

Part II

Density-Functional Tight-Binding (DFTB) as fast approximate DFT method



Helmut Eschrig



Gotthard Seifert



Thomas Frauenheim



Marcus Elstner

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Density-Functional Tight-Binding

Part I

1. Tight-Binding

2. Density-Functional Tight-Binding (DFTB)

Part II

3. Bond Breaking in DFTB

4. Extensions

5. Performance and Applications

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Bond breaking

SCC-DFTB and SDFTB Dissociation of H_2^+

$$E_{SDFTB} = \sum_{\sigma=\uparrow,\downarrow} \sum_i^{MO} n_{i\sigma} \langle \psi_{i\sigma} | \hat{H}[\rho_0] | \psi_{i\sigma} \rangle + \frac{1}{2} \sum_{\alpha\beta} \gamma_{\alpha\beta} \Delta q_{\alpha} \Delta q_{\beta} + \frac{1}{2} \sum_{\alpha} \sum_{l \in \alpha'} \sum_{l' \in \alpha} p_{\alpha l} p_{\alpha l'} W_{\alpha l l'} + E_{rep}$$

H^+ : (no electron):

$$E_{SDFTB} = 0 + \frac{1}{2} \gamma_{HH} \Delta q_H \Delta q_H + 0 + 0 = \frac{1}{2} 0.4195 * 1 * 1 = 0.20975 \text{ h} \neq 0$$

Energy without electrons in an electronic structure method

H: (one electron) (spin-polarized SCC-DFTB):

$$E_{SDFTB} = \epsilon_i + \frac{1}{2} * 1 * 1 * W_{Hss} = -0.2386 + \frac{1}{2} * (-0.072) = -0.2746 \text{ h}$$

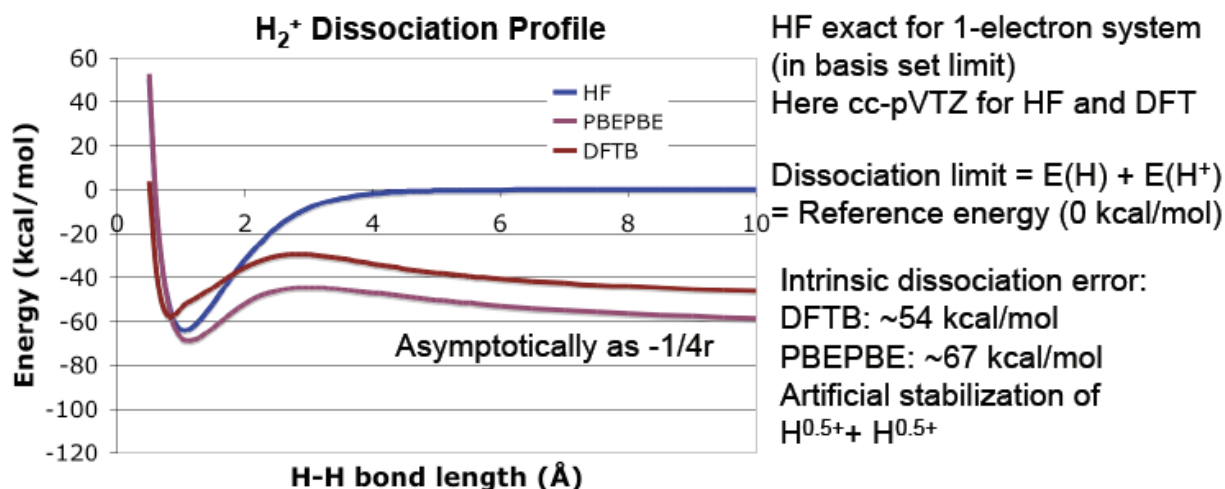
Ionization Potential of H (IP_H) = $E(H^+) - E(H) = 0.5 \text{ h}$

Spin-polarized SDFTB: $IP_H = 0.4821 \text{ h}$ (error of 11.2 kcal/mol)

SCC-DFTB: $IP_H = 0.4461 \text{ h}$ (error of 33.8 kcal/mol)

Spin-polarized SDFTB is qualitatively correct for H^+ , H (and also H^-)

SCC-DFTB and SDFTB Dissociation of H_2^+



In most DFT functionals, the electron unphysically interacts with its own density (i.e. self-interaction error - SIE). This error can contribute to:

- Incorrect dissociation limits for systems with unpaired electrons (e.g. H_2^+)
- Underestimation of transition state barriers (of simple organic reactions)
- Exaggerated delocalization in conjugated systems
- Too low energy of charge-transfer excitations

SCC-DFTB and SDFTB Dissociation of H_2^+

$$E_{SDFTB} = \sum_{\sigma=\uparrow,\downarrow} \sum_i^{MO} n_{i\sigma} \langle \psi_{i\sigma} | \hat{H}[\rho_0] | \psi_{i\sigma} \rangle + \frac{1}{2} \sum_{\alpha\beta}^N \gamma_{\alpha\beta} \Delta q_{\alpha} \Delta q_{\beta} + \frac{1}{2} \sum_{\alpha}^N \sum_{l \in \alpha'} \sum_{l' \in \alpha} p_{\alpha l} p_{\alpha l'} W_{\alpha l l'} + E_{rep}$$

Error in dissociation energy at infinite distance:

$$E_{SDFTB}(H + H^+) = \varepsilon_i + \frac{1}{2} W_{Hss} + \frac{1}{2} \gamma_{HH} \quad (\text{correct})$$

$$E_{SDFTB}(H^{0.5+} + H^{0.5+}) = \frac{1}{2} 2\varepsilon_i + \frac{1}{2} 0.5 * 0.5 W_{Hss} * 2 + \frac{1}{2} 0.5 * 0.5 \gamma_{HH} * 2 = \varepsilon_i + \frac{1}{4} W_{Hss} + \frac{1}{4} \gamma_{HH}$$

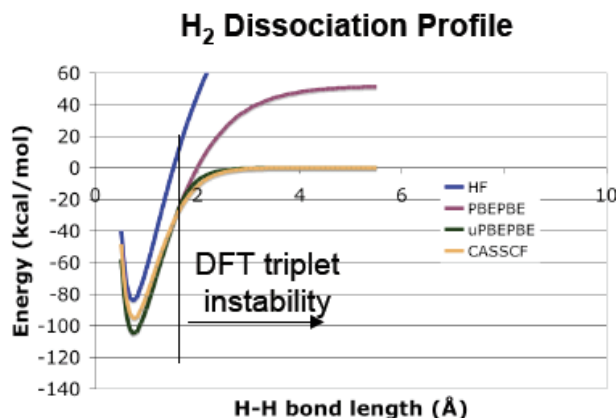
$$\text{Error}(SDFTB) = E_{SDFTB}(H + H^+) - E_{SDFTB}(H^{0.5+} + H^{0.5+}) = \frac{1}{4} W_{Hss} + \frac{1}{4} \gamma_{HH} \quad (\text{wrong})$$

Error in dissociation energy of H_2^+

Spin-polarized DFTB: $(0.4195/4 - 0.072/4) = 0.086875$ h = 54.5 kcal/mol

Normal SCC-DFTB: $0.4195/4 = -0.104875$ = 65.8 kcal/mol

SCC-DFTB and SDFTB Dissociation of H₂

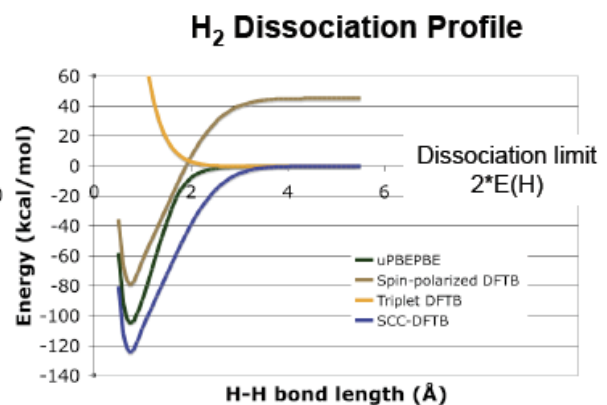


Triples: $\alpha\alpha$ $\beta\beta$ $\alpha\beta+\beta\alpha$

Singlet = $\alpha\beta-\beta\alpha$

UDFT singlet: $\alpha\beta = 1/2[(\alpha\beta - \beta\alpha) + (\alpha\beta + \beta\alpha)]$

Correct energy limit by triplet spin-contamination



Atoms are spin polarized, but H₂ (singlet) does not include spin-polarization at any distance. No UDFTB solution

SCC-DFTB energy is better but has $\alpha/2 + \beta/2$ electrons at each nuclei.

