

Chemical bond in solids



A real space perspective

Outline

1. Why studying chemical bonds?

2. Quantum Chemical topology

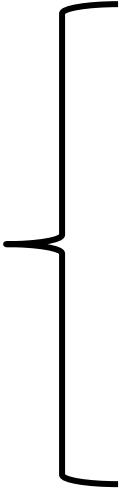
3. Chemical functions

a) electron density

b) ELF

c) NCI

4. Summary

- 
1. The function
 2. The topology
 3. Old concepts
 4. New insight



Outline

5. Applications to high pressure

a) ELF: Predicting compressibility and transitions

b) NCI: He bonds

c) Electron density: the borate family

6. The codes

7. Summary

Outline

1. Why studying chemical bonds?

2. Quantum Chemical topology

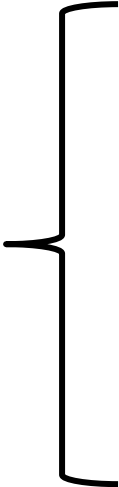
3. Chemical functions

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1. The function
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Motivation

« It's nice to know that the computer understands the problem. But I would like to understand it too. »

E.P. Wigner in Physics Today

The **chemical bond** is a divide-and-conquer approach

Motivation



Desired Properties

Conventional Approach



Materials

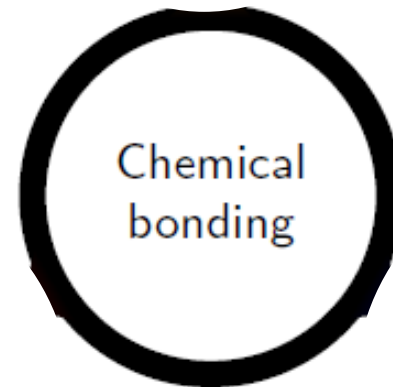
Motivation



Quantum chemical topology

How do we divide and conquer in chemical bonds?

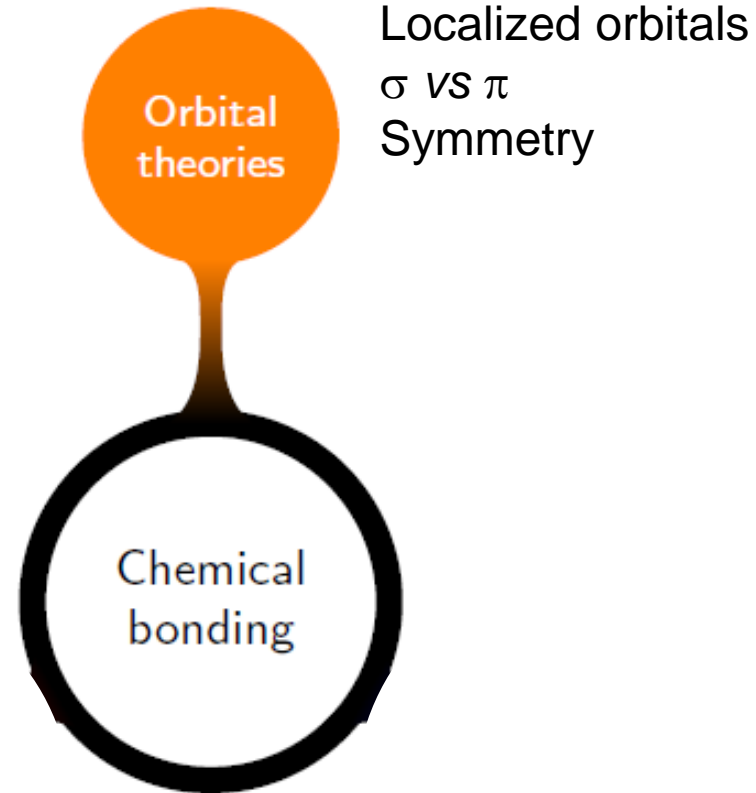
- Quantum chemistry gives us a quantitative description of chemical systems. It is predictive
- Bonds are objects from Classical Chemistry
- We need extra tools to extract quantitative information from Quantum Chemistry
(and reduce the dimensionality)



Quantum chemical topology

How do we divide and conquer in chemical bonds?

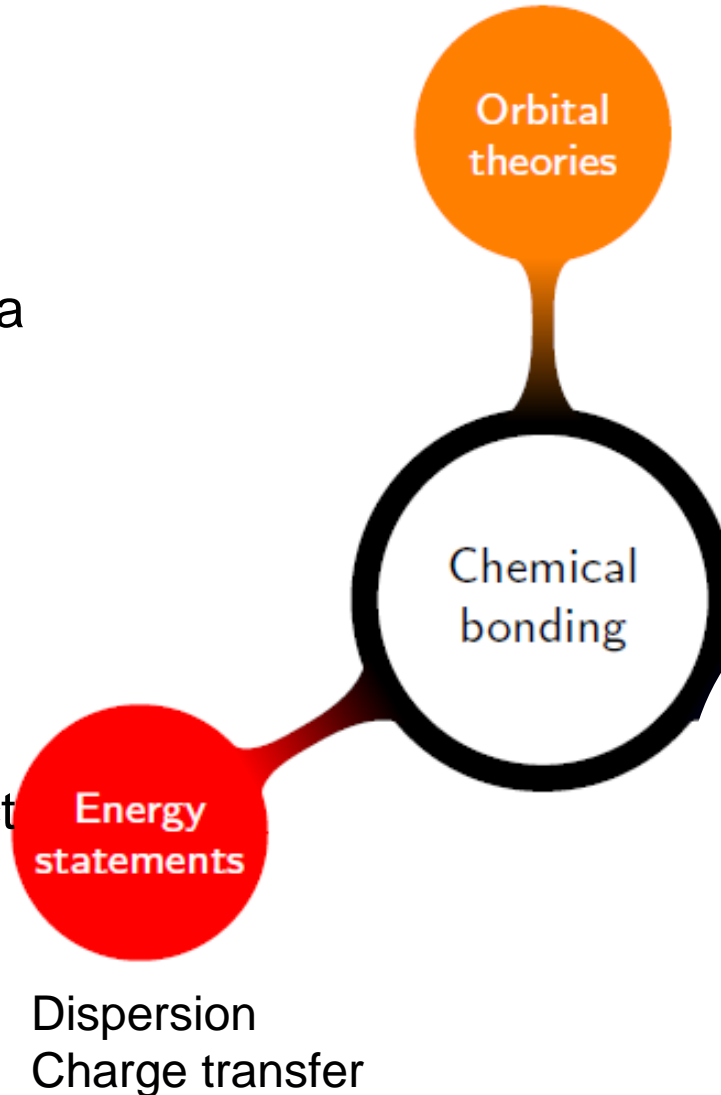
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Quantum chemical topology

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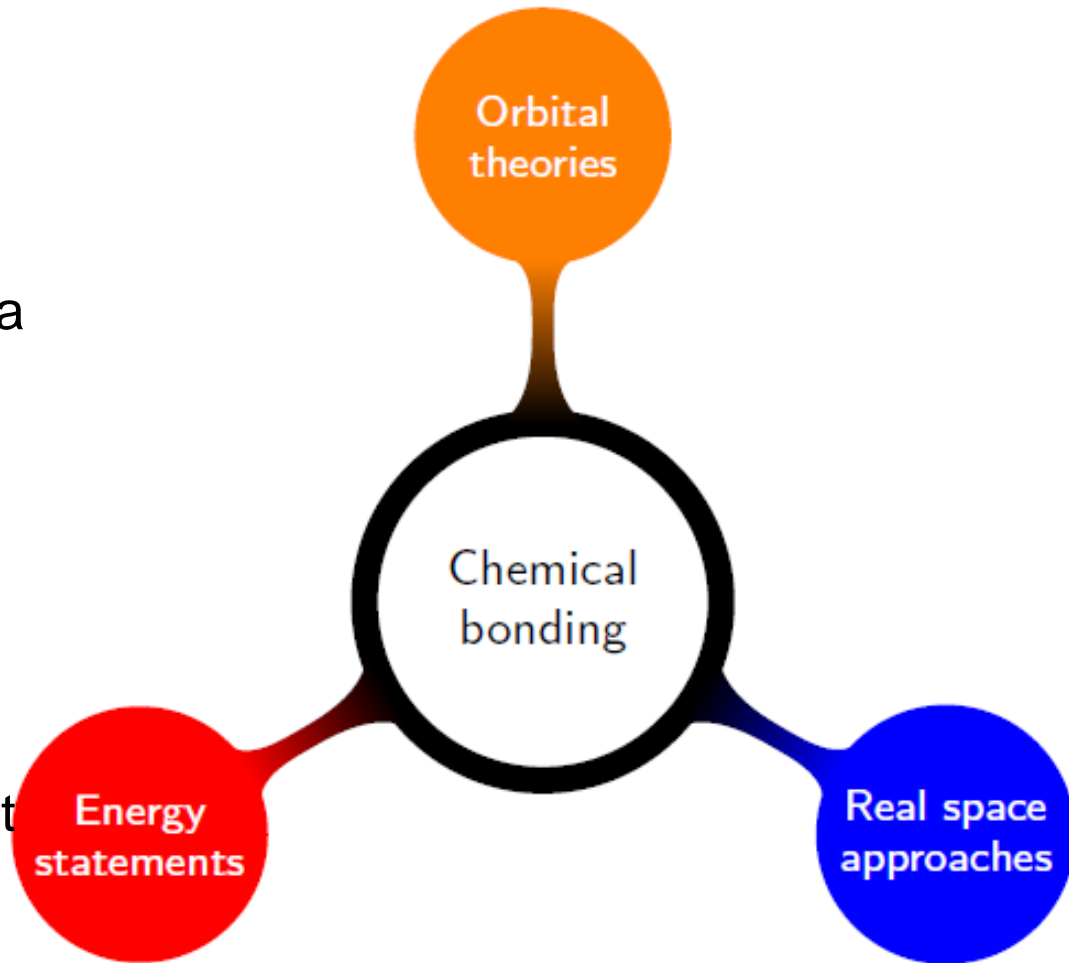
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Quantum chemical topology

How do we divide and conquer in chemical bonds?

- Quantum chemistry gives us a quantitative description of chemical systems. It is predictive
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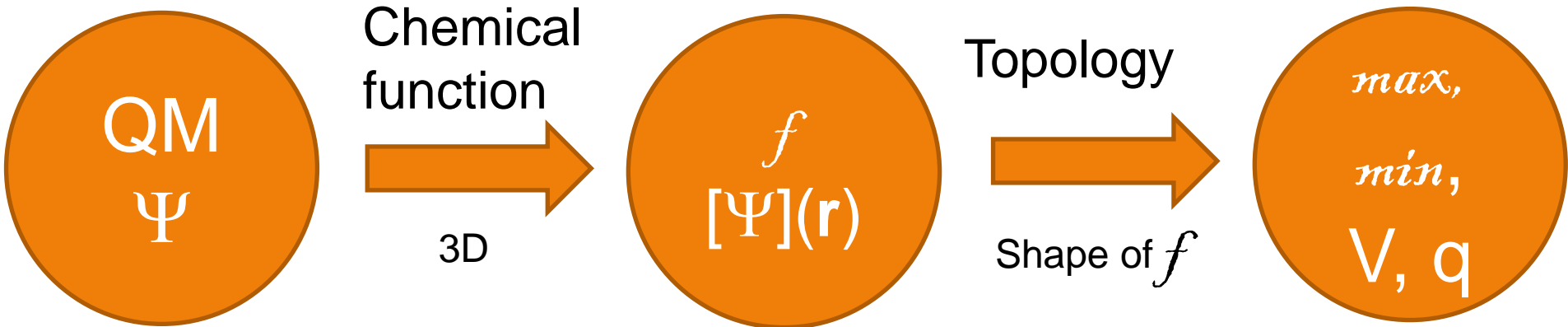
QCT in a nutshell

Quantum topology

Classical Chemistry

3D

Quantum chemistry





QTC in a nutshell

Real space approaches

- Orbital invariant (summation over orbitals)
- Inherently defined in 3D (comprehensible dimensionality)
- Do not depend on the method (applicable to all approximations)
- Usually related to experimental data

- BUT no direct connexion to energetics (lack of the exact functional)

QCT in a nutshell

1

Topology



2

Chemically
sound
functions



3

Quantum
chemical
Topology
(QTC)

QCT in a nutshell

1

Topology

Local information:
critical points
Global information:
basins, integrals



2

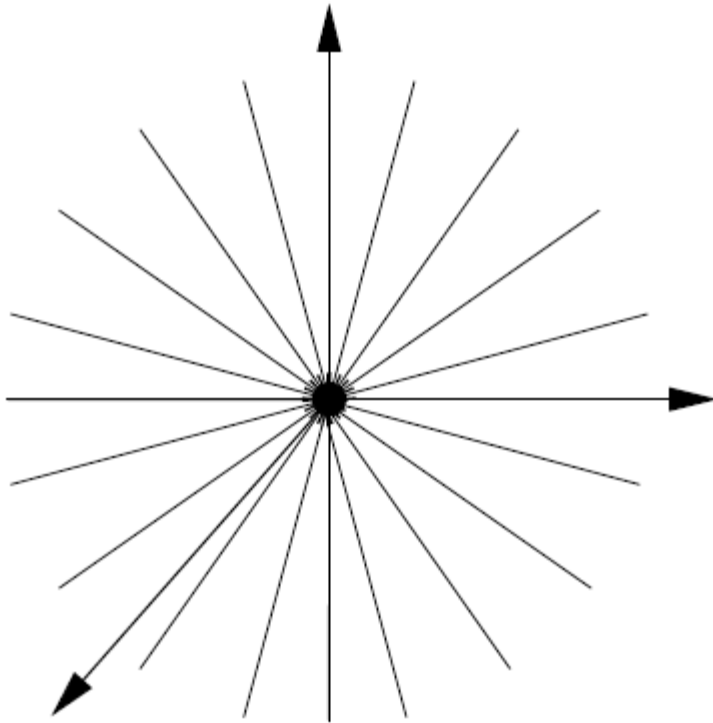
Chemically
sound
functions



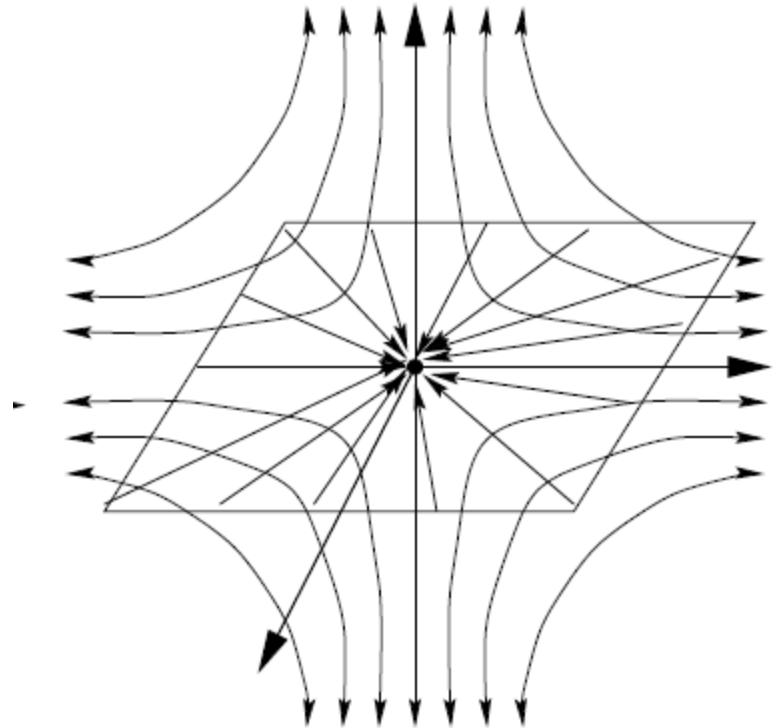
3

Quantum
chemical
Topology
(QTC)

Types of CPs in 3D

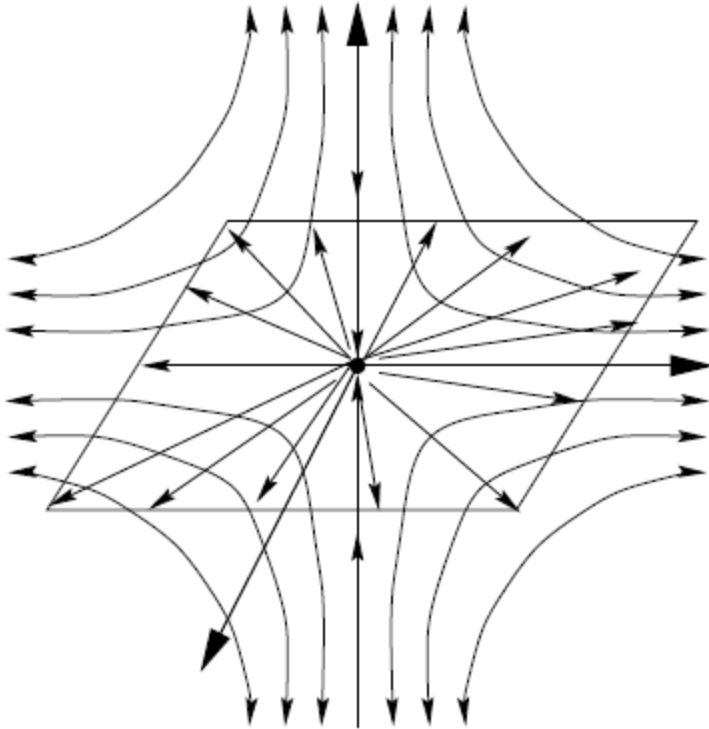


Maximum

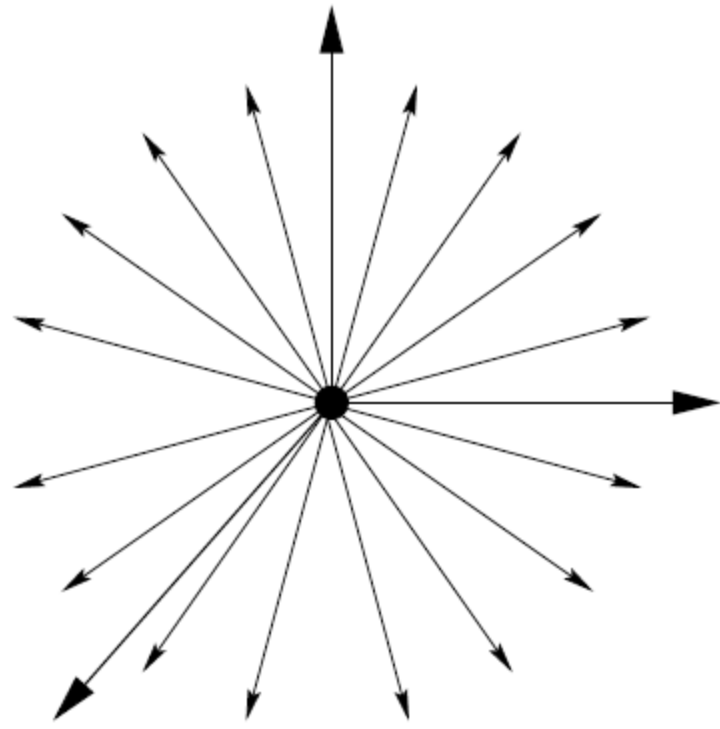


Saddle point of order 1

Types of CPs in 3D

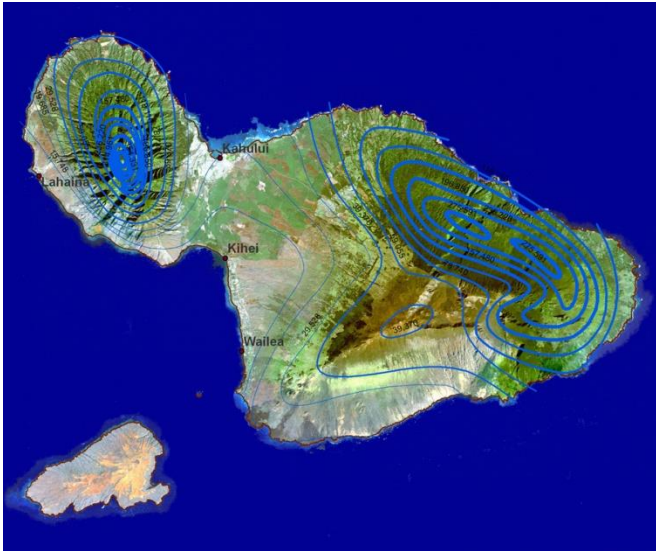


Saddle point of order 2

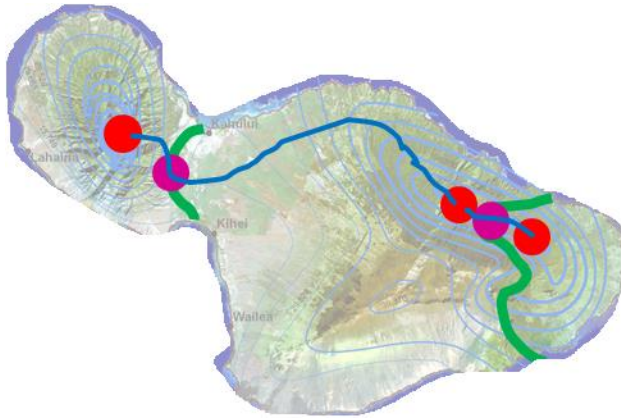


Minimum

Topological partitions are intuitive



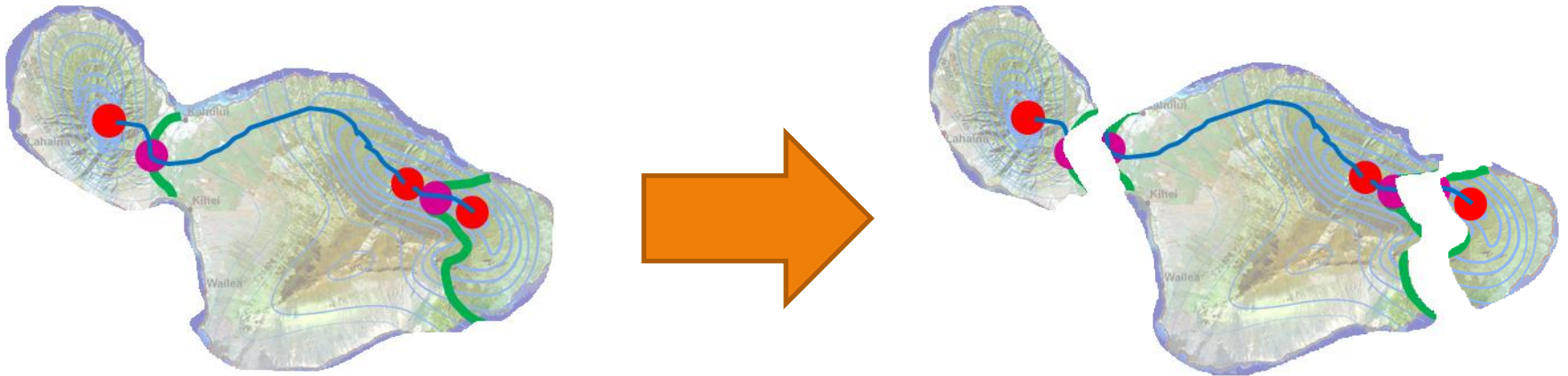
Topological partitions are intuitive



We automatically

- Identify cusps ●
- Divide through the valleys —
- Identify the lowest point in the valley ●

Topological partitions are intuitive



We automatically

- Identify cusps ●
- Divide through the valleys —
- Identify the lowest point in the valley ●

- Use this information to see three regions
- These regions contain orography information

Meaning is inherited

QCT in a nutshell

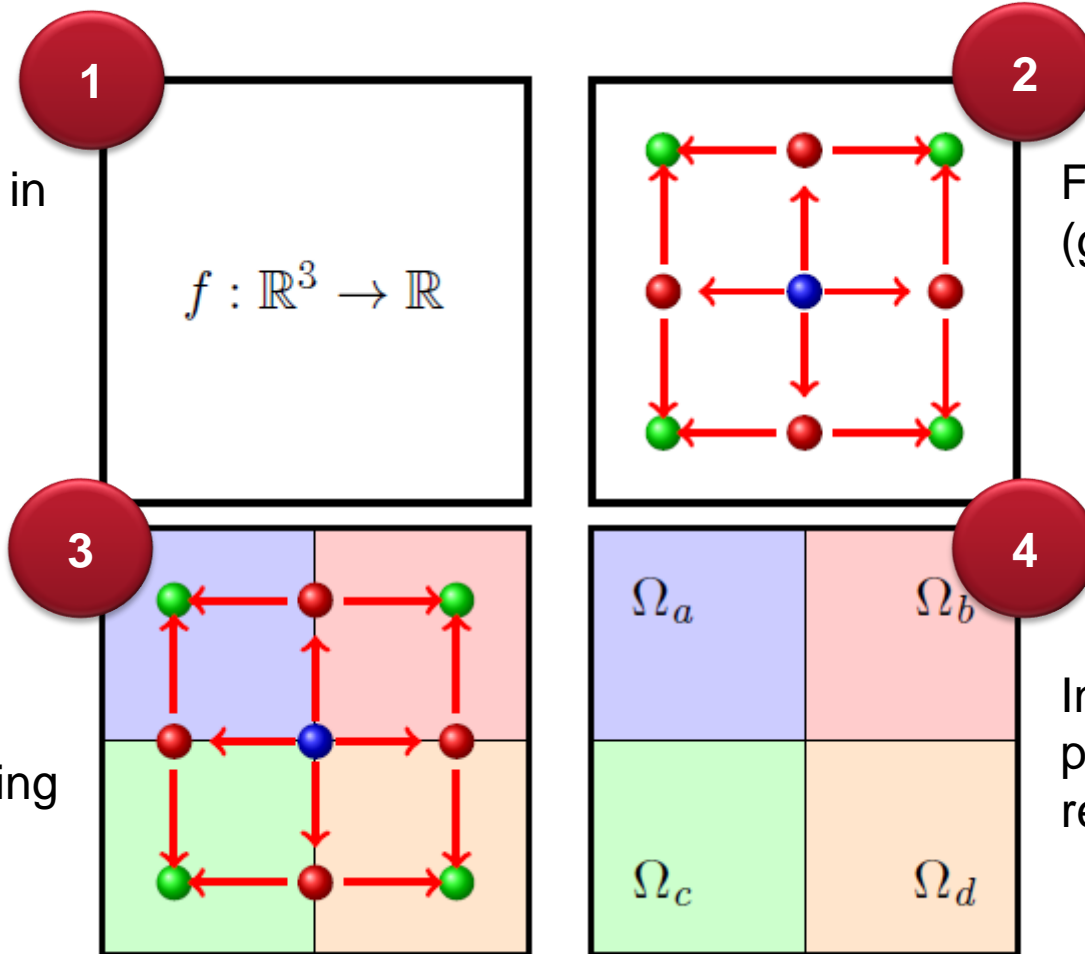
Compute scalar in the molecular /crystal space

$$f : \mathbb{R}^3 \rightarrow \mathbb{R}$$

Find separatrix (flux=0) separating the regions

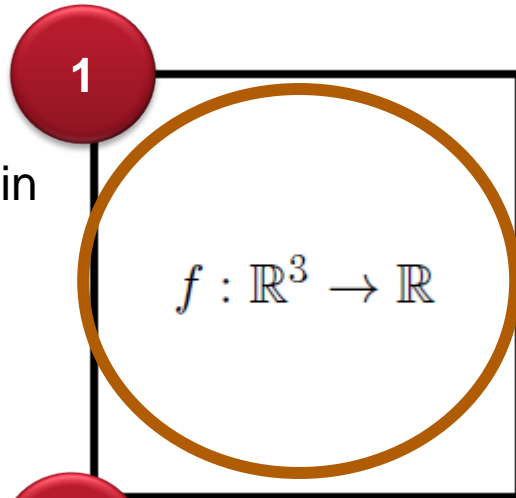
Find critical points (gradient=0)

Integrate properties over regions

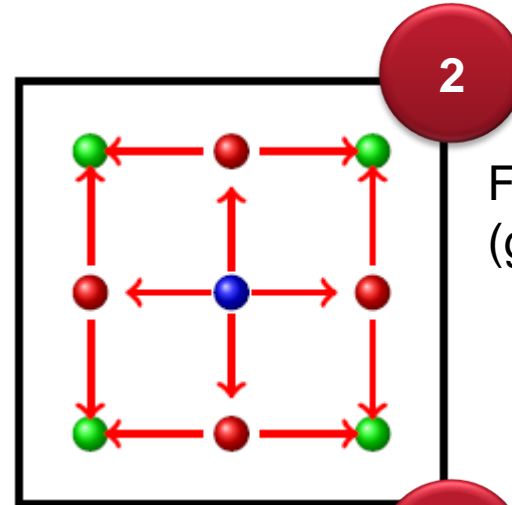


QCT in a nutshell

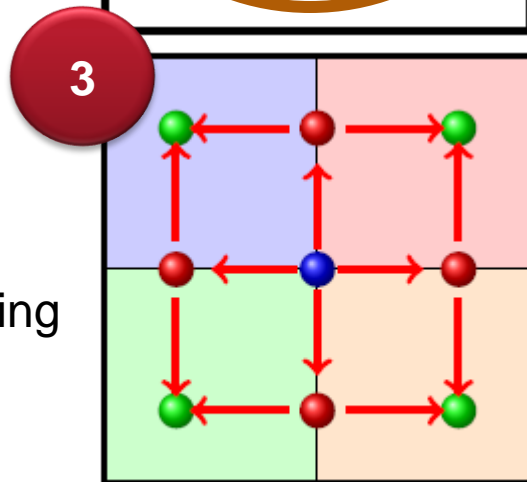
Compute scalar in the molecular /crystal space



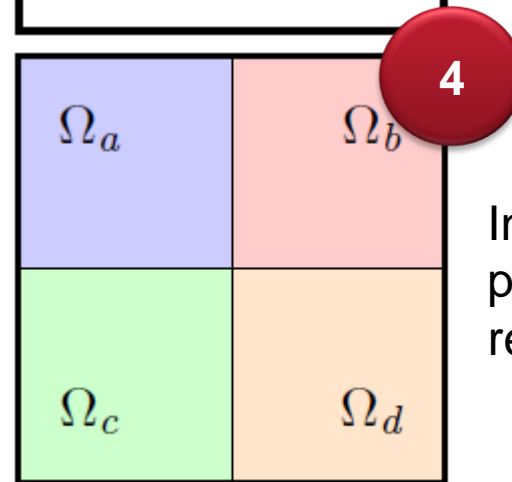
2 Find critical points (gradient=0)



3 Find separatrix (flux=0) separating the regions



4 Integrate properties over regions



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3. Chemical functions
 - a) electron density
 - b) ELF
 - c) NCI
4. Summary

The electron density

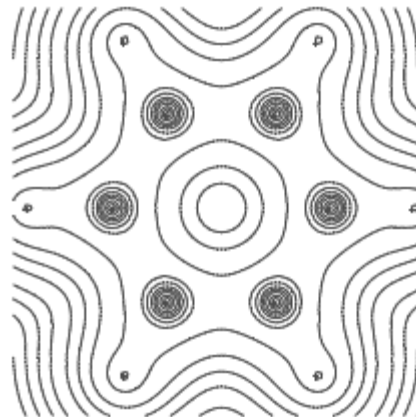
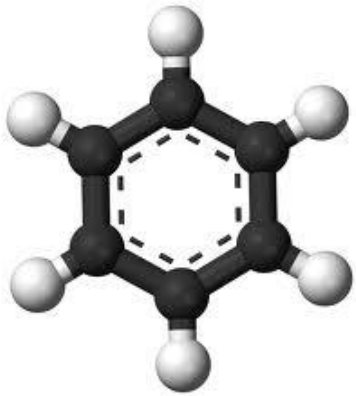
- $\rho(\mathbf{r})$ is a fundamental property of any electronic system

$$\rho(\vec{r}) = N \int \dots \int |\Psi(\vec{x}_1, \vec{x}_2, \dots, \vec{x}_N)|^2 ds d\vec{x}_2 \dots \vec{x}_N$$

- is experimentally accessible
- is defined within the exact many-body theory,
- is supported by the Hohenberg–Kohn theorem

Example: the electron density

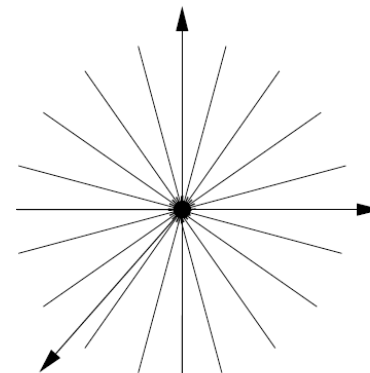
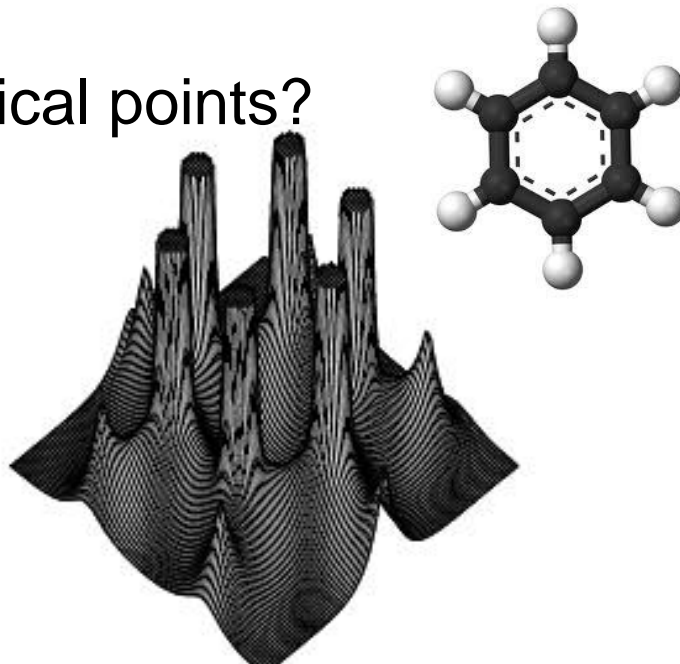
Approxiamately, sum of
exponentials over the
atoms



Example: the electron density

Where are the critical points?

