NMR and EPR spectroscopies with Quantum-Espresso

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Outline

• PART I:

- Basic principles of magnetic resonance spectroscopy
- Introduction to experimental NMR
- Interpretation of NMR spectra
- Solid state NMR
- PART II:
 - Effective NMR spin hamiltonian
 - The GIPAW method
 - Examples
 - Brief introduction to EPR spectroscopy and EPR parameters

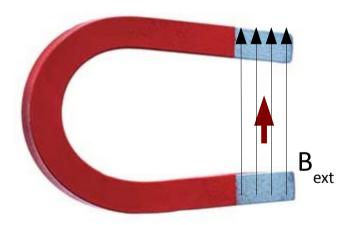
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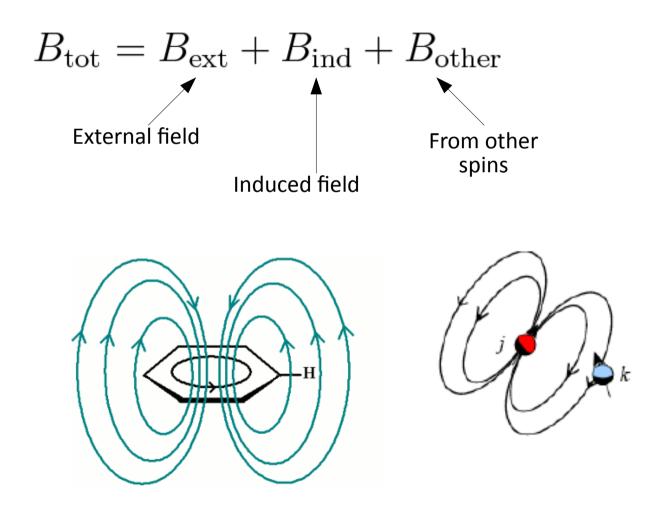
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Principles of magnetic resonance

NMR = <u>N</u>uclear <u>M</u>agnetic <u>R</u>esonance EPR = <u>E</u>lectron <u>P</u>aramagnetic <u>R</u>esonance (also ESR)

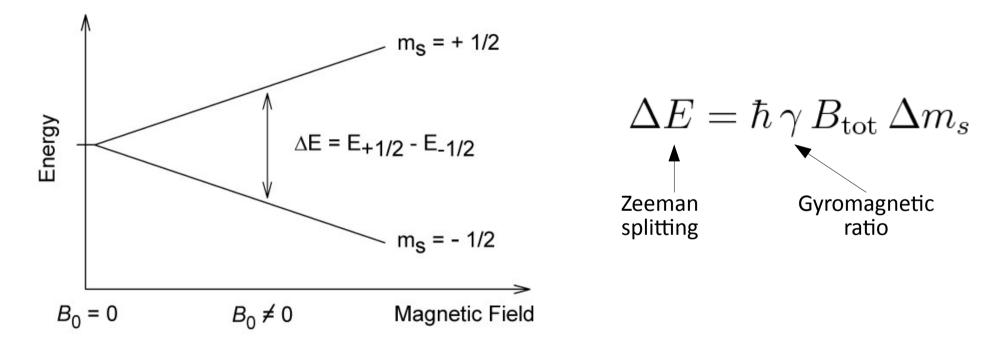


A spin (nuclear or electron), in a magnetic field



Principles of magnetic resonance

A spin in a magnetic field will align parallel or antiparallel to the field. The Zeeman splitting is proportional to the <u>total</u> magnetic field.



The energy splitting can be probed by an electromagnetic wave of frequency ω :

$$\Delta E = \hbar \omega$$

Some useful relations

For a nucleus

$$\gamma = \frac{Q}{2M}g = \frac{g\mu_N}{\hbar}$$

 γ = gyromagnetic ratio q = charge m = mass g = g-factor μ_N = nuclear magneton

For the electron

$$\gamma_e=rac{e}{2m_e}g_e=rac{g_e\mu_B}{\hbar}$$
 $\mu_{\rm B}$ = bohr magneton (>> nuclear magneton)

For example:

g(¹H) = 5.585694 → γ (¹H) = 2.67·10⁸ Hz/T g(elec.) = -2.002319 → γ (elec.) = 1760·10⁸ Hz/T

Resonance frequency is proportional to γ : ¹H @ 9.306 T \rightarrow 400 MHz

NMR active nuclei

Atom	Isotope	Spin	Larmor frequency (MHZ at 9.306 T)	Abundance (%)	Absolute sensitivity	Quadrupole (barn)
Hydrogen	1H	1/2	-400.00	99.98	1.00E+000	0
Deuterium	2D	1	-61.40	0.02	1.45E-006	0.00273
Lithium	6Li	1	-58.86	7.42	6.31E-004	-0.0008
Lithium	7Li	3/2	-155.45	92.58	2.70E-001	-0.045
Boron	10B	3	-42.98	19.58	3.90E-003	0.074
Boron	11B	3/2	-128.34	80.42	1.30E-001	0.0355
Carbon	13C	1/2	-100.58	1.11	1.76E-004	0
Nitrogen	14N	1	-28.90	99.63	1.01E-003	0.016
Nitrogen	15N	1/2	40.53	0.37	3.85E-006	0
Oxygen	170	5/2	54.23	0.04	1.08E-005	-0.026
Fluorine	19F	1/2	-376.31	100.00	8.30E-001	0
Sodium	23Na	3/2	-105.80	100.00	9.25E-002	0.12
Magnesium	25Mg	5/2	24.48	10.13	2.71E-004	0.22
Aluminum	27Al	5/2	-104.23	100.00	2.10E-001	0.149
Silicon	29Si	1/2	79.46	4.70	3.69E-004	0
Phosphorus	31P	1/2	-161.92	100.00	6.63E-002	0
Sulfur	33S	3/2	-30.68	0.76	1.72E-005	-0.055
Chlorine	35Cl	3/2	-39.19	75.53	3.55E-003	-0.08
Chlorine	37Cl	3/2	-32.62	24.47	6.63E-004	-0.0632
Potassium	39K	3/2	-18.67	93.10	4.73E-004	0.055
Potassium	41K	3/2	-10.24	6.88	5.78E-006	0.067
Calcium	43Ca	7/2	26.91	0.15	9.28E-006	-0.05

NMR history

- 1937 **Rabi** predicts and observes nuclear magnetic resonance
- 1946 Bloch, Purcell first nuclear magnetic resonance of bulk sample
- 1953 **Overhauser** NOE (nuclear Overhauser effect)
- 1966 Ernst, Anderson Fourier transform NMR
- 1975 Jeener, Ernst 2D NMR
- 1985 Wüthrich first solution structure of a small protein (BPTI) from NOE derived distance restraints
- 1987 3D NMR + 13C, 15N isotope labeling of recombinant proteins
- 1990 pulsed field gradients (artifact suppression)
- 1996 new *long range* structural parameters: residual dipolar couplings and cross-correlated relaxation

Nobel prizes

1944 *Physics* Rabi
1952 *Physics* Bloch, Purcell
1991 *Chemistry* Ernst
2002 *Chemistry* Wüthrich
2003 *Medicine* Lauterbur, Mansfield

First ¹H NMR spectra of water

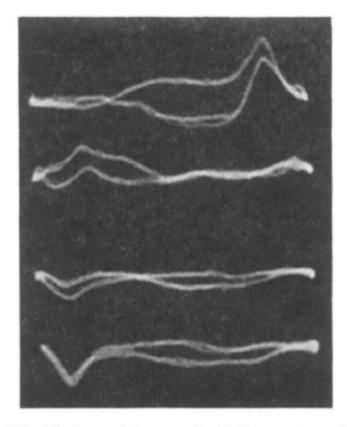
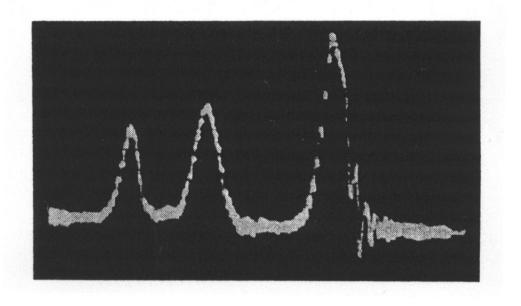


FIG. 10. Photographic record of the proton signal in water. The four traces from top to bottom correspond to the times t_1 , t_2 , t_3 , t_4 of Fig. 9. In the text they are referred to as a, b, c, d, respectively.

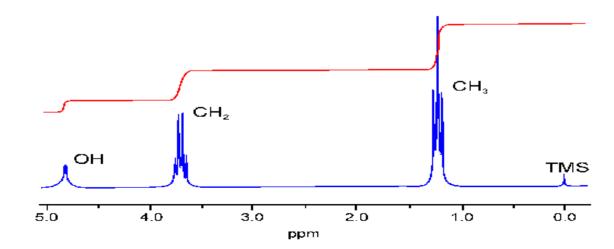
Bloch, Hansen and Packard, **The nuclear induction experiment** Physical Review **70**, 474 (1946),

First observation of the chemical shift



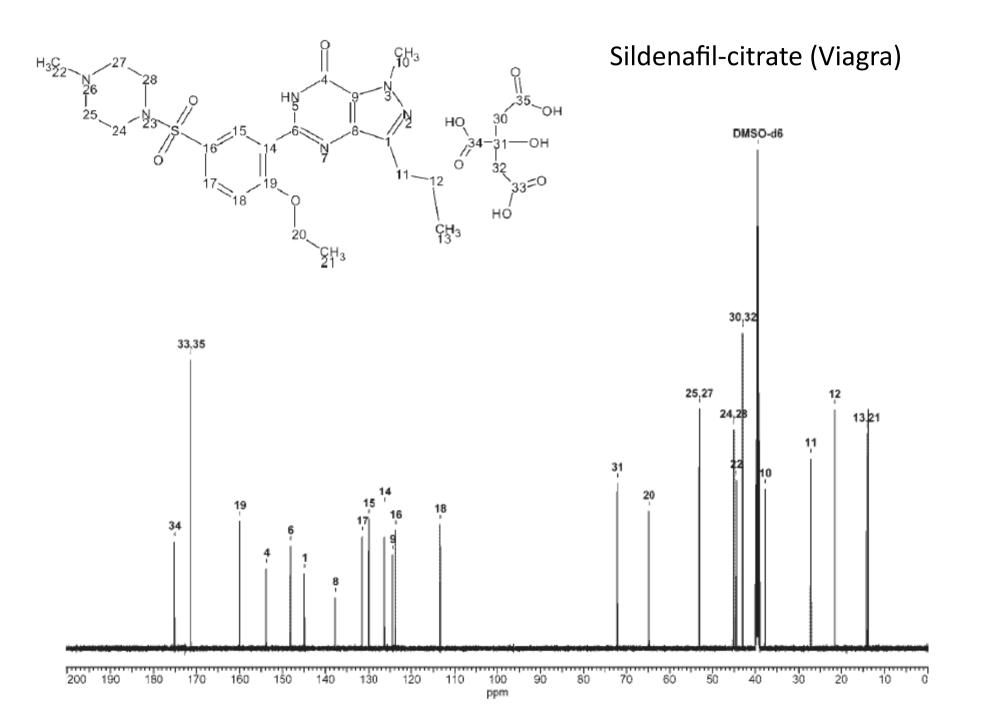
¹H NMR spectra ethanol

Modern ethanol spectra



Arnold, Dhamatti and Packard, J. Chem. Phys. 19, 507 (1951)

Chemical shift as molecular fingerprint



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Typical applications of NMR

1) Structural (chemical) elucidation

Natural product chemistry Synthetic organic chemistry

- analytical tool of choice of synthetic chemists
- used in conjunction with mass spectroscopy and IR

2) Study of dynamic processes

reaction kinetics study of equilibrium (chemical or structural)

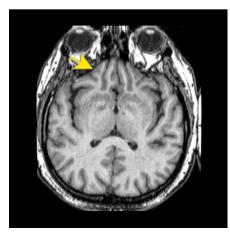
3) Structural (three-dimensional) studies

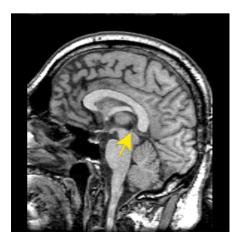
Proteins, Protein-ligand complexes DNA, RNA, Protein/DNA complexes Polysaccharides

4) Drug Design

5) Medicine: MRI

MRI images of the Human Brain





Outline

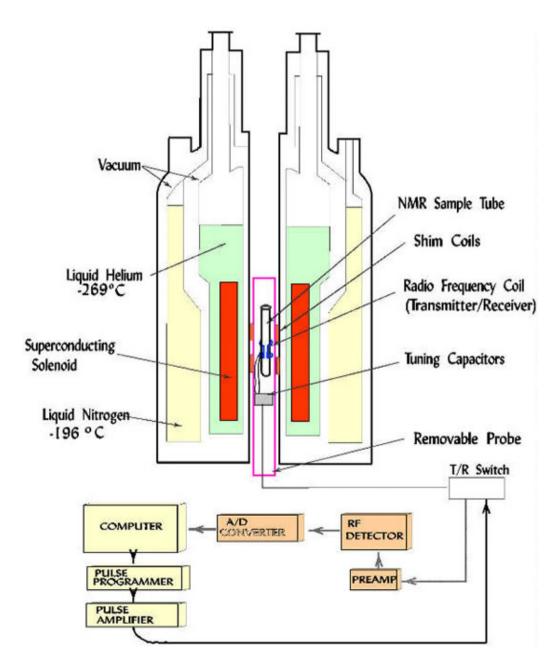
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NMR spectrometer