No DFTB method can give qualitatively correct dissociation curve

M. Lundberg, Y. Nishimoto, *SI, Int. J. Quant. Chem.* **112**, 1701 (2012)

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Analytical Hessian for DFTB and SCC-DFTB

H. A. Witek, S. Irle, K. Morokuma, *J. Chem. Phys.* **121**, 5163 (2005)

$$G_{ab}^{\text{DFTB}} = \frac{\partial^2 E_{\text{rep}}}{\partial a \partial b} + 2 \sum_{im}^{\text{MO}} n_i \sum_{\mu\nu}^{\text{AO}} U_{mi}^b c_{\mu m} c_{\nu i} \left[\frac{\partial H_{\mu\nu}^0}{\partial a} - \varepsilon_i \frac{\partial S_{\mu\nu}}{\partial a} \right] +$$

$$+ \sum_i^{\text{MO}} n_i \sum_{\mu\nu}^{\text{AO}} c_{\mu i} c_{\nu i} \left[\frac{\partial^2 H_{\mu\nu}^0}{\partial a \partial b} - \varepsilon_i \frac{\partial^2 S_{\mu\nu}}{\partial a \partial b} - \frac{\partial \varepsilon_i}{\partial b} \frac{\partial S_{\mu\nu}}{\partial a} \right]$$

$$G_{ab}^{\text{SCC-DFTB}} = \frac{\partial^2 E_{\text{rep}}}{\partial a \partial b} + 2 \sum_{im}^{\text{MO}} n_i \sum_{\mu\nu}^{\text{AO}} U_{mi}^b c_{\mu m} c_{\nu i} \left[\frac{\partial H_{\mu\nu}^0}{\partial a} + \left(\frac{1}{2} \sum_K^{\text{atoms}} (\gamma_{MK} + \gamma_{NK}) \Delta q_K - \varepsilon_i \right) \frac{\partial S_{\mu\nu}}{\partial a} \right] +$$

$$+ \sum_i^{\text{MO}} n_i \sum_{\mu\nu}^{\text{AO}} c_{\mu i} c_{\nu i} \left[\frac{\partial^2 H_{\mu\nu}^0}{\partial a \partial b} + \left(\frac{1}{2} \sum_K^{\text{atoms}} (\gamma_{MK} + \gamma_{NK}) \Delta q_K - \varepsilon_i \right) \frac{\partial^2 S_{\mu\nu}}{\partial a \partial b} \right] -$$

$$- \sum_i^{\text{MO}} n_i \sum_{\mu\nu}^{\text{AO}} c_{\mu i} c_{\nu i} \left[\frac{\partial \varepsilon_i}{\partial b} - \frac{1}{2} \sum_K^{\text{atoms}} \left(\frac{\partial \gamma_{MK}}{\partial b} + \frac{\partial \gamma_{NK}}{\partial b} \right) \Delta q_K - \frac{1}{2} \sum_K^{\text{atoms}} (\gamma_{MK} + \gamma_{NK}) \frac{\partial \Delta q_K}{\partial b} \right] \frac{\partial S_{\mu\nu}}{\partial a} +$$

$$+ \sum_K^{\text{atoms}} \frac{\partial^2 \gamma_{AK}}{\partial a \partial b} \Delta q_A \Delta q_K + \sum_K^{\text{atoms}} \frac{\partial \gamma_{AK}}{\partial a} \frac{\partial \Delta q_A}{\partial b} \Delta q_K + \sum_K^{\text{atoms}} \frac{\partial \gamma_{AK}}{\partial a} \Delta q_A \frac{\partial \Delta q_K}{\partial b}$$

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Coupled Perturbed SCC-DFTB Equations

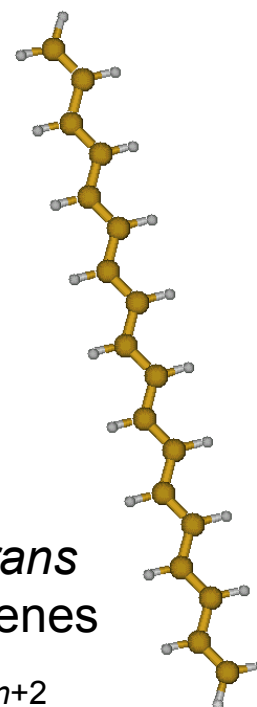
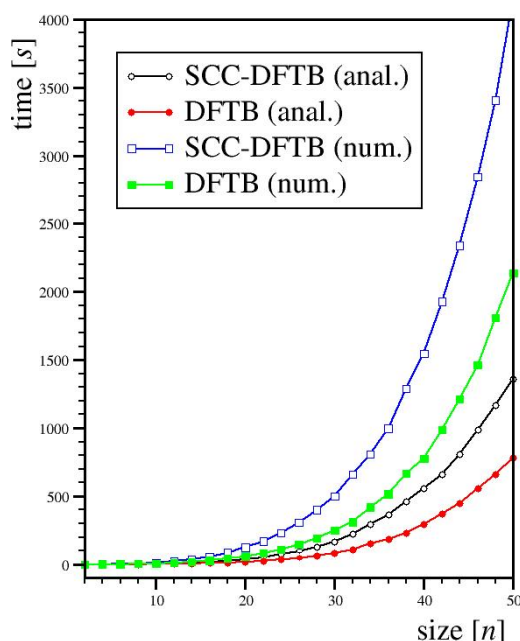
- ❖ To determine response of molecular orbitals to nuclear perturbation, one has to solve a set of iterative coupled-perturbed SCC-DFTB equations

$$\left\{ \begin{array}{l} U_{ij}^b = \frac{1}{\varepsilon_j - \varepsilon_i} \sum_{\mu\nu}^{\text{AO}} c_{\mu i} c_{\nu j} \left[\frac{\partial H_{\mu\nu}^0}{\partial b} + \left(\frac{1}{2} \sum_K^{\text{atoms}} (\gamma_{MK} + \gamma_{NK}) \Delta q_K - \varepsilon_i \right) \frac{\partial S_{\mu\nu}}{\partial b} + \right. \\ \quad - \\ \quad \left. + \frac{1}{2} S_{\mu\nu} \sum_K^{\text{atoms}} \left(\frac{\partial \gamma_{MK}}{\partial b} + \frac{\partial \gamma_{NK}}{\partial b} \right) \Delta q_K + \frac{1}{2} S_{\mu\nu} \sum_K^{\text{atoms}} (\gamma_{MK} + \gamma_{NK}) \frac{\partial \Delta q_K}{\partial b} \right] \\ \\ \frac{\partial \Delta q_K}{\partial b} = \sum_i^{\text{MO}} n_i \sum_{\mu \in K}^{\text{AO}} \sum_{\nu}^{\text{AO}} \left[c_{\mu i} c_{\nu i} \frac{\partial S_{\mu\nu}}{\partial b} + \sum_m^{\text{MO}} U_{mi}^b (c_{\mu m} c_{\nu i} + c_{\mu i} c_{\nu m}) S_{\mu\nu} \right] \end{array} \right.$$

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Frequency calculations

performance test



How to treat van der Waals interactions cheaply?

Addition of empirical **London dispersion** term!

$$E_{\text{tot}} = E_{\text{QM}} - \sum_{\alpha\beta} f(R_{\alpha\beta}) C_6^{\alpha\beta} / R_{\alpha\beta}^6 (+1/R_{\alpha\beta}^8 \text{ etc. terms})$$

$f(R_{\alpha\beta})$: damping function



Fritz London
1/R⁶: 1922

- Ahlrichs *et al.*, Chem. Phys. **19**, 119 (1977): **HFD** (Hartree-Fock Dispersion), C_i parameters from **expt.**
- Mooij *et al.*, J. Phys. Chem. A **103**, 9872 (1999): PES with London dispersion, C_6 parameters from **MP2** calculations
- Elstner *et al.*, J. Chem. Phys. **114**, 5149 (2001): **SCC-DFTB-D** (dispersion-augmented SCC-DFTB), C_6 parameters via Halgren, J. Am. Chem. Soc. **114**, 7872 (1992) from expt.
- Yang *et al.*, J. Chem. Phys. **116**, 515 (2002): **DFT+vdW** (DFT including van der Waals interactions), C_6 parameters from expt.
- Grimme, J. Comput. Chem. **25**, 1463 (2004): **DFT-D**, C_6 parameters from expt.
- ... several more plus many reviews and benchmarks

London Dispersion

- No overlapping densities
- Interaction of two polarisable objects

$$E_{London} = -\frac{2}{3} \frac{I_1 I_2}{I_1 + I_2} \alpha_1 \alpha_2 \frac{1}{r^6}$$

London, F.; Z. Physik 1930, 63, 245.