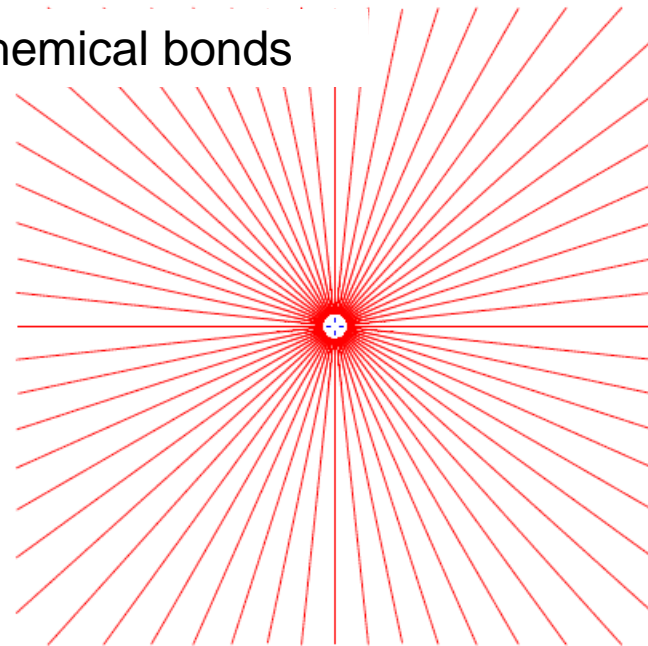
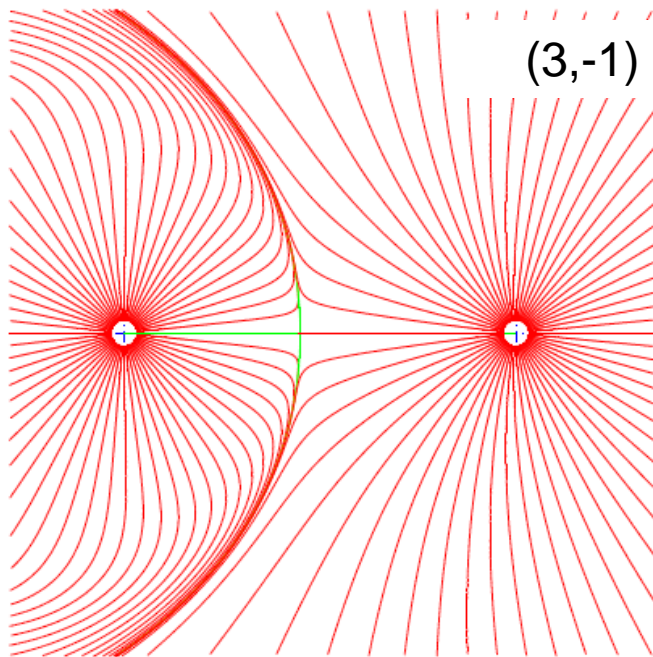
- Maxima = nuclei



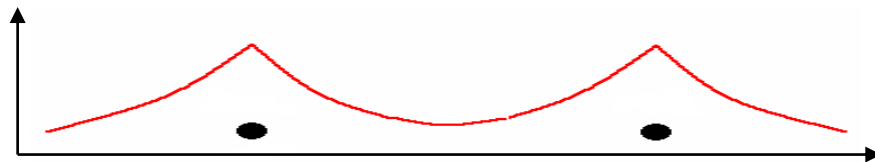
(a) Sumidero de líneas de campo.

The electron density

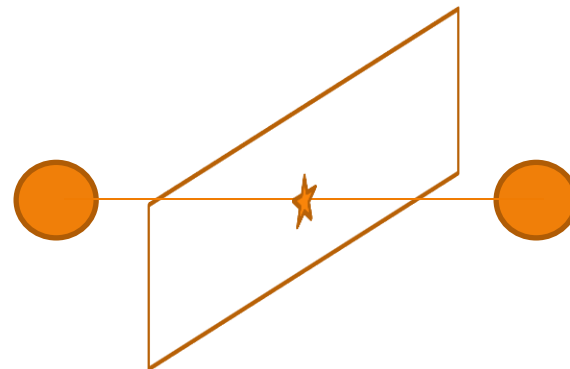
(3,-1) occur at chemical bonds



Plane that contains the nuclei



Plane perpendicular to the internuclear line at the critical point that contains the nuclei

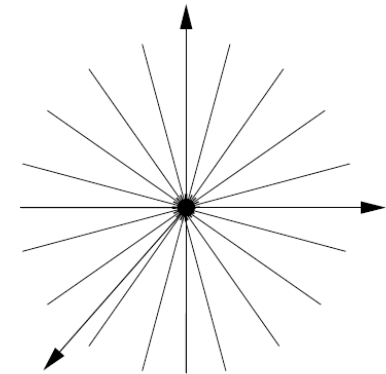
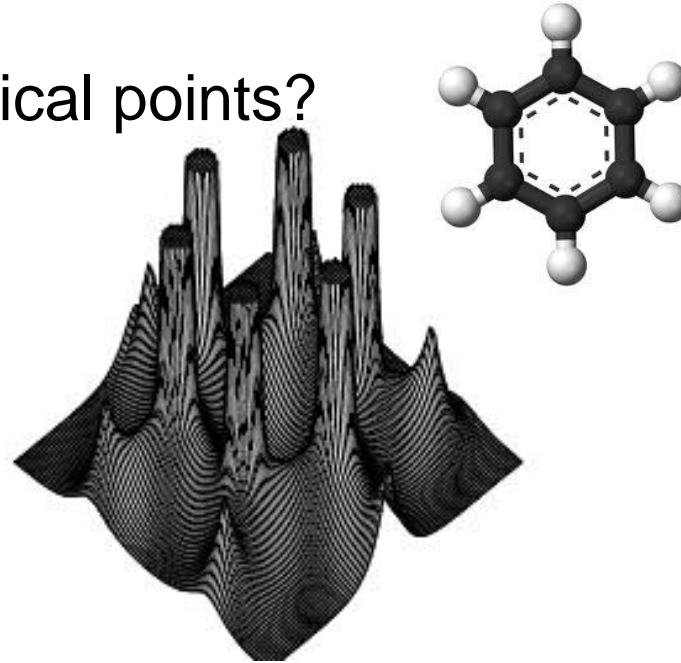


Example: the electron density

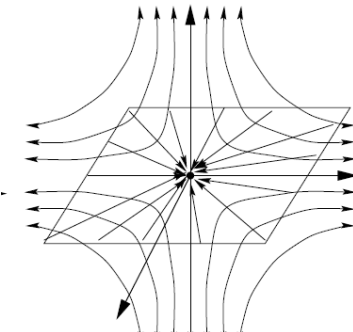
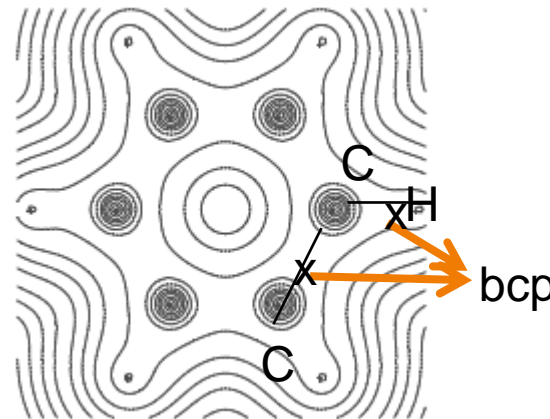
Where are the critical points?

- Maxima = nuclei

- 1st order saddle points=bonds



(a) Sumidero de líneas de campo.



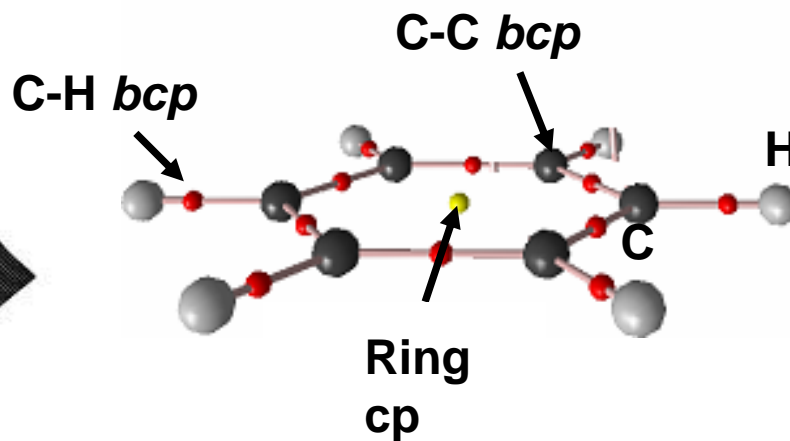
(b) Silla tipo 1.

The electron density

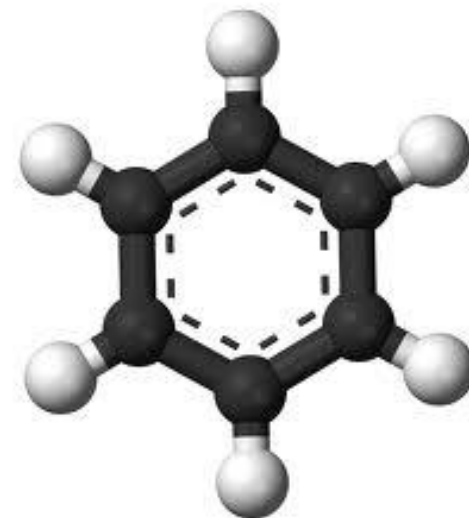
We can know from ab initio methods the chemical structure
i.e. **Where do we draw lines between atoms?**



Density



ρ topology



Chemical structure



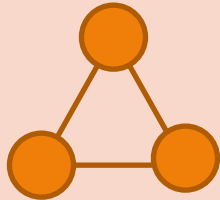
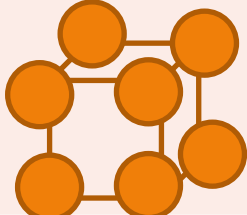
- Maxima = nuclei
- Bond = 1st order saddle point (*bcp*)

QCT in a nutshell

Directions along which the field grows

Directions along which the field decreases

Sum of signs of eigenvalues

CP	$\lambda > 0$	$\lambda < 0$	Signature (s)	(r,s)	name	acronym	Figure
Maximum	0	3	-3	(3,-3)	maximum		
1 st order saddle point	1	2	-1	(3,-1)	Bond critical point	bcp	
2 nd order saddle point	2	1	+1	(3,+1)	Ring critical point	rcp	
Minimum	3	0	+3	(3,+3)	Cage critical point	ccp	

Local information: CPs

- In a periodic system, you will always have the 4 types of non-degenerated critical points
- Morse relationship MUST hold:

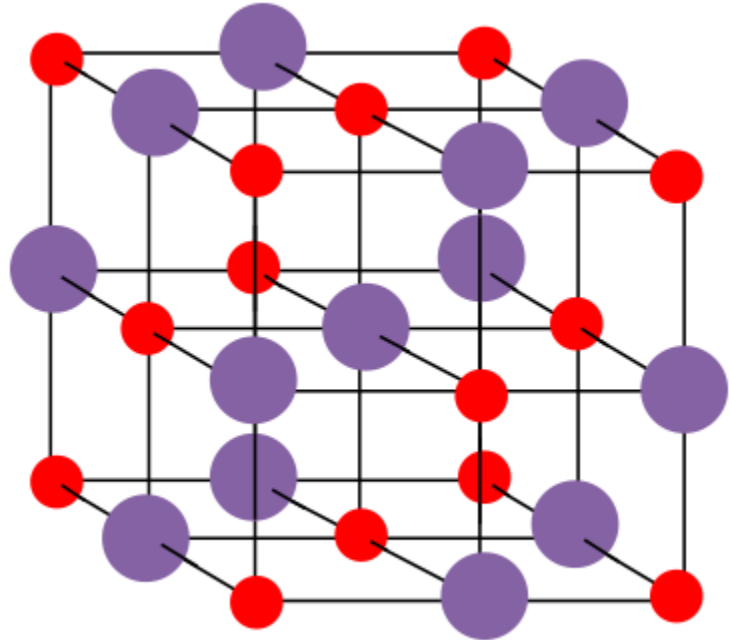
$$n_{(3,-3)} - n_{(3,-1)} + n_{(3,+1)} - n_{(3,+3)} = 0$$

$$\left\{ \begin{array}{l} n_{(3,-3)} \geq 1 \\ n_{(3,-1)} \geq 3 \\ n_{(3,+1)} \geq 3 \\ n_{(3,+3)} \geq 1 \end{array} \right.$$

- If $\lambda_i = 0$, these are degenerated critical « points » (we will see an example later).

E.g. (2,-2)

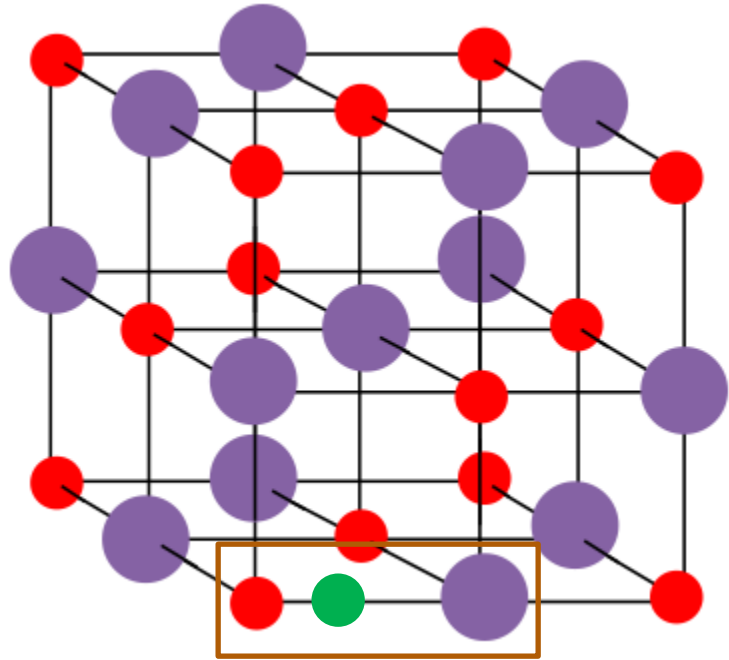
Local information: CPs



- Attractors are at the nuclei positions

Simet.	Clase	Tipo	x	y	z
Oh	(3,-3)	Nucl. Na	0.00000	0.00000	0.00000
Oh	(3,-3)	Nucl. Cl	0.50000	0.50000	0.50000

Local information: CPs

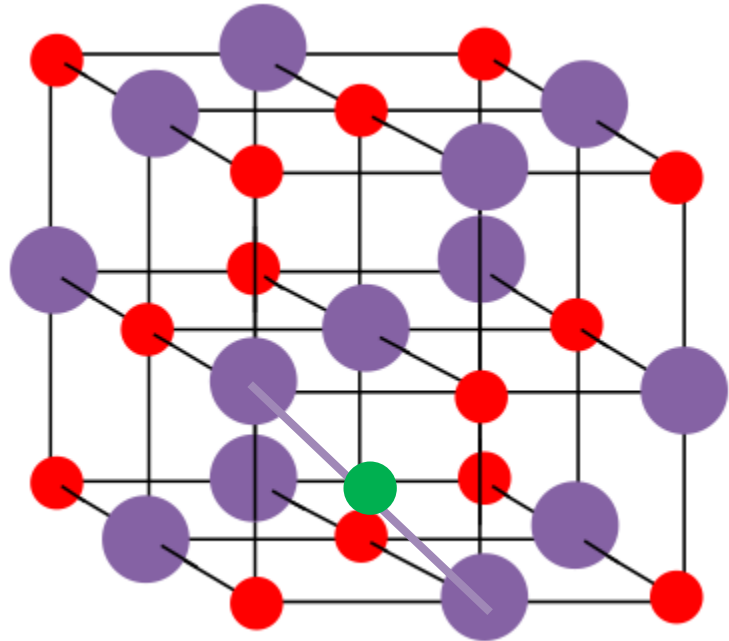


- Attractors are at the nuclei positions
- $(3,-1)$ signal interactions

Simet.	Clase	Tipo	x	y	z
Oh	(3,-3)	Nucl. Na	0.00000	0.00000	0.00000
Oh	(3,-3)	Nucl. Cl	0.50000	0.50000	0.50000
C3v	(3,-1)	Enlace	0.20618	0.20618	0.20618

Cation-anion

Local information: CPs

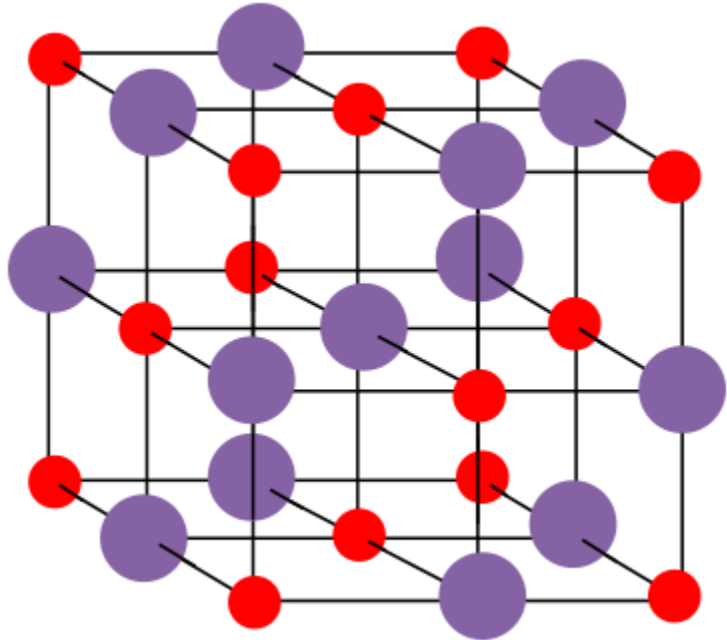


- Attractors are at the nuclei positions
- $(3,-1)$ signal interactions

Anion-anion interactions are quite common in crystals

Simet.	Clase	Tipo	x	y	z	
Oh	(3,-3)	Nucl. Na	0.00000	0.00000	0.00000	
Oh	(3,-3)	Nucl. Cl	0.50000	0.50000	0.50000	
C3v	(3,-1)	Enlace	0.20618	0.20618	0.20618	Cation-anion
D4h	(3,-1)	Enlace	0.00000	0.50000	0.50000	Anion-anion

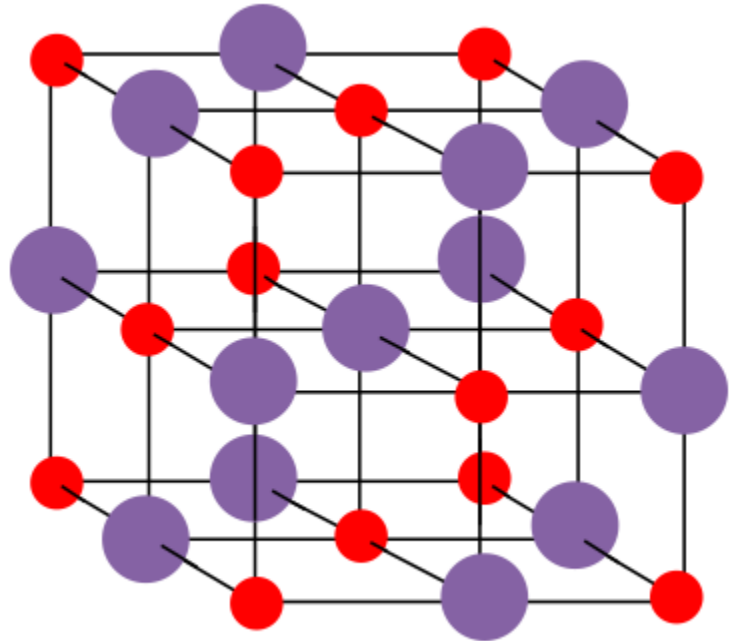
Local information: CPs



- We have all types of critical points

Simet.	Clase	Tipo	x	y	z
Oh	(3,-3)	Nucl. Na	0.00000	0.00000	0.00000
Oh	(3,-3)	Nucl. Cl	0.50000	0.50000	0.50000
C3v	(3,-1)	Enlace	0.20618	0.20618	0.20618
D4h	(3,-1)	Enlace	0.00000	0.50000	0.50000
C2v	(3, 1)	Anillo	0.00000	0.28136	0.28136
D4h	(3, 3)	Caja	0.00000	0.00000	0.50000

Local information: CPs



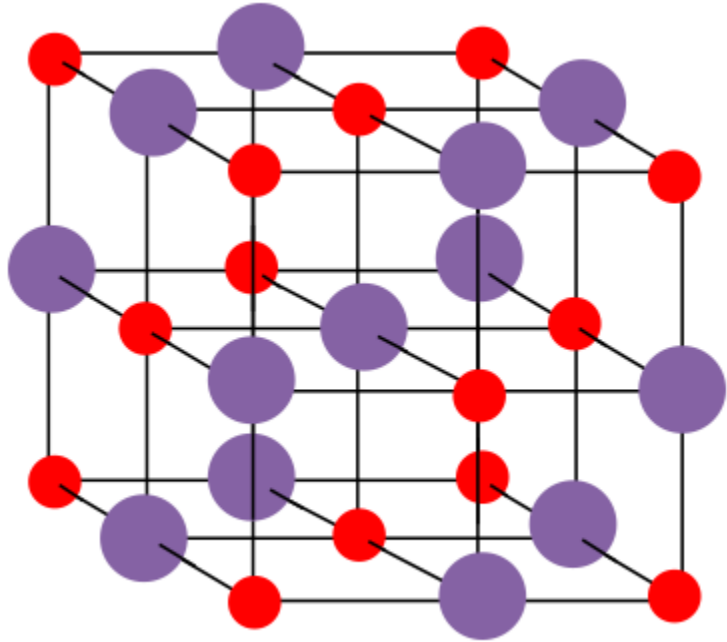
- We have all types of critical points
- Morse sum is working

$$2 - 11 + 12 - 3 = 0$$




Simet.	Clase	Tipo	x	y	z	Mult.
Oh	(3,-3)	Nucl. Na	0.00000	0.00000	0.00000	+ 1
Oh	(3,-3)	Nucl. Cl	0.50000	0.50000	0.50000	+ 1
C3v	(3,-1)	Enlace	0.20618	0.20618	0.20618	- 8
D4h	(3,-1)	Enlace	0.00000	0.50000	0.50000	- 3
C2v	(3, 1)	Anillo	0.00000	0.28136	0.28136	+ 12
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Local information: CPs



- We have all types of critical points
- Morse sum is working

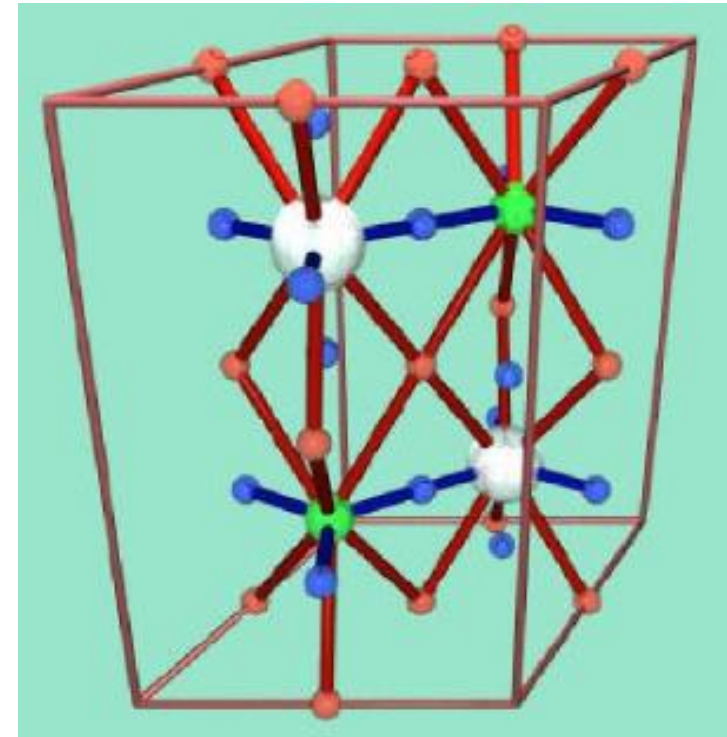
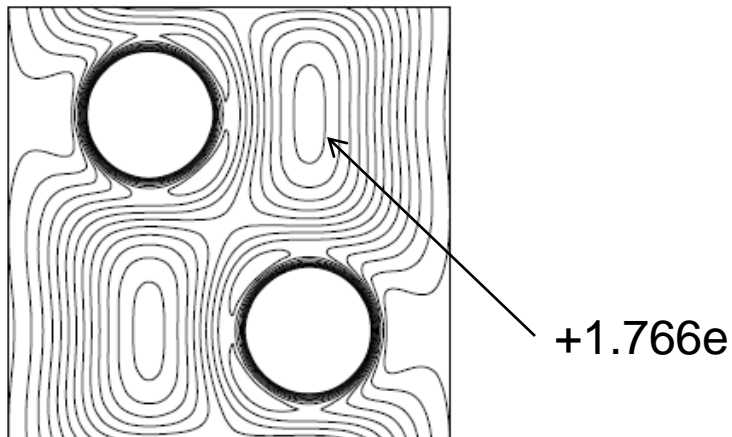
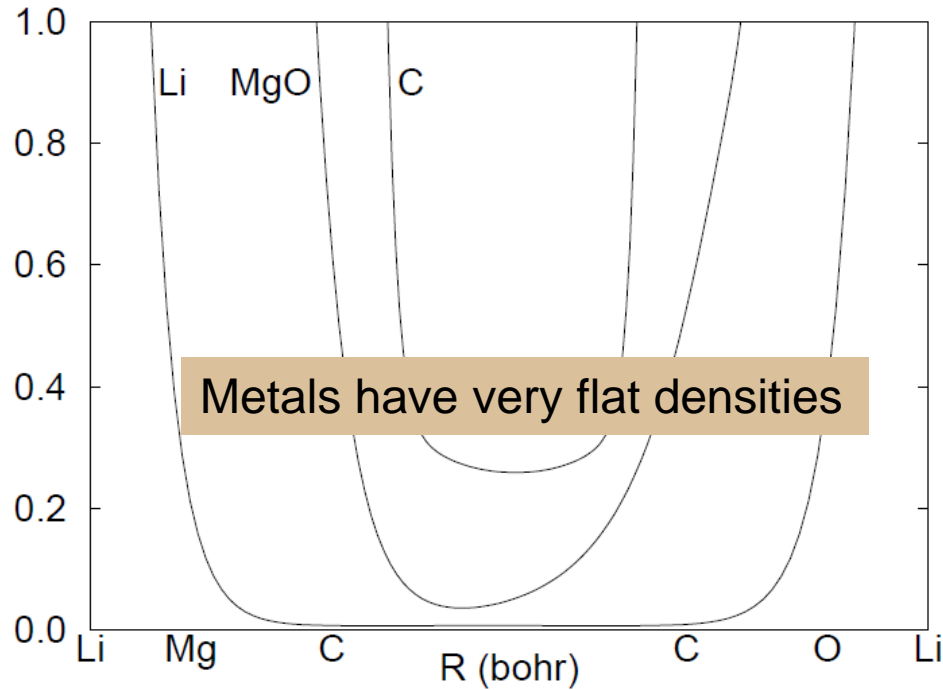
$$2 - 11 + 12 - 3 = 0$$

- We only have (3,-3) at the nuclei?

Simet.	Clase	Tipo	x	y	z	Mult.
Oh	(3,-3)	Nucl. Na	0.00000	0.00000	0.00000	+ 1
Oh	(3,-3)	Nucl. Cl	0.50000	0.50000	0.50000	+ 1
C3v	(3,-1)	Enlace	0.20618	0.20618	0.20618	- 8
D4h	(3,-1)	Enlace	0.00000	0.50000	0.50000	- 3
C2v	(3, 1)	Anillo	0.00000	0.28136	0.28136	+ 12
D4h	(3, 3)	Caja	0.00000	0.00000	0.50000	- 3

Local information: CPs

Berillium HCP

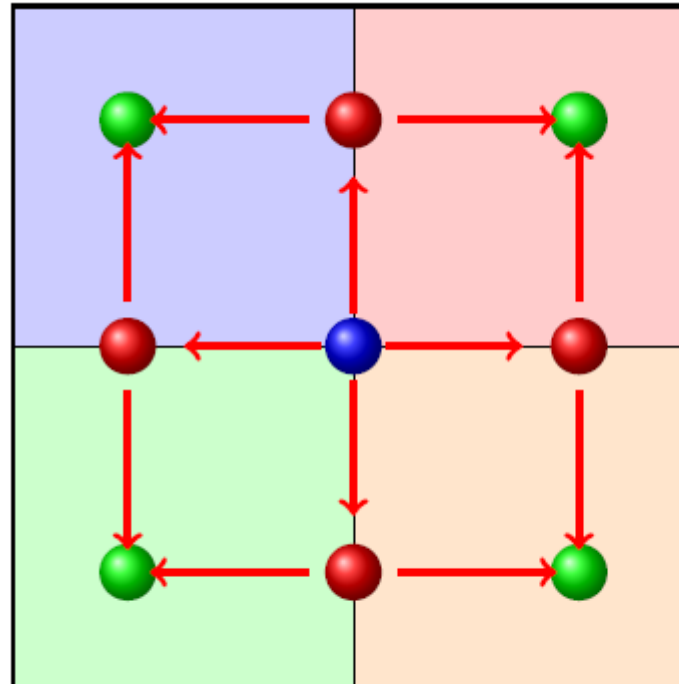
PC	x	y	z	$\rho(\text{PC})$
$n(\text{Be})$	0.3333	0.6667	0.2500	32.6222
$n(\text{MNN})$	0.3333	0.6667	0.7500	0.0488



QCT in a nutshell

3

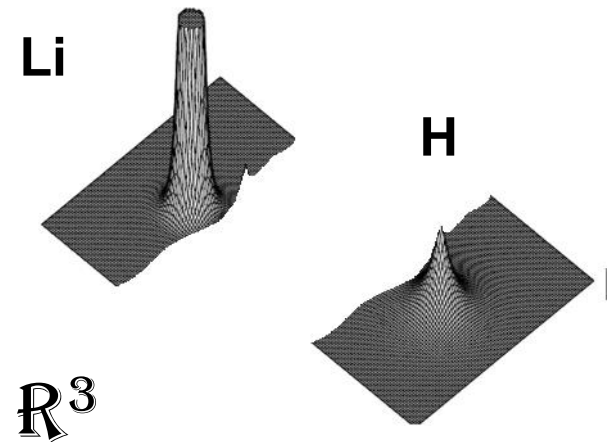
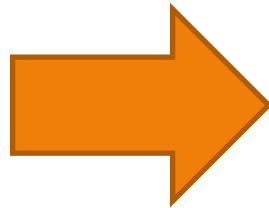
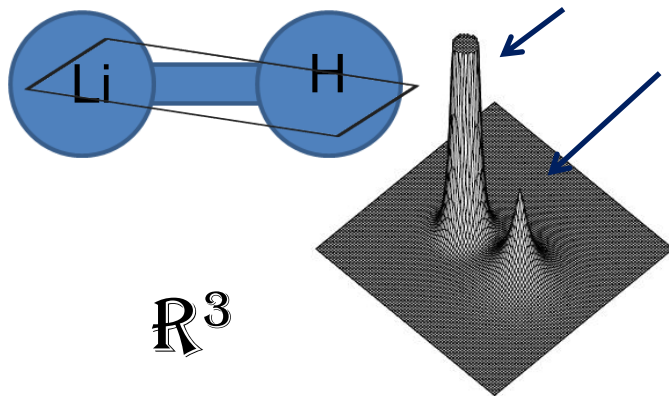
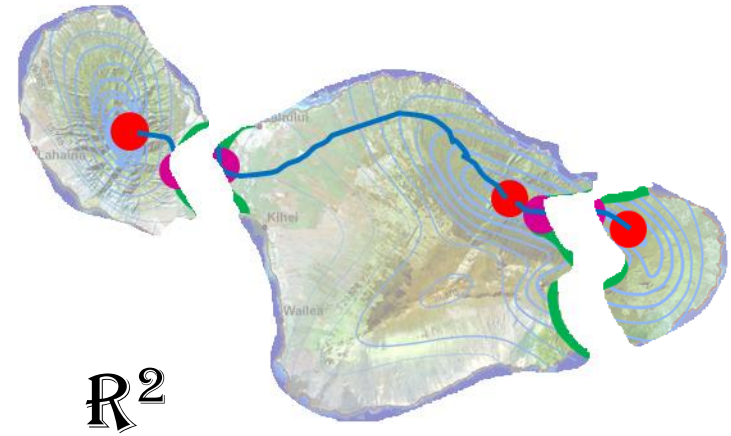
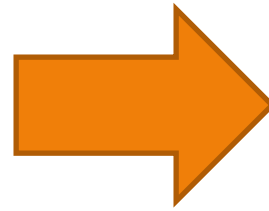
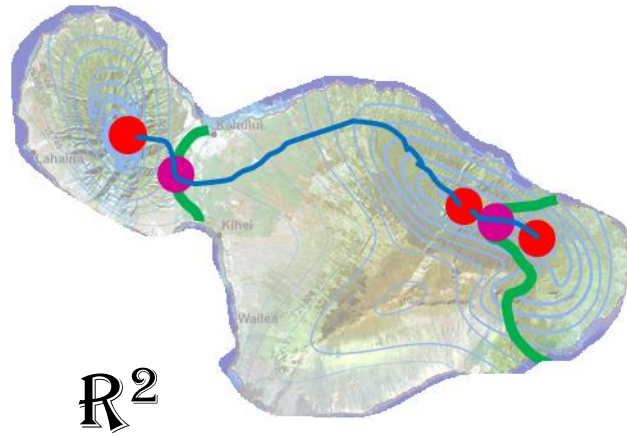
Computing the Morse complex of f



