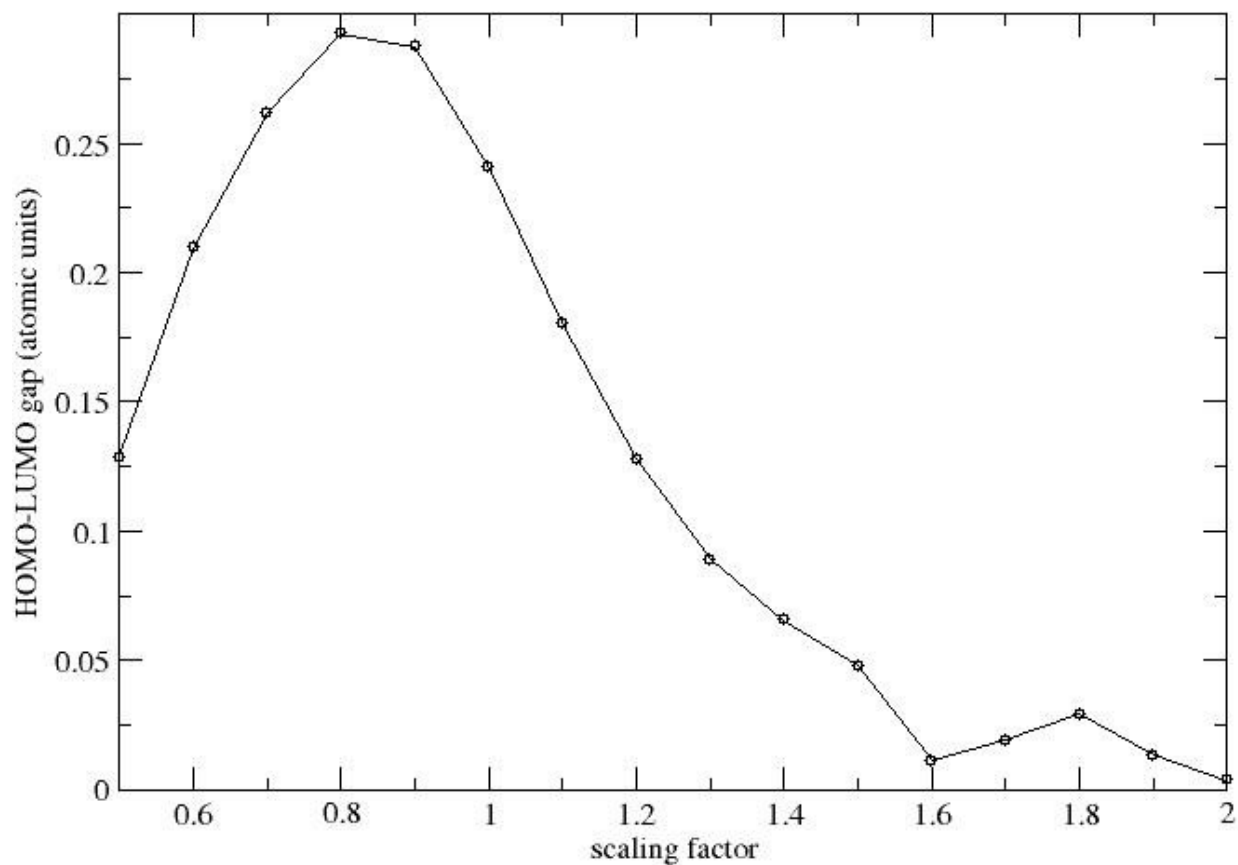
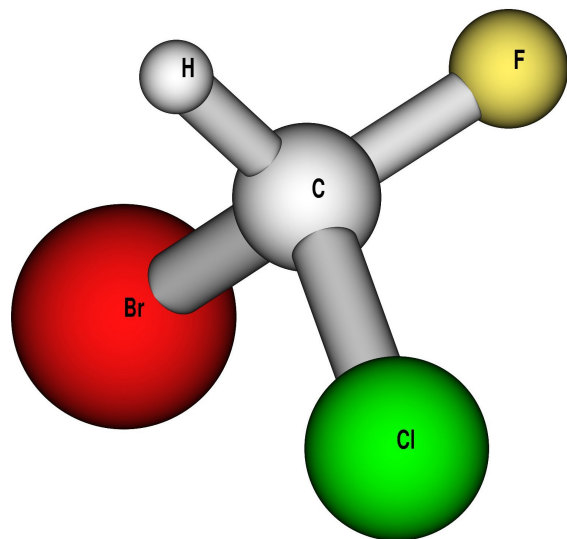
The total E_{PV} energy is not the most interesting quantity since the change in the chiral field for all atoms except the central carbon atom is rather complicated.

Exploring the chiral field: CHFCIBr



More interesting is the contribution to the E_{PV} energy from the central carbon atom. Surprisingly it increases within the range of validity of the model and goes through a minimum around $s=0.82$.

Exploring the chiral field: CHFCIBr



The bond stretching will inevitably lead to a breakdown of the SCF model. Indeed one observes that the HOMO-LUMO gap closes and the SCF did not converge beyond scaling factor 1.4. Further studies requires the calculation of E_{PV} using a method which can break bonds, e.g. MCSCF.