Well-known application: Lennard-Jones potential:

$$E_{Lennard-Jones} = \left(\frac{C_{12}}{r^{12}} - \frac{C_6}{r^6} \right)$$

short-range repulsion long-range attraction

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DFTB+Dispersion

Elstner *et al.*, JCP **114**, 5149 (2001)

$$E_{tot} = E_{DFTB} + \sum_{i < j} E_{dispersion}(i, j)$$

$$E_{dispersion}(i, j) = -\frac{1}{r_{ij}^6} f * C_6(i, j),$$

$$f = \left(1 - \exp \left(-3.0 * \left(\frac{r_{ij}}{R_{ij, vdw}} \right)^7 \right) \right), \text{damping function}$$

$$R_{ij, vdw} = \frac{R_{i, vdw}^3 + R_{j, vdw}^3}{R_{i, vdw}^2 + R_{j, vdw}^2}, \text{cubic mean rule}$$

$$C_6(i, j) = \frac{3}{2} * \frac{\alpha_i \alpha_j}{\left(\alpha_i / N_i \right)^{1/2} + \left(\alpha_j / N_j \right)^{1/2}}$$

α_i, α_j : atomic polarizability parameter

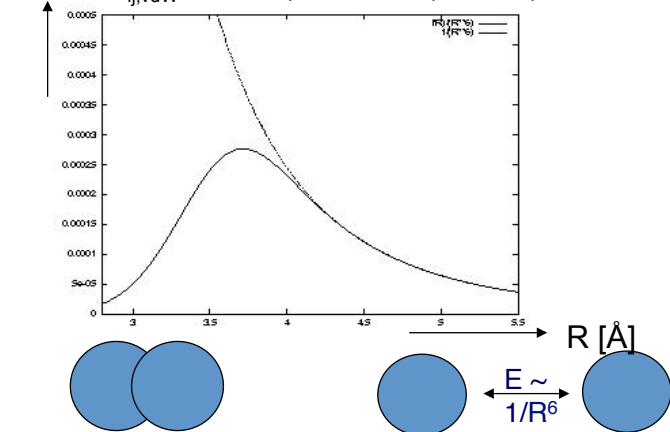
N_i, N_j : Slater Kirkwood effective number of electrons

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DFTB+Dispersion

Elstner *et al.*, JCP **114**, 5149 (2001)

damping $f(R_{\alpha\beta}) = [1 - \exp(-3(r_{ij}/R_{ij,vdW})^7)]^3$
 $R_{ij,vdW} = 3.8 \text{ \AA}$ (for 1st row), 4.8 \AA (for 2nd row)

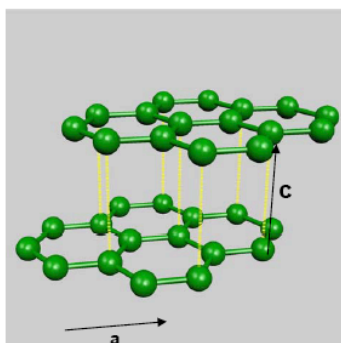


DFTB-D choice of C_6 parameters:

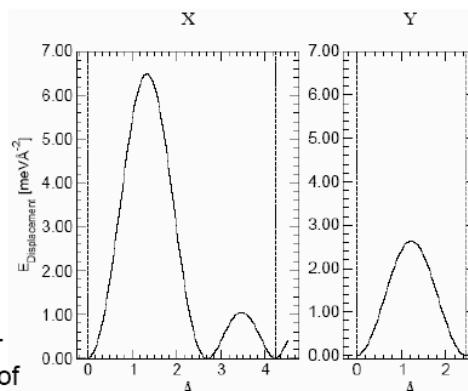
- generally hybridization dependent (i.e. not simply atomic values)
- use “empirical” values for parameters to match BSSE-corrected MP2 interaction energies

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Application to Graphite



	DFTB-D	Experiment
a	2.45Å	2.45Å
c	3.38Å	3.35Å
bulk modulus	330 GPa	318 ±11 GPa

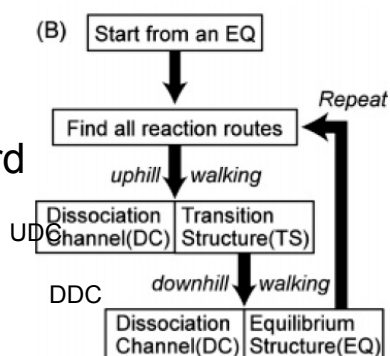
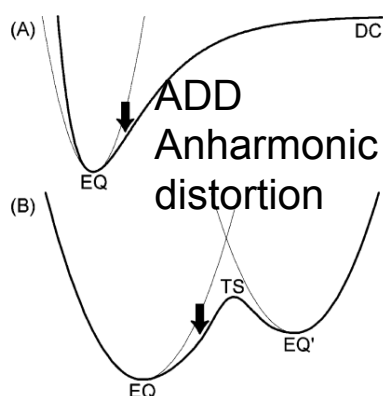


Energy change for lateral movement of graphite layers

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GRRM with DFTB

Principles of GRRM



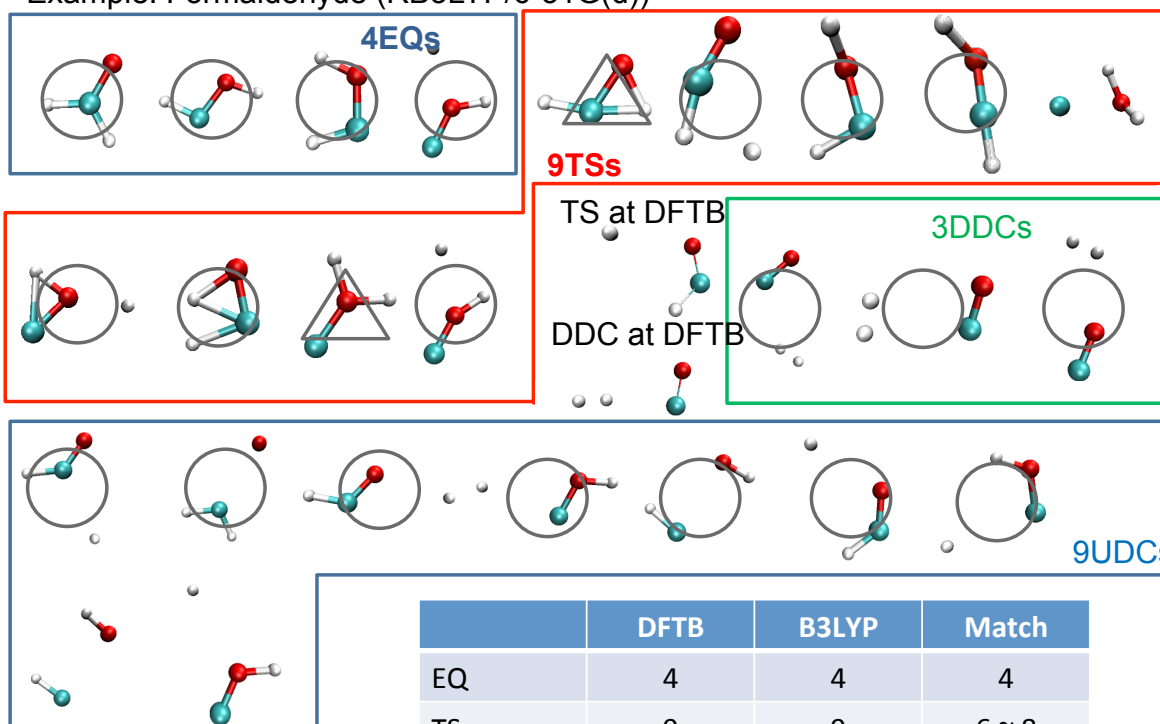
S. Maeda and K. Ohno, *J. Phys. Chem. A*, **2005**, *109*, 5742.

One of the most powerful and reliable methods to find reaction pathways.

Y. Nishimoto and S. Irlé, in preparation

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Example: Formaldehyde (RB3LYP/6-31G(d))



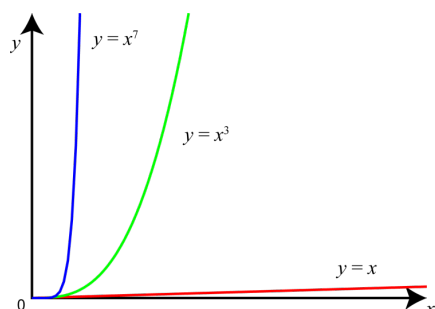
○ : Reproduced by DFTB
 △ : Found

	DFTB	B3LYP	Match
EQ	4	4	4
TS	9	9	6 ~ 8
DDC	4	3	3
UDC	7	9	7

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Want to perform quantum mechanical (QM) calculation of **large systems**

However, problem is the scaling of QM calculations ... QM calculations scale $N^3 \sim N^7$



We need a method which scales **linearly** with **cheap** computational cost (yet QM)

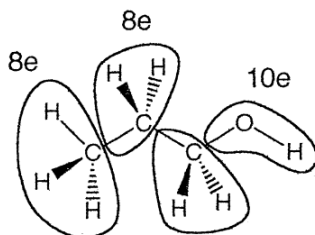
What can we do?