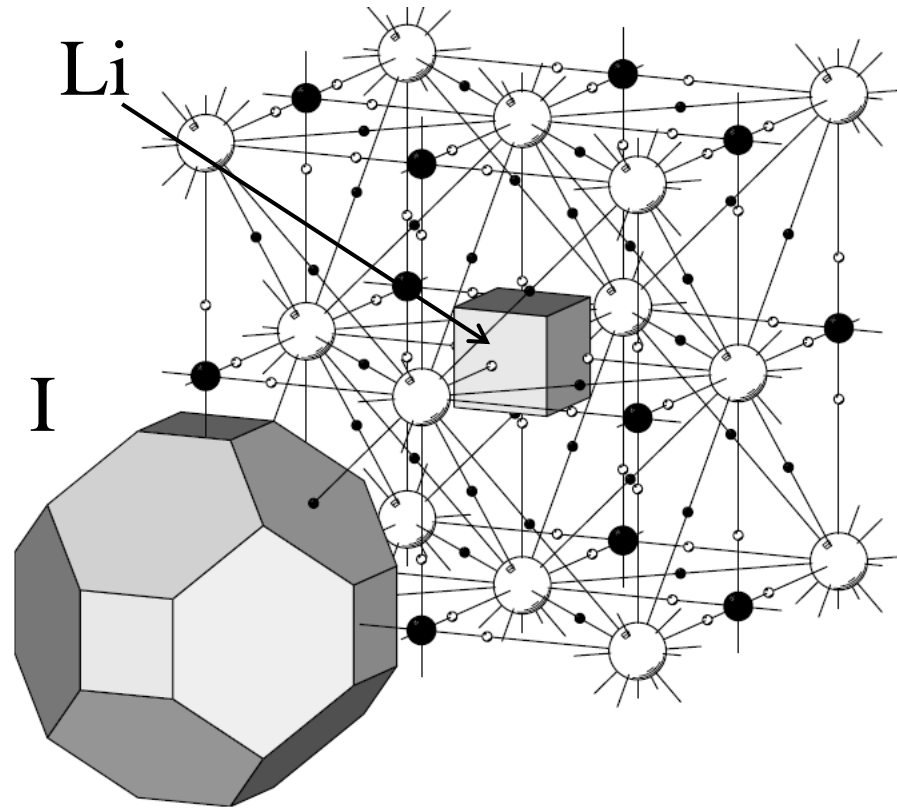
● Maximum ● (3,-1) ● (3,+1)

Each maximum has an associated region of space (basin)

Example: the electron density

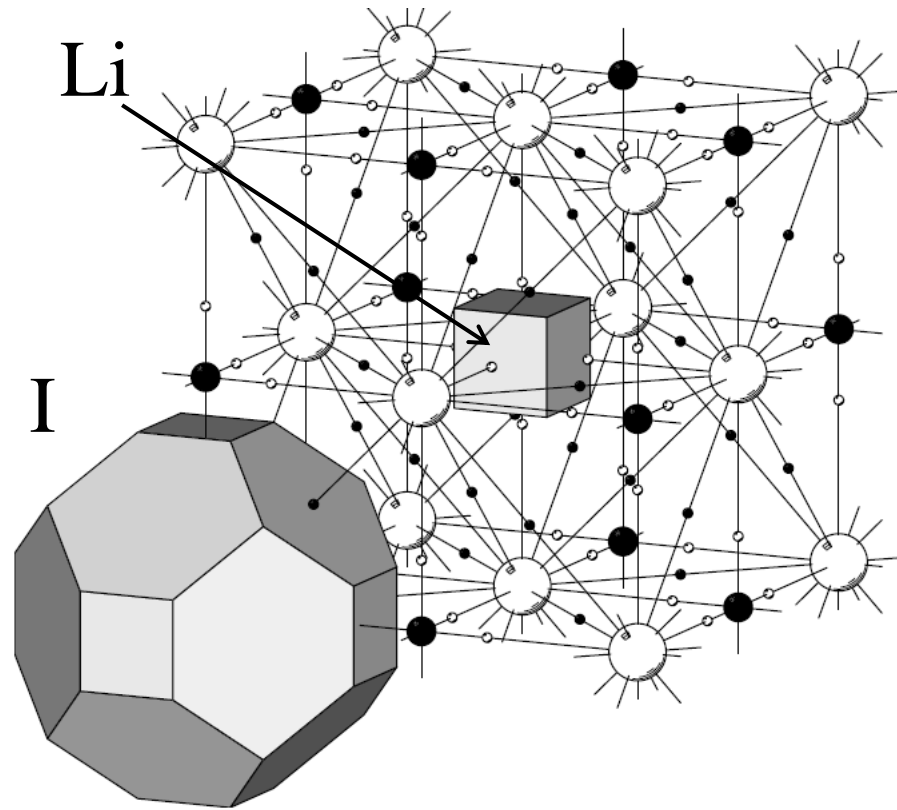


Old concepts

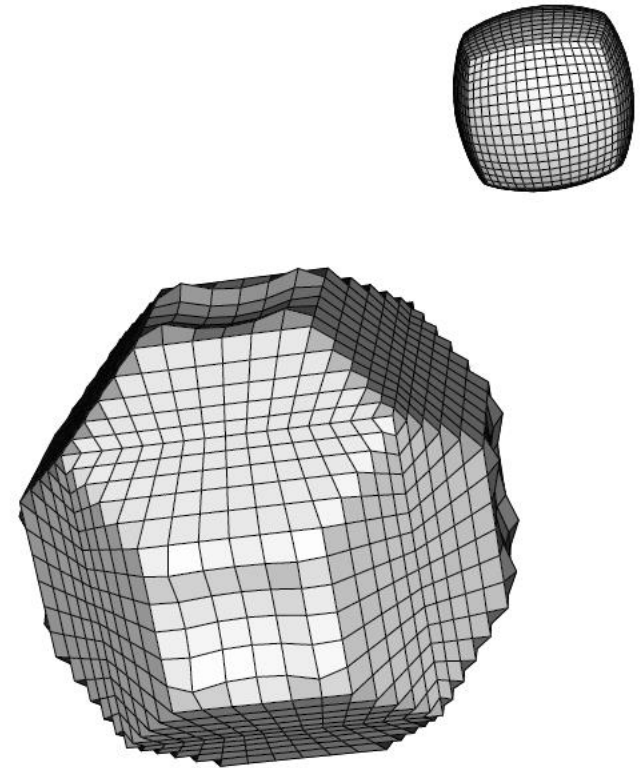


Contact Polyhedra

Old concepts



Contact Polyhedra



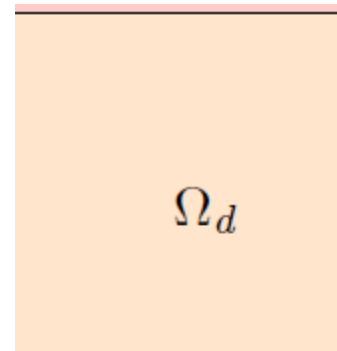
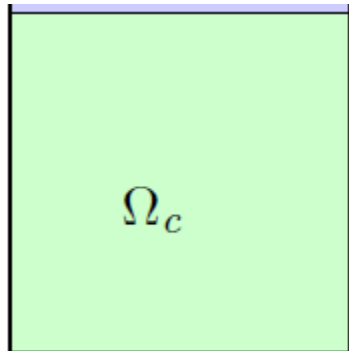
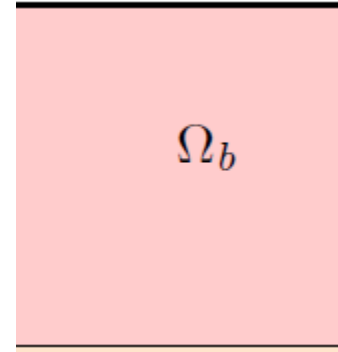
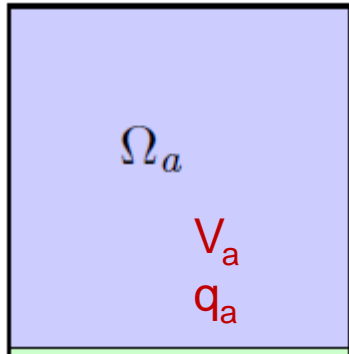
Electron density basins

QCT in a nutshell

4

Calculate properties within the basins of $f(\Omega)$

$$q_A = \int_{\Omega_A} \rho dV$$

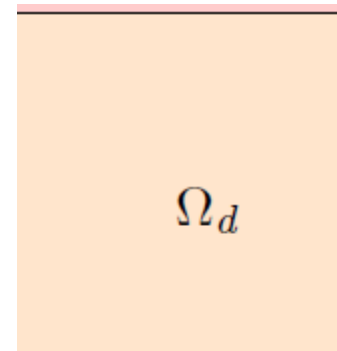
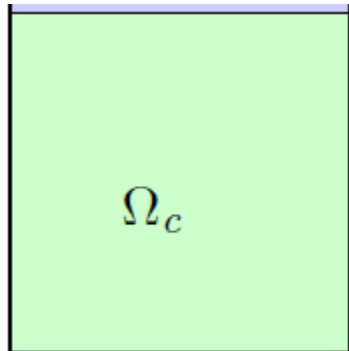
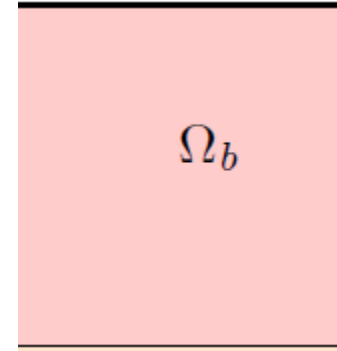
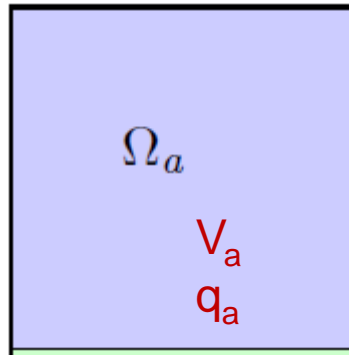


QCT in a nutshell

4

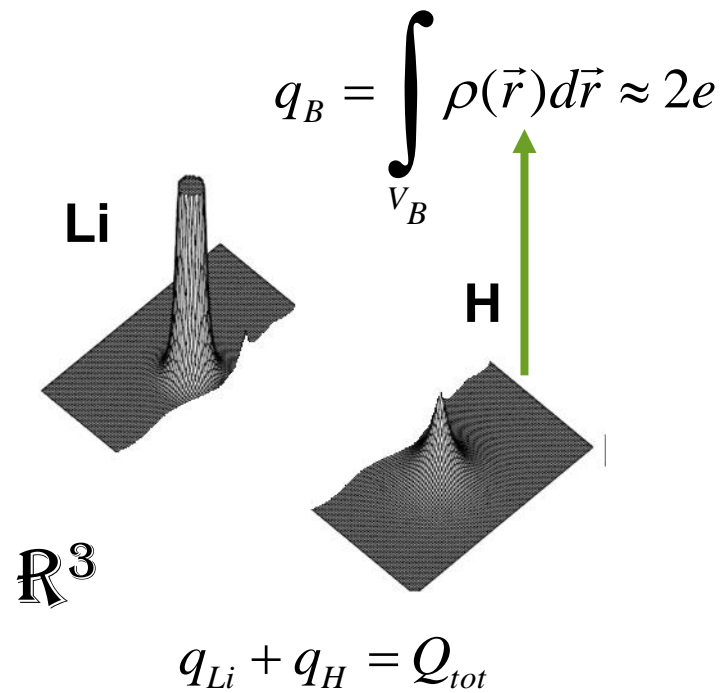
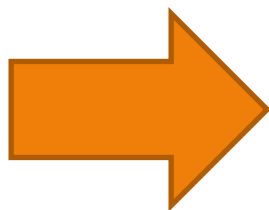
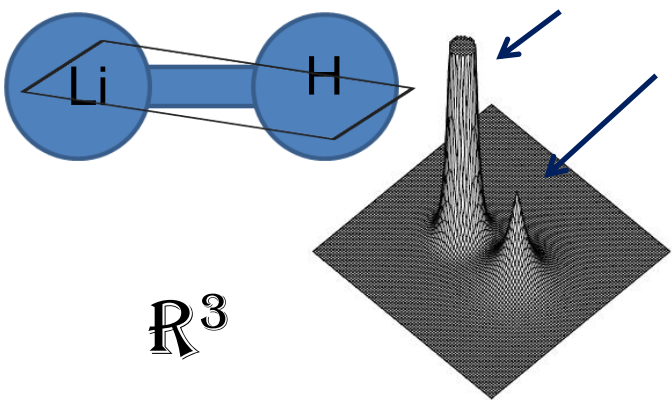
Calculate properties within the basins of $f(\Omega)$

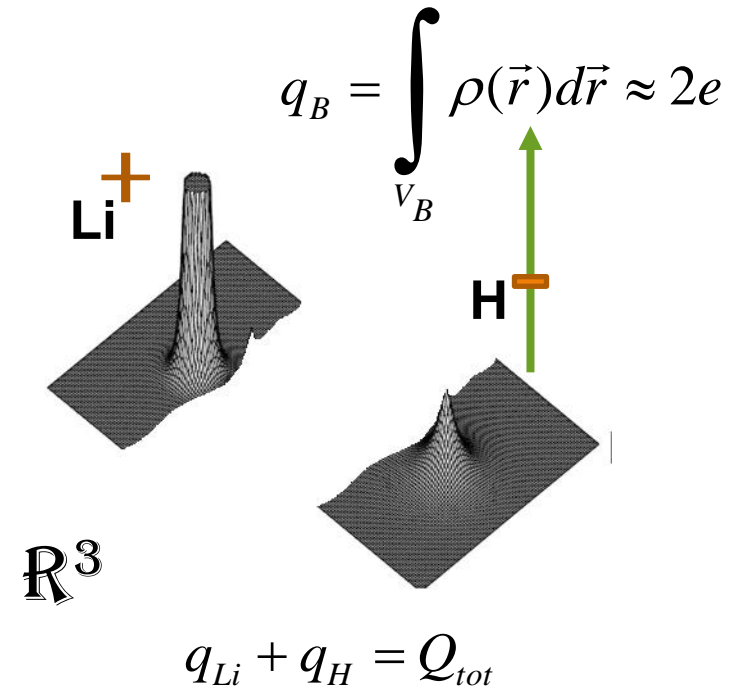
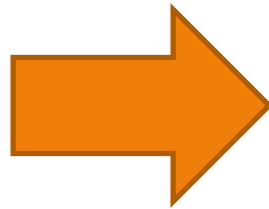
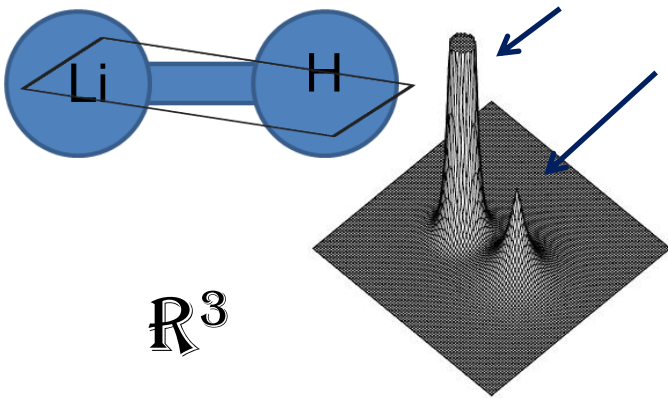
$$q_A = \int_{\Omega_A} \rho dV$$



$$N = \sum_A q_A$$

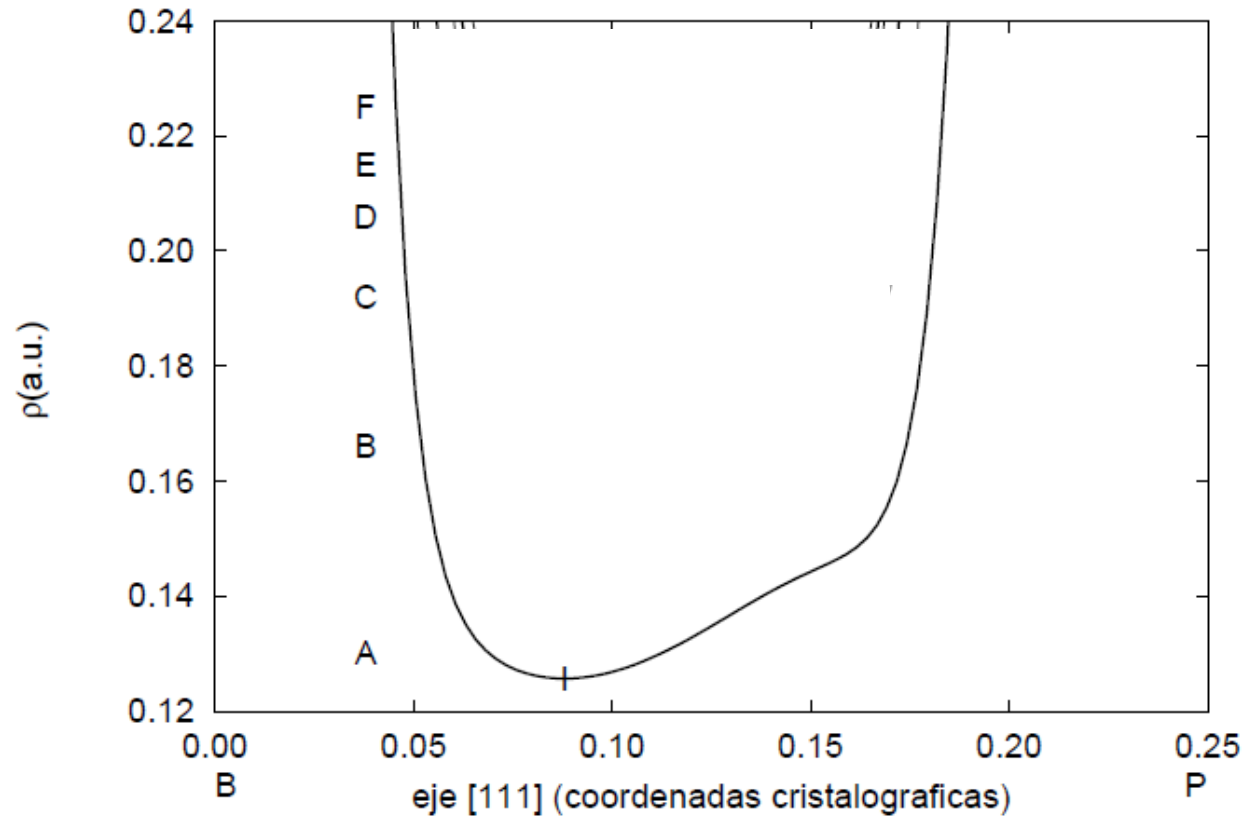
Properties are additive





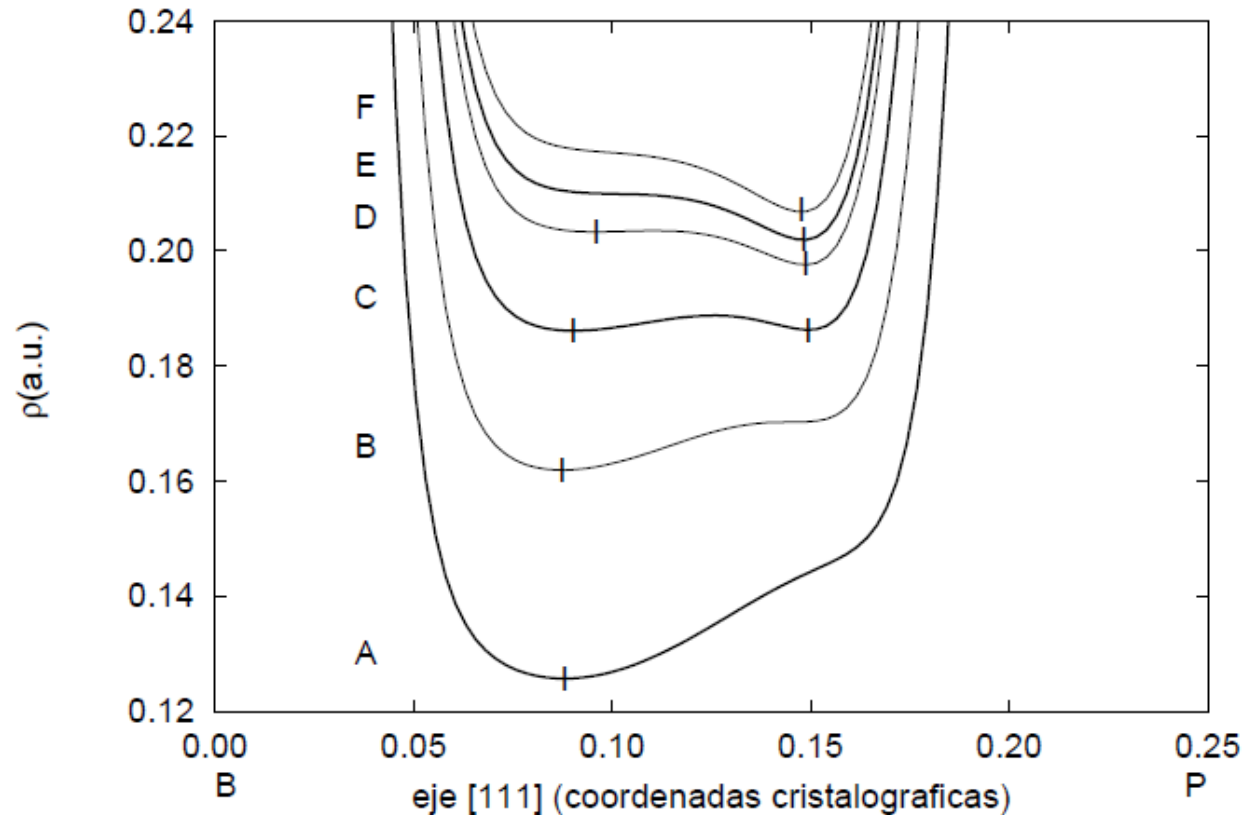
Properties

- BP



Properties

- BP



When the crystal is compressed, the electron density is displaced, leading to a change in polarity from $B^{\delta+}P^{\delta-}$ at V_0 to $B^{\delta-}P^{\delta+}$ at $V=0.57V_0$

Outline

1. Why studying chemical bonds?

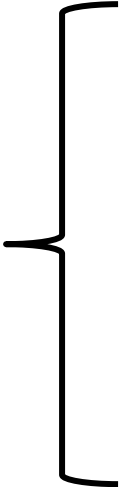
2. Quantum Chemical topology

3. Chemical functions

a) electron density

b) ELF

c) NCI

- 
1. The function
 2. The topology
 3. Old insight
 4. New insight

4. Summary

The electron localization function

Fermionic information

- ELF can be interpreted as an excess of local kinetic energy density due to Pauli repulsion.

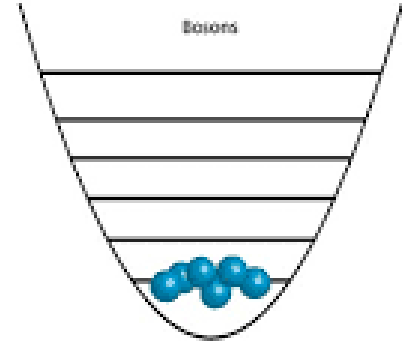
$$t_p(\vec{r}) = t(\vec{r}) - \frac{1}{8} \frac{|\nabla \rho(\vec{r})|^2}{\rho(\vec{r})}$$

The electron localization function

Bosonic system (iso-orbital)

If we are in a region that can be described by a solely orbital j :

$$\rho = \sum_i |\varphi_i|^2 = |\varphi_j|^2 \quad \left\{ \begin{array}{l} \varphi_j = \sqrt{\rho} \\ \nabla \varphi_j = \frac{\nabla \rho}{2\sqrt{\rho}} \end{array} \right.$$



Kinetic energy densities

$$t = \frac{1}{2} \sum_i \nabla \varphi_i^* \nabla \varphi_i \quad \longrightarrow \quad t_w = \frac{1}{2} \frac{\nabla \rho}{2\sqrt{\rho}} \frac{\nabla \rho}{2\sqrt{\rho}} = \frac{1}{8} \frac{(\nabla \rho)^2}{\rho}$$

$$t_{bose}(\mathbf{r}) = \frac{\tau_w(\mathbf{r})}{\tau_{TF}(\mathbf{r})}$$

Its a scaled reference

- with the same density
- Pauli principle has been turned off

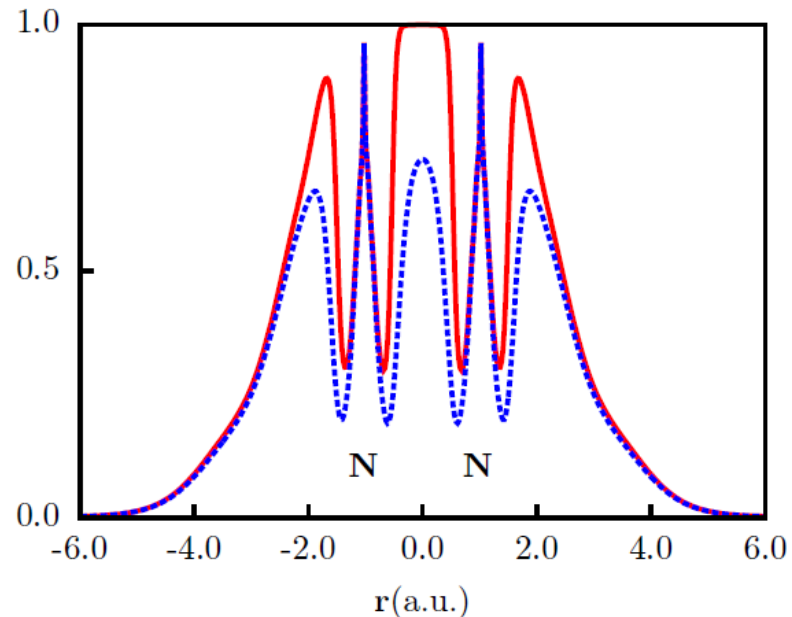
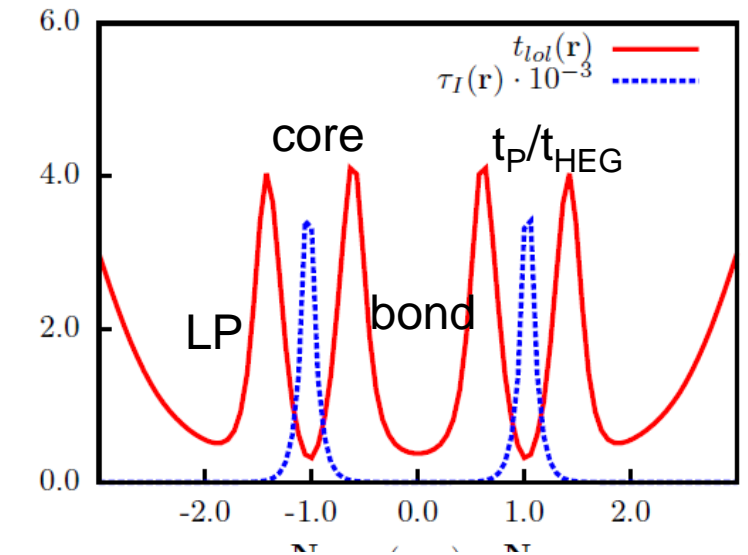
The electron localization function

Fermionic information

- It is divided by TF to get rid of electro density dependence
- Re-scaled to go from 0 to 1

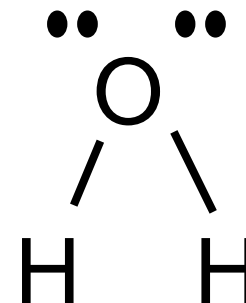
$$t_p(\vec{r}) = t(\vec{r}) - \frac{1}{8} \frac{|\nabla \rho(\vec{r})|^2}{\rho(\vec{r})}$$

$$\chi(\vec{r}) = \frac{t_p(\vec{r})}{c_F \rho(\vec{r})^{5/3}} \quad ELF = \frac{1}{(1 + \chi^2(\vec{r}))}$$



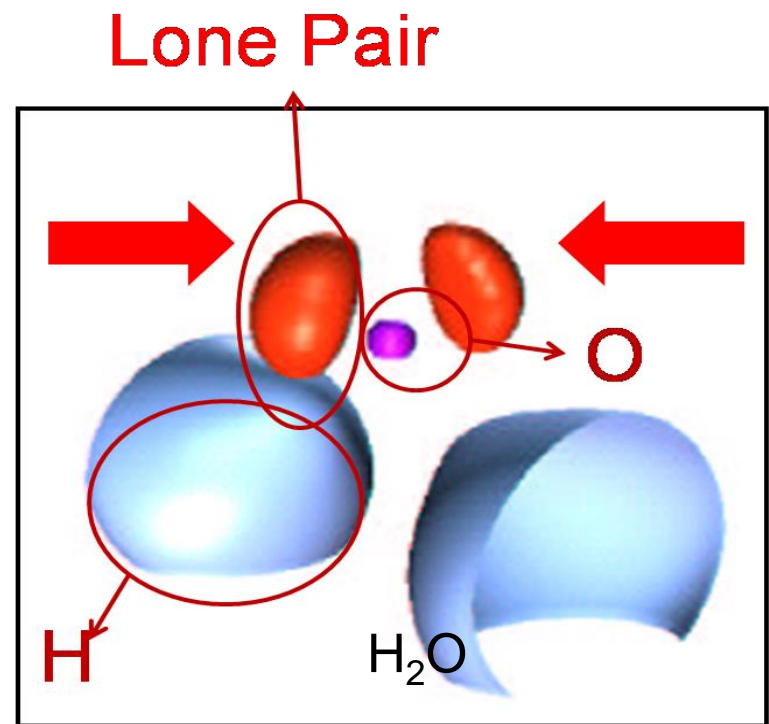
The electron localization function

- It recovers the Lewis picture of a system
- ELF is close to one in:
 - Lone pairs



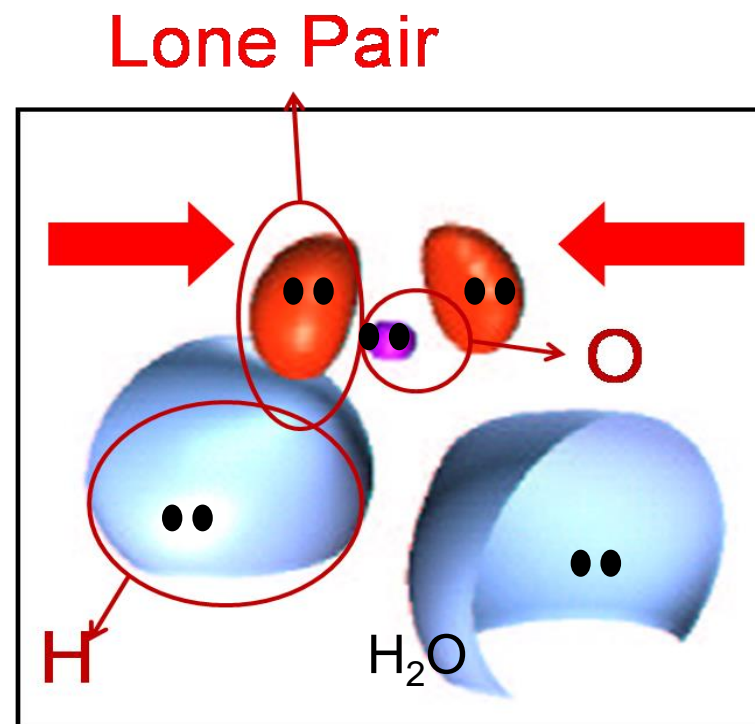
The electron localization function

- It recovers the Lewis picture of a system
- ELF is close to one in:
 - Lone pairs



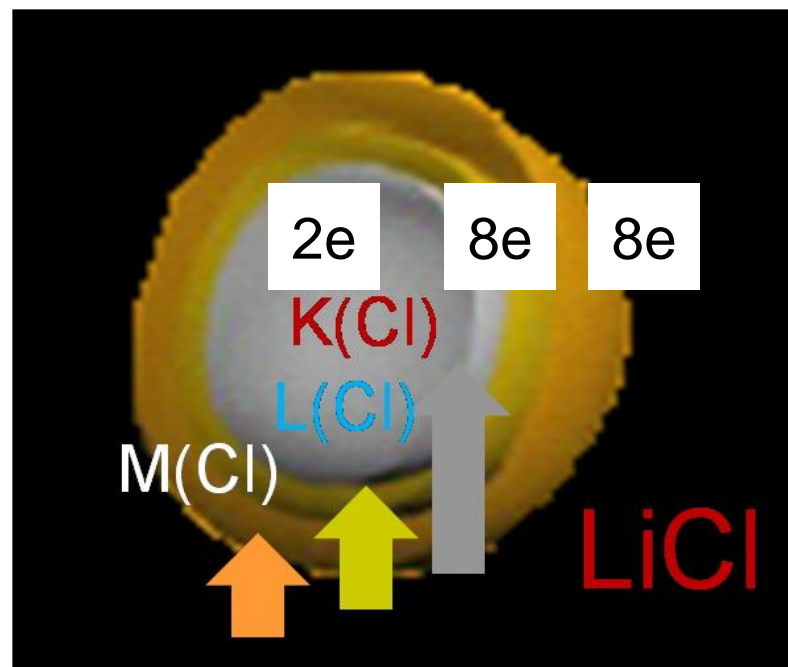
The electron localization function

- It recovers the Lewis picture of a system
- ELF is close to one in:
 - Lone pairs



The electron localization function

- It recovers the Lewis picture of a system
- ELF is close to one in:
 - Lone pairs
 - Atomic shells