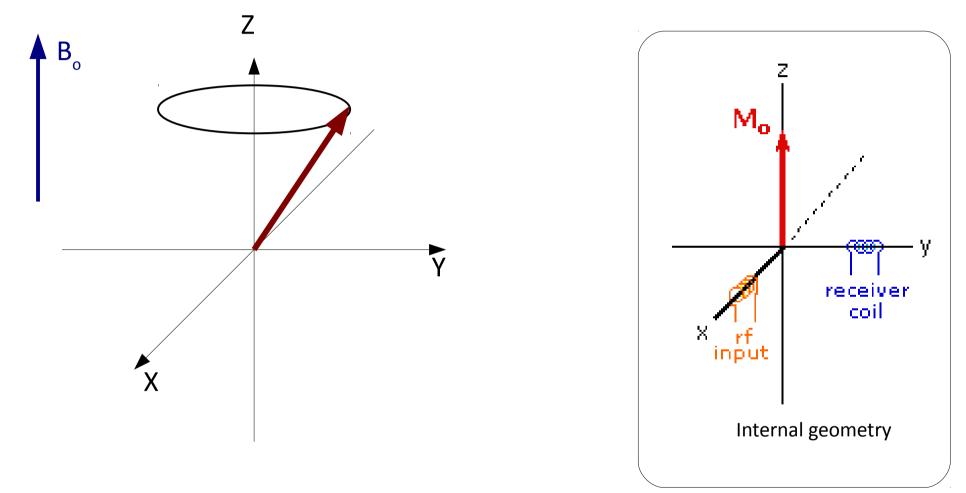


Typical fields~14 TResonance~600 MHzCost~800 k\$



Classical description of NMR

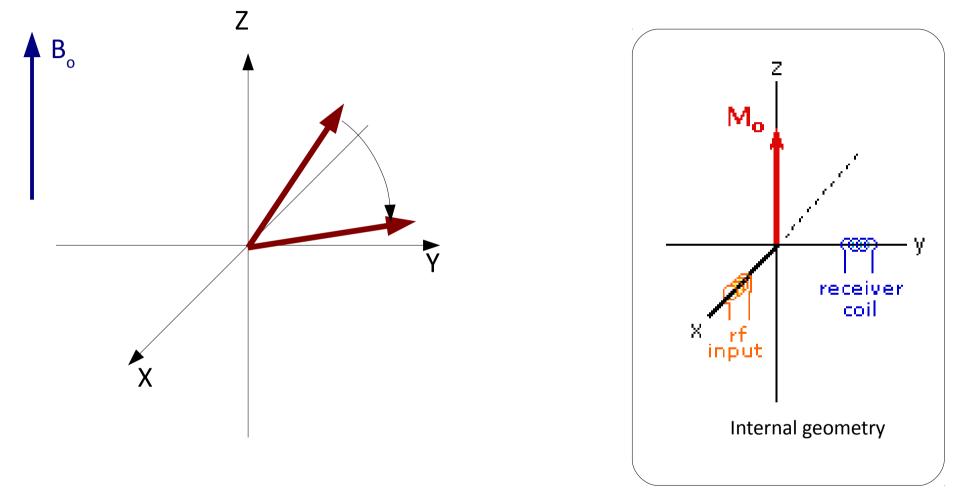
Spin in a magnetic field B₀ precesses forever around the magnetic field



On average, the magnetization lies only along the z-direction. In order to observe NMR, we must perturb the system!

Classical description of NMR

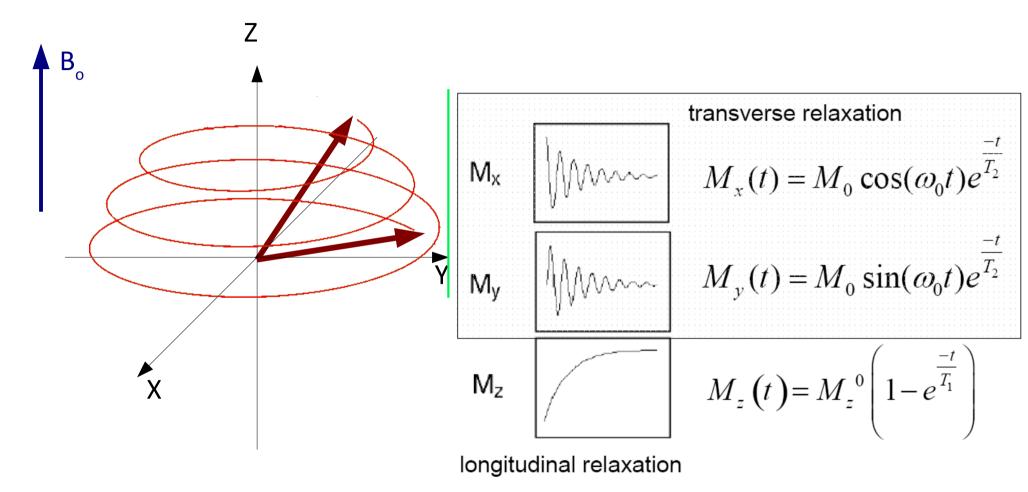
We send a RF pulse to by 90° the precession axis ...



... and we start recording the spin relaxing back to the initial state.

Classical description of NMR

... and we start recording the spin relaxing back to the initial state.

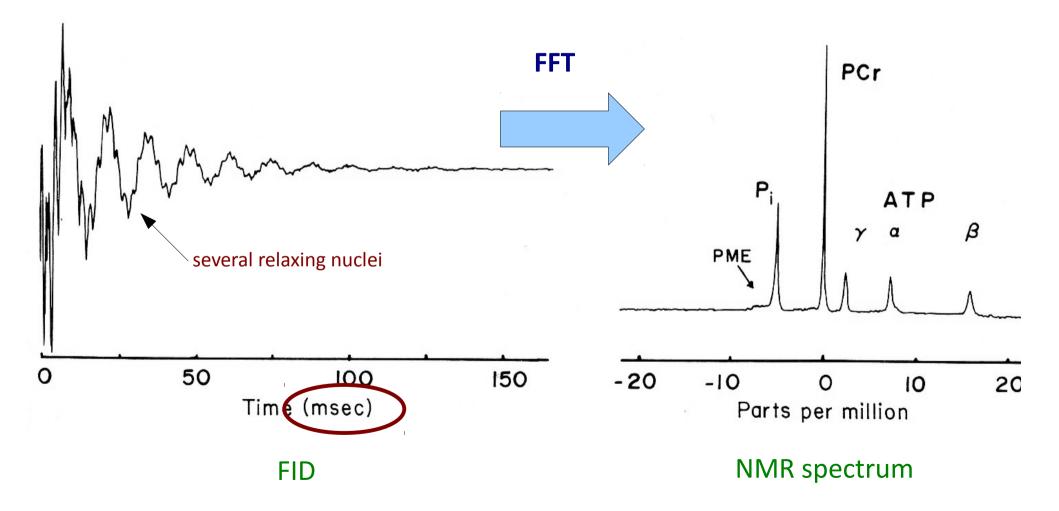


 $T_1 =$ longitudinal (spin-lattice) relaxation time $T_2 =$ transverse (spin-spin) relaxation time

Free Induction Decay

Signal recorded after the pulse is called Free Induction Decay (FID)

The Fourier transform of the FID yields the NMR spectrum.

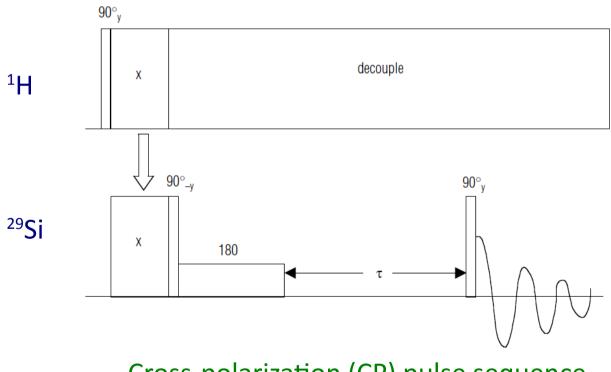


Pulse sequences

The NMR spectrometer is a programmable machine!

Pulse sequences have been designed to:

- increase resolution and sensitivity, decouple spins
- find correlations between neighboring nuclei (2D and 3D NMR)



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Chemical shift

- Different nuclei/isotopes resonate at different Larmor frequencies.
- The resonance frequency is modified by the chemical environment.
- Every nucleus experiences a different local magnetic field.

Definition: shielding tensor $\boldsymbol{\sigma}$

$$\mathbf{B}_{\rm eff} = \mathbf{B}_{\rm ext} (1 - \overset{\leftrightarrow}{\sigma})$$

Definition: chemical shift $\boldsymbol{\delta}$

$$\delta = -(\sigma - \sigma_{\rm ref})$$

- σ independent of magnetic field
- usually $\sigma \ll 1$
- measured in ppm (1 ppm = 10⁻⁶)

- δ given as deviation from a <u>reference</u> compound
- measured in ppm (1 ppm = 10⁻⁶)

Reference compounds can be liquids, solutions, solids.

Chemical shift

In practice $\boldsymbol{\delta}$ is obtained from the resonance frequencies:

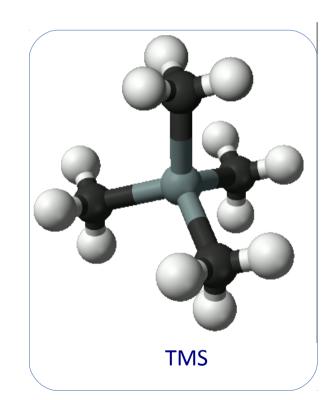
$$\delta = \frac{\omega - \omega_{\rm ref}}{\omega_{\rm ref}} \cdot 10^6$$

Some reference compounds:

- ¹H, ¹³C, ²⁹Si \rightarrow tetramethylsilane (TMS)
- ¹⁵N \rightarrow liquid NH₃
- ${}^{17}\text{O} \rightarrow \text{liquid H}_{2}\text{O}$
- ¹⁹F \rightarrow liquid CFCl₃
- ${}^{27}\text{Al} \rightarrow \text{AlCl}_3 \text{ in } \text{D}_2\text{O}$
- ${}^{43}Ca \rightarrow CaCl_2(aq) 1 mol/L$

It is possible to use a secondary reference if you know its chemical shift from the primary reference.

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Chemical shift

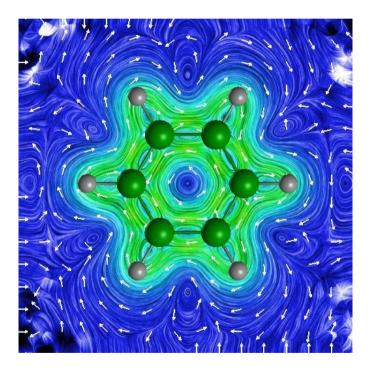
The origin of the chemical shift are the orbital currents induced by the external magnetic field.

The shielding s can be decomposed into a diamagnetic $(\sigma_{d} > 0)$ and paramagnetic $(\sigma_{p} < 0)$ term:

 σ_d depends on the density of circulating electrons increases with electronegativity of substituents

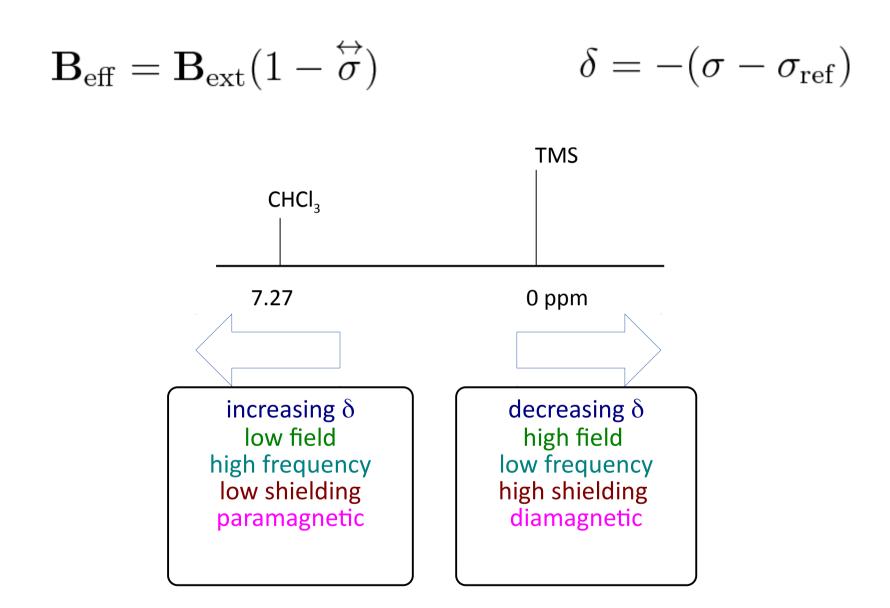
 σ_n depends on the hindering free circulation