- 1) Linear scaling method: **fragment molecular orbital (FMO)**, divide-and-conquer (DC), generalized energy-based fragmentation (GEBF) ...
- 2) Cheap QM method → semi-empirical QM
MNDO, PM3, PM6, AM1, ZINDO, **DFTB** ...

- 1) Divide a large system to a set of fragments

Fragment molecular orbital method (FMO)

Kitaura, K.; Ikeo, E.; Asada, T.; Nakano, T.; Uebayashi, M.
Chem. Phys. Lett. **1999**, *313*, 701-706.



- 2) Use a semi-empirical method

Density-functional tight-binding method (DFTB)

Elstner, M.; Porezag, D.; Jungnickel, G.; Elsner, J.; Haugk, M.; Frauenheim, T.; Suhai, S.; Seiferg, G. *Phys. Rev. B* **1998**, *58*, 7260-7268.

$$E^{\text{SCC}} = \sum_i \sum_{\mu\nu} n_i c_{\mu i}^* c_{\nu i} H_{\mu\nu}^0 + \sum_{A>B} E_{AB}^{\text{rep}} + \frac{1}{2} \sum_{AB} \gamma_{AB} \Delta q_A \Delta q_B$$

FMO-DFTB

Calculate the molecular energy by the sum of internal energies of monomers and dimers:

$$E = \sum_I^N E'_I + \sum_{I>J}^N (E'_{IJ} - E'_I - E'_J) + \sum_{I>J}^N \Delta E_{IJ}^V \quad (\text{FMO2})$$

where internal energy is defined by

$$E'_X = \sum_{i \in X} \sum_{\mu\nu \in X} n_i^X c_{\mu i}^{X*} c_{\nu i}^X H_{\mu\nu}^{0,X} + \sum_{A>B \in X} E_{AB}^{\text{rep}} + \frac{1}{2} \sum_{AB \in X} \gamma_{AB} \Delta q_A^X \Delta q_B^X$$

Additionally, we need the coupling of the charge transfer to the embedding potential:

$$\Delta E_{IJ}^V = \sum_{A \in IJ} \sum_{K \notin I, J} \sum_{C \in K} \gamma_{AC} \Delta q_A^{IJ} \Delta q_C^K$$

FMO-DFTB Hamiltonian is defined by

$$H_{\mu\nu}^X = H_{\mu\nu}^{\text{SCC},X} + V_{\mu\nu}^X + P_{\mu\nu}^X$$

where $V_{\rho\sigma}^X = \frac{1}{2} S_{\rho\sigma}^X \sum_{K \neq X} \sum_{D \in K} (\gamma_{AD} + \gamma_{BD}) \Delta q_D^K$, and $P_{\mu\nu}^X$ is the projection operator.

Nishimoto, Y.; Fedorov, D. G.; Irle, S. submitted.

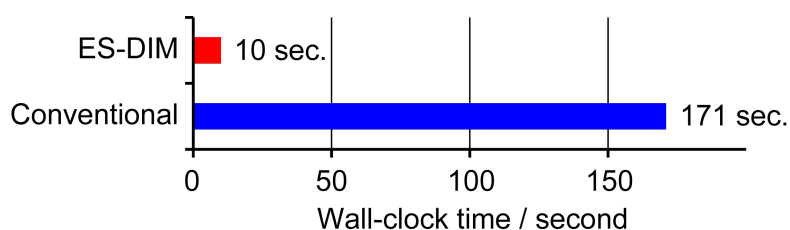
Electrostatic Dimer (ES-DIM)* Approximation

* Nakano, T. *et al. Chem. Phys. Lett.* **2002**, 351, 475-480.

For separated pairs (dimers), we introduce the electrostatic dimer approximation:

$$E'_{IJ} \approx E'_I + E'_J + \sum_{A \in I} \sum_{B \in J} \gamma_{AB} \Delta q_A^I \Delta q_B^J \quad \longrightarrow \quad \text{No SCF calculation}$$

The loss of energy with ES-DIM approximation is only 1.02×10^{-3} kcal/mol for extended COMe-(Ala)₂₀₀-NHMe (2,012 atoms)



ES-DIM approximation saves computational cost dramatically without loss of accuracy.

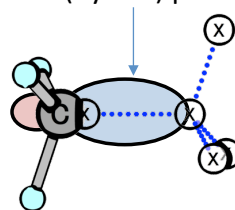
Projection operator works for DFTB too!

$$\begin{aligned}
 P_{\mu\nu} &= B \sum_i \langle \mu | \theta_i \rangle \langle \theta_i | \nu \rangle \\
 &= \sum_i \sum_{\lambda\sigma} B \tilde{c}_{\lambda i} \langle \mu | \lambda \rangle \langle \sigma | \nu \rangle \tilde{c}_{\sigma i}^* \\
 &= B \sum_{\lambda\sigma} \tilde{D}_{\lambda\sigma} \underbrace{S_{\mu\lambda} S_{\sigma\nu}}
 \end{aligned}$$

DFTB has overlap matrix parameters!

T. Nagata *et al. Chem. Phys. Lett.* **2010**, 492, 302.

This (hybrid) p orbital should be removed



$\tilde{D}_{\lambda\sigma}$: weighted density matrix of hybrid orbital



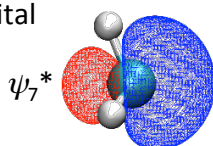
We need to prepare localized orbitals

In DFTB, **Pipek-Mezey localization** is the logical choice.

Pipek-Mezey localization: maximize the sum of Mulliken charges

Pipek, J. *et al. J. Chem. Phys.* **1989**, 90, 4916.

“projected out” orbital



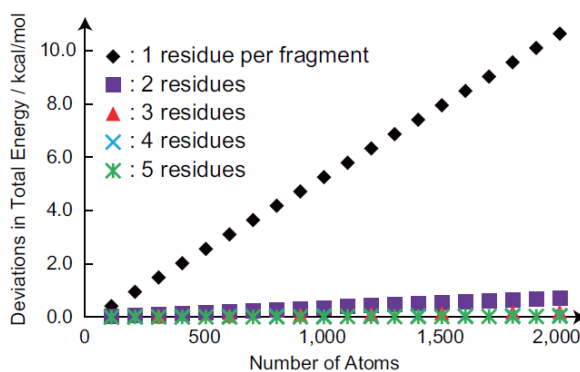
$\epsilon_7 = +2.721 \cdot 10^7$ eV \approx “B” constant value

Accuracy of FMO-DFTB: Energy

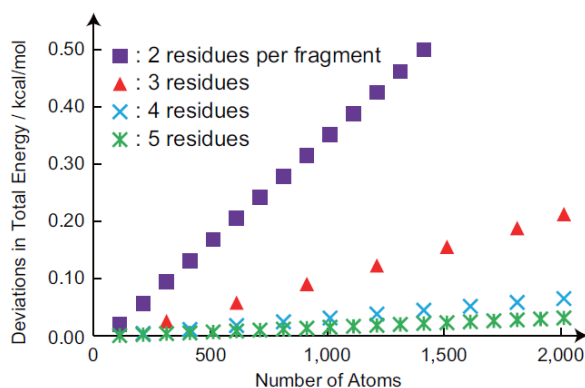
The energy difference between full SCC-DFTB and FMO-DFTB with extended poly-alanine



(A)



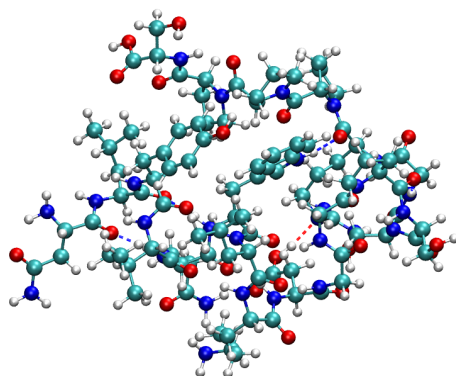
(B) Magnified figure



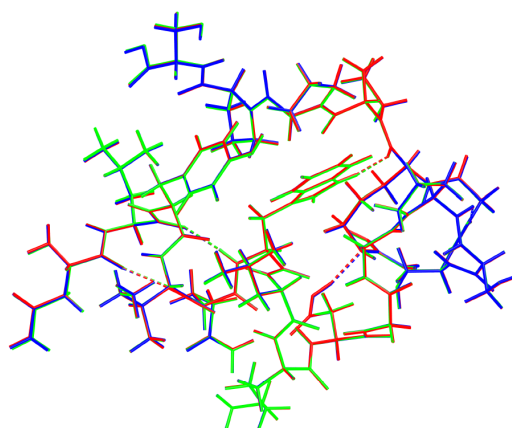
Error is less than 1 kcal/mol for 2,000 atom system (with more than 2 residues per fragment)

Accuracy of FMO-DFTB: Geometries

Geometry optimization of neutralized 1L2Y protein (Trp-cage / 20 residues)



Blue : Full calculation
 Red : FMO-DFTB (1), RMSD = 0.067 Å
 Green : FMO-DFTB (2), RMSD = 0.044 Å