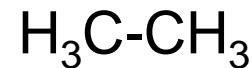
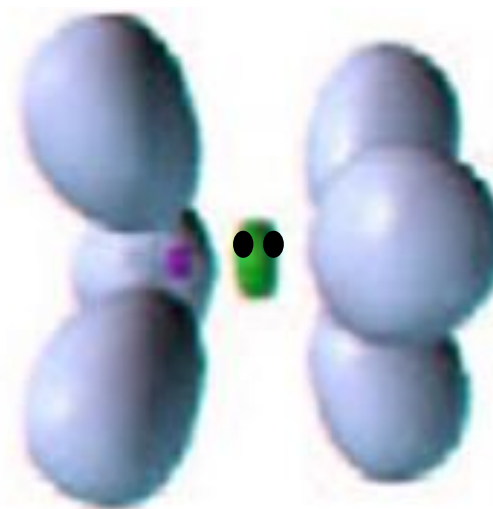
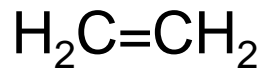
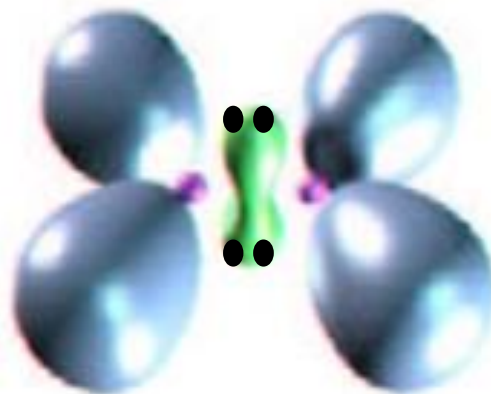
Electron numbers and
charge transfer

The electron localization function

● It recovers the Lewis picture of a system

● ELF is close to one in:

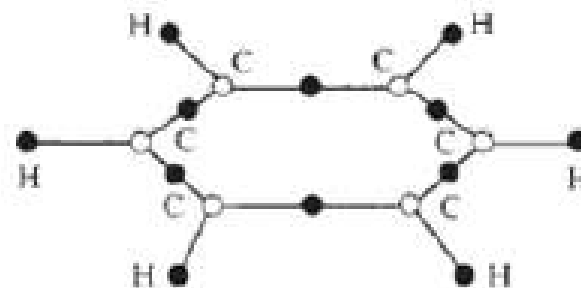
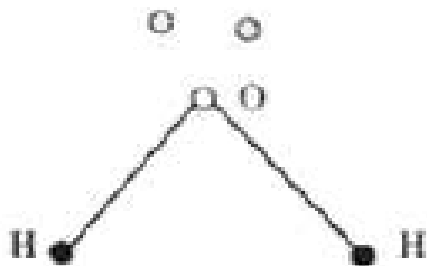
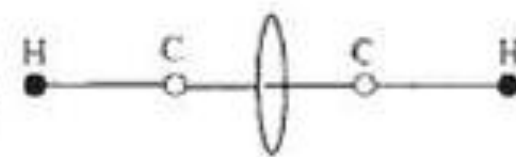
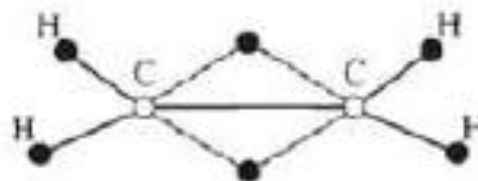
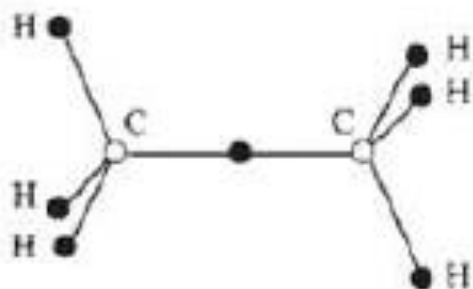
- Lone pairs
- Atomic shells
- Bonds



Bond order

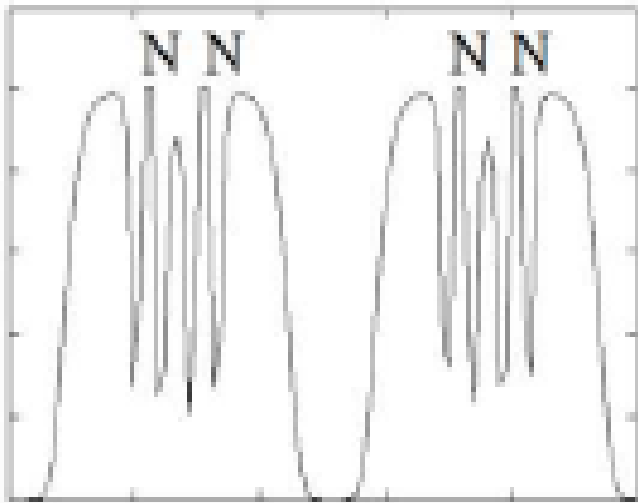
The electron localization function

Critical points

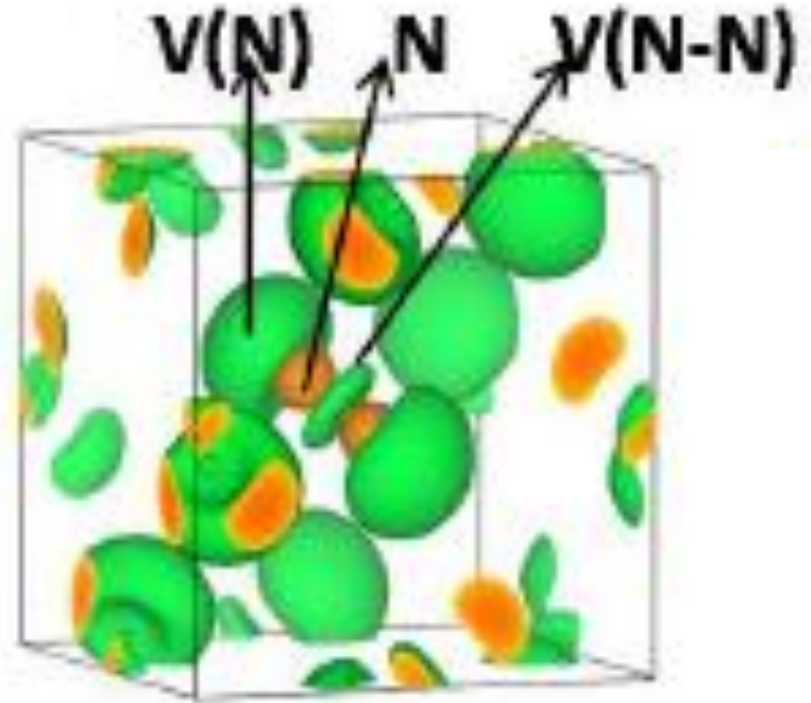


ELF in solids

Molecular solids

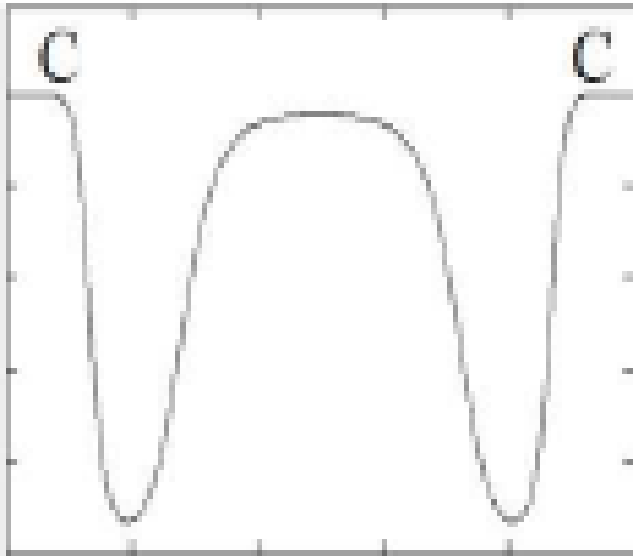


$\epsilon(N_2 \cdots N_2)$

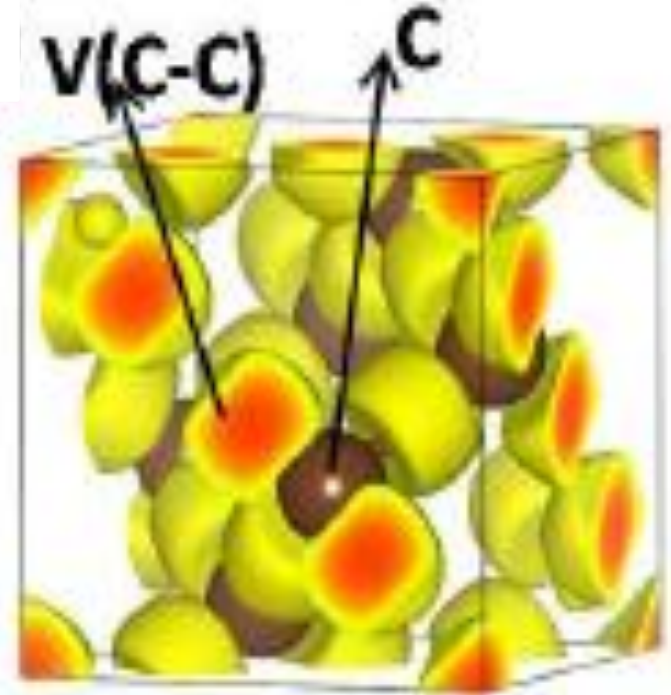


ELF in solids

Covalent solids

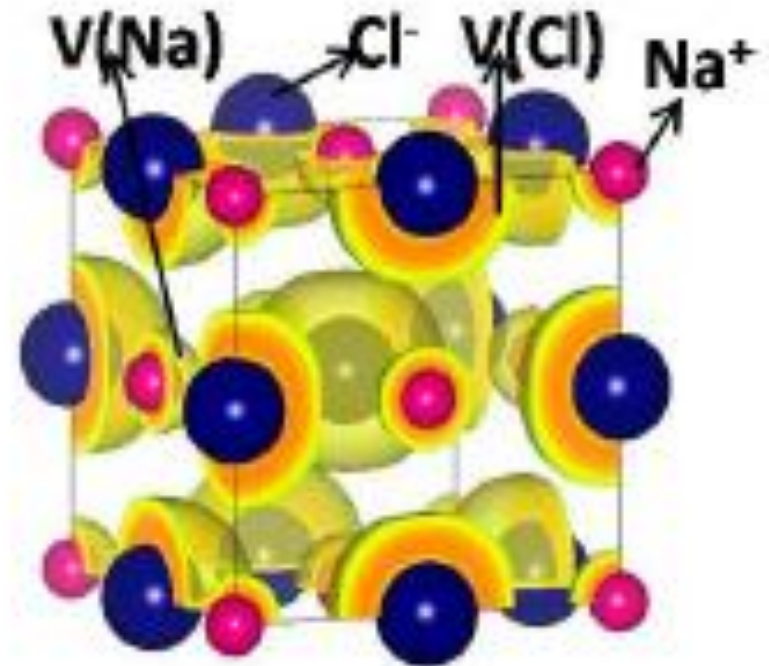
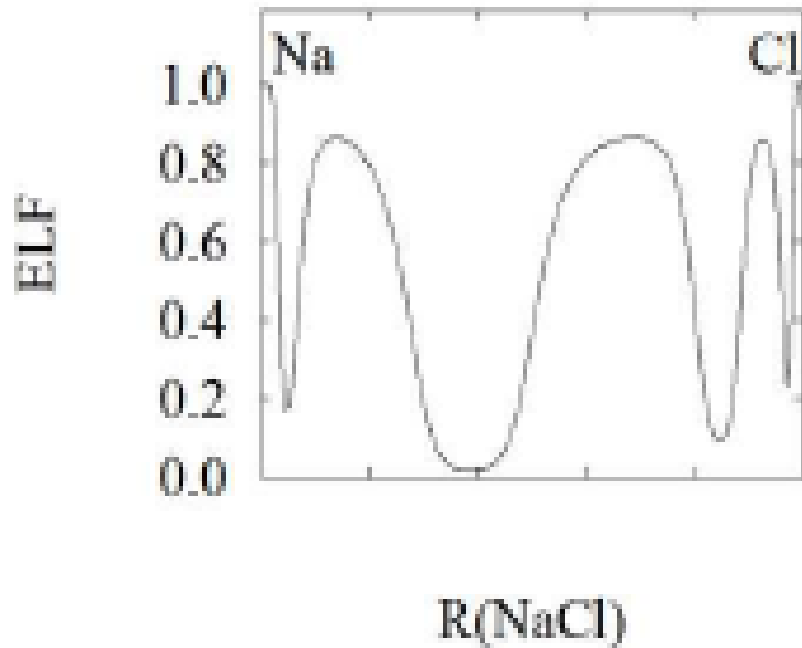


$R(\text{C}\cdots\text{C})$



ELF in solids

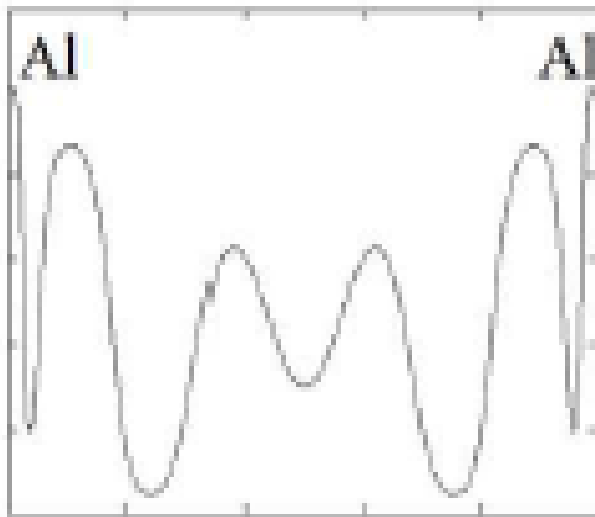
Ionic solids



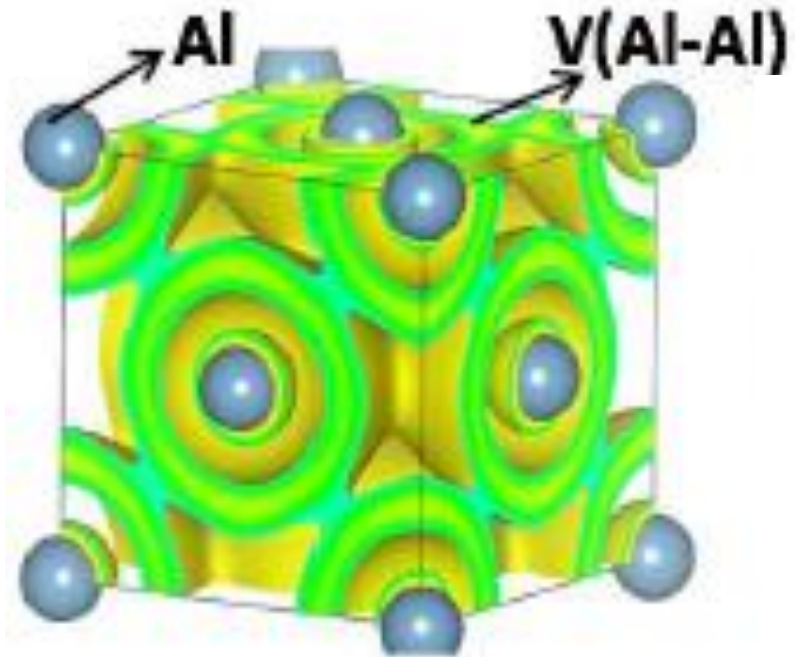
Charge transfer is verified
 $\text{Na}=10 e$
 $\text{Cl}=18 e$

ELF in solids

Metals

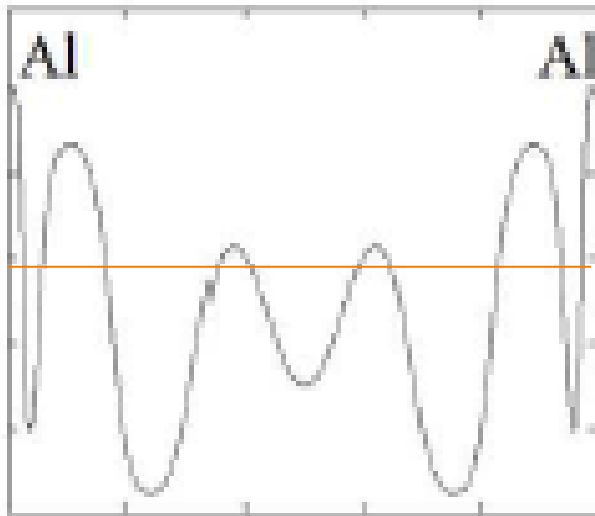


$R(\text{Al})$



ELF in solids

Metals



R(Al)

$$\chi(\vec{r}) = \frac{t_P(\vec{r})}{c_F \rho(\vec{r})^{5/3}} \quad ELF = \frac{1}{(1 + \chi^2(\vec{r}))}$$

$$t_P(\vec{r}) = t_{HEG}(\vec{r}) \quad ELF = 0.5$$

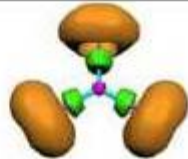
Metallic valence close to HEG

ELF pictures recover VSEPR

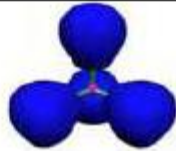
ab₂ BeCl₂



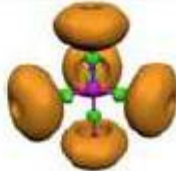
ab₃ BCl₃



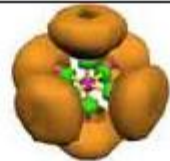
ab₄ CH₄



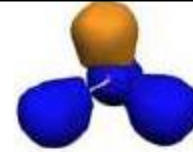
ab₅ PCl₅



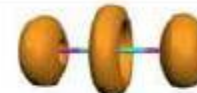
ab₆ SCl₆



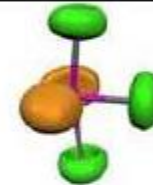
ab₃e NH₃



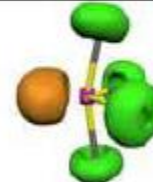
ab₂e₃ XeCl₂



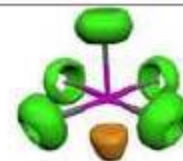
ab₃e₂ ClF₃



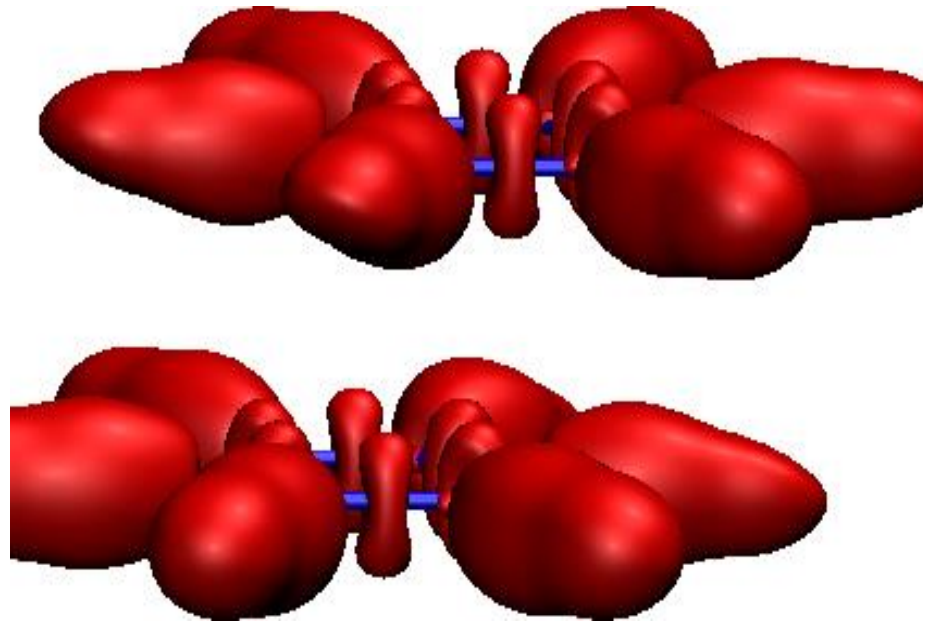
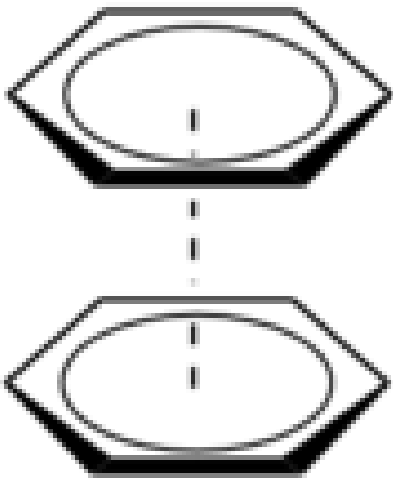
ab₄e SF₄



ab₅e BrF₅



Still missing something...



Outline

1. Why studying chemical bonds?

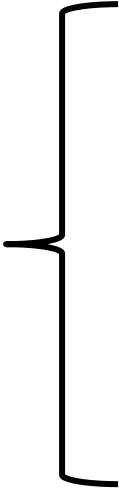
2. Quantum Chemical topology

3. Chemical functions

a) electron density

b) ELF

c) NCI

- 
1. The function
 2. The topology
 3. Old insight
 4. New insight

4. Applications

Critical point : $\nabla\rho = 0$

Let's look at the region $\nabla\rho \rightarrow 0$

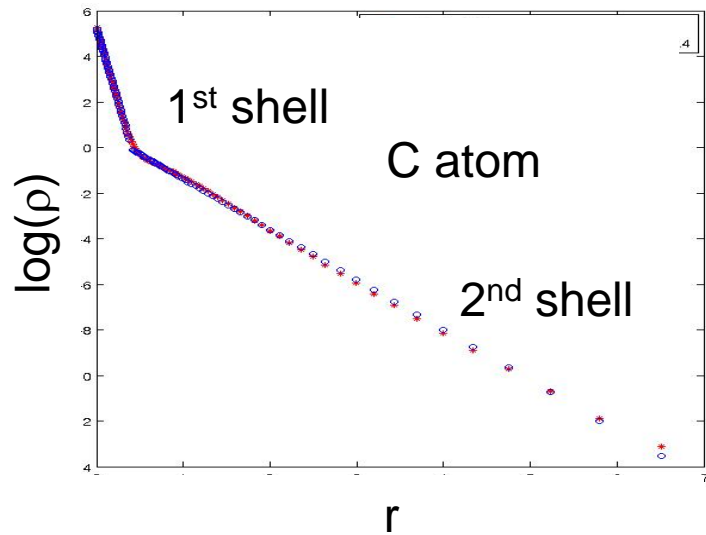
NCI:

analysis of the reduced density gradient at low densities

$$s = \frac{1}{c_s} \frac{|\nabla\rho|}{\rho^{4/3}} \quad \text{with } c_s = 2(3\pi^2)^{1/3}$$

- How does it work?

Model densities



Atoms

- Atomic densities can be mimicked like a sum of N_{shells} exponentials

$$\rho^{at}(r) = \sum_i^{N_{shells}} c_i e^{-\zeta_i r}$$

- 1st shell can be approximated by a 1s orbital

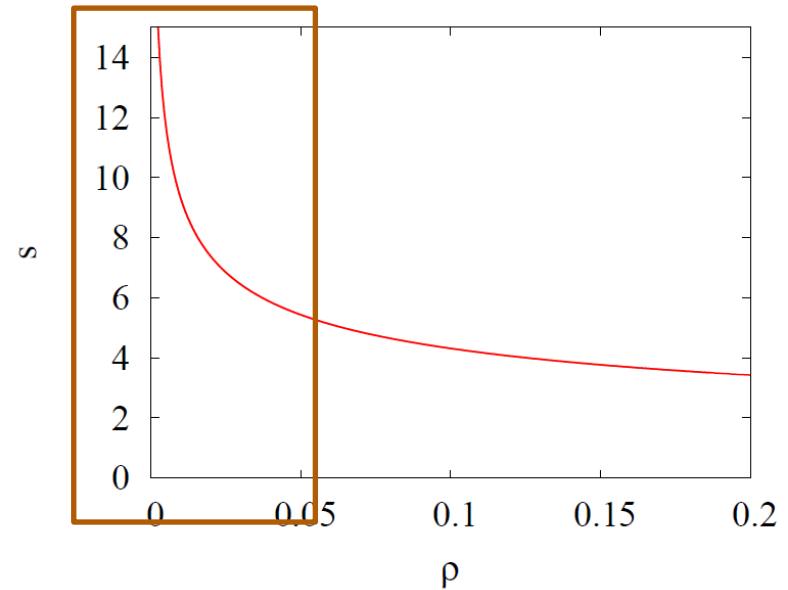
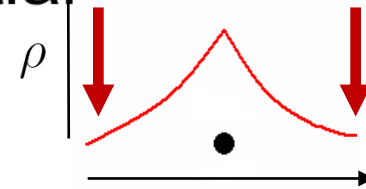
$$\rho_{1s} = \frac{n_{1s}}{\pi} \left(\frac{Z}{a_0} \right)^3 e^{-2Zr/a_0}$$

Atoms

At ∞ Taking the outermost exponential

- $\rho(r) = ce^{-\zeta r}$
- $|\nabla\rho(r)| = c\zeta e^{-\zeta r}$

- $$s = \frac{1}{c_s} \frac{|\nabla\rho|}{\rho^{4/3}} = \frac{1}{c_s} \frac{\zeta}{(ce^{-\zeta r})^{1/3}} = \frac{\zeta}{c_s} \rho^{-1/3}$$



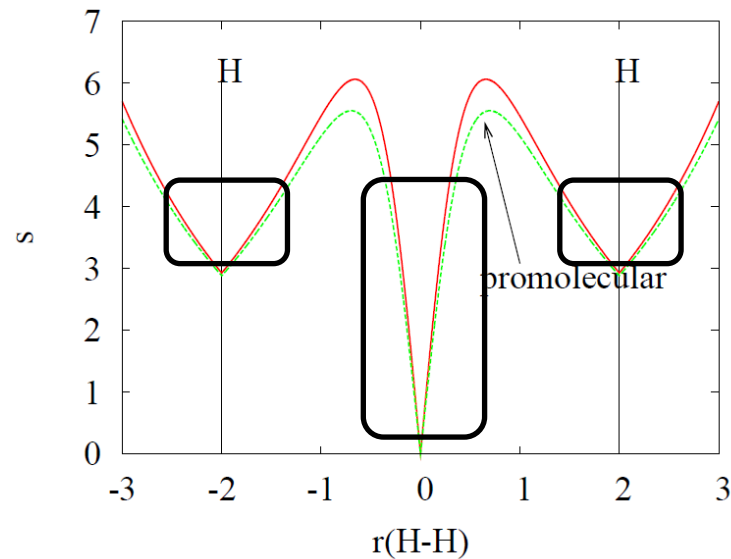
- $\lim_{r \rightarrow \infty} s(r) = \infty$ (i.e. when $\rho \rightarrow 0$)

Model densities

Molecules

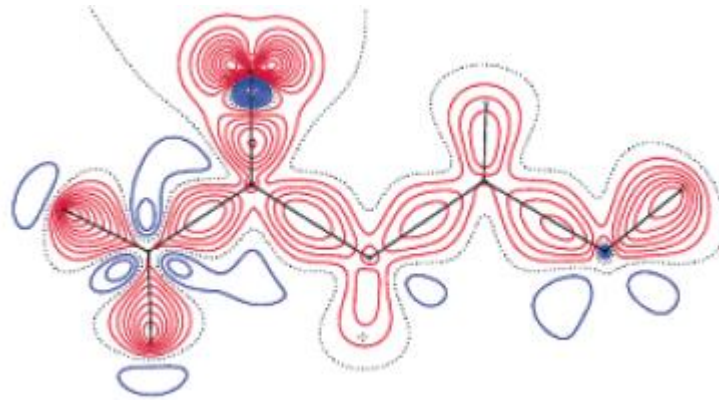
- Promolecular densities: no interference (SCF cycle 0)

$$\rho_{molec}(r) = \sum_j^{N_{atoms}} \rho_j^{at}(r) = \sum_j^{N_{atoms}} \sum_i^{N_{shells}} c_{i,j} e^{-\zeta_{i,j} r}$$



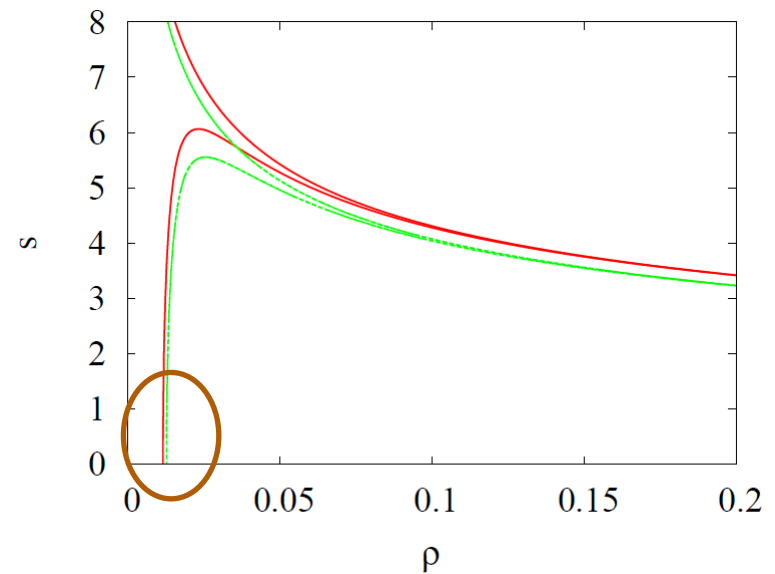
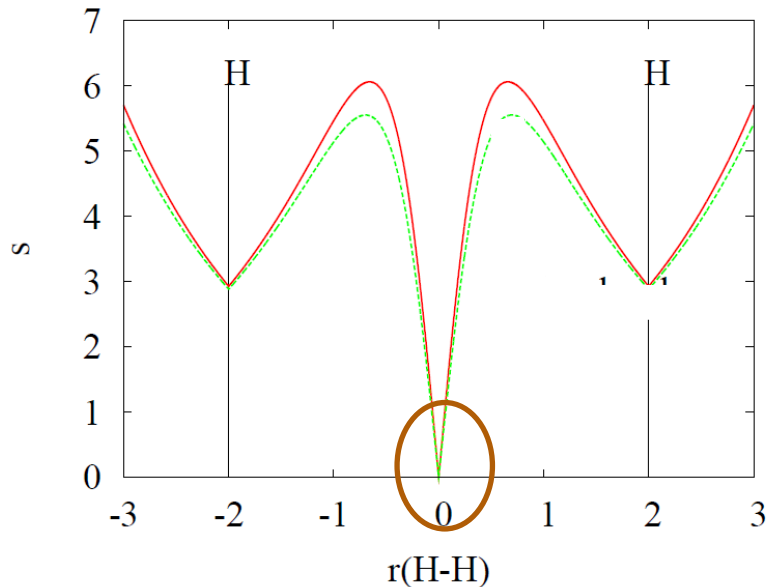
- Works specially well at low densities ($s \rightarrow 0$) at atomic positions (1s orbital)

Promolecular densities



Crystallography: promolecular densities are used in refinement of high-resolution X-Ray data of large systems

Interactions



- Promolecular approach

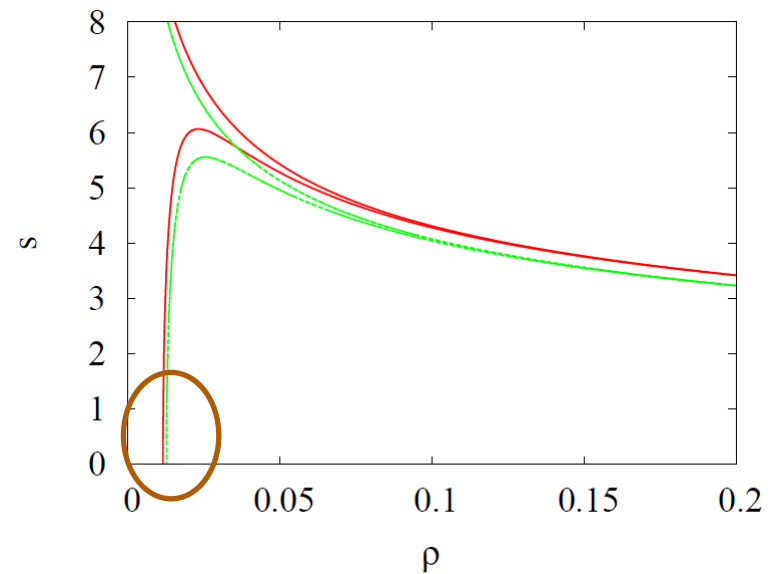
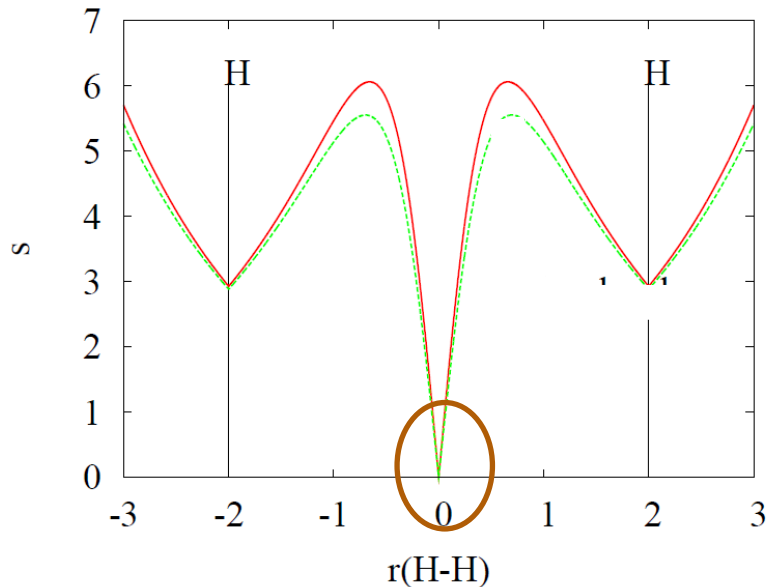
$$\rho(x, y, z) = \rho_{1s}^A + \rho_{1s}^B = \left(c e^{-\zeta \sqrt{x^2 + y^2 + (z - R/2)^2}} \right)^2 + \left(c e^{-\zeta \sqrt{x^2 + y^2 + (z + R/2)^2}} \right)^2$$

- s goes to 0 in the interactions

$$\lim_{z \rightarrow 0} s(\rho) = \lim_{z \rightarrow 0} (e^{R/3} r + O(r^3)) = 0$$

- When we plot $s(\rho)$ at low densities, the behavior is very characteristic

Interactions



- Promolecular approach

$$\rho(x, y, z) = \rho_{1s}^A + \rho_{1s}^B = \left(c e^{-\zeta \sqrt{x^2 + y^2 + (z-R/2)^2}} \right)^2 + \left(c e^{-\zeta \sqrt{x^2 + y^2 + (z+R/2)^2}} \right)^2$$

- s goes to 0 in the interactions

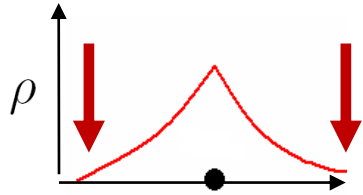
$$\lim_{z \rightarrow 0} s(\rho) = \lim_{z \rightarrow 0} (e^{R/3} r + O(r^3)) = 0$$

- When we plot $s(\rho)$ at low densities, the behavior is very characteristic

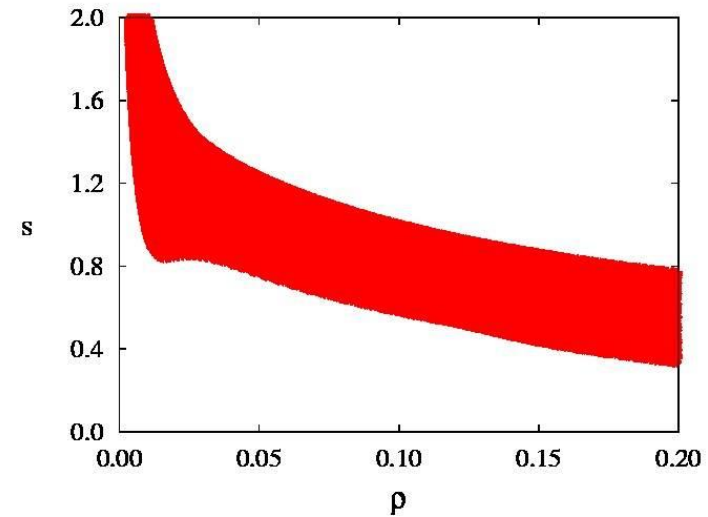
Let's see an example!