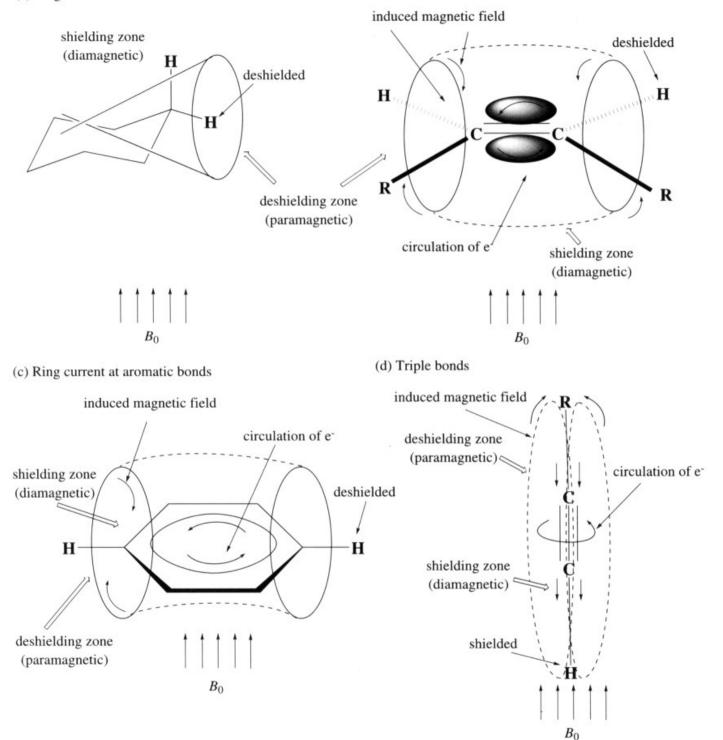
dominates in presence of p and d orbitals, anysotropic chemical bond, positive charges, low lying electronic states, high atomic number elements



Electronic current in benzene, induced by a perpendicular magnetic field

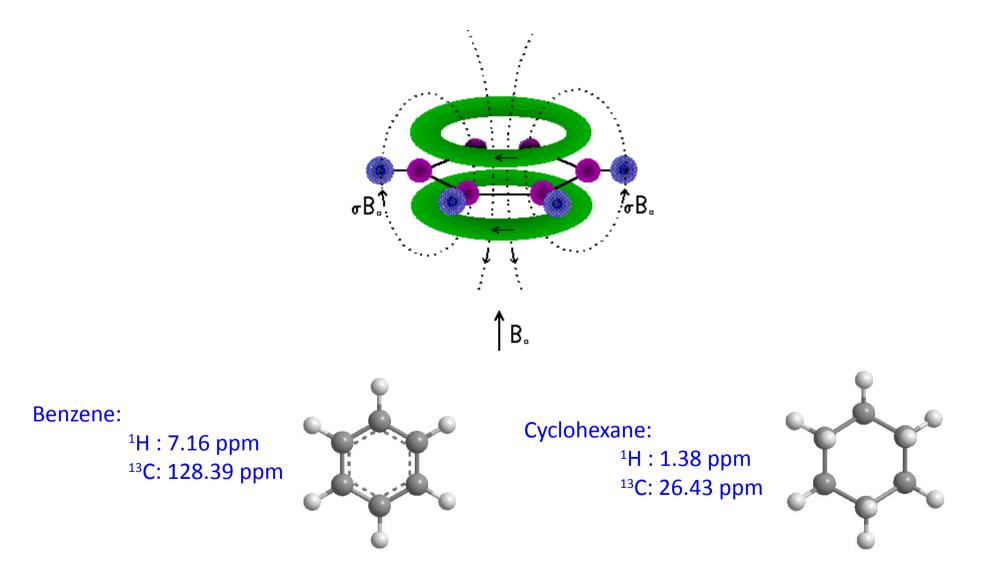
NMR terminology



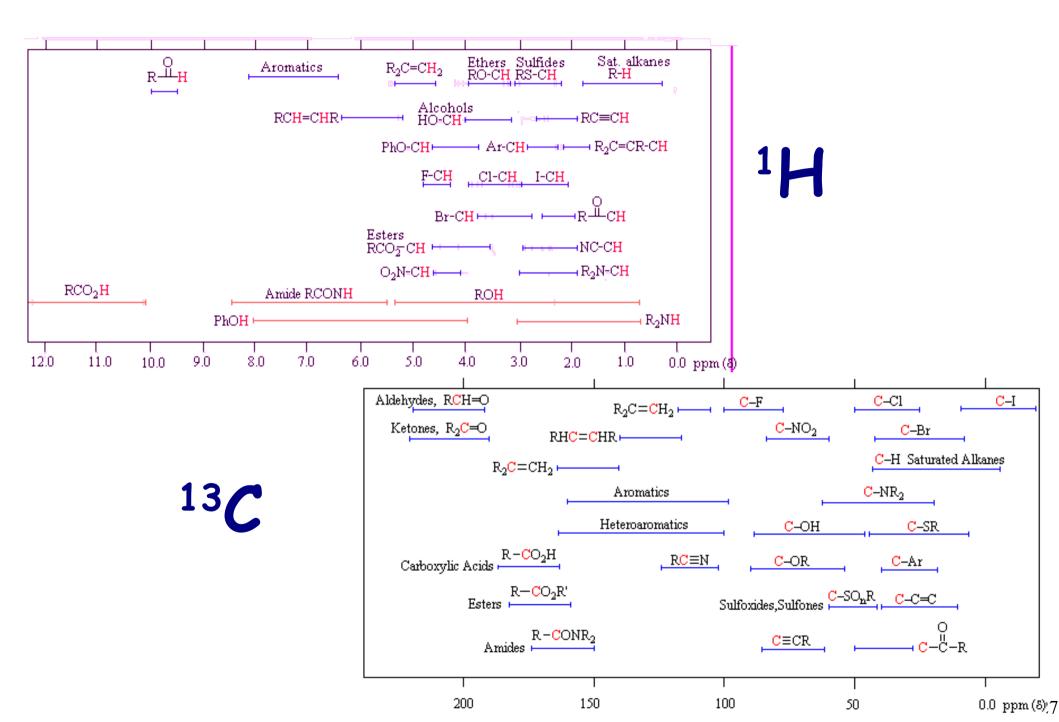


Ring current effect

Atoms on the equatorial plane are strongly de-shielded (higher δ).

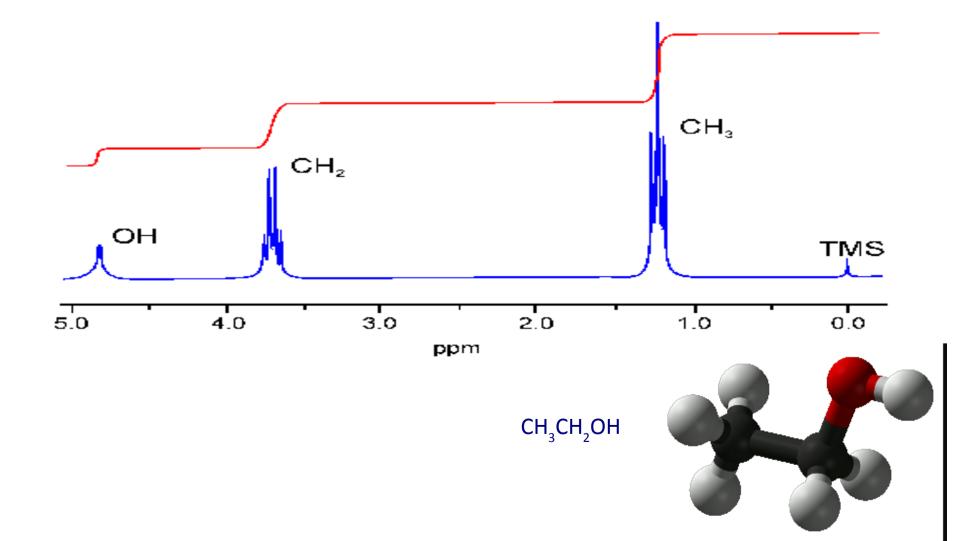


Chemical shift tables



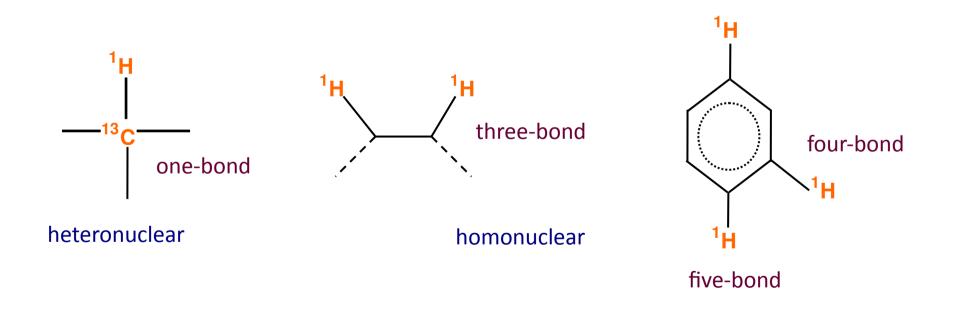
Interpretation of NMR spectra

Area of the peaks (red curve) \rightarrow number of equivalent nuclei Chemical shift table \rightarrow possible chemical group Splitting (J-coupling) \rightarrow connectivity between groups



J coupling

Equivalent nuclei are coupled Heisenberg-like: $-J S_i \cdot S_j$ The "exchange" coupling J decays as $1/r_{ij}^3$ J is independent of the magnetic field (higher B \rightarrow better resolution) Coupling can be homonuclear or heteronuclear



Homonuclear J coupling

Splitting intensities follow Pascal's triangle Example: $-CH_3$ in ethanol (1:2:1 triplet) 1 1 2 3 3 1 1 14641 15101051 ppm 1615201561 172135352171 -**I** 3 Pascal's triangle

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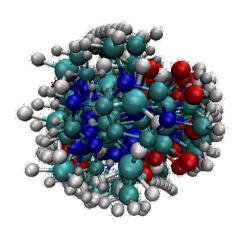
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Solid-state NMR

Molecules in liquids move very fast w.r.t. NMR time, averaging over all orientations: \rightarrow sharp NMR lines (isotropic chemical shift)

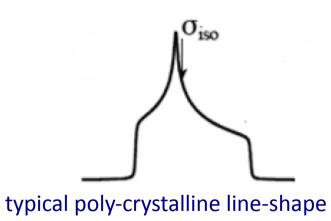
In solids nuclei are fixed in the lattice, no averaging:

- \rightarrow broad NMR peaks (anysotropy)
- \rightarrow interactions between spins are not averaged-out (broader lines)
- \rightarrow long T₁ relaxation time (which implies longer experiments)



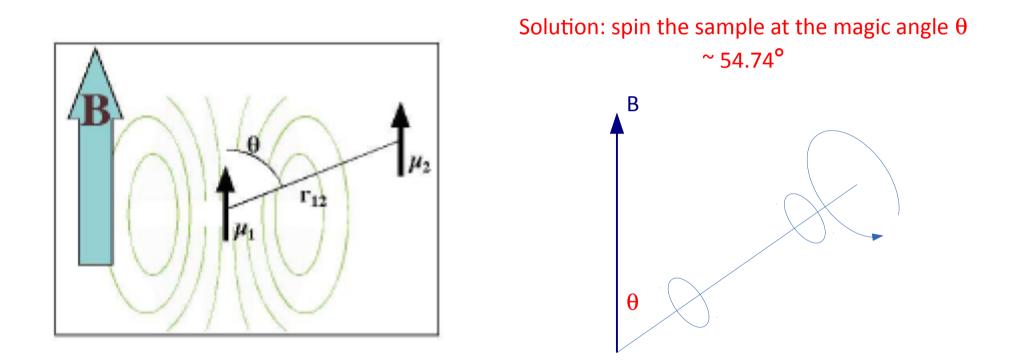
liquid state: rapid molecular tumbling

polycrystalline solid: all possible orientations



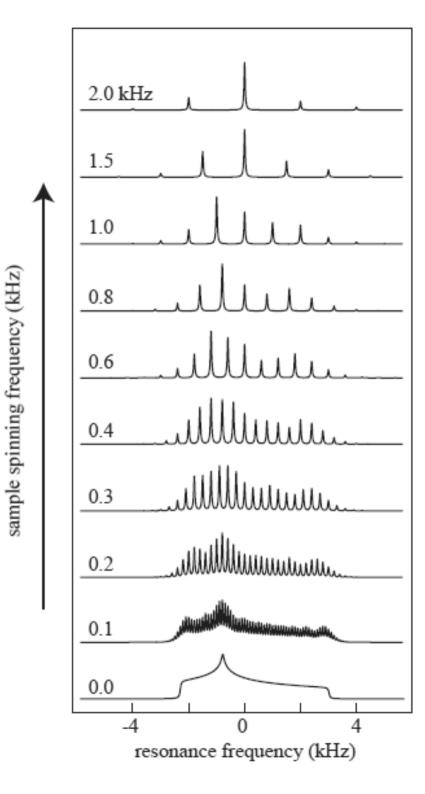
Magic Angle Spinning

The magnetic dipole interaction goes like $(3\cos^2 \theta - 1)$, where θ is the angle with the magnetic field



MAS reduces effect of chemical shift anisotropy and dipolar interactions.