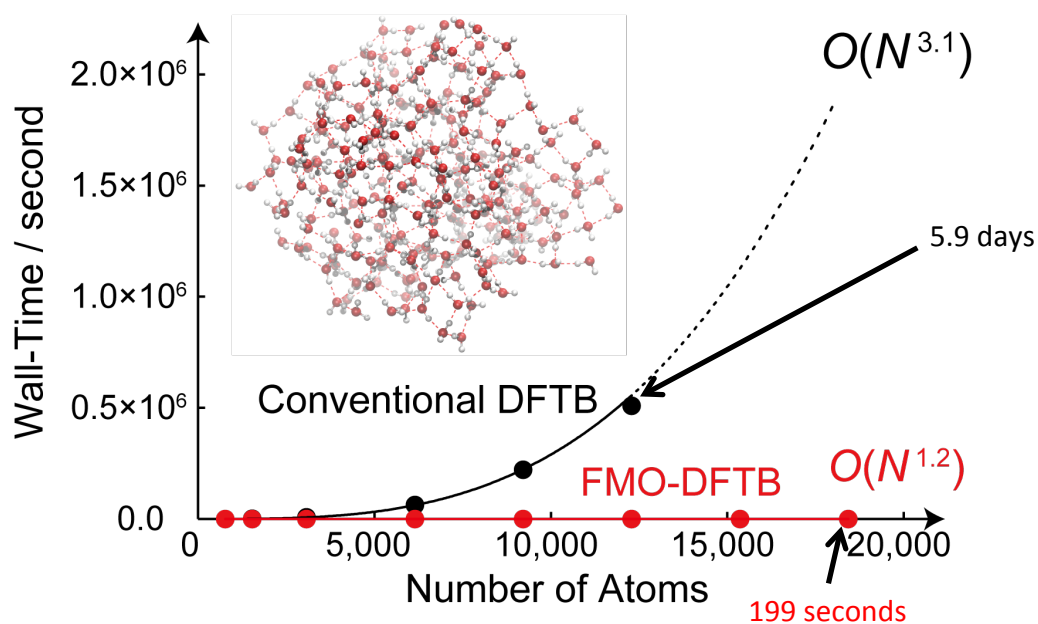
FMO-DFTB (1) :: 1 residue per fragment
 FMO-DFTB (2) :: 2 residues per fragment

Error of total energy is 1.20 and 0.82 kcal/mol

FMO-DFTB well reproduced the optimized geometry of full calculation

Wall-clock time and scaling of FMO-DFTB vs. DFTB

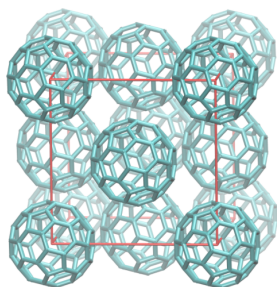
Wall-clock time with 1 CPU core (Xeon E5-2650)



Scaling of FMO-DFTB (for 3D system) is close to linear

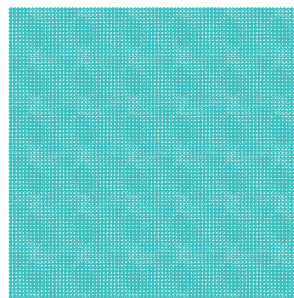
Inter-Node Parallel Calculation

FMO-DFTB is parallelized similar to conventional FMO with generalized distributed data interface (GDDI)



Face centered cubic (FCC)
 $a = 14.04 \text{ \AA}$
 Unit cell : $(C_{60})_{14} = 840$ atoms

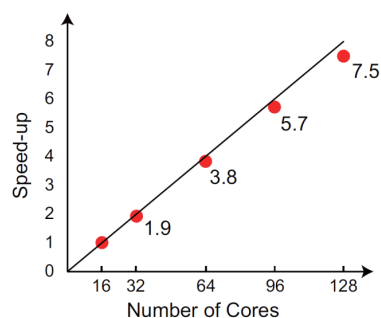
40 x 40



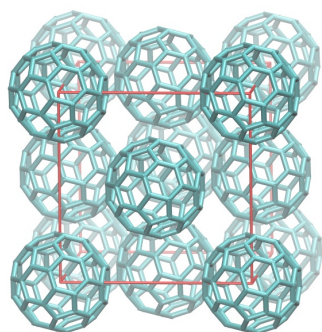
2 dimensional 40x40 fullerite
 $(C_{60})_{9842} = 590,520$ atoms

Calculation on 1 node (16 CPU cores) took **4.5** hours
 → Finished in **0.6** hour with 8 nodes (128 CPUs)

Parallel efficiency (up to 8 nodes) = 93.8 %

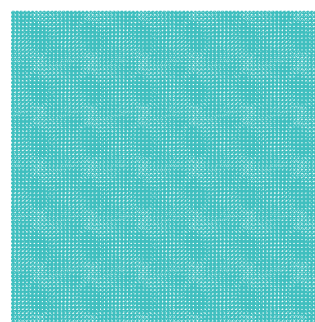


Demonstration with FMO-DFTB



Face centered cubic (FCC)
 $a = 14.04 \text{ \AA}$
 Unit cell : $(C_{60})_{14} = 840$ atoms

53 x 53



2 dimensional 53x53 fullerite
 $(C_{60})_{17174} = 1,030,440$ atoms

Geometry optimization at **FMO-DFTB** ... Converged after 16 cycles in about 4 days
 Total energy = -1772910.025319218 Hartree (1.8 million Hartree)

Geometry optimization at **FMO-DFTB-D** (+1.5 % wall time) ... Converged after 60 cycles
 Total energy = -1764778.120171755 Hartree (1.8 million Hartree)

Geometry optimization of million atom system is manageable

- Geometry optimization of **million atom system** is manageable
- FMO-DFTB for **MD simulations** with ~thousands of atoms becomes possible → prediction of reaction mechanism pathways, proton transfer channels, etc.
- FMO interaction energy decomposition allows the analysis of **ligand-protein interactions**
- Non-adiabatic **electron/hole transfer studies** following FMO-LCMO approach by Tsuneyuki *et al.* (CPL 476, 104 (2009)) becomes possible for **extended systems**

Special thanks to:



Dmitri G. Fedorov
(AIST)

Nishimoto, Y.; Fedorov, D. G.; Irle, S. submitted.

Density-Functional Tight-Binding

Part I

1. Tight-Binding

2. Density-Functional Tight-Binding (DFTB)

Part II

3. Bond Breaking in DFTB

4. Extensions

5. Performance and Applications

SCC-DFTB: general comparison with experiment

*Performance for small organic molecules
(mean absolut deviations)*

- Reaction energies: ~ 5 kcal/mole
- Bond-lenghts: ~ 0.014 Å
- Bond angles: ~ 2°
- Vib. Frequencies: ~6-7 %

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Accuracy of DFTB Geometries and Energies for Fullerene Isomers

Fullerene Isomers Geometries and Energies vs B3LYP/6-31G(d)

•102 Fullerene Isomers

• **small cage non-IPR C₂₀-C₃₆ (35)**, **large cage IPR C₇₀-C₈₆ (67)**

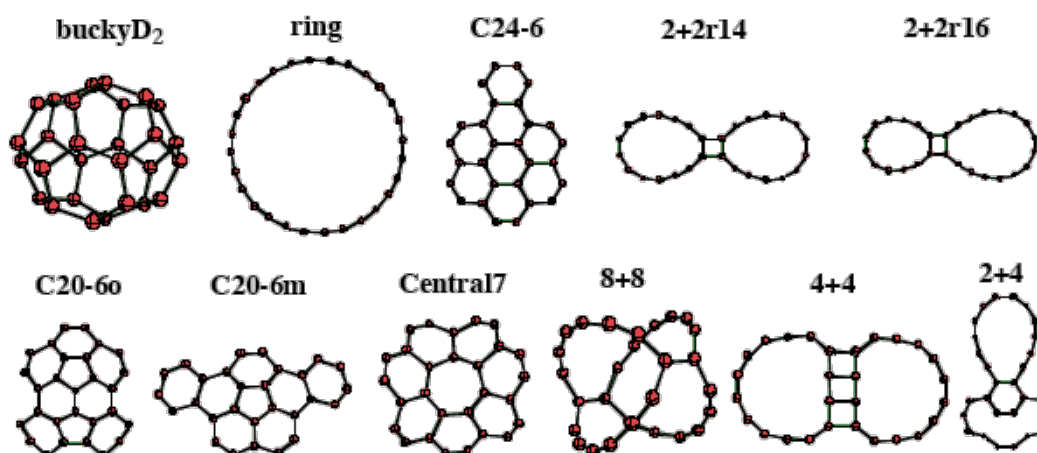
G. Zheng, S.I. M. Elstner, K. Morokuma, *Chem. Phys. Lett.* **412**, 210 (2005)

RMS [Å]	NCC-DFTB	SCC-DFTB	AM1	PM3	
C20-C36	0.025	0.019	0.035	0.030	Geometries
C60-C86	0.014	0.014	0.016	0.015	
R ² (lin. reg.)	NCC-DFTB	SCC-DFTB	AM1	PM3	
C20-C36	0.88	0.93	0.77	0.73	Energetics
C60-C86	0.97	0.98	0.86	0.84	

R²: Energy linear regression between ΔE(Method) and ΔE(B3LYP)

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What about non-cage carbon cluster structures? Some C_{28} isomers as example (from Scuseria, CPL 301, 98 (1999))



Structures of C_{28} used in the benchmark calculations presented in Table 1.