The reduced density gradient

Non-interacting densities

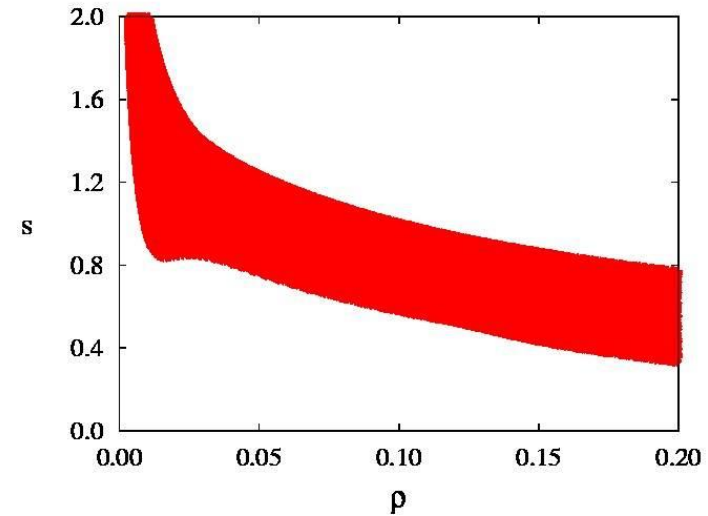
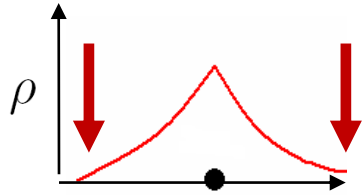


$$s(\rho) \propto \rho^{-1/3} \xrightarrow{\rho \rightarrow 0} \infty$$

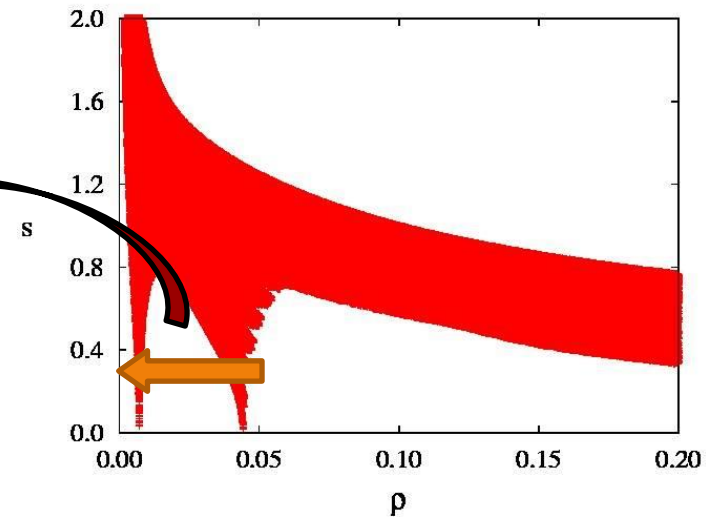
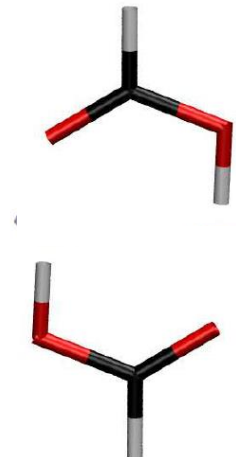
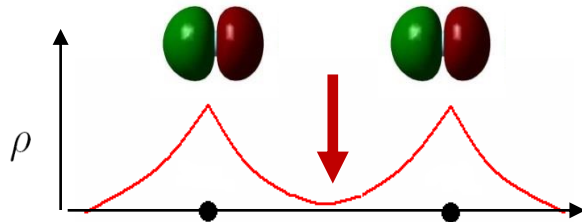


The reduced density gradient

Non-interacting densities



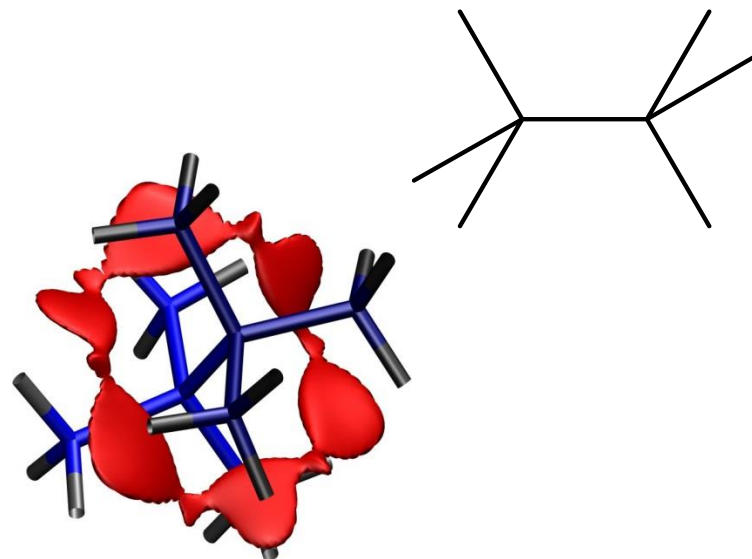
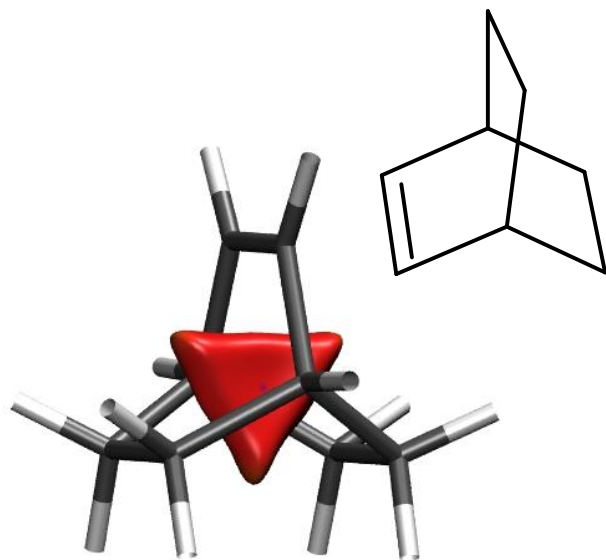
Interacting densities



The reduced density gradient

Repulsive interactions

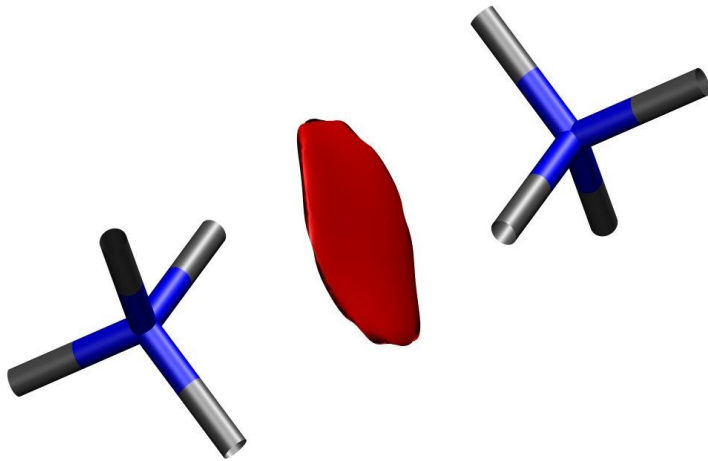
- Steric clashes



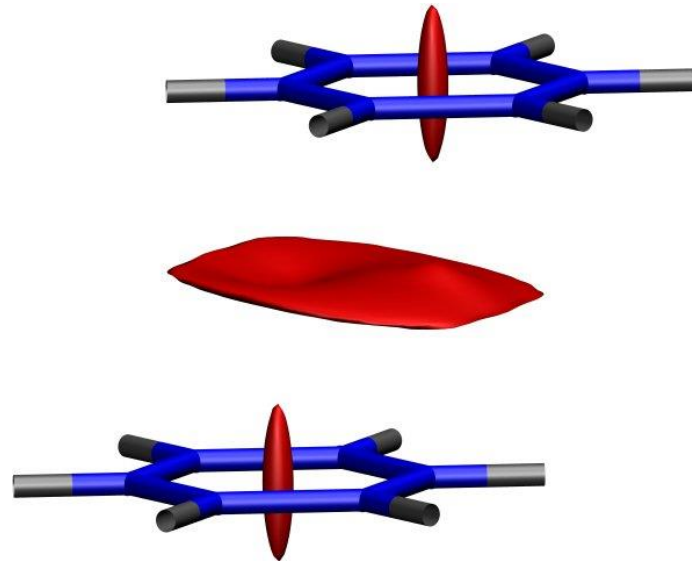
The reduced density gradient

Weak interactions

- Van der Waals



Methane dimer

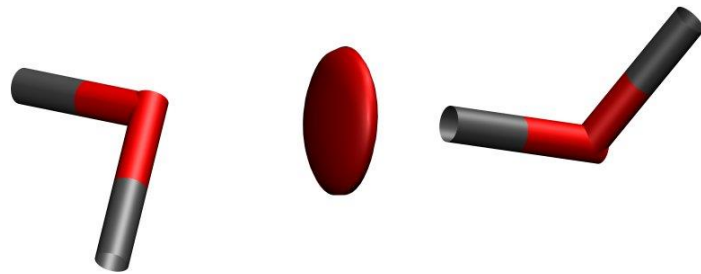


Benzene dimer

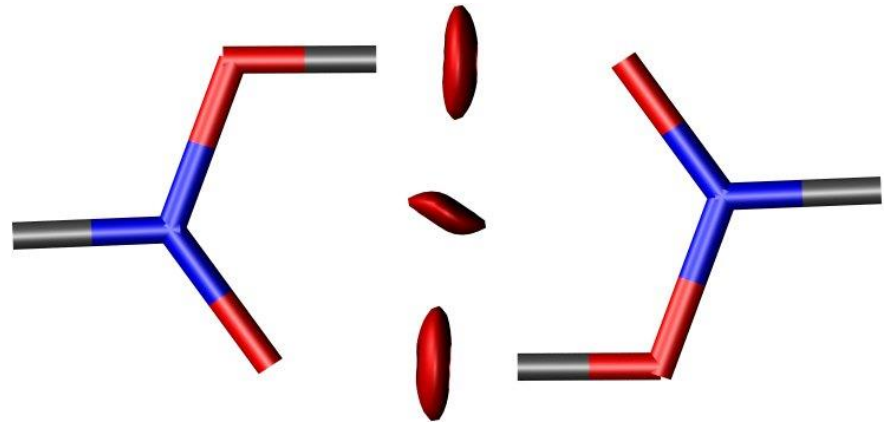
The reduced density gradient

Strongly attractive interactions

- Hydrogen bonds



Water dimer



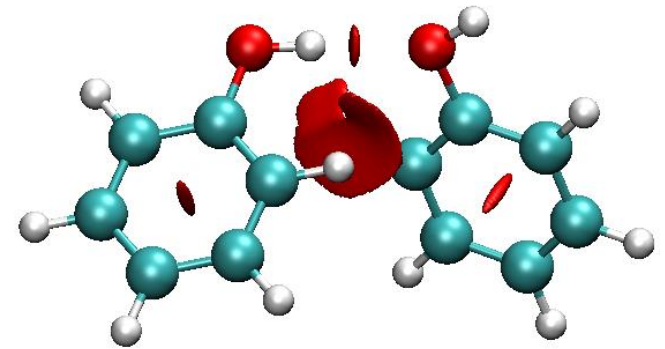
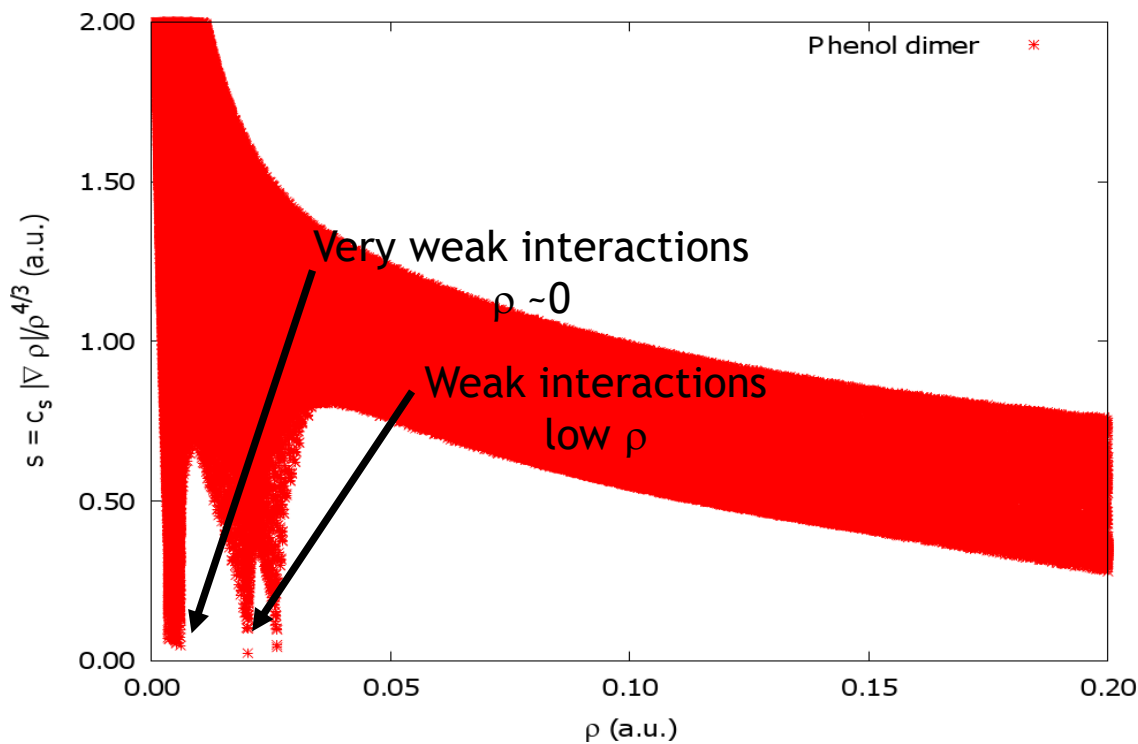
Formic acid dimer

The reduced density gradient

Differentiating interaction types

1

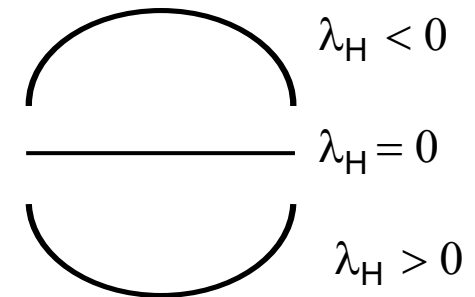
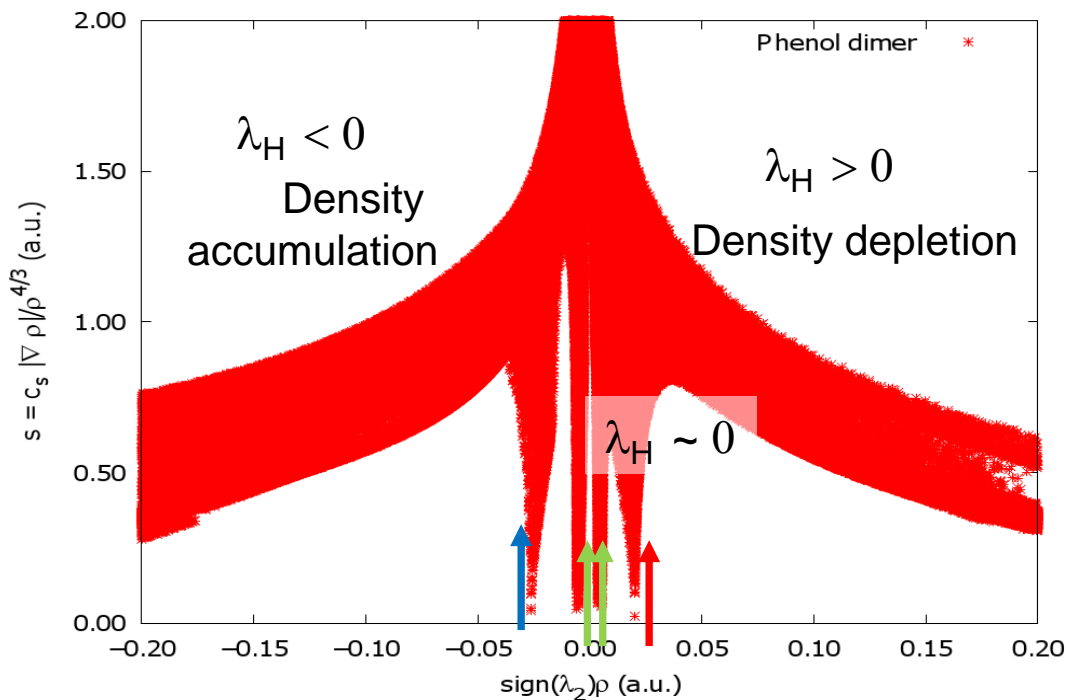
- Density is proportional to the strength of the interaction



The reduced density gradient

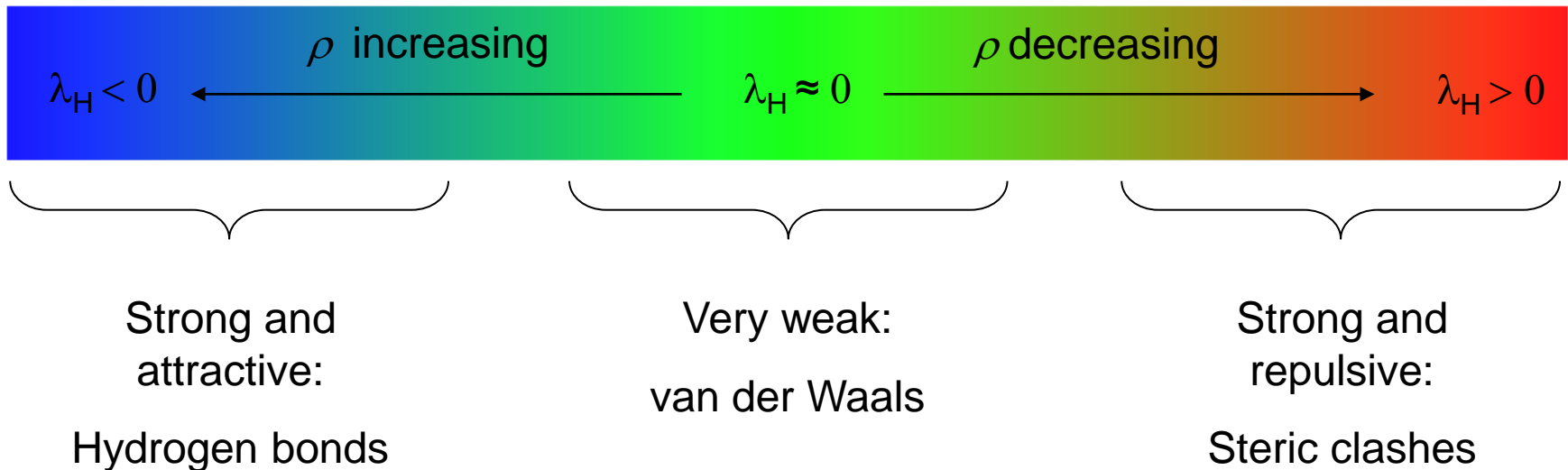
Differentiating interaction types

- 2 {
- Bonding interactions \rightarrow charge accumulation ($\lambda_H < 0$)
 - Antibonding interactions \rightarrow charge depletion ($\lambda_H > 0$)

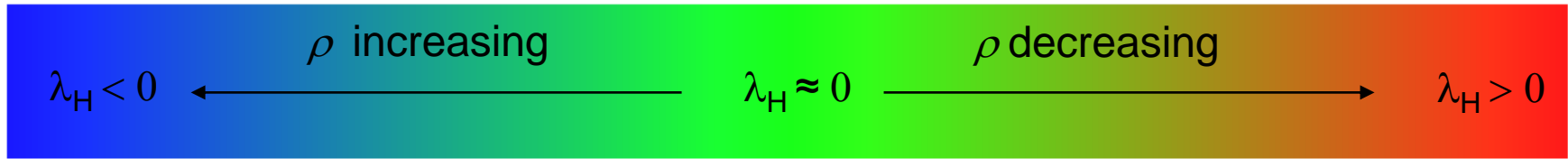
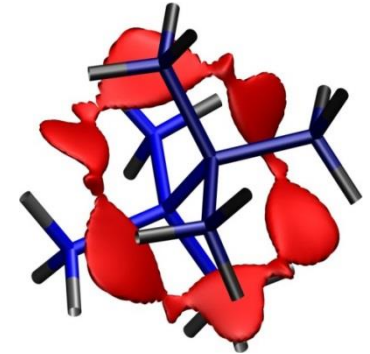
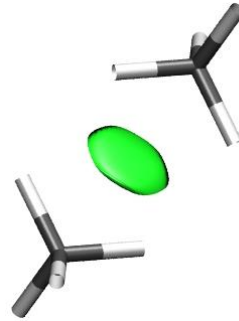
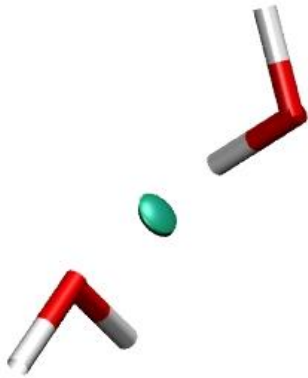


The reduced density gradient

- 1) Represent s isosurfaces in real space
- 2) Colour them in terms of $\text{sign}(\lambda_H) \times \rho$



The reduced density gradient

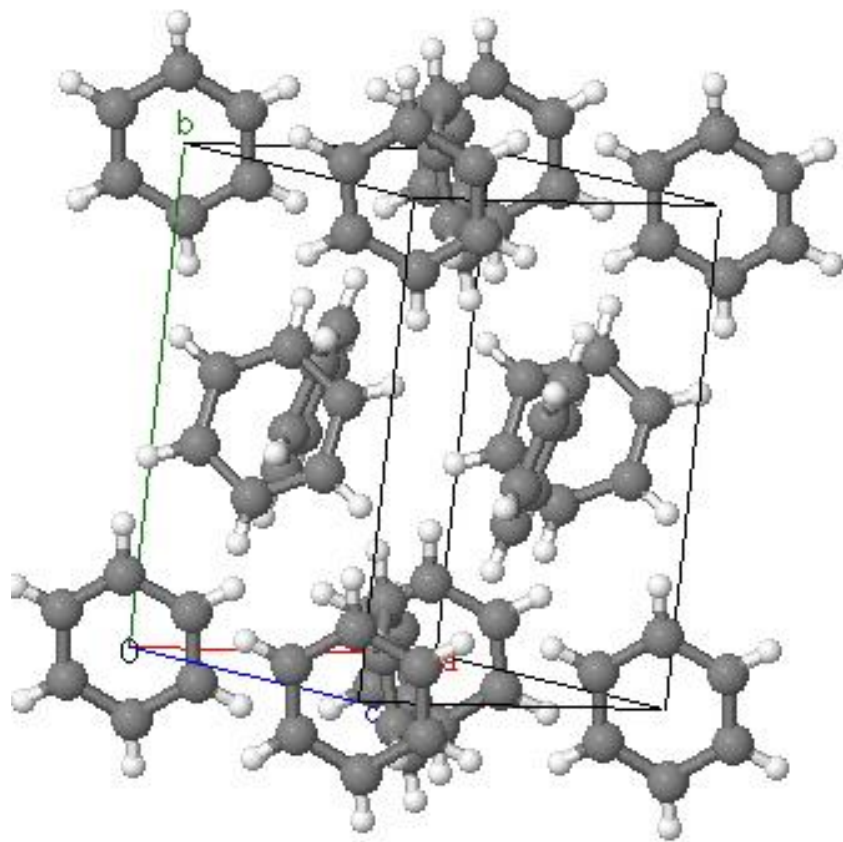


Strong and attractive:
Hydrogen bonds

Very weak:
van der Waals

Strong and repulsive:
Steric clashes

Delocalized interactions



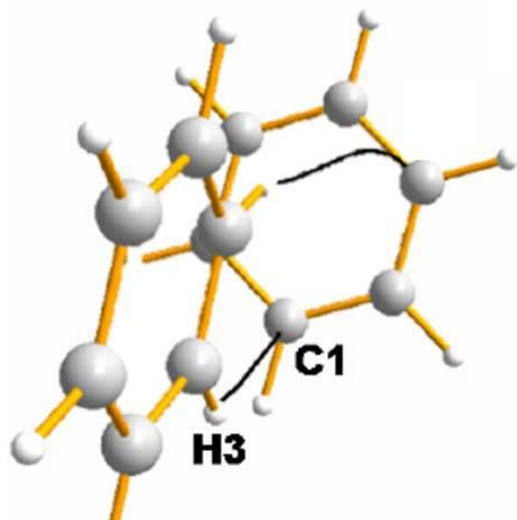
Benzene packing maximizes the number of **C-H... π** and **C-H...C** contacts.

Delocalized interactions

AIM

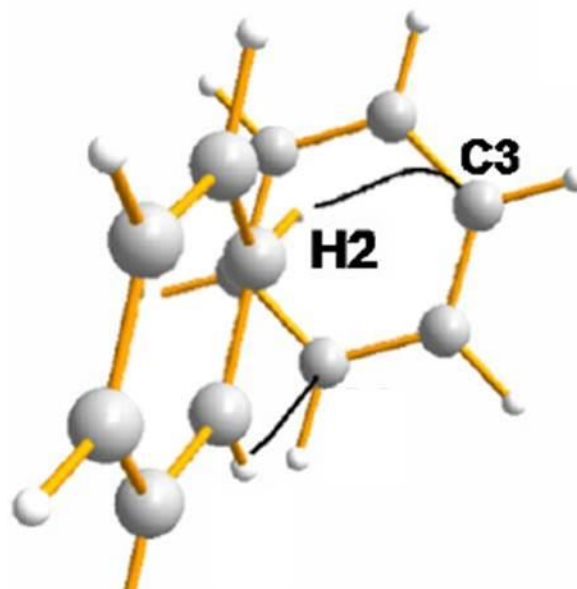
CH-C

Straight bond path



CH- π

H roughly equidistant to the whole ring
Bond path significantly bent

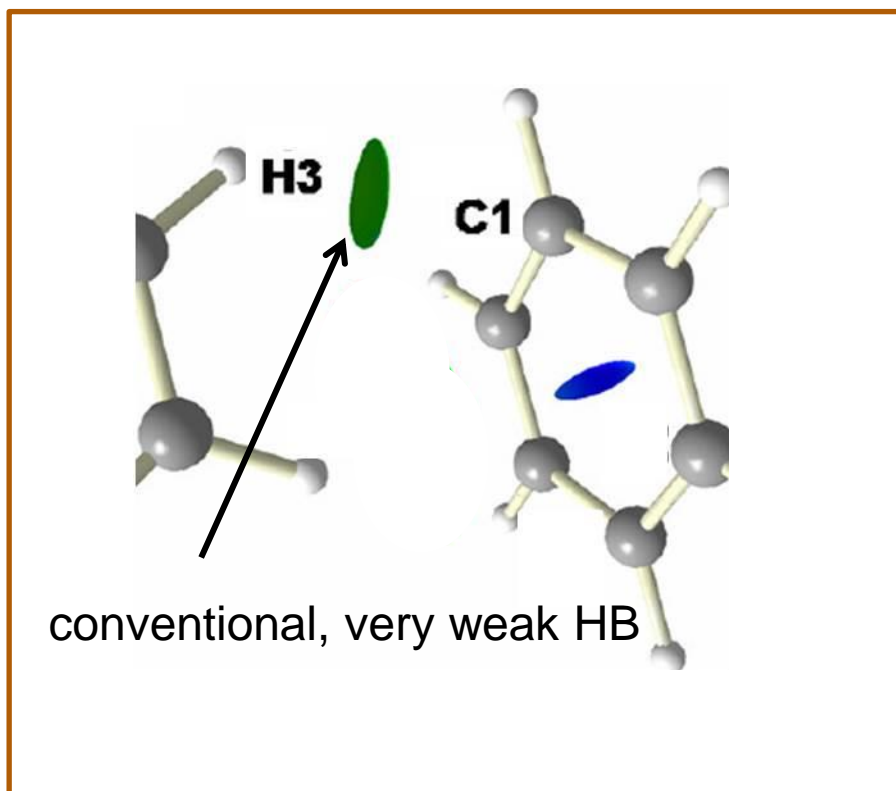


Delocalized interactions

NCI

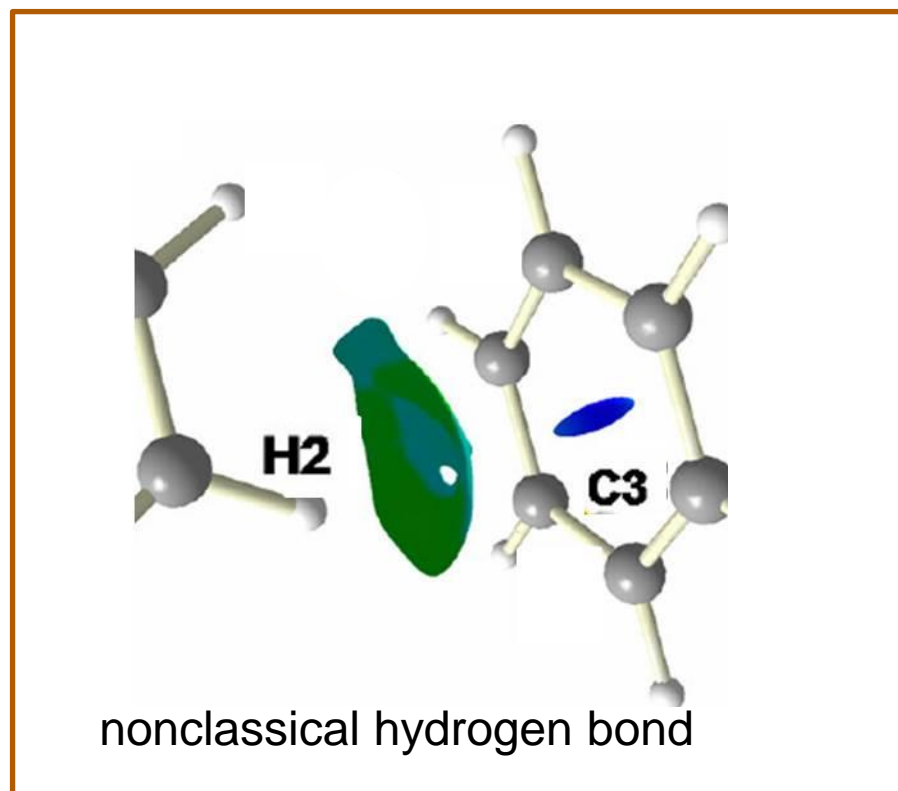
CH-C

Disc-shaped and localized



CH- π

Large isosurface involving the whole π electron cloud



Big systems

- One of the major areas of application of weak interactions are biomolecules
- HOWEVER, wavefunctions are not available
- In the non covalent region, there is very little density reconstruction, so we can use an independent atom model (non relaxed or promolecular densities)

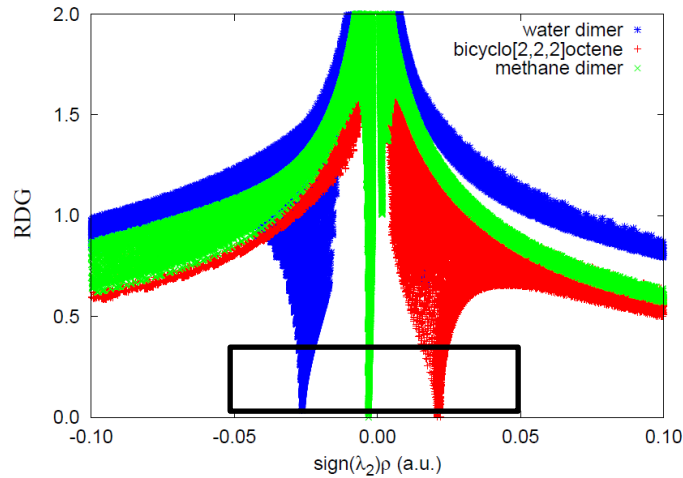
Big systems

How can we apply NCI to such big systems ?