Magic Angle Spinning



ss-NMR:

- sophisticated pulse sequences
- peak assignment more difficult
- need for calculations and simulations

Summary

- NMR uses radio-frequency radiation to induce transitions between different nuclear spin states of samples in a magnetic field.
- NMR can be used for quantitative measurements, but it is most useful for determining the structure of molecules.
- Different atoms in a molecule experience slightly different magnetic fields and therefore resonate at different frequencies (chemical shift).
- Splittings of the spectra lines (J-coupling) provide information about the proximity of different atoms in a molecule.

Advanced NMR topics (not covered in these notes):

- multidimensional NMR
- relaxation time, nuclear Overhauser effect

Questions?



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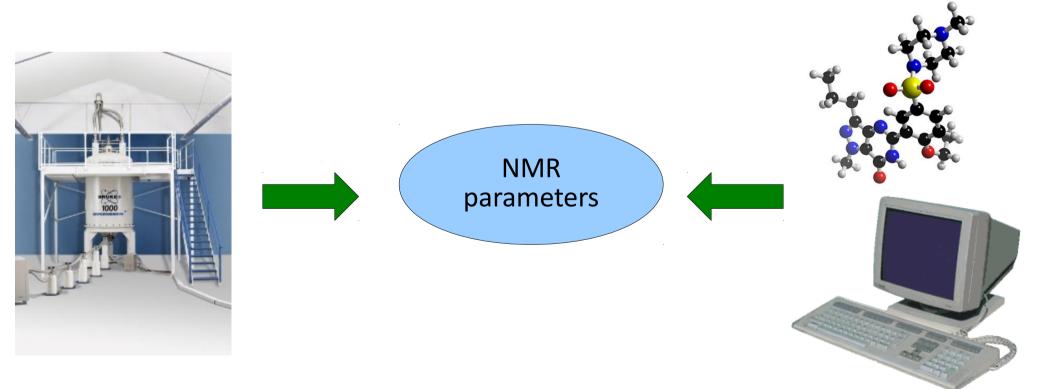
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NMR parameters

NMR parameters are:

- \rightarrow chemical shift
- \rightarrow nuclear quadrupole
- \rightarrow J coupling

They can be extracted by fitting the experimental spectrum or can be calculated from from first-principles, given the atomistic structure.



Effective NMR Hamiltonian

$$\mathcal{H}_{S}(\text{NMR}) = -\hbar \sum_{I} \gamma_{I} \mathbf{B}_{\text{ext}} \left(\bar{\bar{\mathbf{1}}} - \bar{\bar{\sigma}}\right) \mathbf{I}_{I} \quad \stackrel{\text{chemical shift}}{=} \\ + \frac{1}{2} \hbar^{2} \sum_{I} \sum_{J \neq I} \gamma_{I} \gamma_{J} \mathbf{I}_{I} \left(\bar{\bar{\mathbf{D}}}_{IJ} + \bar{\bar{\mathbf{J}}}_{IJ}\right) \mathbf{I}_{J} \\ + \sum_{I, |\mathbf{I}_{I}| \geq 1} \mathbf{I}_{I} \bar{\bar{\mathbf{Q}}}_{I} \mathbf{I}_{I} \quad \stackrel{\text{nuclear}}{\stackrel{\text{quadrupole}}{\stackrel{(\mathbb{I} > 1/2)}{=}}$$

$$\begin{split} \mathbf{I}_{I} &= \text{Nuclear spin}; \ \mu_{I} = \gamma_{I} \hbar \mathbf{I}_{I} \ ; \ \sigma = \text{Nuclear chemical shielding tensor} \\ & \bar{\bar{\mathbf{D}}}_{IJ} = \text{Nuclear magnetic dipolar coupling tensor} \\ & \bar{\bar{\mathbf{J}}}_{IJ} = \text{Indirect nuclear spin} - \text{spin coupling tensor} \\ & \bar{\bar{\mathbf{Q}}}_{II} = \text{Nuclear quarupolar coupling tensor} \end{split}$$

The chemical shift

From the NMR Hamiltonian the shielding tensor is defined as:

$$\mathbf{B}_{ ext{ind}} = -ar{ar{\sigma}} \mathbf{B}_{ ext{ext}}$$

It is the second derivative of energy w.r.t. field and nuclear moment:

$$\overset{\leftrightarrow}{\sigma}_{I} = \frac{\partial^{2} E}{\partial \boldsymbol{\mu}_{I} \partial \mathbf{B}}$$

The chemical shift is then defined by:

$$\delta = - \left(\sigma - \sigma_{
m ref}
ight)$$

 $\sigma_{_{\!ref}}$ is a reference value in a well-characterized material.

"Direct" and "converse" methods

Direct approach (traditional):

- linear response to external magnetic field
- calculate the induced current, then the induced field
- Mauri, Louie (1996); GIPAW: Pickard, Mauri (2001)

$$\left. \stackrel{\leftrightarrow}{\sigma}_{I} = - \left. \frac{\partial \mathbf{B}_{\text{ind}}}{\partial \mathbf{B}_{\text{ext}}} \right|_{r=r_{I}} \right.$$

$$\mathbf{B}_{\text{ind}}(\mathbf{r}) = \frac{1}{c} \int d^3 \mathbf{r}' \, \frac{\mathbf{j}_{\text{ind}}(\mathbf{r}') \times (\mathbf{r} - \mathbf{r}')}{|\mathbf{r} - \mathbf{r}'|^3}$$

Converse approach:

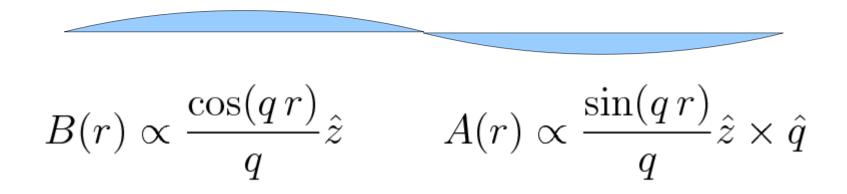
- no linear response, no magnetic field, no gauge-origin problem
- calculate the change of orbital magnetization due to nuclear magnetic moment
- based on the "Modern Theory of the Orbital Magnetization"

$$\overset{\leftrightarrow}{\sigma}_{I} = -\Omega \frac{\partial \mathbf{M}_{\text{orb}}}{\partial \boldsymbol{\mu}_{I}}$$

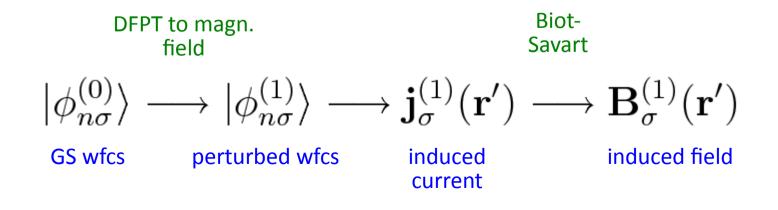
$$\mathbf{M}_{\rm orb} = \frac{\alpha}{2} \operatorname{Im} \sum_{n\mathbf{k}} f_{n\mathbf{k}} \left\langle \partial_{\mathbf{k}} u_{n\mathbf{k}} \right| \times \left(H_{\mathbf{k}} + E_{n\mathbf{k}} - 2\mu \right) \left| \partial_{\mathbf{k}} u_{n\mathbf{k}} \right\rangle$$

Outline of linear response

Vector potential $A(r)=(1/2)B \times r$ is incompatible with crystal periodicity Solution: apply a long-wavelength magnetic field (q << 1)



The response to an incommensurate perturbation is obtained by Density Functional Perturbation Theory (DFPT):



EFG: electric field gradient tensor

Quadrupolar nuclei (I>1/2); non-zero only when no cubic symmetry:

$$\bar{\bar{\mathbf{Q}}}_{\alpha\beta}^{I} = \frac{eQ}{h} \left\langle \psi_{0} \right| \frac{\delta \mathbf{E}_{\alpha}}{\delta \mathbf{r}_{\beta}} \left| \psi_{0} \right\rangle = \frac{eQ}{h} \bar{\bar{\mathbf{V}}}_{\alpha\beta}$$

 $\mathbf{E} = ext{electric field}$; $eQ = ext{electric quadrupole moment}$

Principal axis system: Eigenvectors and -values of Convention: $|V_{zz}| \ge |V_{yy}| \ge |V_{xx}|$

Observables:

- Quadrupolar coupling constant
- Asymmetry parameter

$$Cq = \frac{eQV_{zz}}{h}$$
$$\eta = \frac{V_{xx} - V_{yy}}{V_{zz}}$$

 $ar{ar{V}}_{lphaeta}$

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GIPAW

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All-electron magnetic response with pseudopotentials: NMR chemical shifts

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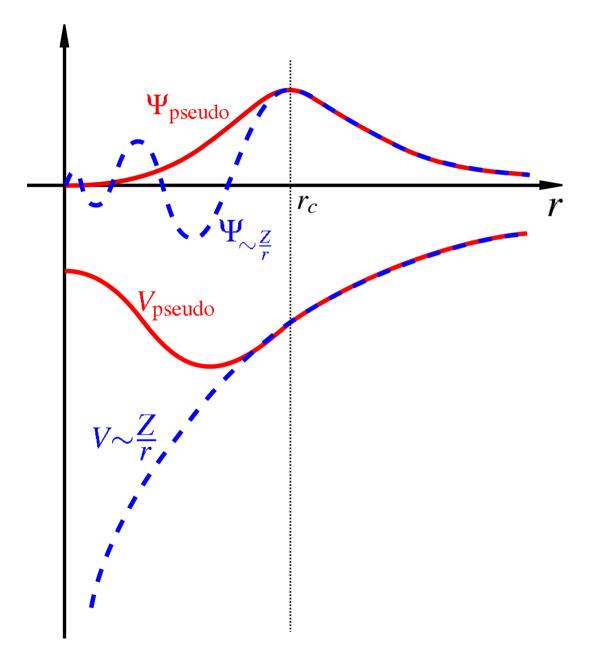
A theory for the *ab initio* calculation of all-electron NMR chemical shifts in insulators using pseudopotentials is presented. It is formulated for both finite and infinitely periodic systems and is based on an extension to the projector augmented-wave approach of Blöchl [P. E. Blöchl, Phys. Rev. B **50**, 17 953 (1994)] and the method of Mauri *et al.* [F. Mauri, B. G. Pfrommer, and S. G. Louie, Phys. Rev. Lett. **77**, 5300 (1996)]. The theory is successfully validated for molecules by comparison with a selection of quantum chemical results, and in periodic systems by comparison with plane-wave all-electron results for diamond.

DOI: 10.1103/PhysRevB.63.245101

PACS number(s): 71.45.Gm, 76.60.Cq, 71.15.-m

Need to reconstruct the wavefunction near the nuclei!

Pseudopotential / all-electron

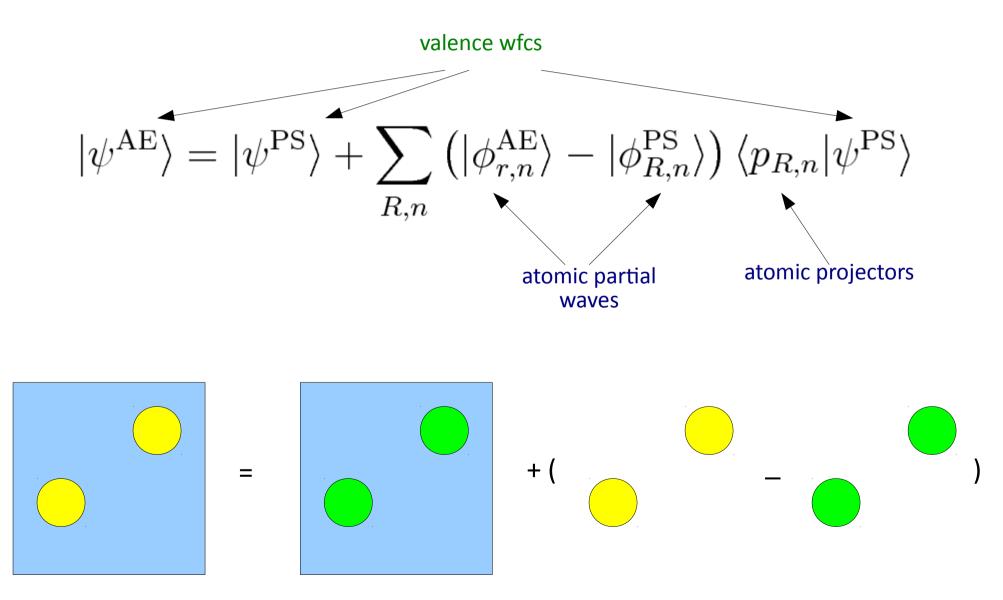


AE wfc (blue) oscillates rapidly near the nucleus

PS wfc (red) smooth, no nodes in the core region

PAW idea

PAW (Blöchl, 1994) = projector augmented wave Idea: "reconstruct" the AE wfc from the PS wfc



Gauge Including PAW

Translation in magnetic field yields a gauge phase factor:

$$H' = \frac{1}{2} \left[\mathbf{p} + \frac{1}{c} \mathbf{A} \left(\mathbf{r} \right) \right]^2 + V \left(\mathbf{r} - \mathbf{t} \right)$$
$$\psi'_i \left(\mathbf{r} \right) = e^{i/(2c)\mathbf{r} \cdot \mathbf{t} \times \mathbf{B}} \psi_i \left(\mathbf{r} \right)$$