What about **non-cage** carbon cluster structures?
Some C_{28} isomers as example *TACC'04 Symp. Proceedings*

Table 1. Relative energies of C_{28} isomers in [eV] and corresponding linear regression coefficient R^2 for correlation between B3LYP/6-31G(d), and DFTB, B3LYP/6-31G, AM1, and PM3 methods. All energetics are obtained using individually optimized molecular structures at respective levels.

	B3LYP/6-31G(d)	DFTB	B3LYP/6-31G	AM1	PM3
buckyD2	0.00	0.00	0.00	0.00	0.00
ring	3.32	8.10	0.78	-7.69	-2.15
c24-6	3.17	3.56	1.99	0.43	1.77
2+2r14	5.08	9.66	2.90	-3.34	0.91
2+2r16	6.01	10.25	3.87	-3.37	0.90
c20-6o	5.41	5.52	4.34	3.42	4.23
c20-6 m	5.57	5.62	4.48	3.43	4.24
2 + 4	7.97	10.28	6.00	0.10	3.60
central7	5.86	6.07	4.84		
8 + 8	7.43	9.43	5.31	-3.24	0.79
4 + 4	9.91	14.27	8.52	1.52	4.97
R^2		0.7571	0.5390	0.2079	0.2964

Wrong minimum structures!

AM1 and PM3 are performing **very bad!**

DFTB includes effects of polarization functions through parameterization

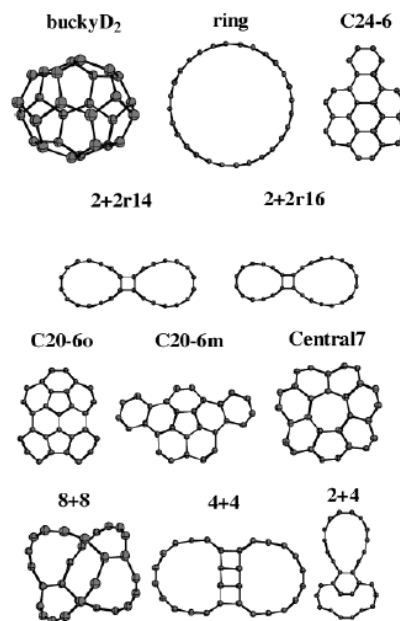
Accuracy of DFTB Geometries

Table 3. The relative energies (in eV) of 11 different isomeric singlet structures of C_{28} calculated with B3LYP/6-31G(d), and DFTB.

Structure ^a	B3LYP/6-31G(d)	DFTB	B3LYP/6-31G(d) // DFTB
buckyD2	0.00	0.00	0.00
ring	3.32	8.10	3.43
c24-6	3.17	3.56	3.66
2+2r14	5.08	9.66	5.22
2+2r16	6.01	10.25	6.13
c20-6o	5.41	5.52	5.96
c20-6m	5.57	5.62	6.09
2+4	7.97	10.28	8.52
central7	5.86	6.07	6.47
8+8	7.43	9.43	7.41
4+4	9.91	14.27	10.20
R^2 ^b		0.75	0.99

^a Structures illustrated below, with the labels taken from "Portmann, S.; Galbraith, J. M.; Schaefer, H. F.; Scuseria, G. E.; Lüthi, H. P. *Chem. Phys. Lett.* **1999**, *301*, 98-104."

^b Squared correlation coefficients R^2 in the linear regression analysis with B3LYP/6-31G(d) energies.

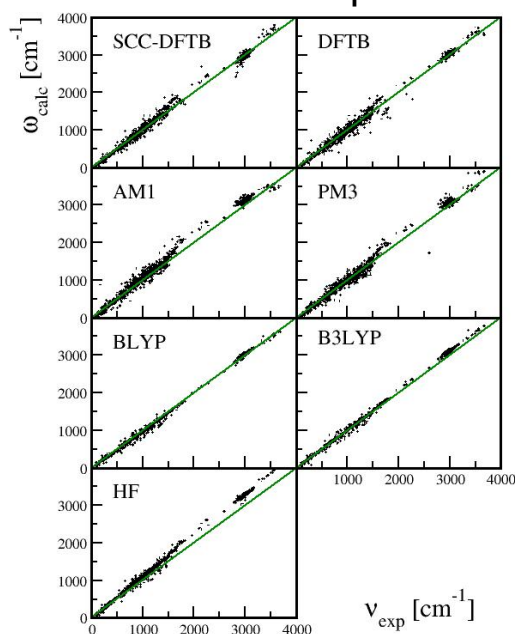


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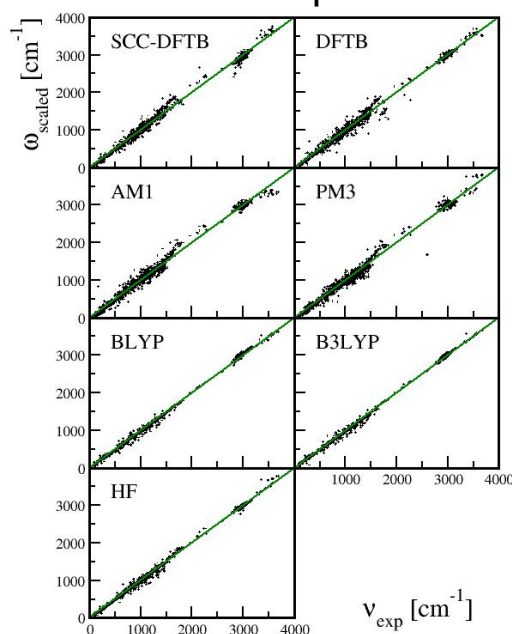
Accuracy of DFTB frequencies

Testing set of 66 molecules, 1304 distinct vibrational modes

calculated frequencies



scaled frequencies



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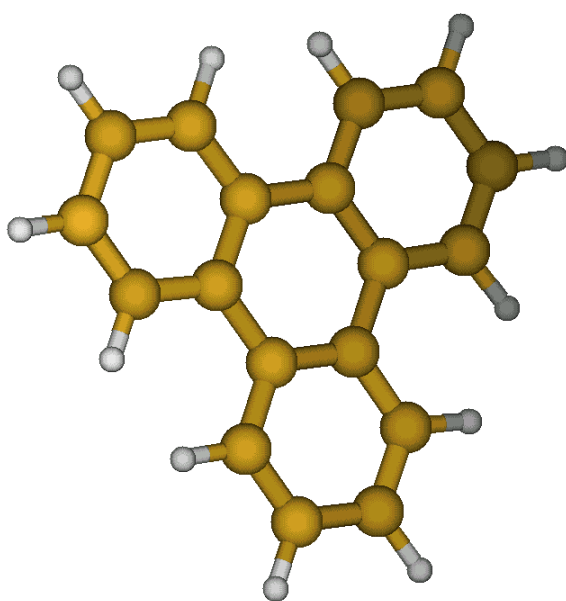
Accuracy of DFTB frequencies

Testing set of 66 molecules, 1304 distinct vibrational modes

	mean absolute deviation	standard deviation	maximal absolute deviation	scaling factor
SCC-DFTB	56 cm ⁻¹	82 cm ⁻¹	529 cm ⁻¹	0.9933
DFTB	60 cm ⁻¹	87 cm ⁻¹	536 cm ⁻¹	0.9917
AM1	69 cm ⁻¹	95 cm ⁻¹	670 cm ⁻¹	0.9566
PM3	74 cm ⁻¹	102 cm ⁻¹	918 cm ⁻¹	0.9762
HF/cc-pVDZ	30 cm ⁻¹	49 cm ⁻¹	348 cm ⁻¹	0.9102
BLYP/cc-pVDZ	34 cm ⁻¹	47 cm ⁻¹	235 cm ⁻¹	1.0043
B3LYP/cc-pVDZ	29 cm ⁻¹	42 cm ⁻¹	246 cm ⁻¹	0.9704