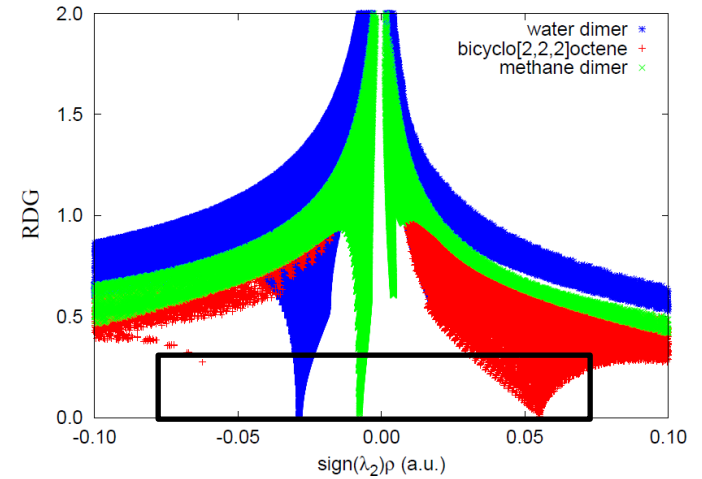
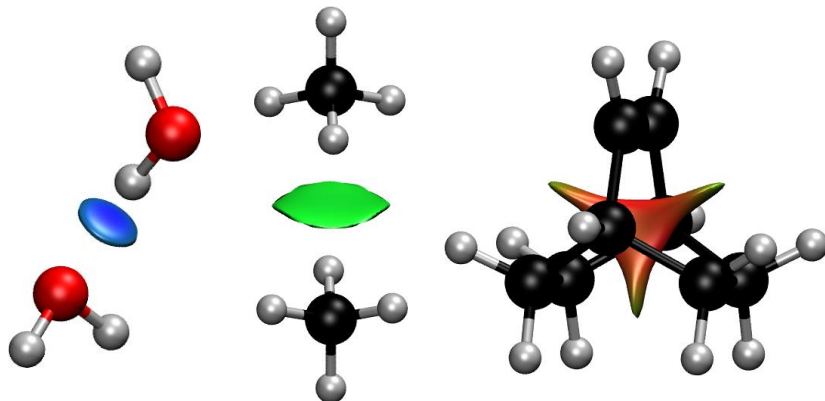
- Since promolecular works well for small densities, we use the promolecular approximation
- Parameters ζ_i and c_i for each atom are internally stored in the code
- No need for wavefunction
- Sum on a grid (very fast)

Big systems

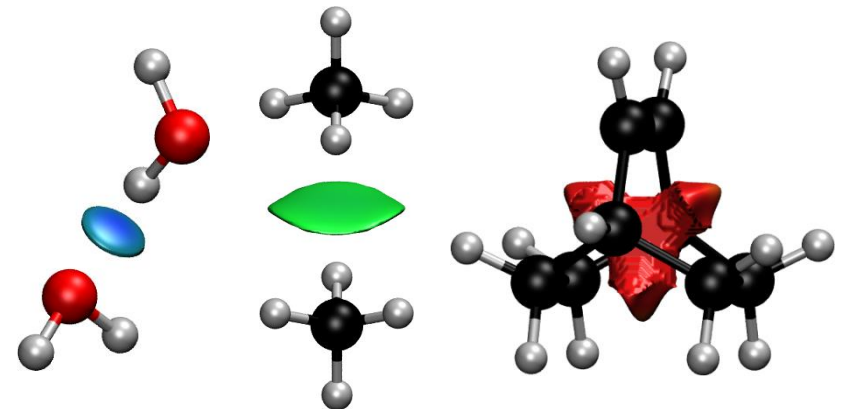
Are characteristics really preserved?



SCF

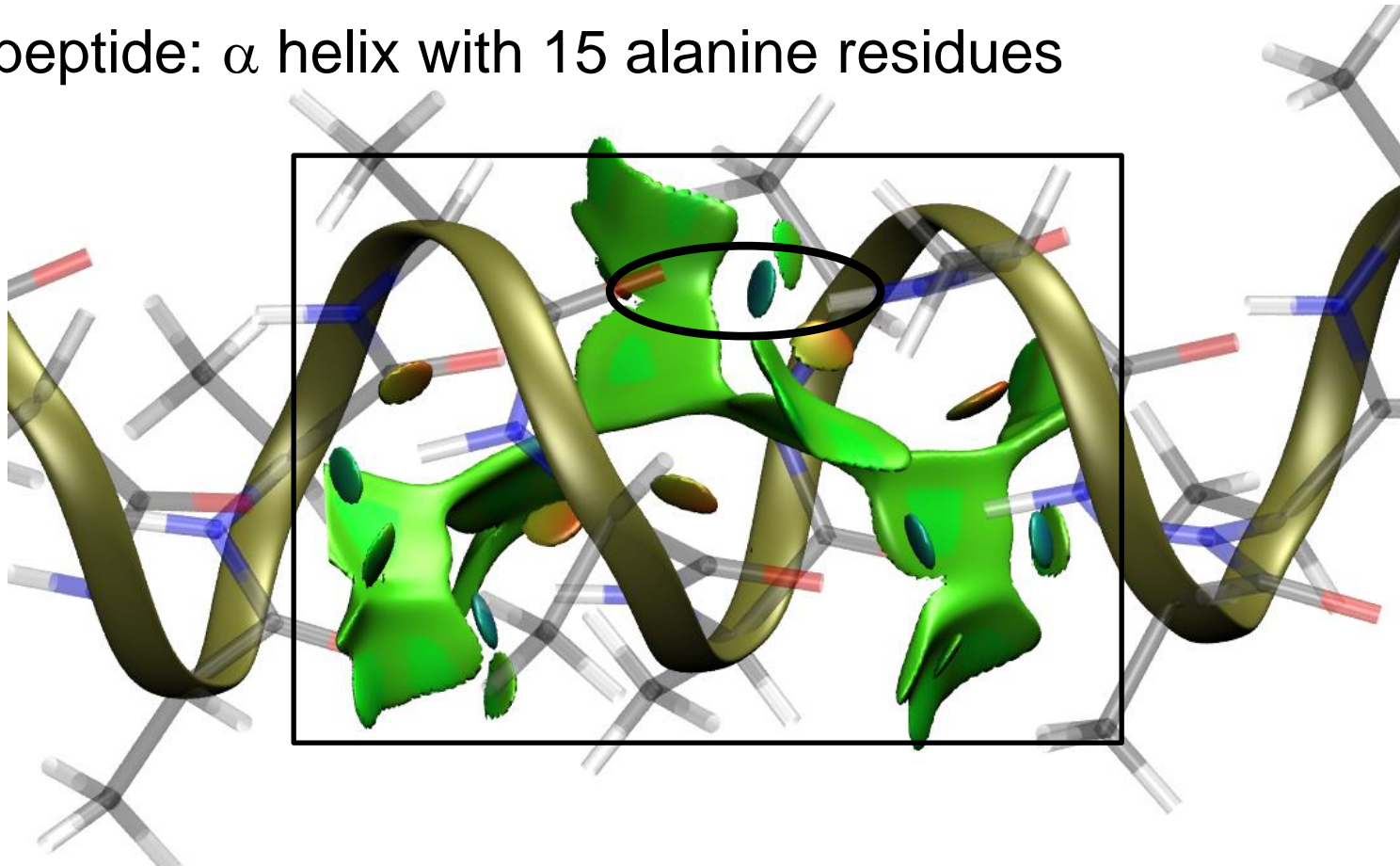


PROMOLECULAR



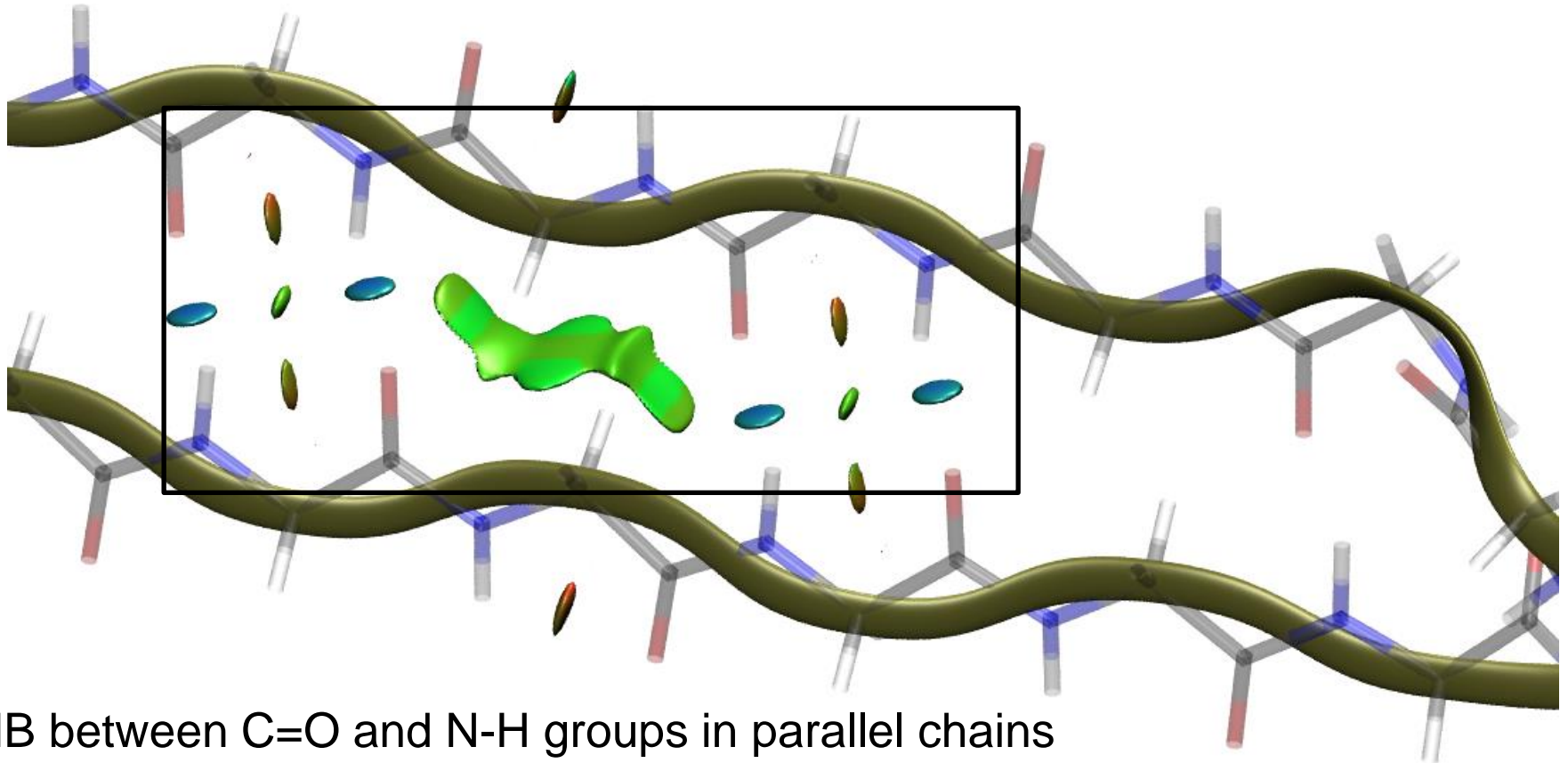
Proteins

Polipeptide: α helix with 15 alanine residues



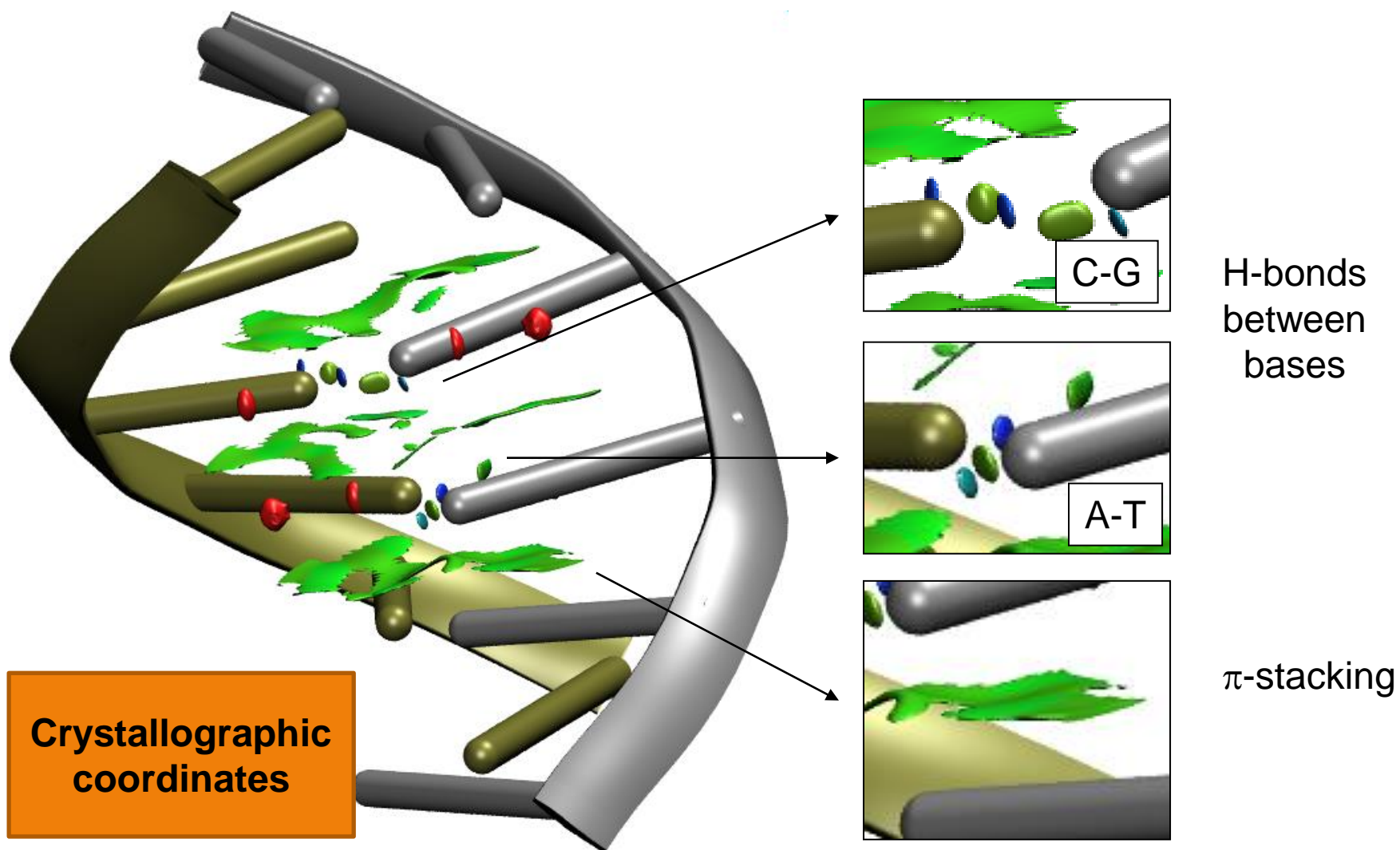
- Hydrogen bonds stabilize the helix
- Big region of van der Waals interaction inside the helix and between methyle lateral chains one step away

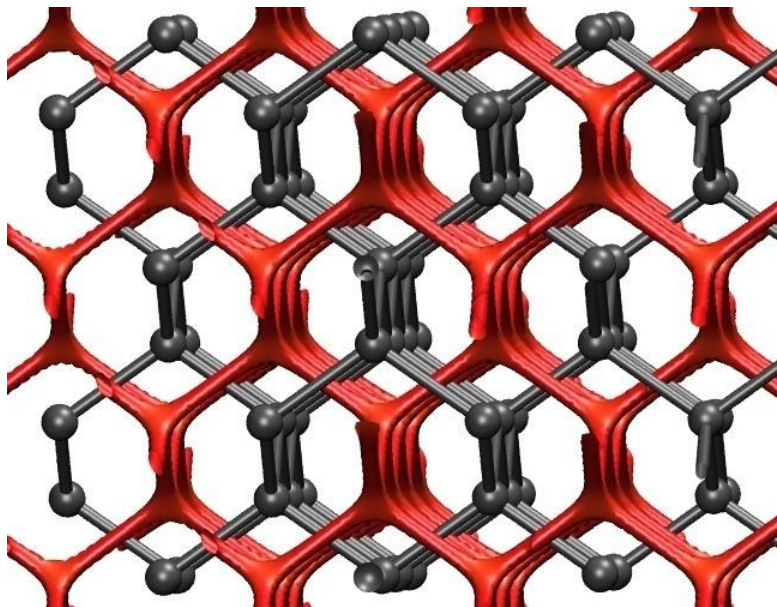
Proteins



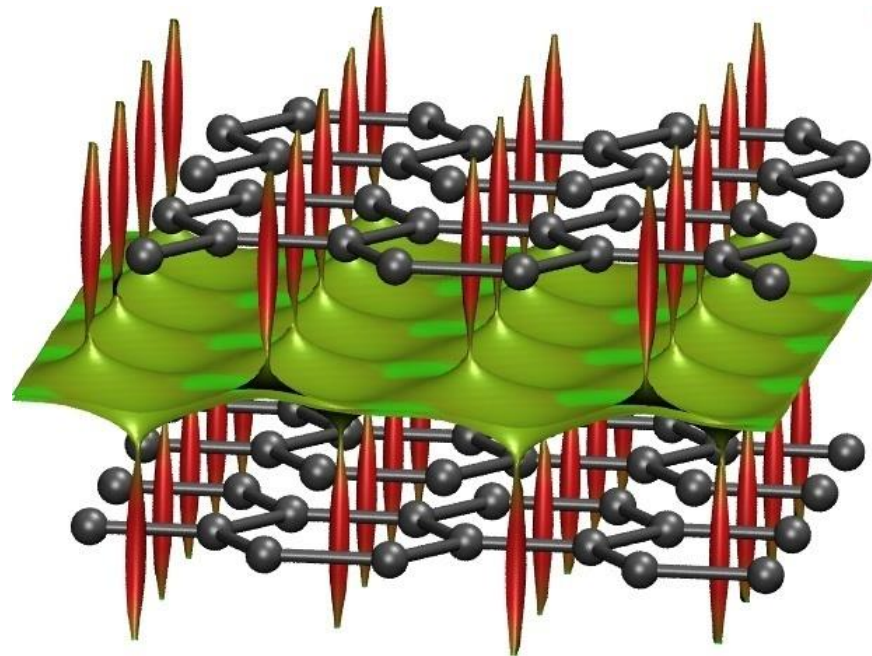
- HB between C=O and N-H groups in parallel chains
- Van der Waals interactions between CH₂ groups

DNA



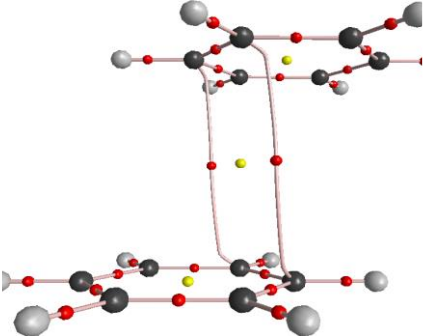
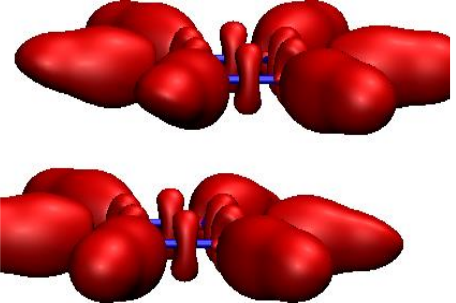
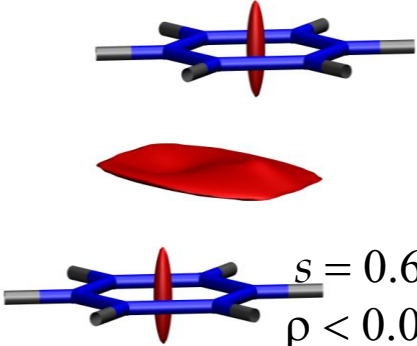


Diamond



Graphite

Summary

What are you looking for	Atoms and bonds Atomic charges	Bonds, lone pairs Charges and volumes in Lewis entities	Non covalent interactions (even in big systems)
You should use...	Electron density	ELF	NCI
Example		 <p>ELF = 0.9</p>	 <p>$s = 0.6,$ $\rho < 0.05$</p>

Summary

- It is important to choose the correct tool
- ...or even to mix them all: for example, in chemical transformations, where we usually go from non covalent to covalent interactions

Outline

5. Applications to high pressure

a) ELF: high pressure metals

b) NCI: He bonds

6. The codes

7. Summary



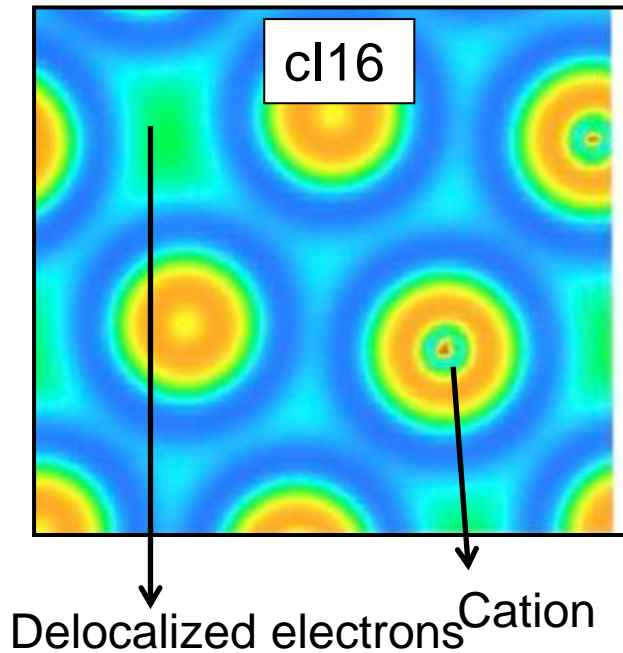
Metals under pressure

New materials : Electronic structure of high pressure metals

- Under pressure, solids exhibit increasingly shorter interatomic distances. Intuitively, this response is expected to be accompanied by an increase in the widths of the valence and conduction bands and hence a more pronounced free-electron-like behavior.
- However, recent experiments have shown a pressure-induced transformation of Na into an optically transparent and insulating phase at 200 GPa (5.0-fold compression)
- What is the electronic structure behind this new state of matter?

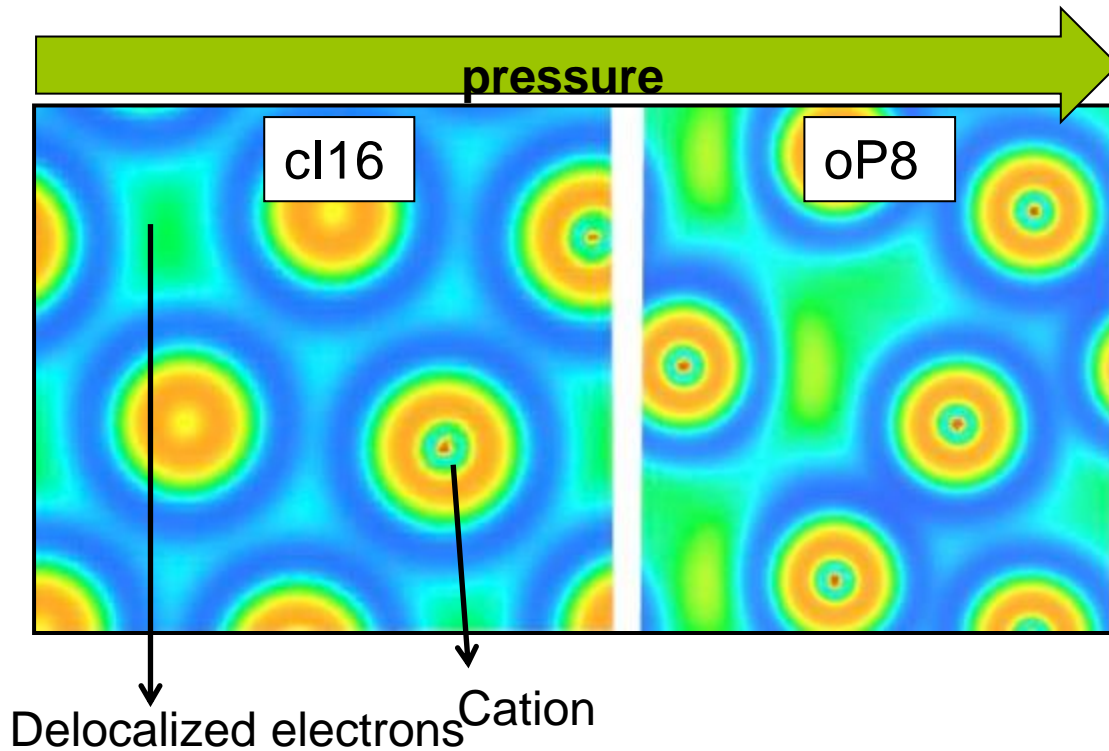
New bonding patterns: new properties

Na



New bonding patterns: new properties

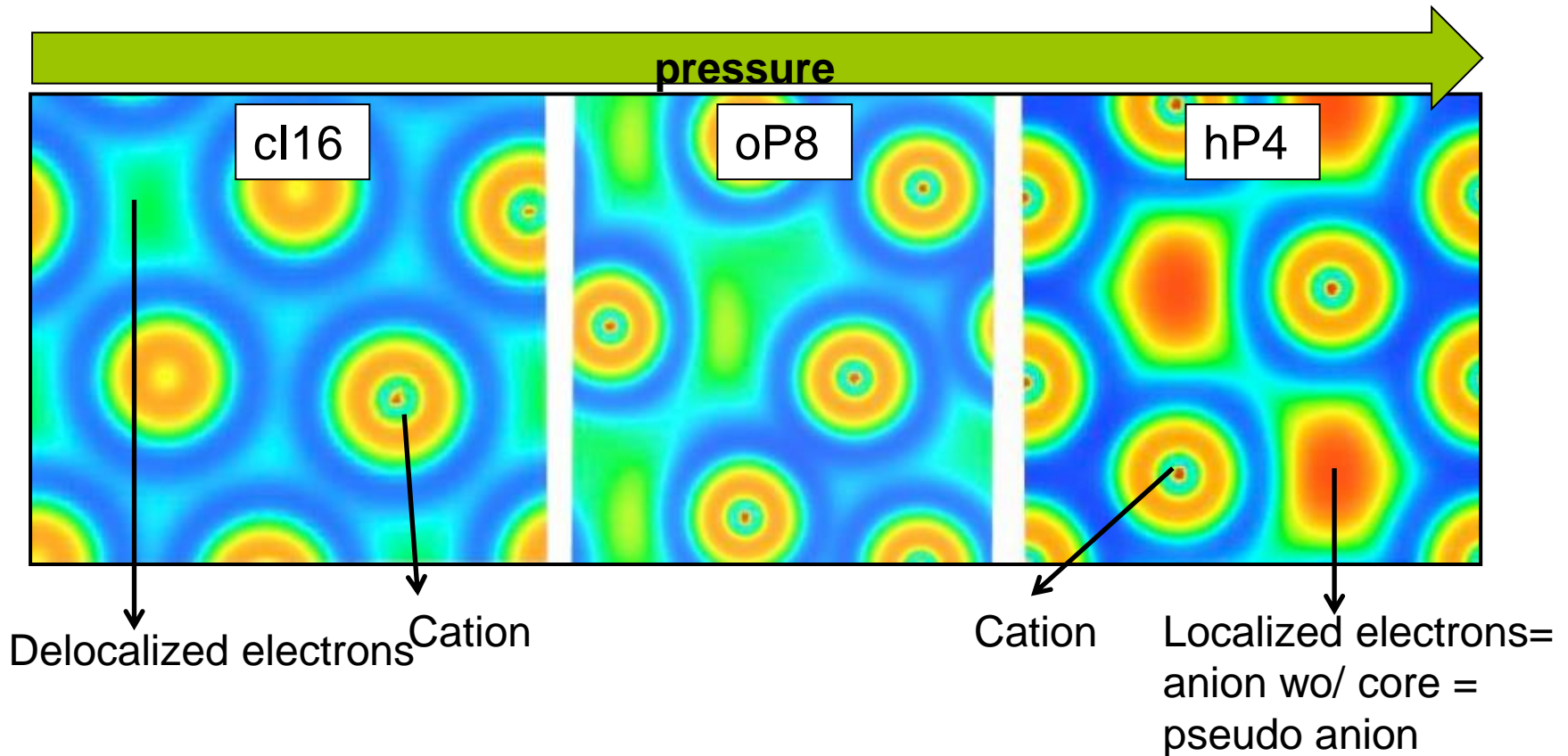
Na



New bonding patterns: new properties

Na

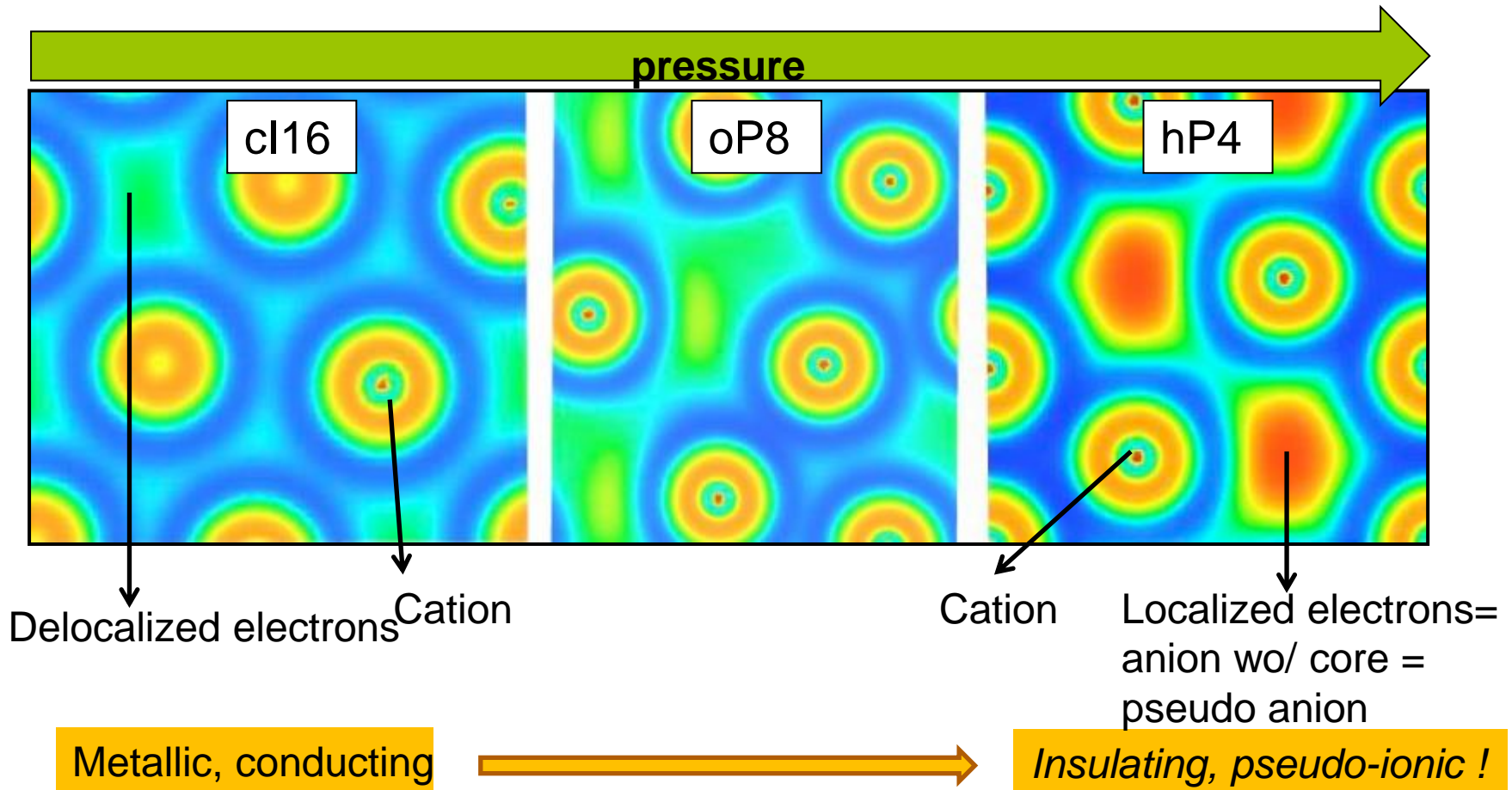
Localization of valence electrons is again observed under pressure



New bonding patterns: new properties

Na

Localization of valence electrons is again observed under pressure



New bonding patterns: new properties

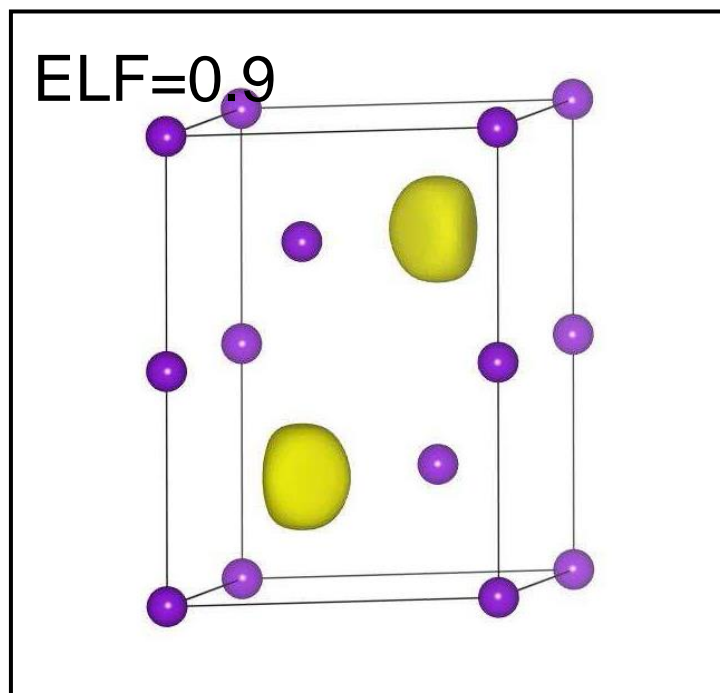
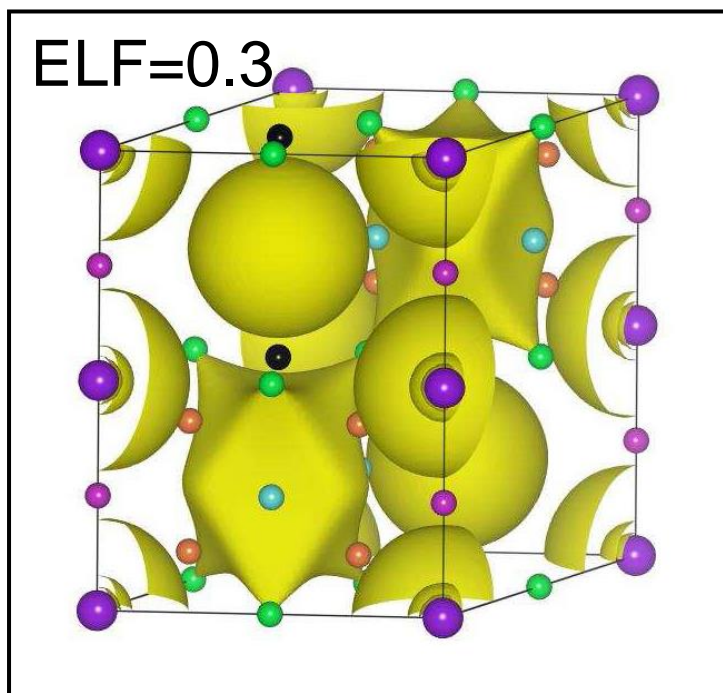
K

Same structure as A_2X and AX_2 compounds!
Electrons in same position as anions!