PAW + magnetic field = GIPAW

$$\mathcal{T}_{\mathbf{B}} = \mathbf{1} + \sum_{\mathbf{R},n} e^{i/(2c)\mathbf{r} \cdot \mathbf{R} \times \mathbf{B}} \left[\left| \phi_{\mathbf{R},n} \right\rangle - \left| \tilde{\phi}_{\mathbf{R},n} \right\rangle \right] \left| \tilde{p}_{\mathbf{R},n} \right\rangle e^{-i/(2c)\mathbf{r} \cdot \mathbf{R} \times \mathbf{B}}$$

Take-home message: (1) wfc reconstruction (2) gauge-invariance

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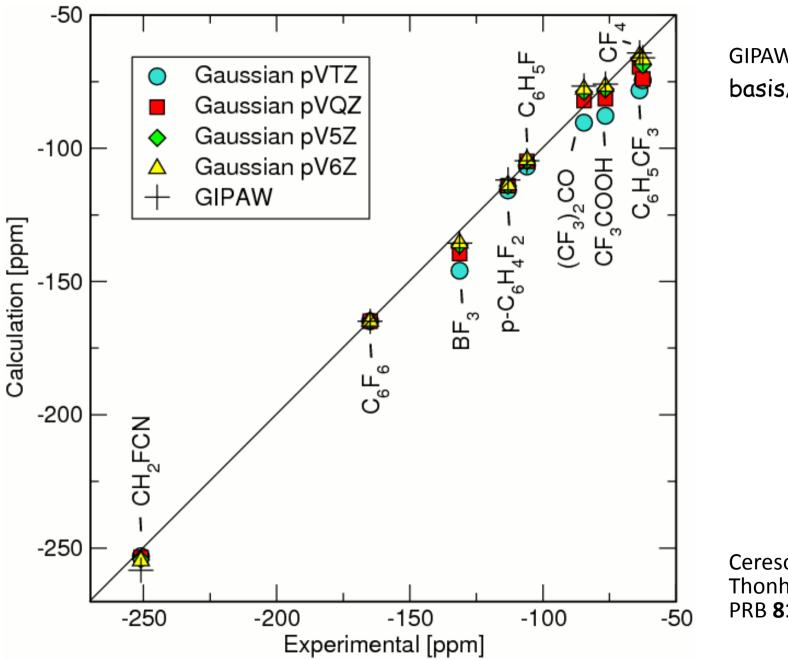
Example: molecules

Molecule			$\sigma_{\rm GIPAW}$			σ_{IGAIM}
	Core	bare	Δd	Δp	Total	Total
H atom						
CH ₄	0.00	30.47	0.40	0.00	30.87	30.99
CH₃F	0.00	25.71	0.41	0.00	26.13	26.50
C ₆ H ₆	0.00	22.33	0.41	0.00	22.74	23.25
TMS	0.00	30.41	0.40	0.00	30.80	31.02
SiH₃F	0.00	24.92	0.38	0.00	25.30	25.13
Si_2H_4	0.00	24.53	0.36	0.00	24.90	24.78
SiH ₄	0.00	26.96	0.37	0.00	27.33	27.28
C atom	15					
CO	198.88	- 126.25	4.59	- 100.15	-22.93	-21.16
CH ₄	198.88	16.86	3.97	-28.76	190.96	191.22
CH₃F	198.88	- 49.64	3.93	- 54.70	98.47	99.66
CH ₃ NH ₂	198.88	-13.98	3.91	- 39.05	149.77	150.44
C ₆ H ₆	198.88	- 89.51	4.07	-77.32	36.12	39.52
CF ₄	198.88	-92.12	3.51	-76.05	34.22	35.29
TMS	198.88	9.12	3.97	-32.65	179.33	182.08
Si atom	1s2s2p					
SiF ₄	832.39	- 19.43	5.28	-408.26	409.97	409.69
SiH₃F	832.39	- 19.50	5.70	- 510.30	308.29	305.45
Si ₂ H ₄	832.39	-9.04	5.80	- 622.45	206.70	202.99
SiH ₄	832.39	-0.21	5.98	-410.20	427.97	424.37
TMS	832.39	-17.39	5.70	- 518.00	302.70	304.39
P atom	1s2s2p					
PF ₃	902.47	-32.94	6.08	- 697.61	178.00	172.52
P ₂	902.47	-33.84	7.58	- 1236.95	- 360.75	- 375.45
P ₄	902.47	49.84	7.42	- 126.79	832.94	826.62

red = GIPAW blue = Gaussian[™]

Pickard, Mauri PRB **63**, 245101 (2001)

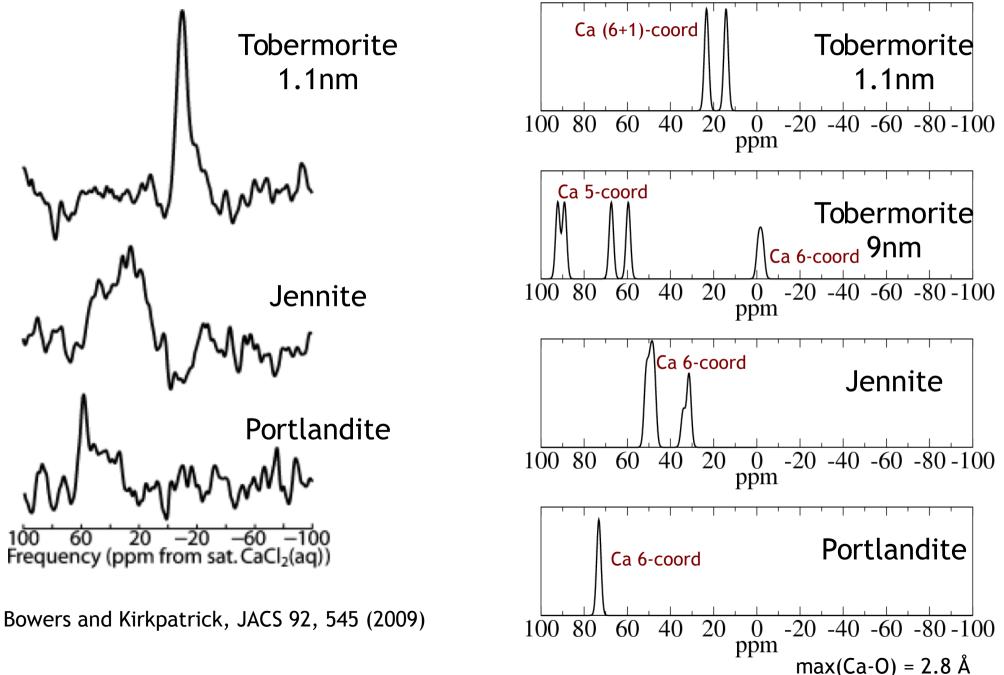
Example: ¹⁹F



GIPAW ~ Gaussian 6Z (140 basis/atom)

Ceresoli, Marzari, Lopez, Thonhauser PRB **81**, 184424 (2010)

Example: ⁴³Ca cement models



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Outline

• PART I:

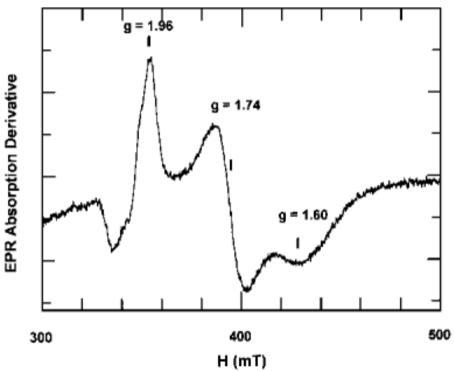
- Basic principles of magnetic resonance spectroscopy
- Introduction to experimental NMR
- Interpretation of NMR spectra
- Solid state NMR
- PART II:
 - Effective NMR spin hamiltonian
 - The GIPAW method
 - Examples
 - Brief introduction to EPR spectroscopy and EPR parameters

EPR/ESR spectroscopy

<u>Electron Paramagnetic Resonance / Electron Spin Resonance</u>

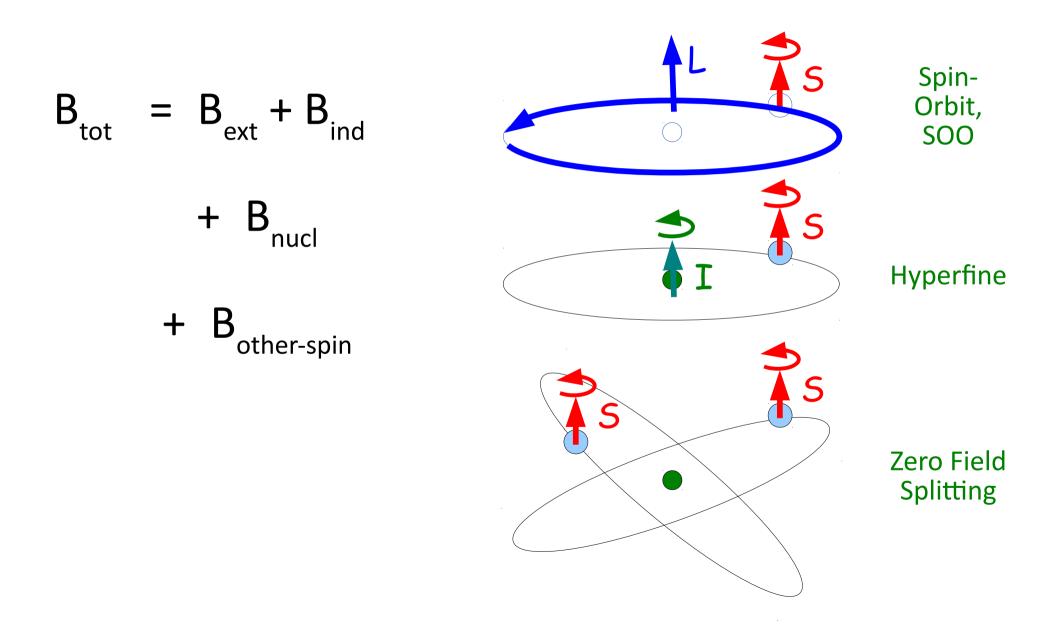


Tipical fields ~0.5 T Resonance ~14 GHz

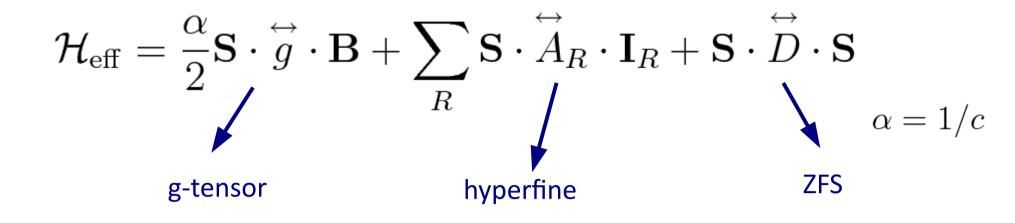


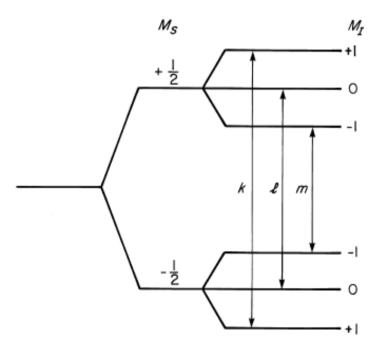
- Paramagnetic defects in solids, radicals in proteins
- Sensitive to local geometry and electronic structure (charge, hybridization)
- Non destructive, small samples, dilute spins

Interpretation of EPR spectra



Effective Spin Hamiltonian





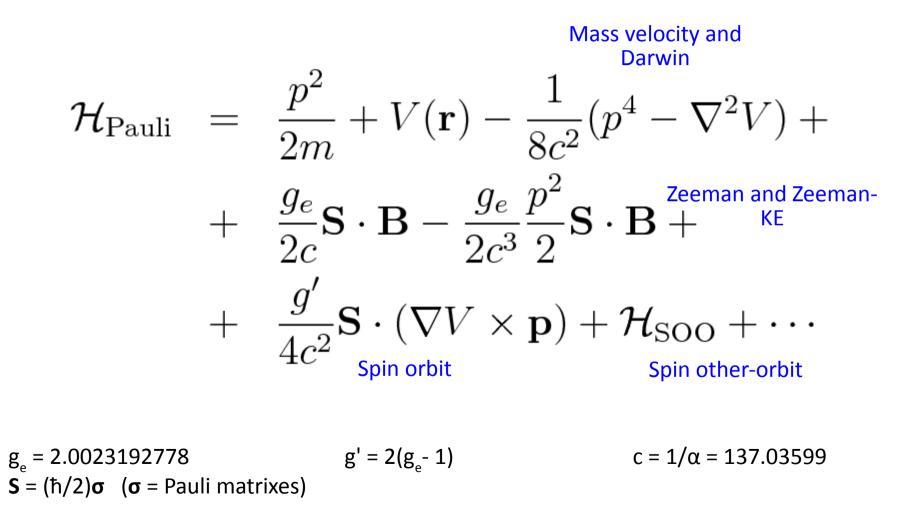
Selection rules:

$$|\Delta M_{s}| = 1$$
$$|\Delta M_{l}| = 0$$

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Theory of EPR parameters

The effective spin Hamiltonian allows to extract EPR parameters from the experiments, but in order to calculate EPR parameters, we need a "physical" Hamiltonian. The simplest non-relativistic Hamiltonian that accounts for spin is the Pauli Hamiltonian:



... plus other terms

Label	Description	Operator
	Scalar relativistic corrections	
h^{mv}	Mass-velocity term	$-\frac{1}{8}\alpha^2 \sum_i \nabla_i^4$
$h^{\operatorname{Dar}(1)}$	Electron-nuclear Darwin term	$\frac{\pi}{2} \alpha^2 \sum_{i,K} Z_K \delta(\mathbf{r}_{iK})$
$h^{\operatorname{Dar}(2)}$	Electron-electron Darwin term	$-\frac{\pi}{2} \alpha^2 \sum_{i,j}^{\prime} \delta(\mathbf{r}_{ij})$
h^{OO}	Electron-electron orbital interaction	$\frac{\frac{2}{-\frac{\pi}{2}}\alpha^{2}\sum_{i,j}^{\prime}\delta(\mathbf{r}_{ij})}{\frac{1}{4}\alpha^{2}\sum_{i,j}^{\prime}\frac{\nabla_{i}\cdot\nabla_{j}}{r_{ij}}-\frac{(\nabla_{i}\cdot\mathbf{r}_{ij})(\mathbf{r}_{ij}\cdot\nabla_{j})}{r_{ij}^{3}}}$
	Spin-dependence, no field-dependence Electron spin	
$h^{\mathrm{SO}(1)}$	Spin-orbit interaction	$\frac{1}{4} \alpha^2 g_e \sum_{i,K} \frac{Z_K}{r_{iK}^3} \mathbf{s}_i \cdot \mathbf{I}_{iK}$
$h^{\mathrm{SSO}(2)}$	Electron-electron spin-orbit interaction	$-\frac{1}{4}\alpha^2 g_e \sum_{i,j} s_i \cdot \frac{\mathbf{I}_{ij}}{r_{ij}^3}$
$h^{\mathrm{SOO}(2)}$	Electron-electron spin-other-orbit interaction	$-\frac{1}{2}\alpha^2 g_e \sum_{i,j}' \mathbf{s}_j \cdot \frac{\mathbf{l}_{ij}'}{r^3}$
$h^{\mathrm{SSD}(2)}$	Spin-spin dipolar interaction	$\frac{1}{4} \alpha^2 g_e^2 \sum_{i,j}' \frac{r_{ij}^2 (\mathbf{s}_i \cdot \mathbf{s}_j) - (\mathbf{s}_i \cdot \mathbf{r}_{ij}) (\mathbf{r}_{ij} \cdot \mathbf{s}_j)}{r_{ij}^5}$
$h^{\mathrm{SSC}(2)}$	Spin-spin contact interaction	$-\frac{\pi g_e^2}{3}\alpha^2 \sum_{i,j}' \mathbf{s}_i \cdot \mathbf{s}_j \delta(\mathbf{r}_{ij})$
$h_K^{ m PSO}$	Orbital hyperfine interaction	$\alpha^2 \gamma_K \sum_i \mathbf{I}_K \cdot \frac{\mathbf{I}_{iK}}{r_{iK}^2}$
	Electron and nuclear spins	16
$h_K^{ m SD}$	Dipolar hyperfine interaction	$\frac{1}{2}\alpha^2 g_e \gamma_K \sum_i \mathbf{s}_i \cdot \frac{3\mathbf{r}_{iK}\mathbf{r}_{iK} - 1r_{iK}^2}{r_{iK}^5} \cdot \mathbf{I}_K$
$h_K^{ m FC}$	Fermi contact hyperfine interaction	$\frac{4\pi}{3}\alpha^2 g_e \gamma_K \sum_i \delta(\mathbf{r}_{iK}) \mathbf{s}_i^{\prime \kappa} \mathbf{I}_K$

From: Pekka Manninen PhD thesis, University of Oulo, Finland (2004). http://herkules.oulu.fi/isbn9514274318/

Zero Field Splitting

In case: S > 1/2

$$\mathcal{H}_{\text{ZFS}} = \frac{g_e \,\mu_B}{r^5} \left[\left(\mathbf{s}_1 \cdot \mathbf{s}_2 \right) r^2 - 3 \left(\mathbf{s}_1 \cdot \mathbf{r} \right) \left(\mathbf{s}_2 \cdot \mathbf{r} \right) \right]$$

- Two electron integrals
- Usually small (?)
- Not implemented in Espresso

 $\mathbf{r} \equiv \mathbf{r}_1 - \mathbf{r}_2$

g-tensor: the easy part

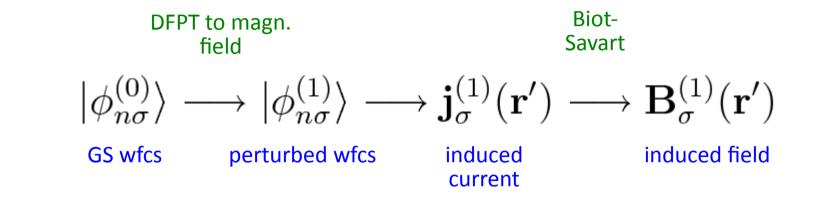
From the effective spin Hamiltonian, the g-tensor is defined as:

$$\stackrel{\leftrightarrow}{g} = \frac{2}{\alpha} \frac{\partial^2 \mathcal{H}}{\partial \mathbf{S} \partial \mathbf{B}}$$

The Zeeman and Zeeman-Kinetic Energy term yield:

g-tensor: SO and SOO

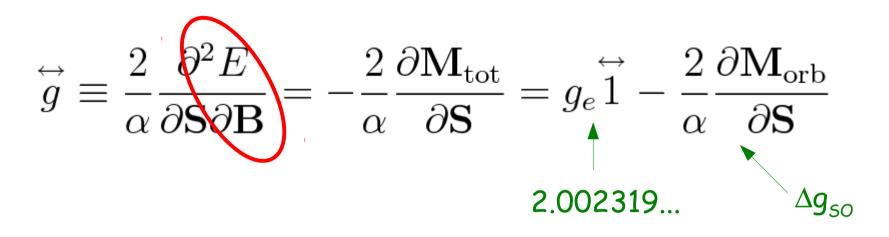
The SO and SOO don't depend explicitly on the magnetic field, but implicitly through the wavefunctions. Their contribution can be calculated in perturbation theory:



Finally:

$$\Delta \overset{\leftrightarrow}{g}_{SO} = \alpha^2 g' \sum_{S=\pm 1/2} S \int d^3 \mathbf{r}' \nabla V(\mathbf{r}') \times \overset{\leftrightarrow}{j}_{S}^{(1)}(\mathbf{r}')$$
$$\Delta \overset{\leftrightarrow}{g}_{SOO} \simeq \alpha \sum_{S=\pm 1/2} S \int d^3 \mathbf{r}' \overset{\leftrightarrow}{B}_{S}^{(1)}(\mathbf{r}') \rho_S(\mathbf{r}')$$

Converse approach to the g-tensor

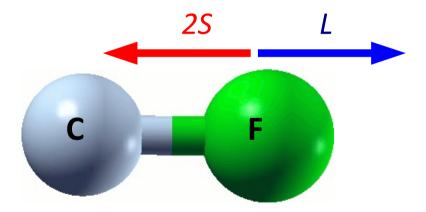


- "Converse" method: $\partial M_{orb} / \partial S \approx 1/(2S) [M_{orb}(S=\uparrow) M_{orb}(S=\downarrow)]$
- Modern Theory of the Orbital Magnetization
- GS calc. including Spin-Orbit; no magnetic field, no linear response
- Implemented in an experimental version of Espresso (ask me)

EPR of diatomic radicals

Molecule		Expt.	LR	This work	
CN·	g_{\parallel}	2.0023	2.0022	2.0022	L = 0
	g_{\perp}	2.0003	2.0002	2.0002	2 0
CF·	g_{\parallel}	0.0000	N/A	0.0002	L = 1
	g_\perp	2.0002	2.0003	2.0003	

LR = <u>linear response</u> This work = <u>converse method</u>



Important when g very different from 2 (i.e. transition metals impurities)!

Hyperfine coupling

$$\overset{\leftrightarrow}{A}_{R} = \left\langle \frac{\partial^{2} \mathcal{H}}{\partial \mathbf{S} \, \partial \mathbf{I}_{R}} \right\rangle = A_{\mathrm{iso,R}} \overset{\leftrightarrow}{1} + \overset{\leftrightarrow}{A}_{\mathrm{dip,R}}$$

Isotropic (Fermi-contact)

$$A_{\rm iso} = \frac{4\pi}{3} \frac{g_e \,\mu_e \,g_N \,\mu_N}{S} \int d^3 \mathbf{r} \, n_s(\mathbf{r}) \underline{\delta(\mathbf{r})}$$
$$\mathbf{r} \equiv \mathbf{r} - \mathbf{R}$$

Dipolar (traceless)

$$A_{\rm dip}^{\alpha\beta} = \frac{1}{2} \frac{g_e \,\mu_e \,g_N \,\mu_N}{S} \int d^3 \mathbf{r} \, n_s(\mathbf{r}) \frac{3r_\alpha r_\beta - \delta_{\alpha\beta} r^2}{r^5}$$

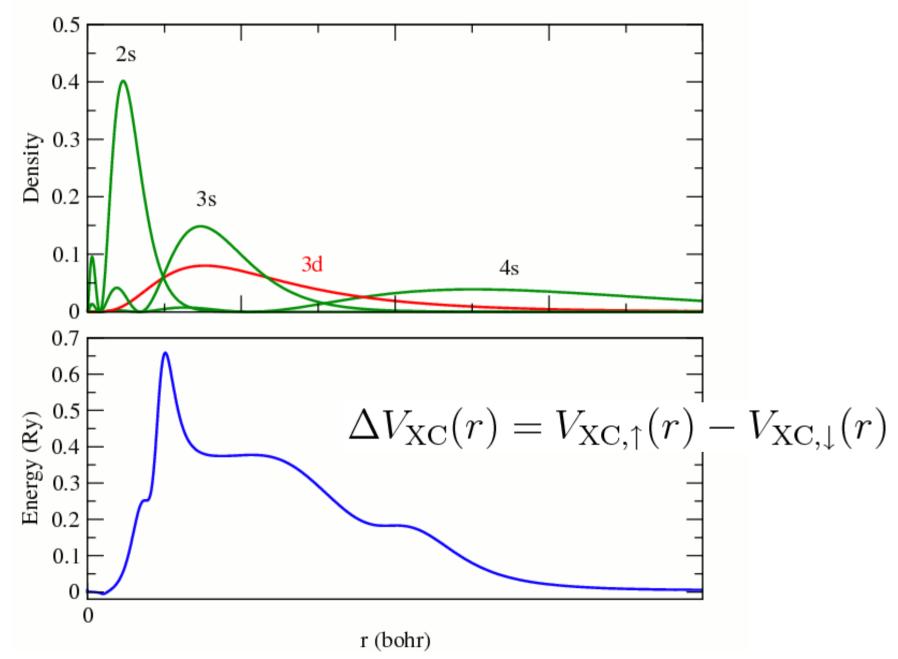
Spin density:
$$n_s({f r})=n_{\uparrow}({f r})-n_{\downarrow}({f r})$$

Spin densities of isolated atoms

Configuration	1 5	2s	3 <i>s</i>	4 5	Total
2s(1↑,0↓)	0.003	0.219			0.222
2s² 2p(2↑,0↓)	-0.199	0.178			-0.021
2s² 2p(3↑,0↓)	-0.440	0.429			-0.011
2s² 2p(3↑,1↓)	-0.417	0.437			0.019
3 <i>s</i> (1↑,0↓)	0.019	-0.030	0.810		0.799
3s² 3p(2↑,0↓)	-0.094	0.056	-0.193		-0.232
4s² 3d(5↑,0↓)	-0.008	-2.203	0.913	1.384	0.086
4s¹ 3d(5↑,0↓)	0.040	-2.215	0.833	7.194	5.852
4sº 3d(5↑,0↓)	-0.018	-2.294	0.888	0.000	-1.424
4sº 3d(4↑,0↓)	-0.018	-2.103	0.942	0.000	-1.179
4sº 3d(3↑,0↓)	-0.017	-1.793	0.904	0.000	-0.907
	$2s(1^{+},0^{+})$ $2s^{2} 2p(2^{+},0^{+})$ $2s^{2} 2p(3^{+},0^{+})$ $2s^{2} 2p(3^{+},1^{+})$ $3s(1^{+},0^{+})$ $3s^{2} 3p(2^{+},0^{+})$ $4s^{2} 3d(5^{+},0^{+})$ $4s^{0} 3d(5^{+},0^{+})$ $4s^{0} 3d(4^{+},0^{+})$	$2s(1\uparrow,0\downarrow)$ 0.003 $2s^2 2p(2\uparrow,0\downarrow)$ -0.199 $2s^2 2p(3\uparrow,0\downarrow)$ -0.440 $2s^2 2p(3\uparrow,1\downarrow)$ -0.417 $3s(1\uparrow,0\downarrow)$ 0.019 $3s^2 3p(2\uparrow,0\downarrow)$ -0.094 $4s^2 3d(5\uparrow,0\downarrow)$ -0.008 $4s^1 3d(5\uparrow,0\downarrow)$ 0.040 $4s^0 3d(5\uparrow,0\downarrow)$ -0.018 $4s^0 3d(4\uparrow,0\downarrow)$ -0.018	$2s(1\uparrow,0\downarrow)$ 0.003 0.219 $2s^2 2p(2\uparrow,0\downarrow)$ -0.199 0.178 $2s^2 2p(3\uparrow,0\downarrow)$ -0.440 0.429 $2s^2 2p(3\uparrow,1\downarrow)$ -0.417 0.437 $3s(1\uparrow,0\downarrow)$ 0.019 -0.030 $3s^2 3p(2\uparrow,0\downarrow)$ -0.094 0.056 $4s^2 3d(5\uparrow,0\downarrow)$ -0.008 -2.203 $4s^1 3d(5\uparrow,0\downarrow)$ 0.040 -2.215 $4s^0 3d(5\uparrow,0\downarrow)$ -0.018 -2.294 $4s^0 3d(4\uparrow,0\downarrow)$ -0.018 -2.103	$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$\begin{array}{cccccccccccccccccccccccccccccccccccc$