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Efficiency of DFTB frequencies



triphenylene

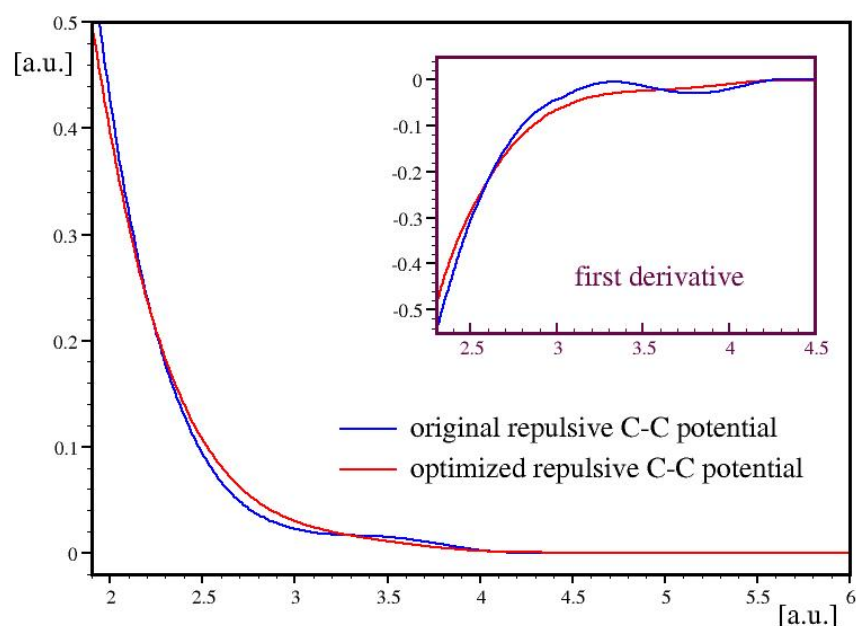
Energy, frequencies, and Raman and IR intensities calculations

DFT BLYP/cc-pVDZ
Linux 2.4GHz machine
32 hours

SCC-DFTB
Linux 333MHz machine
24 seconds

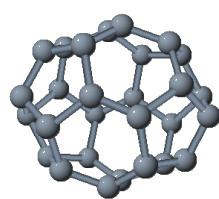
Improvement of DFTB E_{Rep}

H. A. Witek, et al, J. Theor. Comp. Chem. **4**, 639 (2005) and others

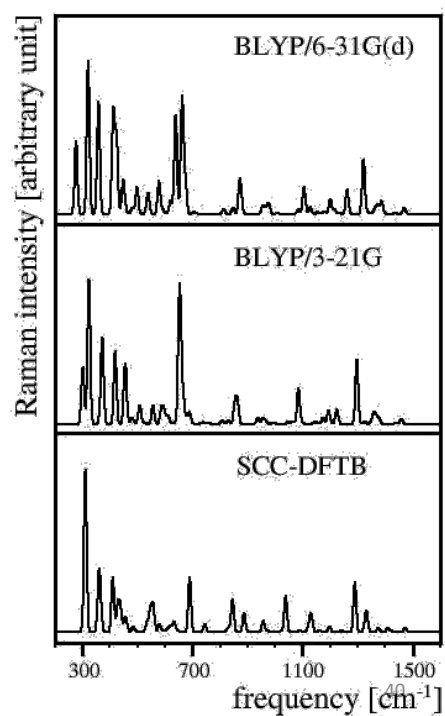
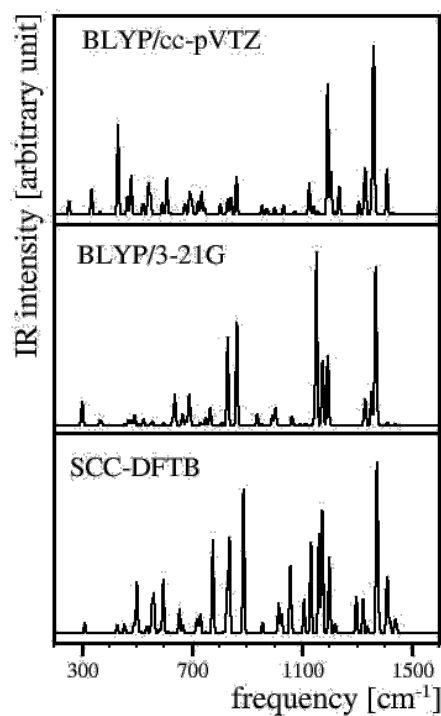