Equivalency external and chemical pressure

COMPOUND → IONIC-LIKE

ELECTRONS → PSEUDOANIONS



27
GPa

Outline

5. Applications to high pressure

a) ELF: predicting compressibility & transformations

b) NCI: He bonds

c) Electron density: the borate family

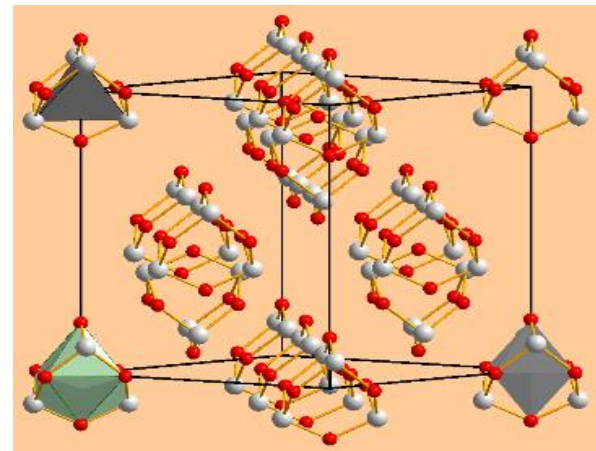
6. The codes

7. Summary

He is noble?

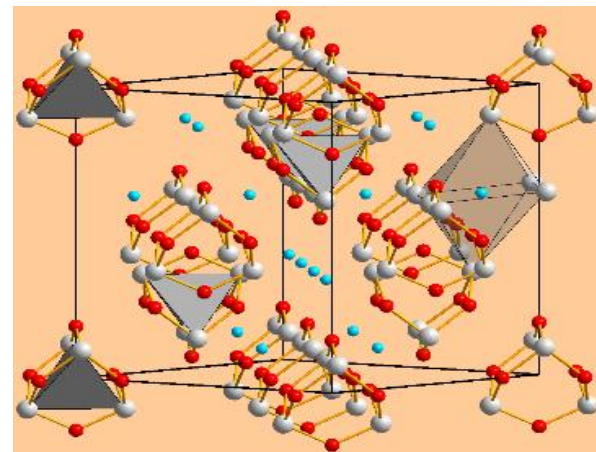
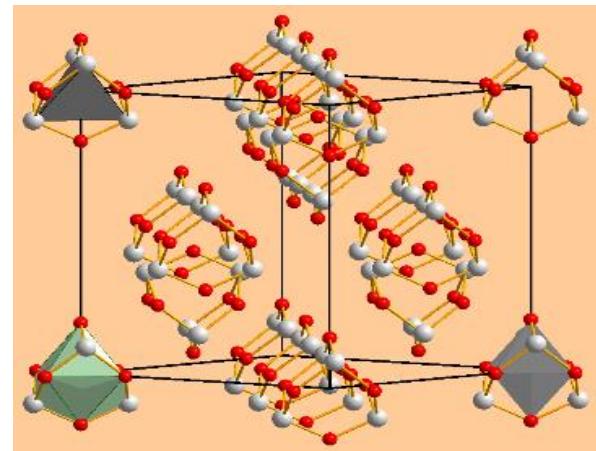
- Arsenolite has a closed-compact As_4O_6 cages
- It amorphizes at 15 GPa when compressed alone
- It does not amorphize when He is used as pressure transmitting medium.

WHY??



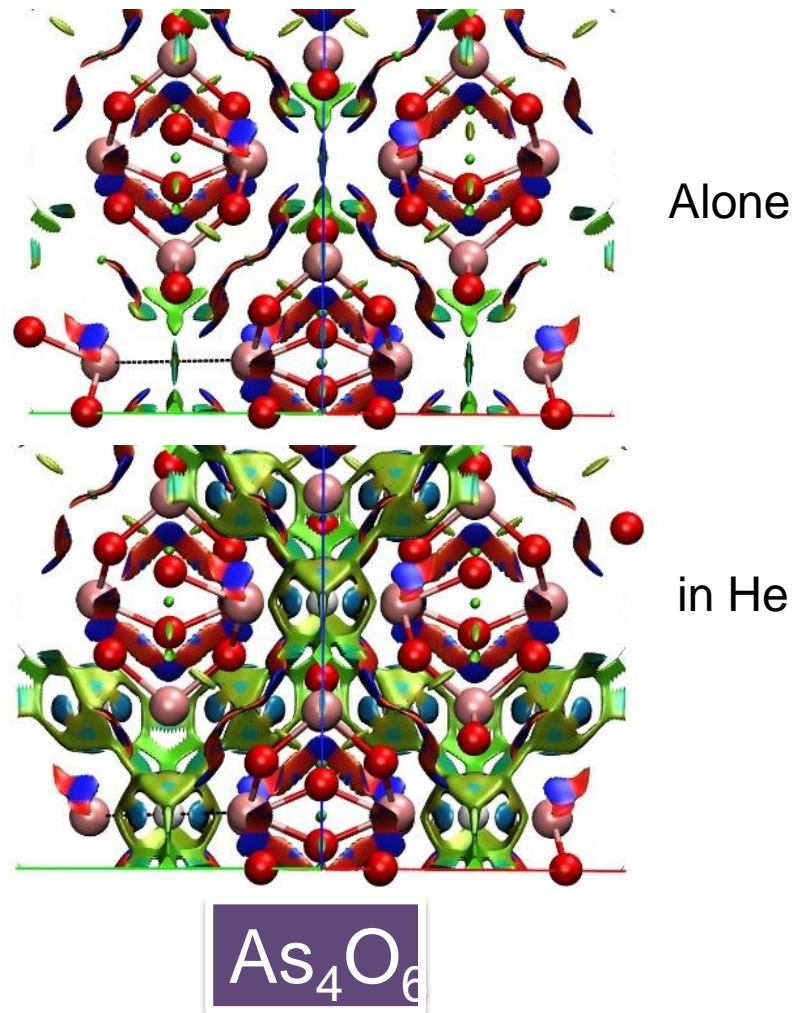
He is noble?

- It gives ordered helium trapping in the voids when compressed in He



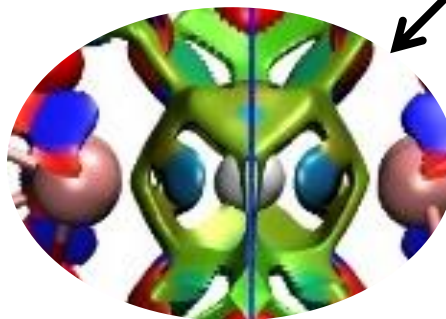
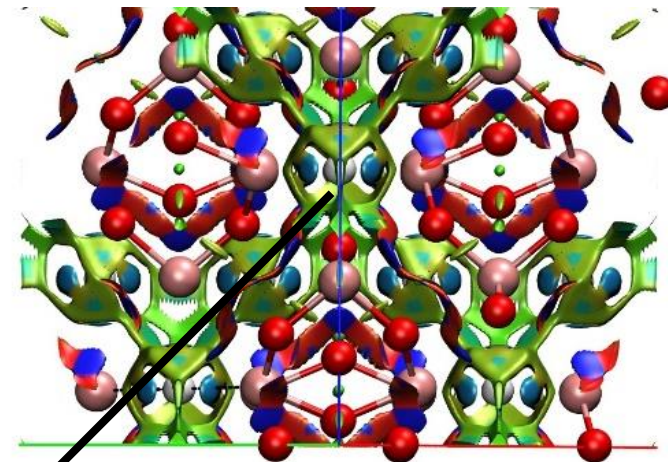
He is noble?

- It gives ordered helium trapping in the voids when compressed in He



He is noble?

- Very localized structural bonds
- They alter the properties of pure As_4O_6 ,
 - e.g. pressure-induced amorphization is hampered



Outline

5. Applications to high pressure

a) ELF

Polarity of ionic compounds

Predicting compressibility

Rationalizing phase transitions

b) NCI: He bonds

c) ELF: superconductivity

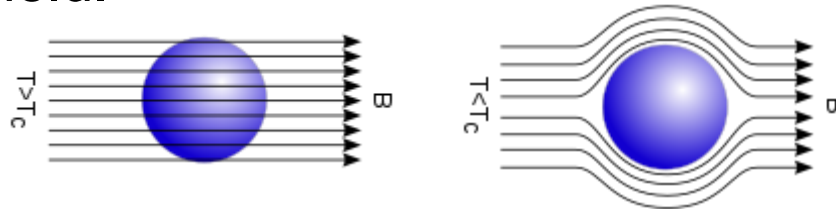
8. The codes

9. Summary

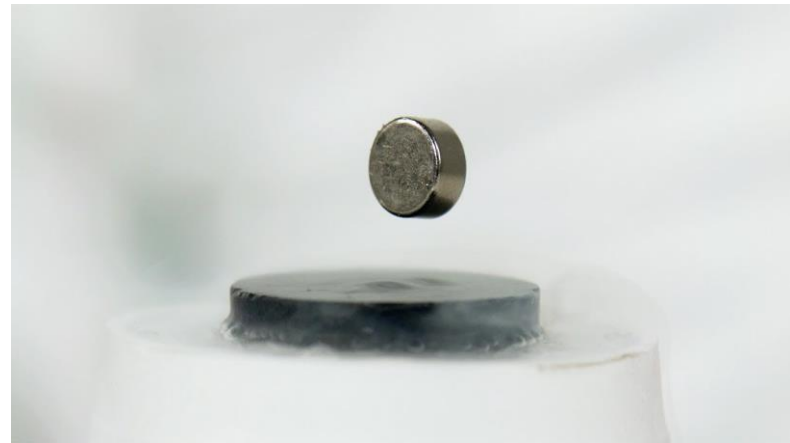
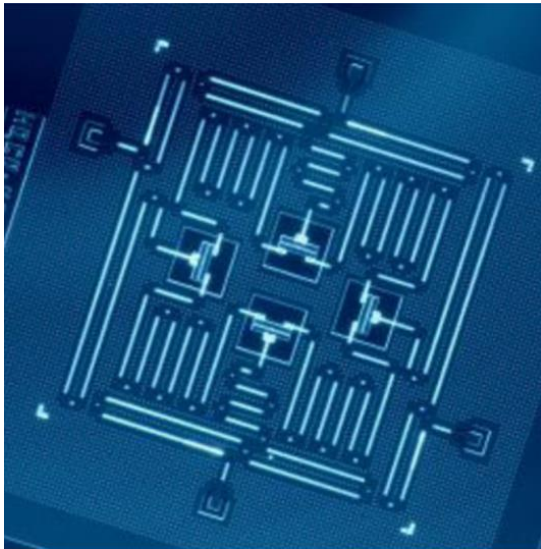
Superconducting materials

Superconductivity is technologically very interesting...

- Two key features: null resistivity and perfect repulsion of magnetic field.



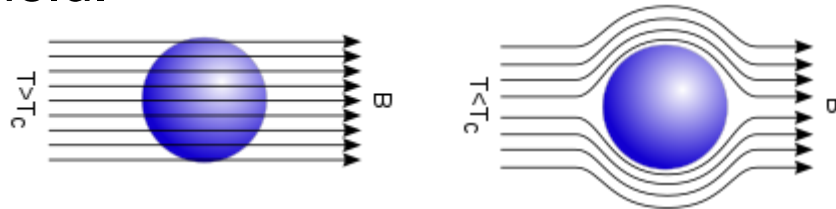
- Interesting applications: electronics, levitation



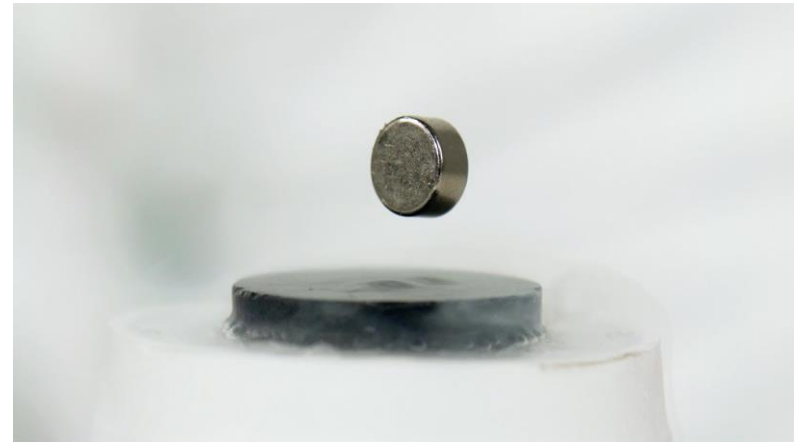
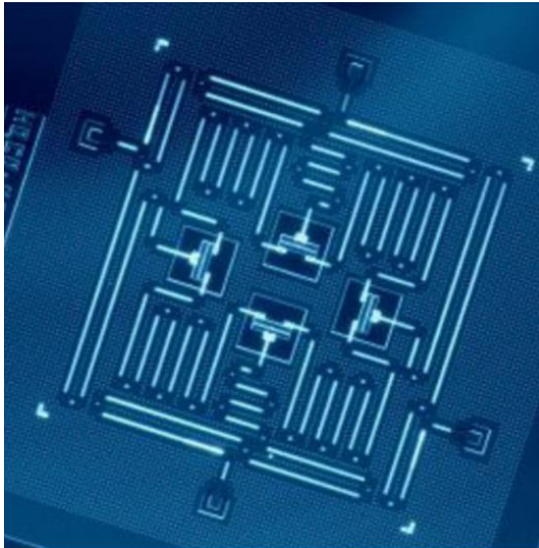
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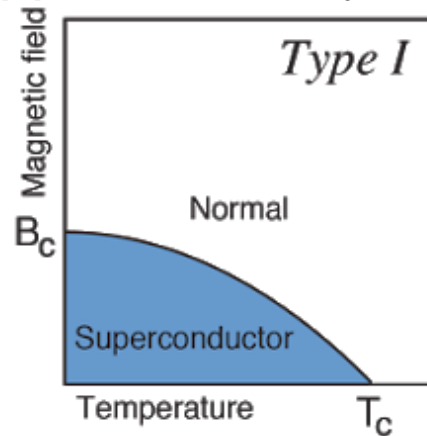
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However...

Superconducting materials

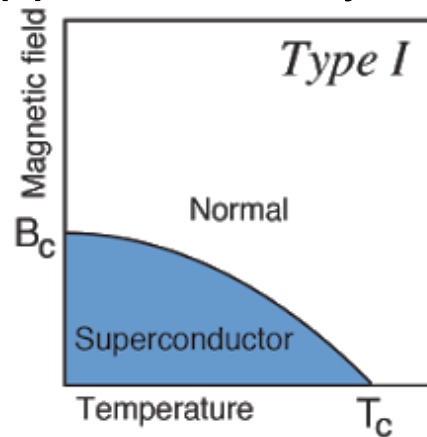
- Superconductivity appears at very low temperatures



- The problem: very low is VERY low: Al (1.2K), Zn (0.88 K)

Superconducting materials

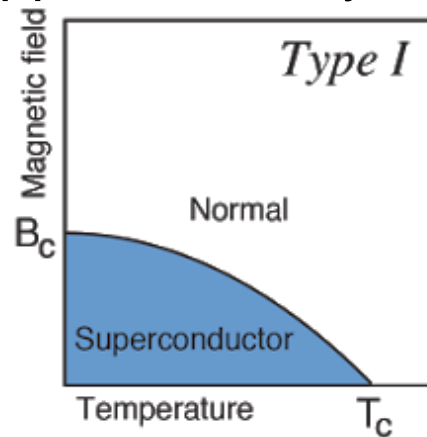
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- Calculations are extremely expensive, and not always work
AlH₃ : T_c theo= 132-146 K ... T_c exp=4 K! (250GPa)

Superconducting materials

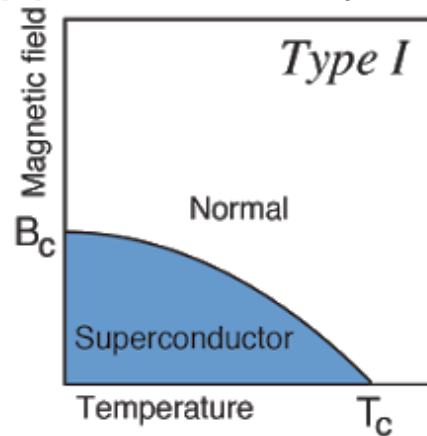
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Superconducting materials

- Superconductivity appears at very low temperatures

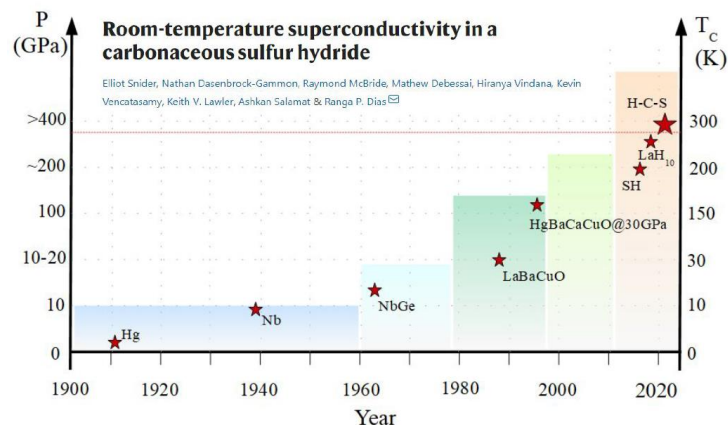


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AlH₃ : T_c theo= 132-146 K ... T_c exp=4 K! (250GPa)
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Let's start from a reasonable starting point...H-based systems!

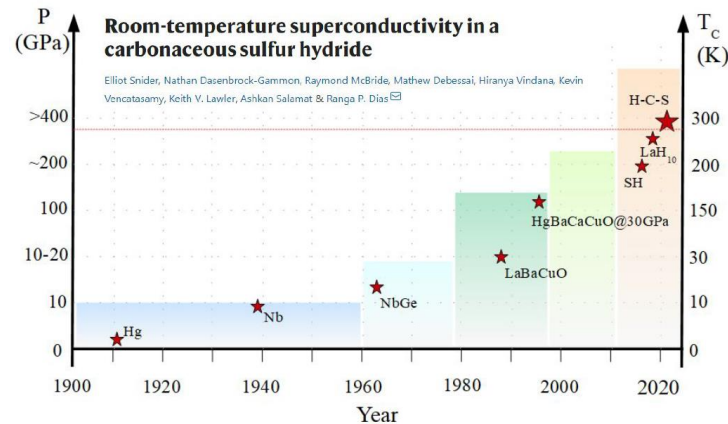
Hydrogen based superconductors

- A new family of high-temperature superconductors: H_3S (203K), LaH_{10} (260K). **Room temperature SC of C-S-H system!**



Hydrogen based superconductors

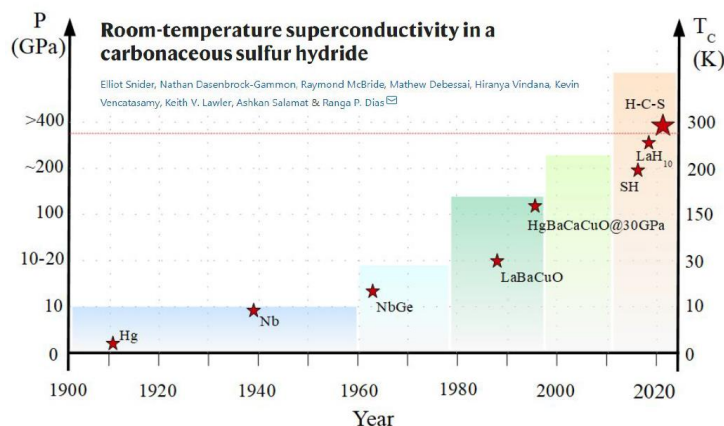
- A new family of high-temperature superconductors: H_3S (203K), LaH_{10} (260K). **Room temperature SC of C-S-H system!**



- Something to keep in mind:
 - High pressures are needed:
 H_3S at 203K and **150GPa**, LaH_{10} at 260K and **190GPa**.
 - Hopefully replaced by chemical pressure! (next step)

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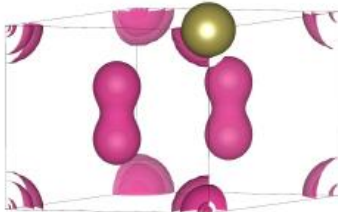


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Let's start!

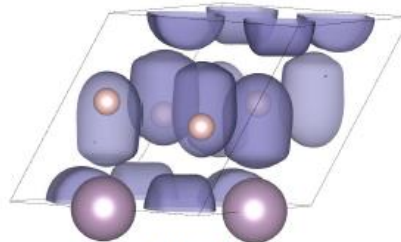
- 178 systems with available Tc values (pure and binary compounds)

Molecular



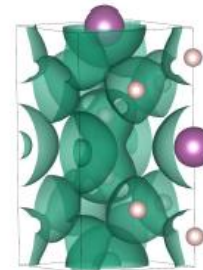
TeH₄ (150)
ELF = 0.9

Covalent



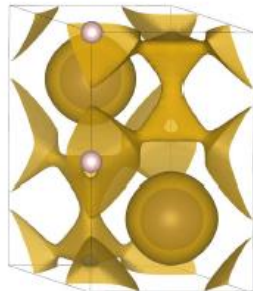
PH₂ (173)
ELF = 0.65

Weak H Interactions



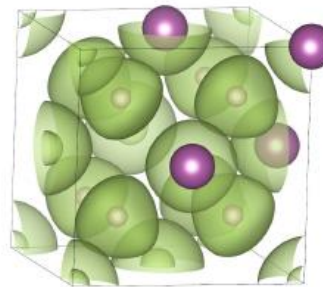
ScH₄ (17)
ELF = 0.45

Electride



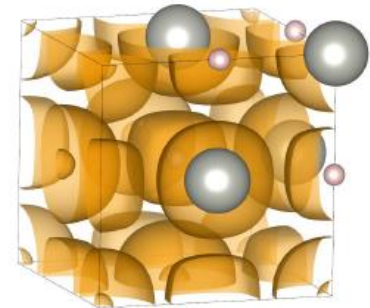
CrH (53)
ELF = 0.39

Ionic

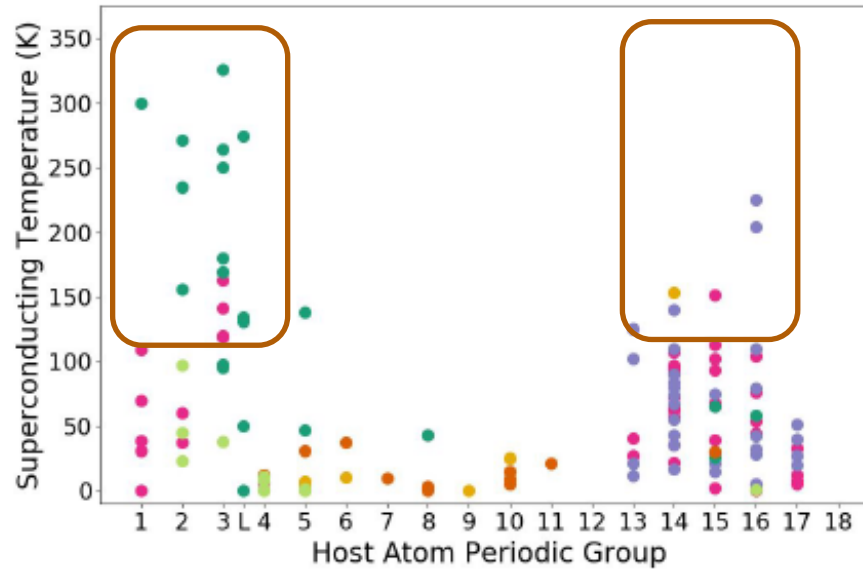
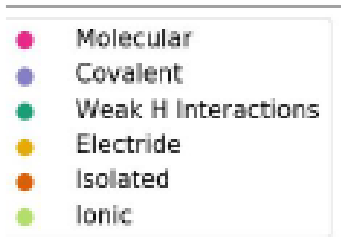


ScH₂ (14)
ELF = 0.52

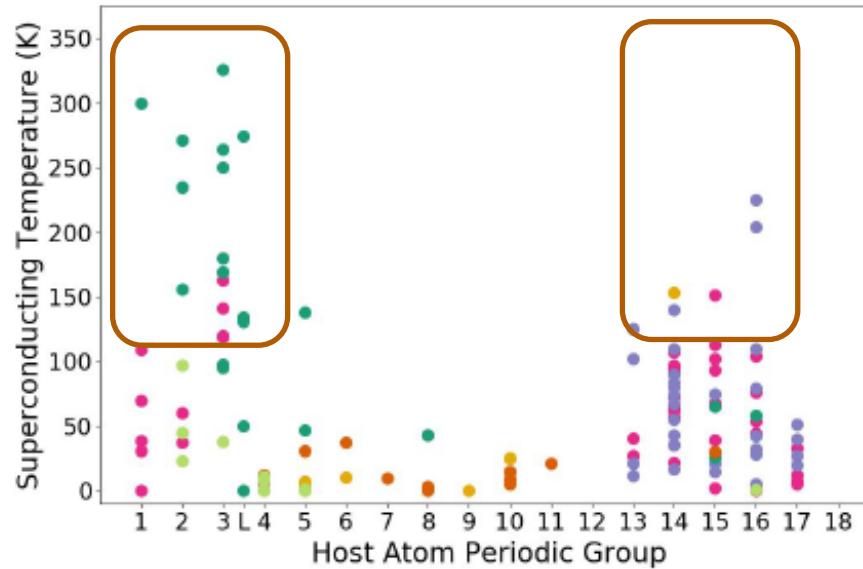
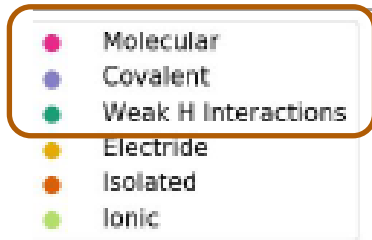
Isolated



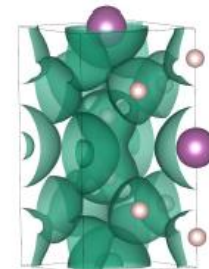
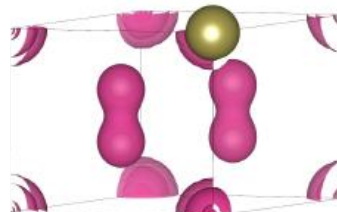
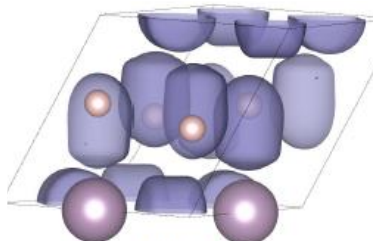
PdH (70)
ELF = 0.25

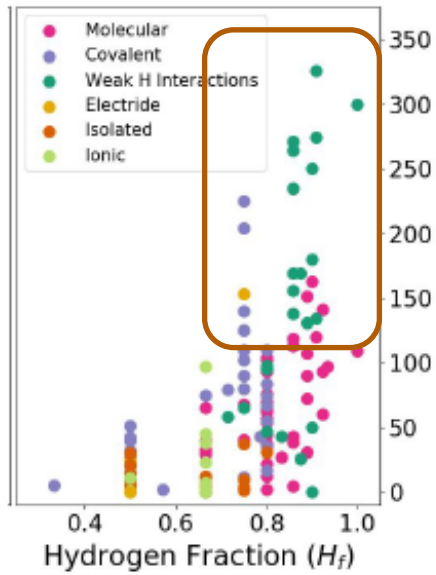


- Characteristics:
 - s, p atoms

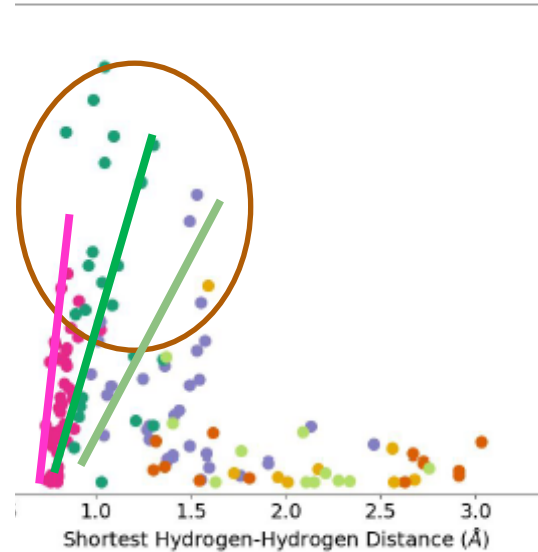
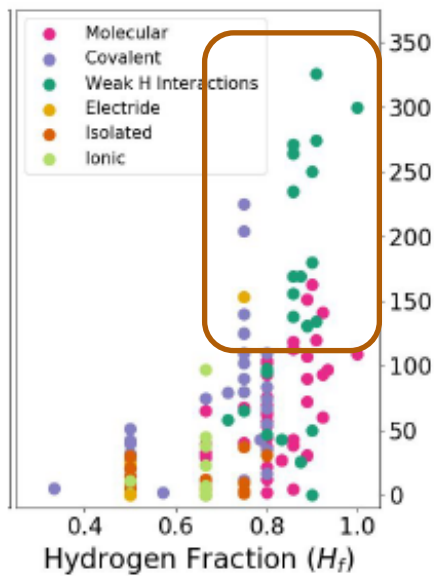


- Characteristics:
 - s, p atoms
 - Non charged (covalent, molecular or weak H interactions)

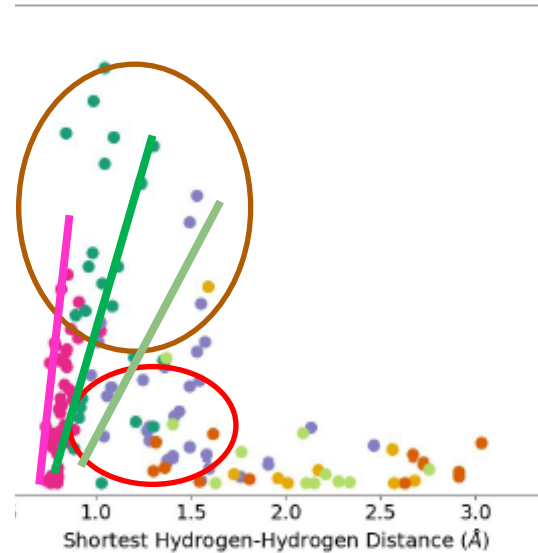
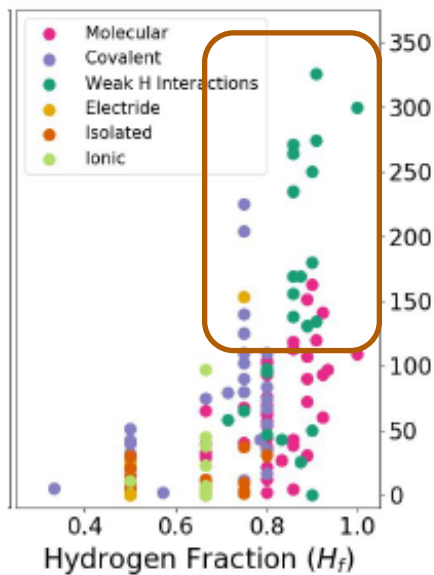