- Calculated with *ld1.x*, extrapolated at the nucleus
- Spin-polarized LDA
- values in elec./bohr³

Mn orbitals and ΔV_{xc}



Core relaxation

• Project valence density around atoms and add GIPAW reconstruction

$$n_{\sigma}(r) = n_{\operatorname{val},\sigma}(r) + n_{\operatorname{rec},\sigma}(r), \qquad r < R_c$$

- Calculate $\Delta V_{\rm xc}$ from projected spherical density

$$\Delta V_{\rm XC} = -\frac{2}{\pi} \frac{n_{\uparrow} - n_{\downarrow}}{\left(n_{\uparrow} + n_{\downarrow}\right)^{2/3}}$$

• Core spin density from Incomplete Perturbation Theory

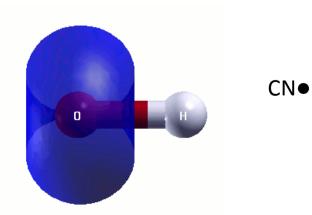
$$n_{\rm s,core}(r) = 4 \sum_{u \in \text{core}} \sum_{v \neq u} \frac{\langle \phi_{us} | \Delta V_{\rm XC} | \phi_{vs} \rangle}{\epsilon_{us} - \epsilon_{vs}} \phi_{us}(r) \phi_{vs}(r)$$

Available since GIPAW 5.4. Highly experimental!

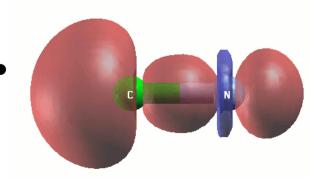
Hyperfine couplings of 2nd row radicals

Molecule	e Atom	no core relax core	relax exper	riment
CH•	С	204	90	47
	Н	-50	-50	-58
OH•	0	-142	-61	-45
	Н	-56	-56	<mark>-73</mark>
CH3•	С	210	122	107
-	Н	-69	-69	-70
CN.	С	636	581	<mark>588</mark>
	N	-7	-18	-13
H ₂ CN•	С	-73	-67	-81
_	N	70	34	26
	Н	230	230	234

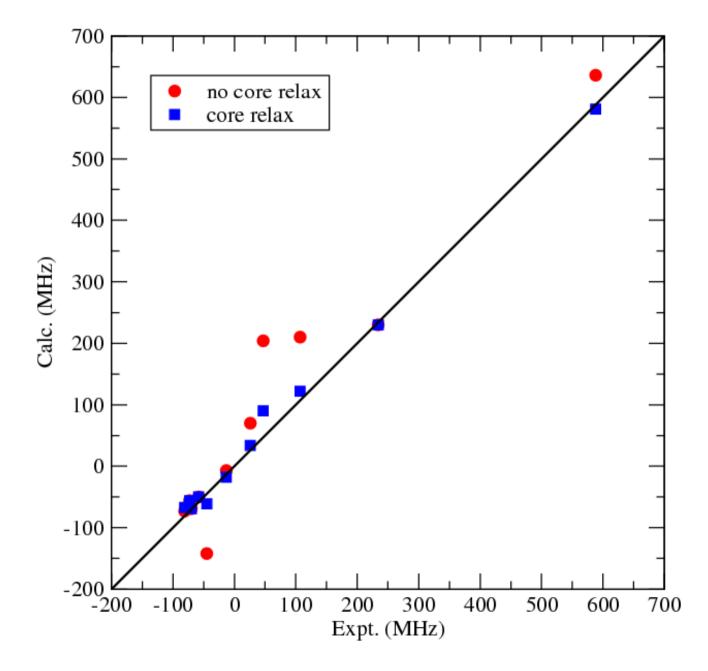
Fermi contact (A_{iso}), values in MHz



OH●

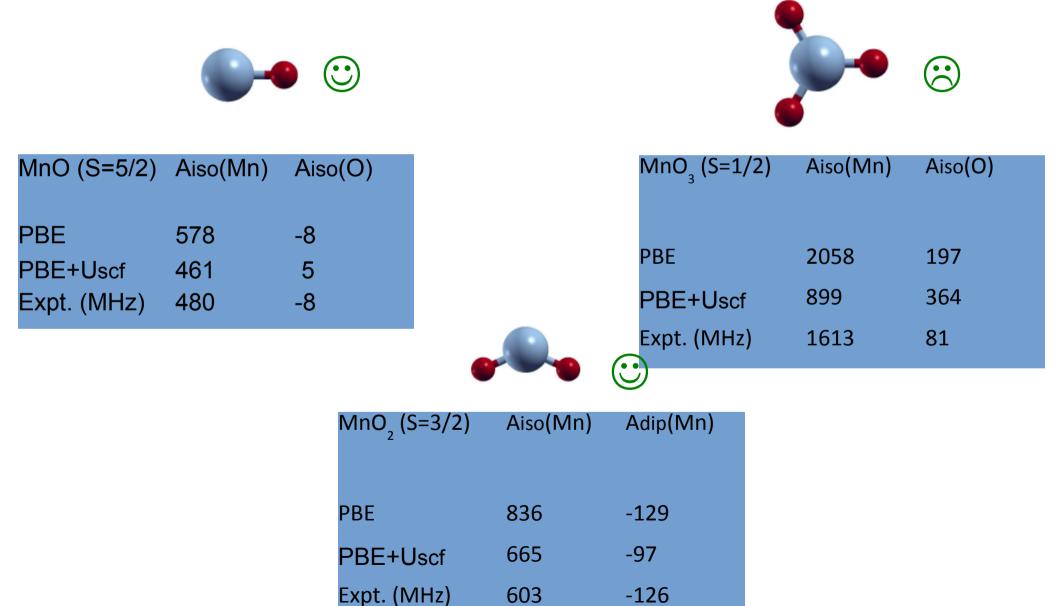


Hyperfine couplings of 2nd row radicals



"Converse" EPR: DFT+U on MnO_x (x=1..3)

- Using experimental geometry (if unavailable, CCSD(T) geometry)
- Comparison only to available experimental data



Bibliography and references

Books

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- T. N. Mitchell and B. Costisella, NMR From Spectra to Structures, Springer
- G. S. Rule and T. K. Hitchens, Fundamentals of Protein NMR Spectroscopy, Springer
- N. Jacobsen, NMR spectroscopy explained, Wiley
- M. Duer, Solid state NMR spectroscopy, Blackwell
- J. A. Weil and J. R. Bolton, *Electron Paramagnetic Resonance*, Wiley
- M. Knaupp, M. Bühl and V. G. Malkin, Calculation of NMR and EPR Parameters, Viley-VCH

Hamiltonians

The simplest Hamiltonian describing all the physics of NMR and EPR is the <u>Dirac-Breit Hamiltonian</u>. The DB Hamiltonian is fully relativistic (four-component) and difficult to solve. In order to make it numerically tractable, it must be reduced to a two-component non-relativistic Hamiltonian by some transformations and approximations.

Physicists apply the <u>Foldy-Wouthuysen</u> transformation to obtain the <u>Pauli</u> Hamiltonian. Chemists prefer the <u>Douglas-Kroll-Hess</u> transformation. Another popular approximation is the <u>ZORA</u> (zeroth-order regular approximation). These transformation lead to different expressions for the Hamiltonian terms, that are numerically very close.

 Pekka Manninen PhD thesis, University of Oulo, Finland (2004). http://herkules.oulu.fi/isbn9514274318/

Theory of EPR parameters

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- C. J. Pickard and F. Mauri, Phys. Rev. Lett. 88, 086403 (2002)
- S. Patchkovskii, R. T. Strong, C. J. Pickard and S. Un, J. Chem. Phys. 122, 214101 (2005)

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- M. S. Barhamy, M. H. F. Sluiter and Y. Kawazoe, Phys. Rev. B 76, 035124 (2007)

Converse approach

- D. Ceresoli, U. Gerstmann, A. P. Seitsonen and F. Mauri, *First-principles theory of the orbital magnetization*, PRB **81**, 060409 (2010)
- T. Thonhauser, D. Ceresoli, A. A. Mostofi, N. Marzari, R. Resta and D. Vanderbilt, A converse approach to the calculation of NMR shielding tensors, JCP **131**, 101101 (2009)
- D. Ceresoli, N. Marzari, M. G. Lopez and T. Thonhauser, *Ab-initio converse NMR for pseudopotentials*, PRB **81**, 184424 (2010).

Modern Theory of the Orbital Magnetization

- T. Thonhauser, D. Ceresoli, D. Vanderbilt and R. Resta, *Orbital magnetization in periodic insulators*, Phys. Rev. Lett. **95**, 137205 (2005)
- D. Ceresoli, T. Thonhauser, D. Vanderbilt and R. Resta, *Orbital magnetization in crystalline solids: multi-band insulators, Chern insulators, and metals,* Phys. Rev. B **74**, 024408 (2006)
- I. Souza and D. Vanderbilt, *Dichroic f-sum rule and the orbital magnetization of crystals*, Phys. Rev. B **77**, 054438 (2008)

Online resources: blogs, codes, lectures

- NMR Wiki: http://nmrwiki.org
- Software: http://edunmrsoft.blogsome.com
- NMR periodic table: http://www.bruker-nmr.de/guide/eNMR/chem/NMRnuclei.html
- Solid state NMR literature blog: http://ssnmr.blogspot.com
- Other blogs:

http://nmr-software.blogspot.com http://u-of-o-nmr-facility.blogspot.com http://scienceblogs.com/scientificactivist/2006/11/nmr_blogs.php

... and of course: www.quantum-espresso.org www.gipaw.net http://qe-forge.org/projects/qe-gipaw



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Uwe Gerstmann



Francesco Mauri

Sapienza Università di Roma condensed matter physics, computational materials science, nanotechnology, quantum chemistry, geochemistry Verified email at uniroma1.it



Emine Kucukbenli

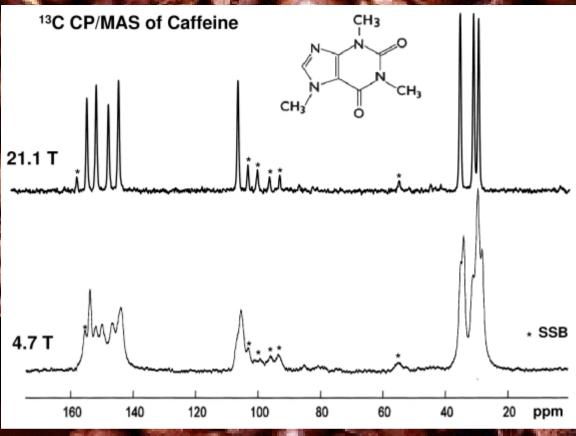


Nicola Varini

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Extra slides

Five practical steps to NMR parameters

0 - Download and compile the code

1 - Write the input file

- 2 Choose the proper pseudopotentials
- 3 Get the ground state structure with PW
- 4 Calculate NMR/EPR parameters with GIPAW

(5 - Simulate NMR/EPR spectra)

(6 - Publish on Science, win Nobel prize)

Step 0 – Download and compile

QE-GIPAW depends on PW code. To compile PW, you first need to configure QE, then to compile. You need at least a UNIX-like environment (Linux, Mac OSX, Cygwin) and a working F90/F95 compiler.