Harmonic IR and Raman Spectra of C_{28}

H. A. Witek, SI, G. Zheng, W. A. de Jong, K. Morokuma, J. Chem. Phys. **125**, 214706 (2005)

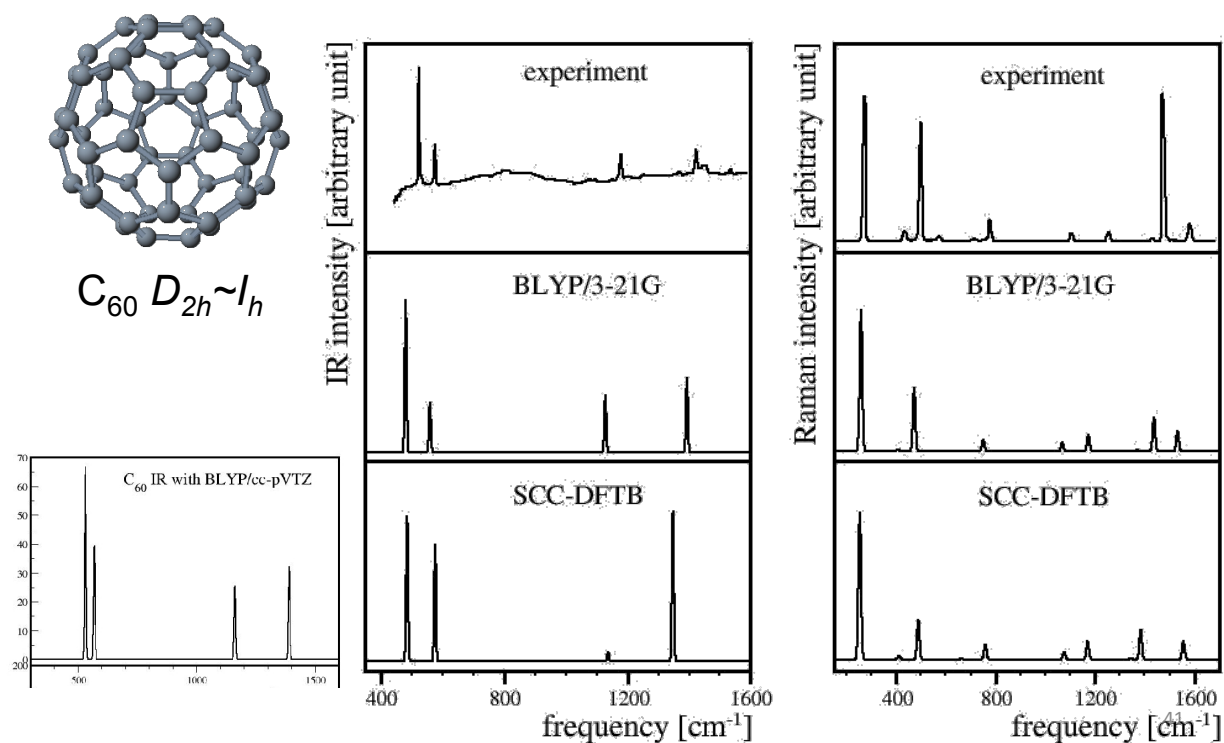


$C_{28} D_2$



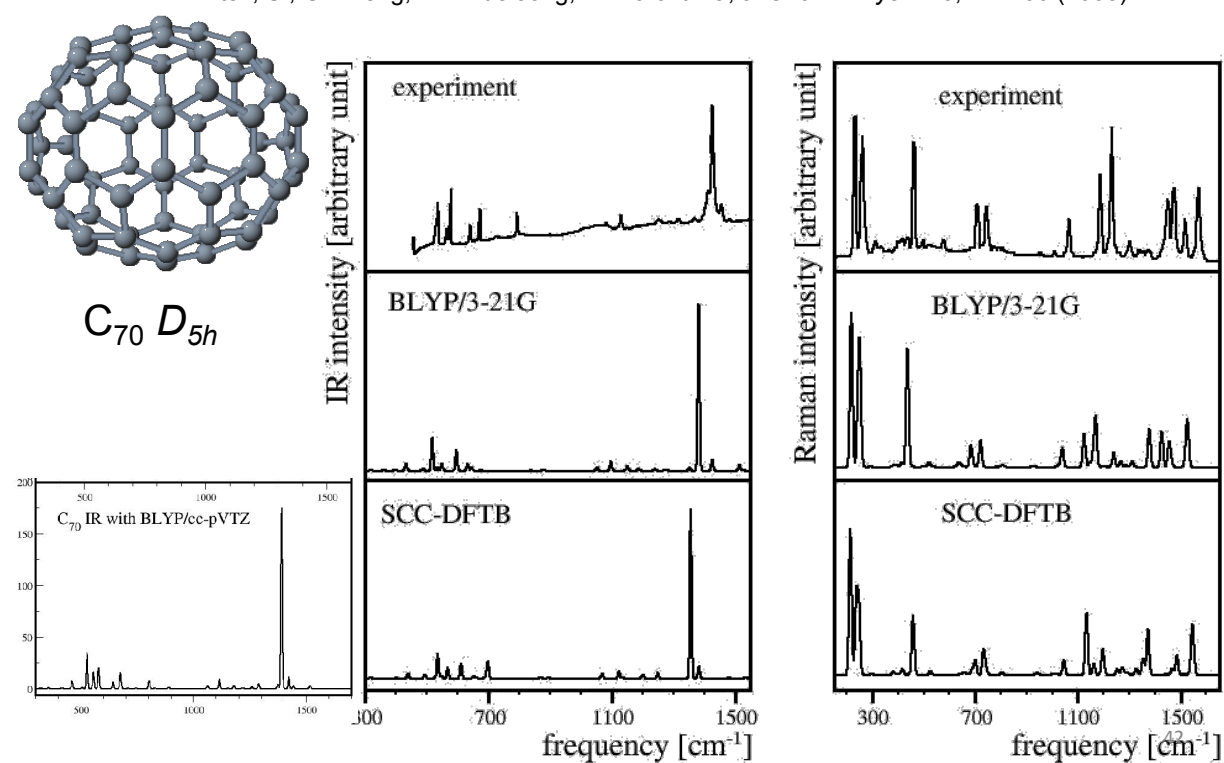
Harmonic IR and Raman Spectra of C₆₀

H. A. Witek, SI, G. Zheng, W. A. de Jong, K. Morokuma, *J. Chem. Phys.* **125**, 214706 (2005)



Harmonic IR and Raman Spectra of C₇₀

H. A. Witek, SI, G. Zheng, W. A. de Jong, K. Morokuma, *J. Chem. Phys.* **125**, 214706 (2005)



SCC-DFTB: Systematic comparison with other methods by Walter Thiel et al.

wavenumbers (compared with the other methods). The excellent performance of SCC-DFTB for geometries and vibrational frequencies has been noted before.^{18–20}

TABLE 3: Mean Absolute Deviations for Heats of Formation (kcal/mol) (Own Test Sets) (reference: exptl. ΔH_0)

	N^a	MNDO	AM1	OM2	DFTB ^b
neutral CHNO molecules	140	6.3	5.5	3.1	7.7
hydrocarbons	57	5.9	4.9	1.7	6.3
CHN compounds	32	6.2	4.6	3.9	6.1
CHO compounds	39	4.8	5.5	4.5	2.7
XNO compounds	8	16.3	11.4	2.9	43.9
anions	24	14.4	11.3	8.4	12.7
cations	33	11.5	9.8	7.2	14.5
radicals	42	11.9	10.6	5.0	17.0

^a N comparisons. ^b $N = 139, 31,$ and 39 in rows 1, 7, and 8, respectively, triplets excluded (see Supporting Information for details).

Thiel and coworkers, *J. Phys. Chem. A* **111**, 5751 (2007)

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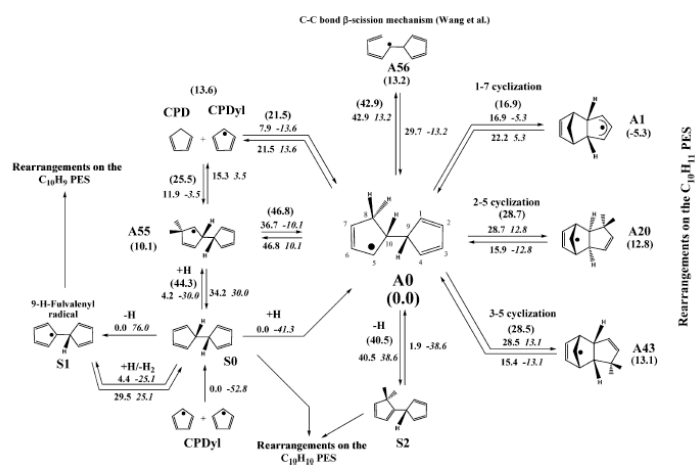
SCC-DFTB: Systematic comparison with other methods by Walter Thiel et al.

approach was also found to perform well.²⁰ Like other semiempirical methods, SCC-DFTB has its strengths and weaknesses. It is excellent for geometries and provides reasonable energetics for many types of compounds, especially for biomolecular systems. It seems less suitable for radicals and excited states, and suffers from occasional outliers (e.g., for molecules with NO bonds).

Thiel and coworkers, *J. Phys. Chem. A* **111**, 5751 (2007)

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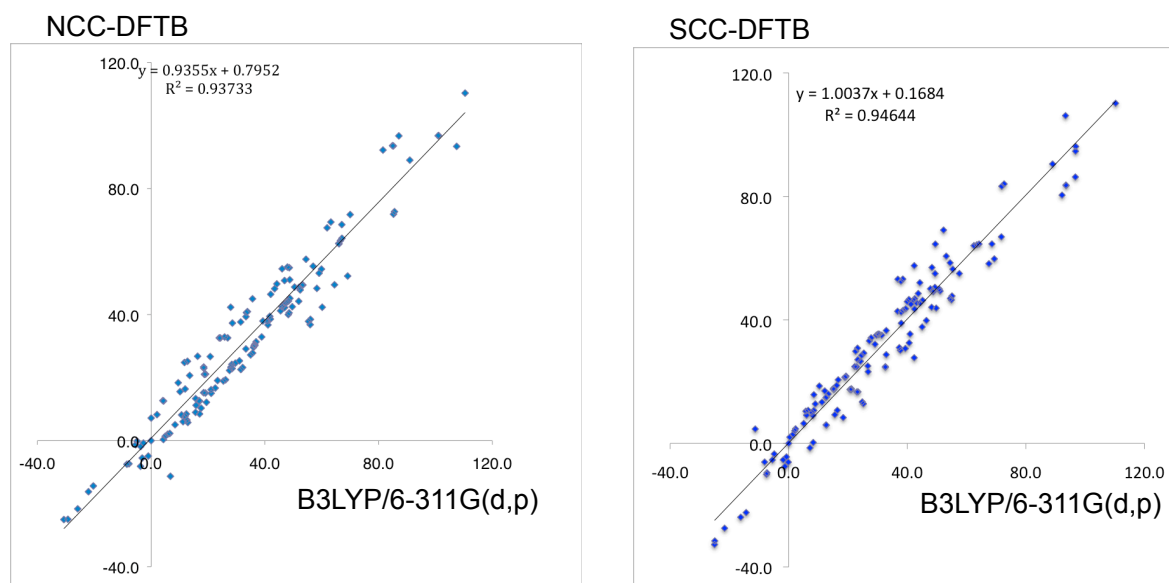
But: NCC- and SCC-DFTB for radicals: own comparison with B3LYP/6-311G**



	NCC-DFTB	SCC-DFTB
Barrier heights	6.60	5.99
Rel. energies	7.96	7.31
Overall RMSD	7.17	6.55
Overall R^2	0.937	0.946

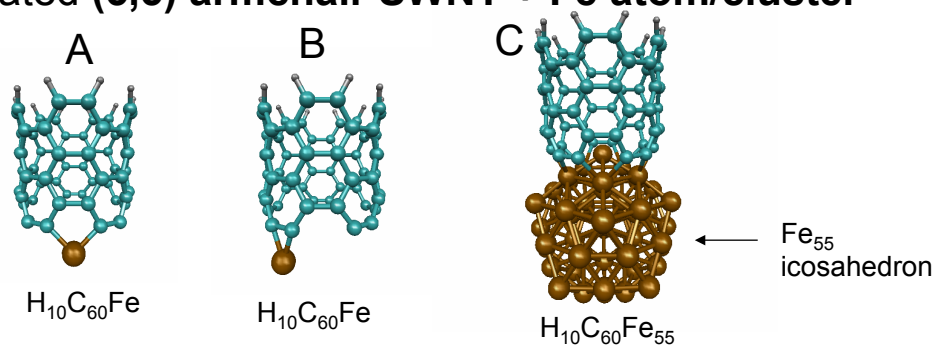
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NCC- and SCC-DFTB for radicals: own comparison



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H-terminated (5,5) armchair SWNT + Fe atom/cluster



Adhesion energies (eV/atom)

	A	B	C
DFT:PW91 ^[1]	-6.24	-5.63	-1.82
SCC-DFTB ^[2]	-5.17	-4.68	-1.86

[1] Phys. Rev. B **75**, 115419 (2007) [2] Fermi broadening=0.13 eV

PW91: An ultrasoft pseudopotential with a plane-wave cutoff of 290 eV for the single metal and the projector augmented wave method with a plane-wave cutoff of 400 eV for the metal cluster

Fe-Fe and Fe-C DFTB parameters from: G. Zheng *et al.*, *J. Chem. Theor. Comput.* **3**, 1349 (2007)

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