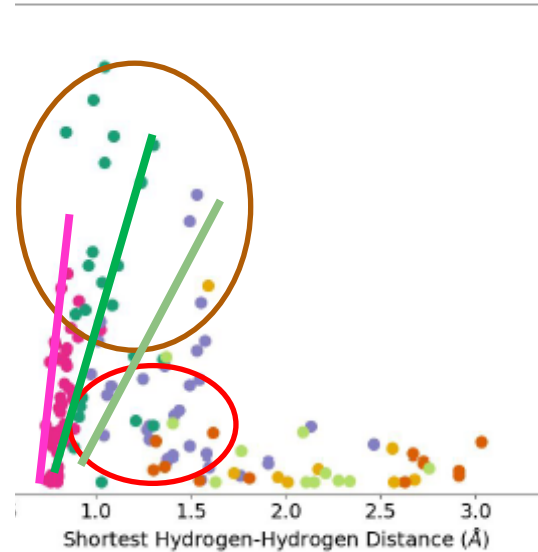
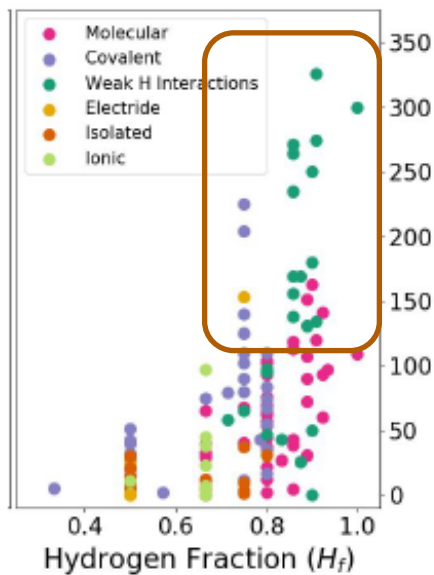
- Characteristics:
 - Hydrogen rich systems



- Characteristics:
 - Hydrogen rich systems
 - Stretched H-H bonds



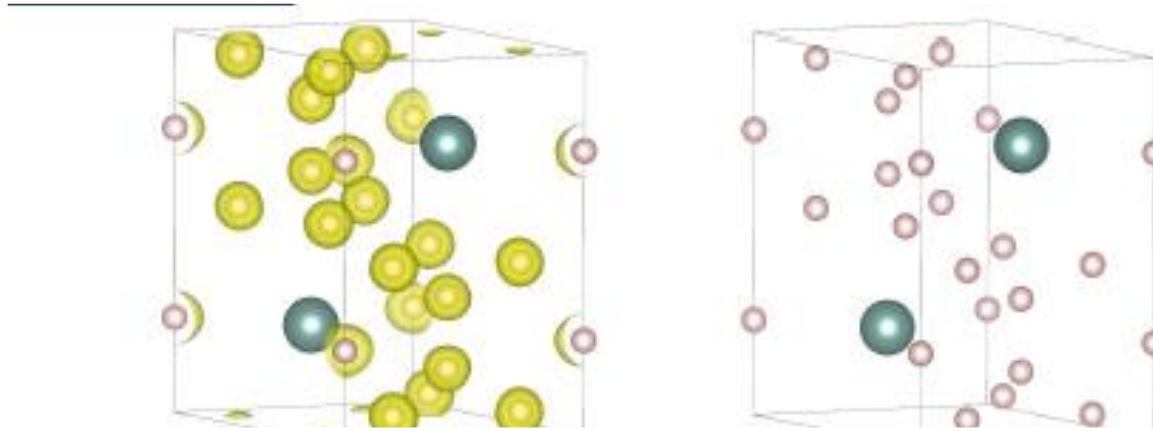
- **Characteristics:**
 - Hydrogen rich systems
 - Stretched H-H bonds
- **Necessary but not sufficient conditions!**



- **Characteristics:**
 - Hydrogen rich systems
 - Stretched H-H bonds
- **Necessary but not sufficient conditions!**

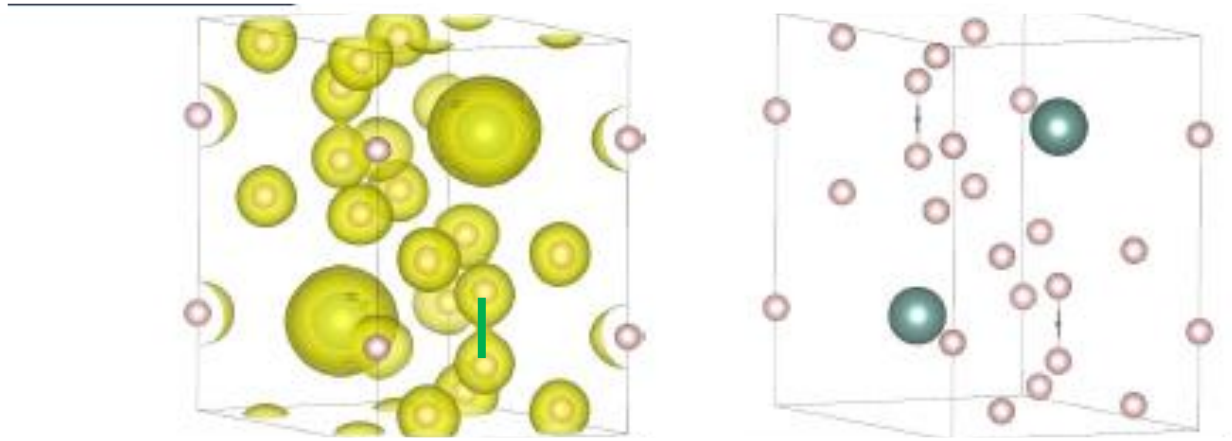
How about the electron localization?

YH_9

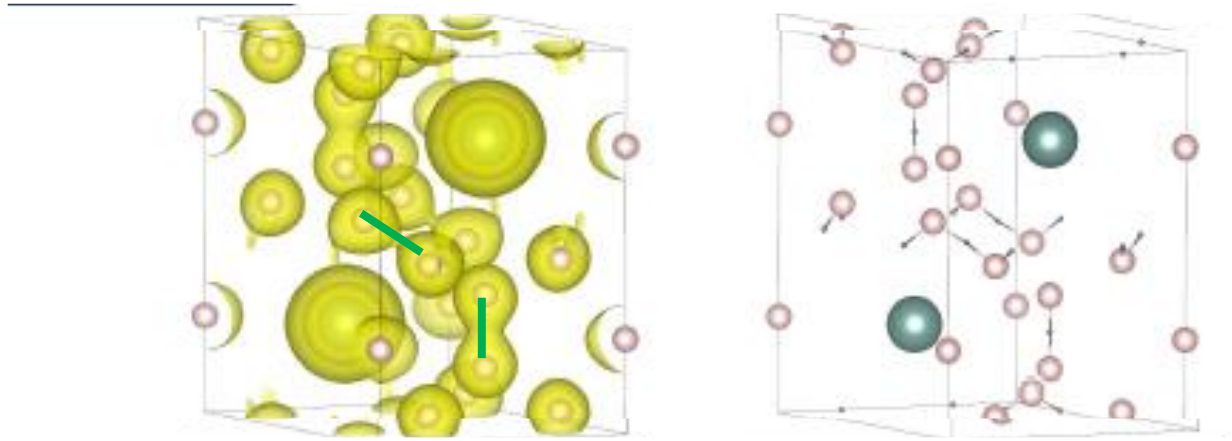


$\text{ELF}=0.9$

YH_9

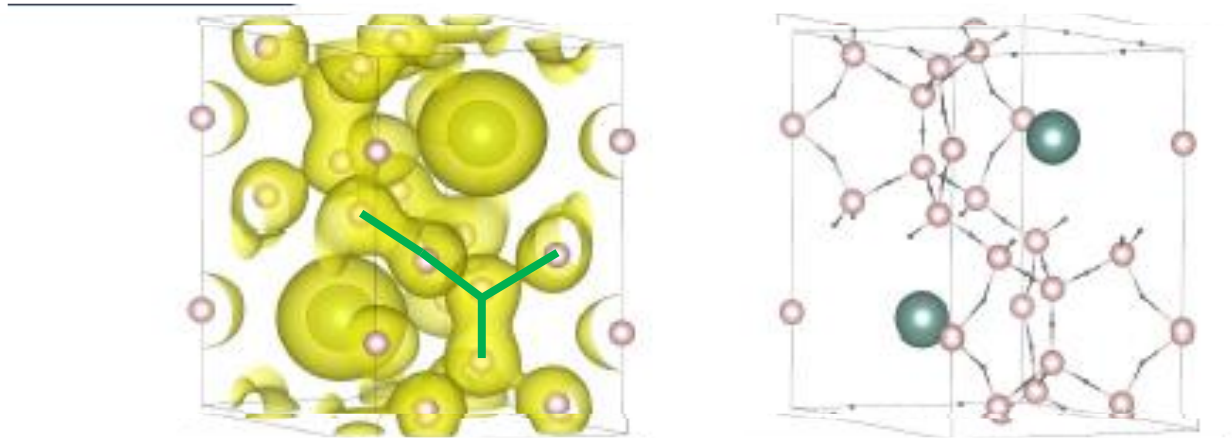


$\text{ELF}=0.78$

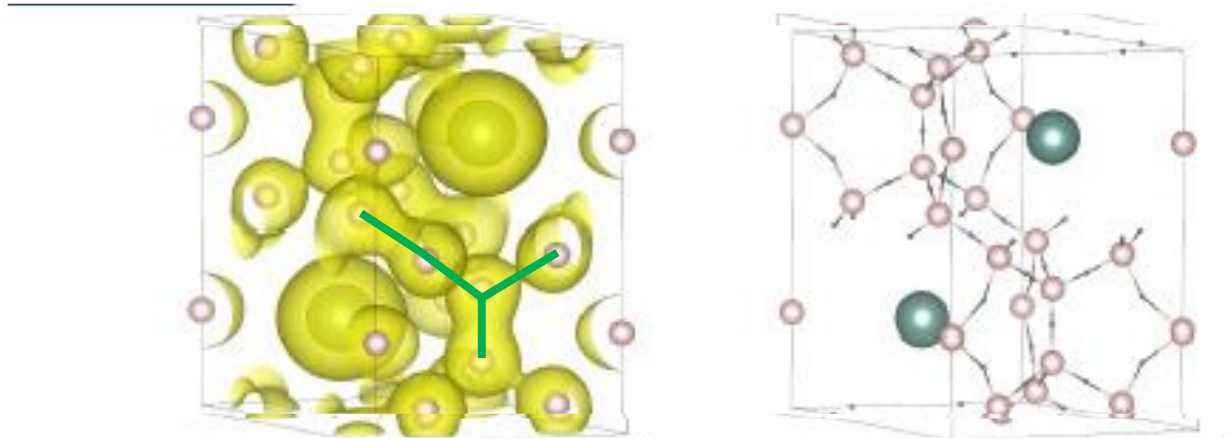


$\text{ELF}=0.72$

YH_9



$\text{ELF}=0.57$



$\text{ELF}=0.57$

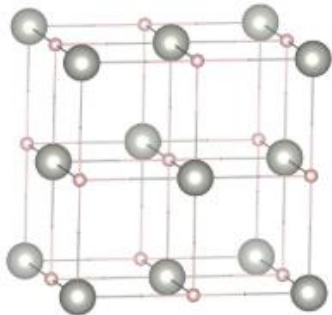
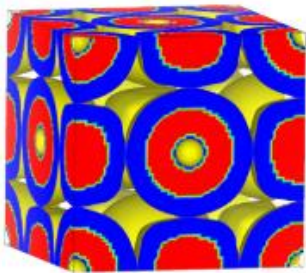
- At $\text{ELF}=0.57$ the hydrogen network is connected
- We call this ELF value the networking value, ϕ

Networking value ϕ

- We determined it in the 178 crystalline structures

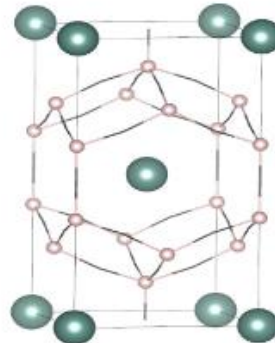
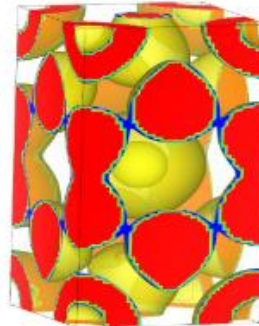
PdH(70)

ELF = 0.19



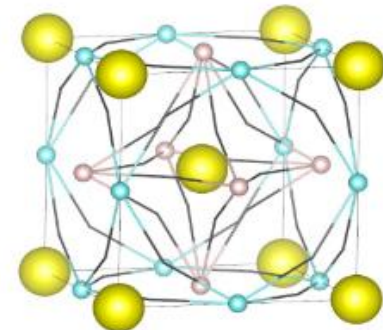
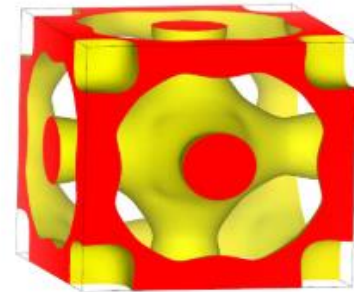
YH4(15)

ELF = 0.43

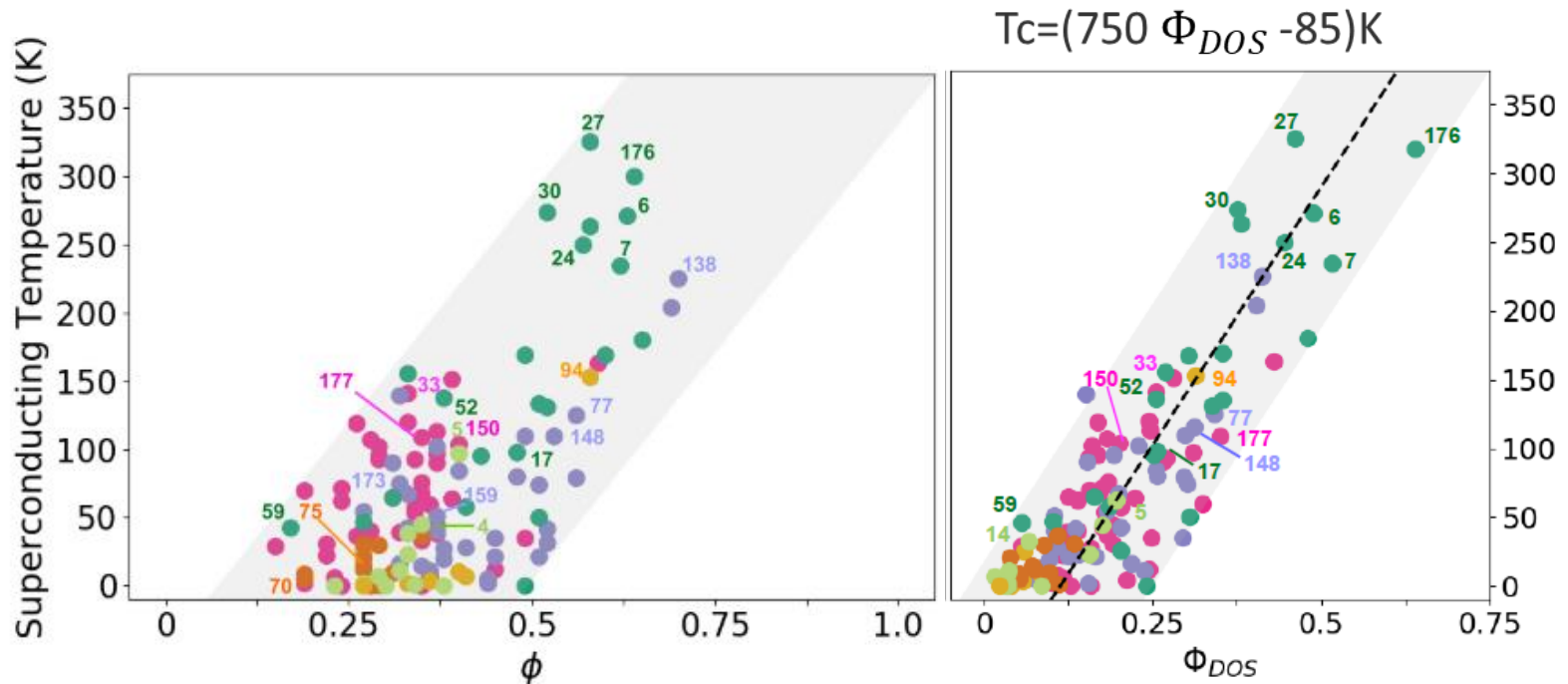


H3S(138)

ELF = 0.68



Correlation between T_c and ϕ



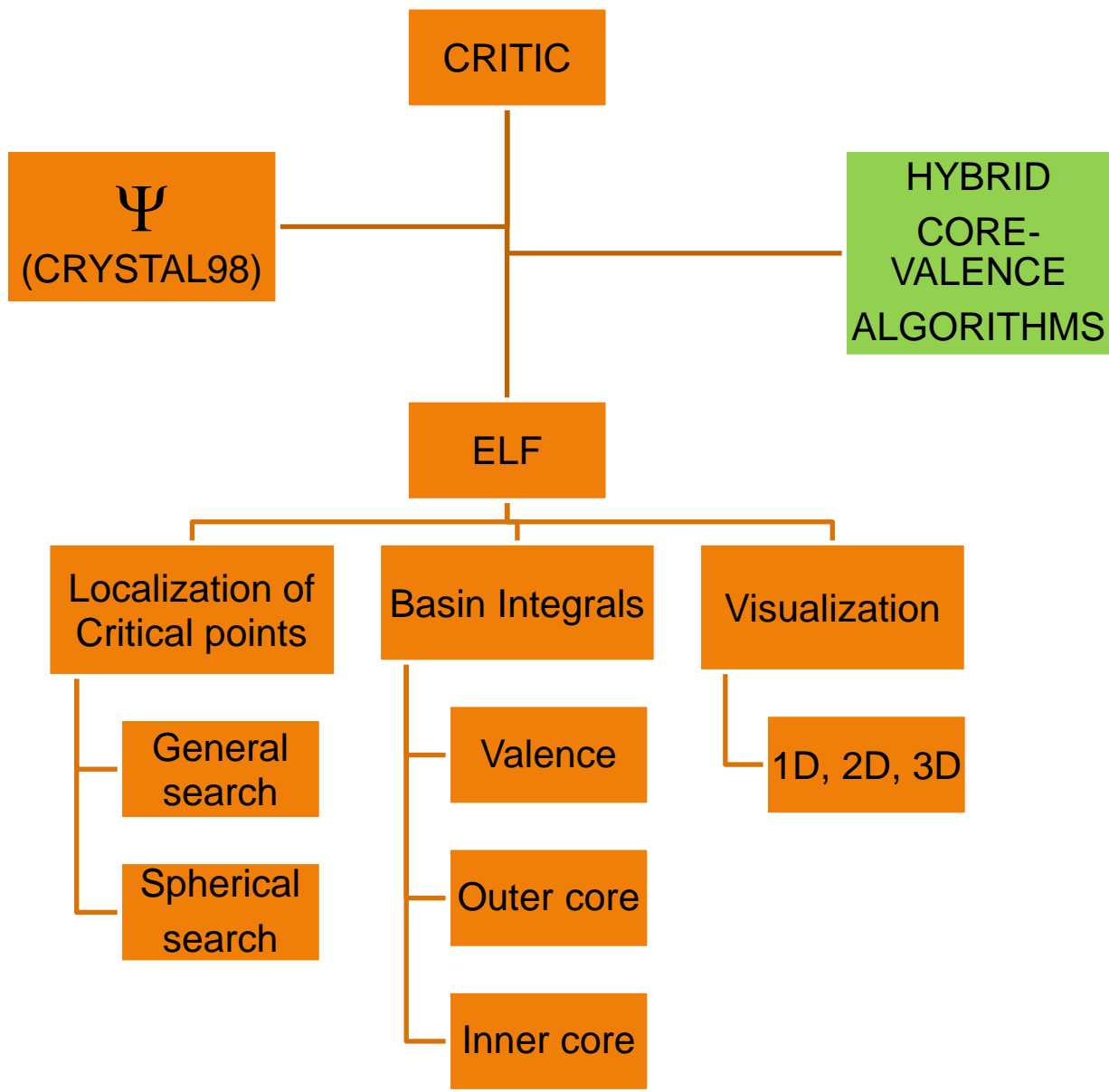
- ϕ provides a first estimate of the superconducting critical temperature
- Including chemical and electronic structure features reduces the dispersion within 60K
- Sufficient condition, avoids very expensive calculations!

Summary

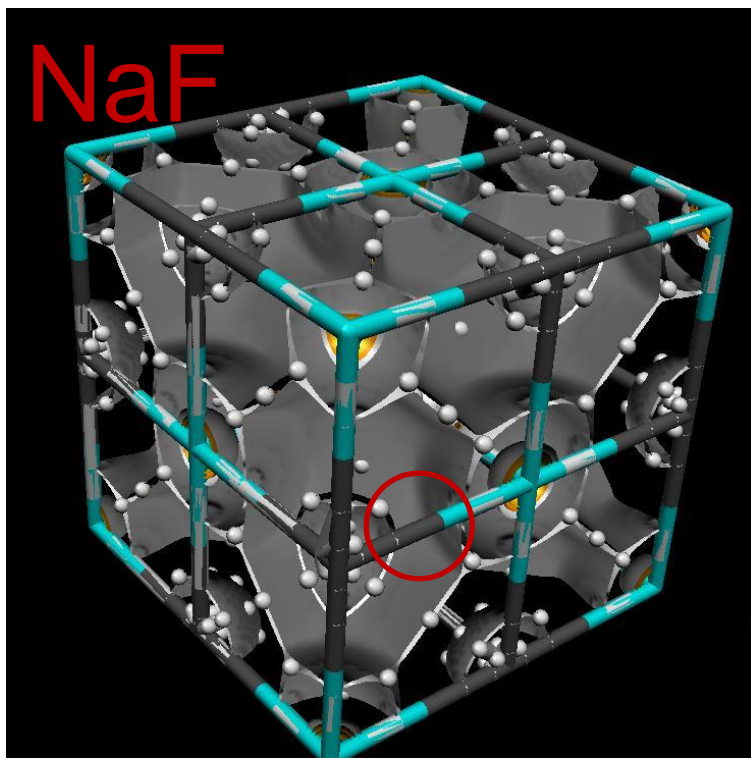
- We can derive macroscopic properties from the electronic structure/bonding
 - It improves all other structural or electronic indexes previously proposed
 - It only requires DFT ground state+ELF calculations
- It offers a simple way of screening new superconductors
 - Points the way to inverse engineering new hydrogen-based superconductors

Programs

- ELF and NCI from periodic calculations:
CRITIC
- NCI from molecular calculations, PDBs
and crystallographic data
- NCI from PDB with PDB treatment:
NCIWeb now also available at:
<https://nciweb.dsi.upmc.fr/index.php>



CRITIC

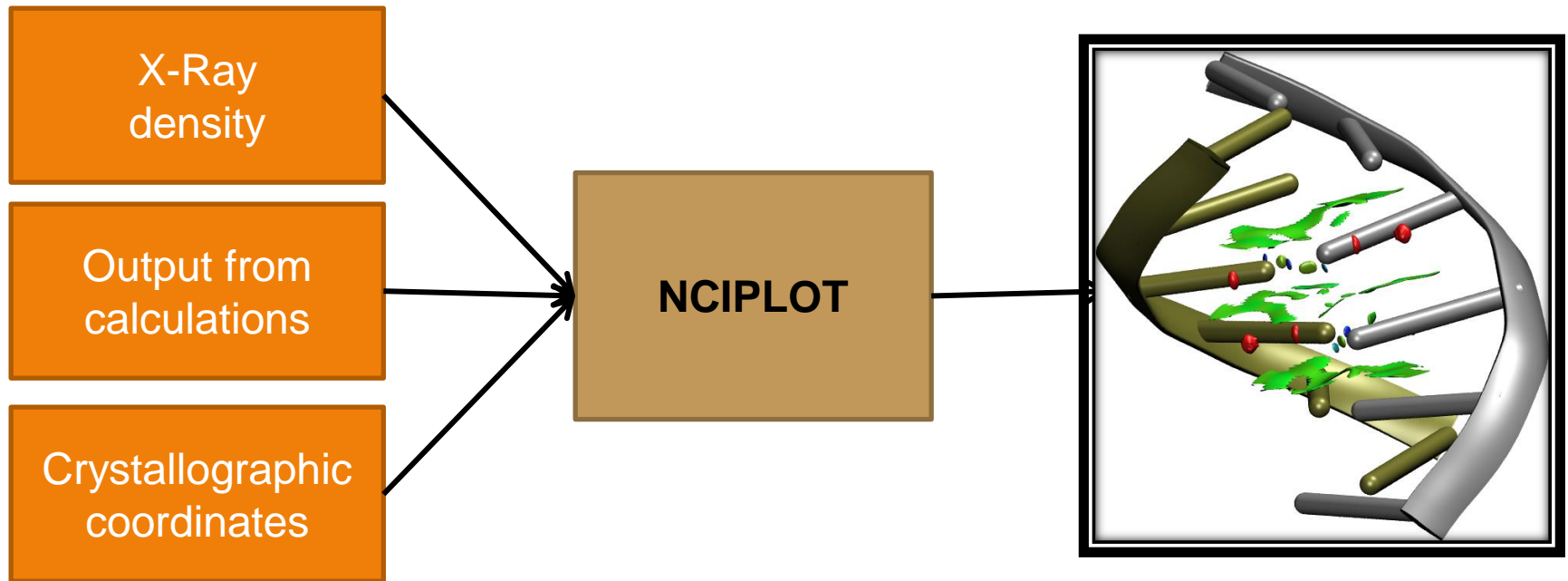


Total	Generic Hybrid	
Maximum number	64	64
Bond number	176	200
Ring number	200	200
Cage number	64	64
Morse sum equals	24	0

N	SYM	CP	TYPE	X	Y	Z	M
1	Oh	(3,-3)	Nucleus	0.000	0.000	0.000	4
2	Oh	(3,-3)	Nucleus	0.000	0.000	0.500	4
3	C3v	(3,-3)	Maximum	0.038	0.462	0.462	32
4	C4v	(3,-3)	Maximum	0.000	0.084	0.500	24
5	D2h	(3,-1)	Bond	0.250	0.500	0.250	24
6	C4v	(3,-1)	Bond	0.790	0.500	0.500	24
7	C2v	(3,-1)	Bond	0.059	0.500	0.941	48
8	C2v	(3,-1)	Bond	0.000	0.453	0.453	48
9	C3v	(3,-1)	Bond	0.487	0.487	0.013	32
10	C4v	(3,-1)	Bond	0.000	0.000	0.530	24
11	C2v	(3,+1)	Ring	0.000	0.320	0.320	48
12	C3v	(3,+1)	Ring	0.048	0.048	0.452	32
13	C4v	(3,+1)	Ring	0.000	0.000	0.934	24
14	C2v	(3,+1)	Ring	0.521	0.021	0.000	48
15	C2v	(3,+1)	Ring	0.516	0.016	0.500	48
16	Td	(3,+3)	Cage	0.250	0.250	0.250	8
17	C3v	(3,+3)	Cage	0.982	0.518	0.018	32
18	C4v	(3,+3)	Cage	0.500	0.500	0.023	24

Morse sum must is zero

NCIPLLOT



NCIWEB

The screenshot displays the NCIweb web interface. At the top, there are logos for NCIweb, LCT (Laboratoire de Chimie Théorique), Sciences Sorbonne Université, and CNRS. The main content is divided into two columns. The left column is titled "Submit NCIweb job" and contains a form with the following fields and options:

- Name:** A text input field with the placeholder "Enter Name".
- Email:** A text input field with the placeholder "Enter Email".
- Upload your structure (PDB or XYZ format):** A file upload button labeled "Parcourir..." and a status message "Aucun fichier sélectionné."
- Choose structure by PDB ID:** A text input field with the example "E.g.: 2RC5".
- Choose the operation mode:** Three radio button options: "Intramolecular" (selected), "Intermolecular", and "Ligand".
- Clean structure:** A checkbox that is checked.
- Protonate protein:** An unchecked checkbox.
- Protonate ligands:** An unchecked checkbox.
- Preselected ligands:** An unchecked checkbox.

The right column is titled "What is NCIweb?" and "What do I need to use NCIweb?".

What is NCIweb?

NCIweb is a web implementation of the popular NCIplot code. At the moment, NCIweb works with promolecular densities. It provides a representation of the non-covalent interactions of a system based on the reduced density gradient of the electron density. More information on the NCIplot code and the theoretical background of the method can be found [here](#).

What do I need to use NCIweb?

To use NCIweb you simply need a PDB or XYZ structure to upload, on which the analysis will be performed. You can also fetch a PDB file directly with its PDB ID. If everything works fine, you will receive an email with your results, including a simple VMD script for visualization. Three operation modes are available:

- Running in **Intramolecular mode** will study all non-covalent interactions in the system.
- Running in **Intermolecular mode** will require manual definition of two fragments, and will only study interactions between them.
- Running in **Ligand mode** will require manual definition of a ligand and a receptor, and will only study interactions between the ligand and the receptor in the proximity.

For **PDB files only**, additional options are available:

- The **clean structure** checkbox will selectively remove non-protein fragments from the structure, enabling more processing options.
- The **protonate protein** checkbox will add hydrogens to the protein using **OpenBabel**.
- The **protonate ligands** checkbox will add hydrogens to the ligands using **OpenBabel**. This may fail for some molecules.
- The **preselected ligands** checkbox will assume that the ligands to be preserved have been signaled with residue id's LIG1, LIG2 etc. in the input PDB file. This will skip visual selection of ligands.

With our preprocessing pipeline, a PDB structure from a data bank may be directly analyzed. Nevertheless, we recommend manual and careful preparation and protonation whenever possible.

1. Visual, interactive interface.
2. Clear instructions with interactive explanations.
3. Both xyz and pdb files are accepted, either from uploaded or from a PDB record.

NCIWEB

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MyPDB

Search

Visualize

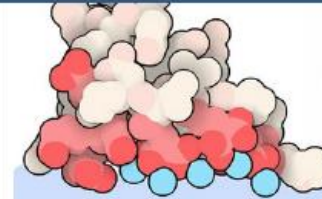
Analyze

Download

Learn

The RCSB PDB builds upon the data by creating tools and resources for research and education in molecular biology, structural biology, computational biology, and beyond.

New Video: Penicillin and Antibiotic Resistance



Proteins and Biomaterials

Latest Entries

As of Tuesday Apr 09 2019



Features & Highlights



Mandatory PDBx/mmCIF format files submission for MX depositions
Submission of PDBx/mmCIF format files for crystallographic depositions to the PDB will be mandatory from July 1st 2019 onward. PDB format files will no longer be accepted for deposition of structures solved by MX techniques.



Join Our Team as a Biocurator
Curate, validate, and standardize macromolecular structures from the PDB community at Rutgers, The State University of New Jersey.



New Architecture and Services Enable Faster Access to More Information
Explore the improved display of PDB Statistics, structure funding information, and 3D views of ligands and electron density.

News

Publications



Take the Molecule of the Month User Survey and Enter to Win
Answer a few questions and enter a drawing for a signed copy of *The Machinery of Life*. - 04/09/2019

High School Students: Submit Antibiotic Resistance Videos Before April 23 - 04/02/2019

Molecular Landscapes and the Art of Science - 03/26/2019

The PDB Archive Reaches a Significant Milestone - 03/19/2019

PDB at a Glance 46689 Distinct Protein Sequences | 42251 Structures of Human Sequences | 10904 Nucleic Acid Containing Structures | More Statistics

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RCSB Partners Nucleic Acid Database

wwPDB Partners RCSB PDB PDBu PDBj BMRB

RCSB PDB (citation) is managed by two members of the Research Collaboratory for Structural Bioinformatics: Rutgers and UCSD-SDSC

RUTGERS | UC San Diego | SDSC

RCSB PDB is a member of the PDB Consortium

NCIWEB

The screenshot displays the NCIWEB web interface. At the top, there are logos for NCIweb, LCT (Laboratoire de Chimie Théorique), Sciences Sorbonne Université, and CNRS. A red warning message reads: "Remember to carefully check the protonation of ligands!". Below this, a white box contains a success message: "Success! Help us clean your file". It lists three prompts with corresponding checkboxes and radio buttons: "A total of 2 ligands were found. By default, no ligands will be preserved. Do you want to select which ligands to preserve?" (with a "YES" radio button selected), "Preserve ligand GTJ, number undefined from chain A1404 ?" (with an unchecked checkbox), "Preserve ligand DMS, number undefined from chain A1403 ?" (with an unchecked checkbox), "A total of 1 chains were found. By default, all chains will be considered. Do you want to select which chains to preserve?" (with a "NO" radio button selected), and "A total of 34 residues with multiple orientations were found. By default, the first orientation will be considered. Do you want to select which orientations to choose?" (with a "NO" radio button selected). A large "VALIDATE" button is at the bottom of this section. To the right, a "Visualize your system" panel shows a 3D ribbon model of a protein-ligand complex. Below the model are controls: "Toggle spin!", "Toggle rock!", "White background", "Add Ball & Stick", "Add Licorice", and "Download image". A "Visualize current selection" button is at the bottom of the visualization panel.

1. All crystallographic molecules and salts have been removed automatically.
2. An **interactive visualization** page lets us select chains, ligands and orientations to properly prepare our structure.

Summary

- Quantum chemistry and topology can be coupled to provide a quantitative picture of bonding
- This picture is very helpful in understanding high pressure behavior

**Keep all this in mind for
this afternoon lab
work!!!**