Example: cd espresso

(serial code, no blas/lapack installed) ./configure --disable-parallel --with-internal-blas --with-internal-lapack CC=gcc FC=gfortran \ F77=gfortran

(using Intel compiler, FFTW3 library path specified explicitely) ./configure CC=icc FC=ifort F77=ifort LIBDIRS="/opt/fftw3/lib"

(assuming you have a working MPI library, compile MPI+OpenMP version with Scalapack) ./configure --enable-parallel --with-scalapack --enable-openmp

Optionally you might want to modify make.sys for fine tuning, and after that: make pw pp or make all

Step 0 - Download and compile

If the previous compilation went well, now change into QE-GIPAW directory and:

```
cd GIPAW
./configure --with-qe-source=absolute_path_to_make.sys
make
```

The executables have the '.x' suffix and they will be placed into bin/ and GIPAW/bin/ directories

ceresoli@ada	ello:~/Cod	es/ge-espresso/trunk/bin\$ ls					
average.x	d3.x	fd_ifc.x	iotk.x	phcg.x	pp.x	pw.x	wannier_plot.x
band_plot.x	dist.x	fd.x	kpoints.x	ph.x	projufc.x	q2qstar.x	wfck2r.x
bands_FS.x	dos.x	feha.x	kvecs_FS.x	plan_avg.x	pw2bgw.x	q2r.x	wfdd.x
bands.x	dynnat.x	generate_vdl_kernel_table.x	landa.x	plotband,x	pm2gm_x	q2trans_fd.x	
bgw2pw.x	epsilon.x	initial_state.x	ld1.x	plotproj.x	pw2wannier90.x	q2trans.x	
Chibia * X	ev.x	iotk	nanypu.x	plotrho.x	pw_export.x	suppos.x	
cp.x	fd_ef.x	iotk_print_kinds.x	natdyn.x	pmu,x	pwi2xsf.x	wannier_ham,x	
ceresoli@adamello:"/Codes/ge-espresso/trunk/bin\$							

ceresoli@adamello:"/Codes/qe-gipaw/trunk/bin\$ ls gipam.x ceresoli@adamello:"/Codes/qe-gipaw/trunk/bin\$

PW input file is composed of Fortran "namelists" followed by "cards":

```
&control
    calculation = 'scf'
    restart_mode = 'from scratch'
    prefix = 'betacrist'
    pseudo dir = './pseudo/'
    outdir = './scratch/'
    tstress = .true.
    tprnfor = .true.
&svstem
    ibrav = 2
    a = 7.1315
    nat = 6
    ntvp = 2
    ecutwfc = 35
    ecutrho = 350
    spline ps = .true.
&electrons
    diagonalization = 'david'
    diago thr init = 1e-4
    mixing mode = 'plain'
    mixing beta = 0.7
    conv thr = 1e-10
/
```

!type of calculation: 'scf', 'relax', !'vc-relax, 'nscf', 'bands' !'from_scracth' or 'restart' !basename for all binary files !directory containing pseudopotentials !scratch directory for large binary files !if .true., print stress tensor !if .true., print forces

```
!see manual, ibrav=2 means fcc
!conventional lattice spacing in Å
!number of atoms
!number of different atomic speciec
!plane wave cutoff in Ry
!density cutoff in Ry
!useful for NMR
```

!'davidson' (faster, more memory) or 'cg'
!eigenvalue accuracy at first diagonalization

```
!0.3-0.7 in insulators, <0.1 in metals
!SCF accuracy</pre>
```

continued on the next slide...

continuing from the previous slide...

ATOMIC_SPECIES

Si 28.086 Si.pbe-tm-new-gipaw-dc.UPF 0 15.999 0.pbe-rrkjus-gipaw-dc.UPF

K_POINTS automatic

4 4 0 0 0

!atom mass pseudopotential

!4x4x4 not-shifted Monkhorst-Pack mesh

ATOMIC POSITIONS alat

Si	0.00000000	0.000000000	0.000000000
Si	0.250000000	0.250000000	0.250000000
0	0.075458128	0.174541872	0.125000000
0	0.424541872	0.325458128	0.125000000
0	0.174541872	0.424541872	0.375000000
0	0.325458128	0.075458128	0.375000000

```
!alat=cartesian in units of 'a'
!other possibilities: crystal (=fractional)
!or angstrom (=cartesian, in Å)
```

- One of the most common source of misunderstanding is how to enter the crystal cell parameters
- Two possibilities: choose a Bravais lattice or enter cell vectors manually

Bravais lattices supported by QE (ibrav != 0):

- ibrav=1 simple cubic
- ibrav=2 cubic F (fcc)
- ibrav=3 cubic I (bcc)
- ibrav=4 hexagonal or trigonal P
- ibrav=-5,5 trigonal R (two settings)
- ibrav=6 tetragonal P
- ibrav=7 tetragonal I (bct)
- ibrav=8 orthorhombic (P)
- ibrav=-9,9 base-centered orthorhombic (two settings)
- ibrav=10 face-centered orthorhombic
- ibrav=11 body-centered orthorhombic
- ibrav=-12,12 monoclinic P (two settings)
- ibrav=13 base-centered monoclinic
- ibrav=14 tricilinic

Please, check the settings (axis) and cell vectors in **PW/Doc/INPUT_PW.def**

If there is a Bravais lattice suitable for your input, then enter the lattice parameters in the namelist &system:

Using Bohr atomic units:	Using Angstrom units:
celldm(1) = a	a = <i>a</i>
celldm(2) = b/a	b = <i>b</i>
celldm(3) = c/a	C = <i>C</i>
celldm(4) = cos(see INPUT_PW.def)	cosAB = cos(<i>gamma</i>)
celldm(5) = cos(see INPUT_PW.def)	$\cos AC = \cos(beta)$
celldm(6) = cos(see INPUT_PW.def)	$\cos BC = \cos(alpha)$

Otherwise, if there is no default lattice matching your crystal structure, set **ibrav=0** and add the following card:

CELL_PARAMETERS [angstrom|bohr] a_x a_y a_z b_x b_y b_z

c x c y c z

Finally, enter the atomic coordinates in the ATOMIC_POSITIONS card:

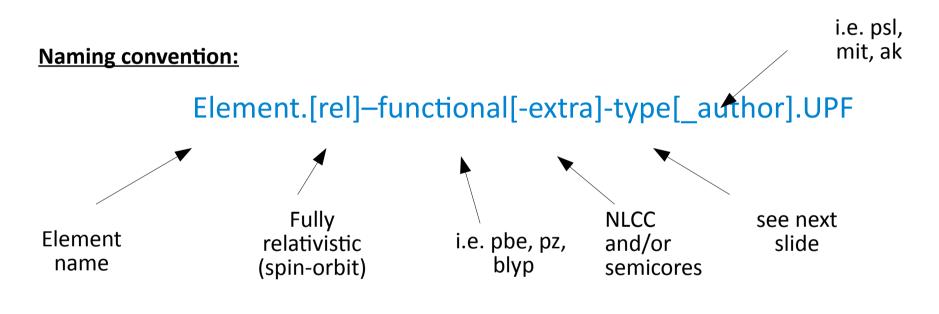
ATOMIC_POSITION [bohr|angstrom|alat|crystal] atom coord coord coord

••••

- The keywords **bohr**, **angstrom** and **alat** mean that the coordinates are cartesian and in units of, respectively bohr, angstrom, a (the length of the first lattice parameter).
- Otherwise, the crystal keyword allows you to enter fractional coordinates (i.e. from a CIF file)
- There is no automatic symmetry generation, you must enter all atoms in the unit and not just the asymmetric unit

2 – Choose pseudopotentials

- For GIPAW calculations you need GIPAW or PAW pseudopotentials
- For sake of clarity, I will illustrate all kinds of pseudopotentials you can find
- The recommended format is UPF (v1 or v2), which is text based. Other formats are deprecated (and probably unsupported)



For example: Fe.pw91-sp-van_ak.UPF Fe.rel-pbe-spn-kjpaw_psl.0.2.1.UPF

2 – Choose pseudopotential

Three PP types are supported in QE, and can be mixed:

- Norm conserving (NCPP)
 - tm = Troullier-Martins
 - rrkj = Rappe-Rabe-Kaxiras-Johannopulos
 - vbc = Van Barth-Car
 - hgh = Hartwigsen-Goedecker-Hütter
 - fhi = converted from FHI table of pseudopotentials
- Ultrasoft (USPP)
 - van = Vanderbilt
 - rrkjus = Rappe-Rabe-Kaxiras-Johannopulos ultrasoft
- PAW
 - kjpaw = Kresse-Joubert
 - there is no official naming for pseudos generated by atompaw (N. Holzwarth)

many elements, well tested few elements few elements most elements, high cutoff most elements, well tested?

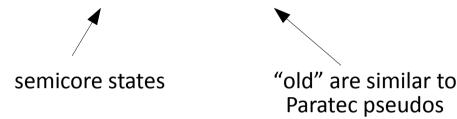
well tested, _ak with problems? well tested, low cutoff

I don't have much experience

2 – Choose pseudopotential

• Slightly different naming convention for GIPAW pseudos, sorry!

Element.functional-type[-semi]-gipaw[-new]-author.UPF



• PAW-GIPAW pseudos generated by atompaw, are marked by '-nh' as author

For example:

O.pbe-tm-gipaw-new-dc.UPF C.pbe-rrkjus-gipaw-dc.UPF K.pbe-tm-semi-gipaw-xy.UPF Fe.pbe-paw-gipaw-nh.UPF

Beware:

gipaw pseudopotentials found on www.quantum-espresso.org are for XANES core-hole calculations, not for NMR/EPR!

2 – Choose pseudopotential

Pseudopotential sources (incomplete):

- www.quantum-espresso.org/pseudopotentials (QE official pseudos)
- sites.google.com/site/dceresoli/pseudopotentials (D. Ceresoli)
- www.impmc.jussieu.fr/~software/gipaw/pseudopotentials.html (A. Seitsonen)
- qe-forge.org/gf/project/pslibrary (A. Dal Corso)
- users.wfu.edu/natalie/papers/pwpaw/newperiodictable (N. Holzwarth)
- www.physics.rutgers.edu/gbrv (for high-trhoughput calculations)
- www.nnin.org/research-support/computation/nnin-computation-nanotechnologyresources/virtual-vaults (searchable database, many formats, many codes)
- did I miss someone?

2 - Choose pseudopotential

Fe bcc ferromagnetic, equation of state, courtesy of D. Dragoni (EPFL)

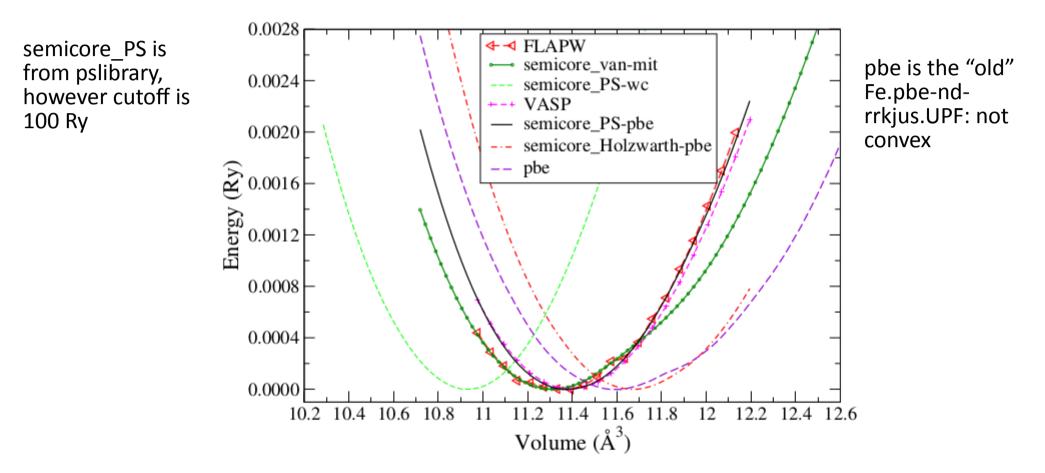


Figure 9: E vs. V comparison of QE ($24 \times 24 \times 24$, smearing of 0.001 Ry), VASP and all-electron calculations. Different QE pseudopotentials are used while for VASP there is a single one. The ewfc cutoff varies according to the pseudopotential considered (60 Ry for the standard pbe and the semicore_van-mit while 100 Ry for the others.)

2 - Chose pseudopotential

semicore_PS is from pslibrary, however cutoff is 100 Ry

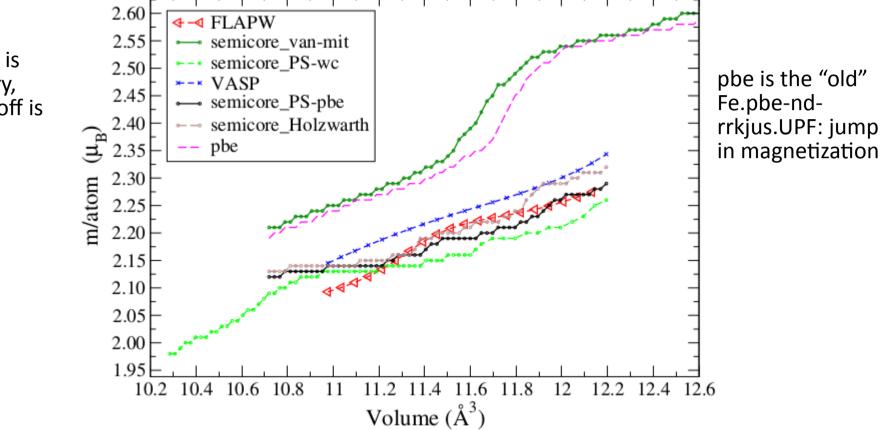


Figure 10: M vs. V comparison of QE (24×24×24, smearing of 0.001 Ry), VASP and all-electron calculations. Details as in Fig. 9)

3 - Ground state and relaxation

- Now you are ready to obtain the ground state wavefunctions and/or relax the structure with PW
- You can relax positions and cell parameters simultaneously
- You can constrain atom positions, bonds, angles, cell shape, ...
- More details during the afternoon lab

Running PW:

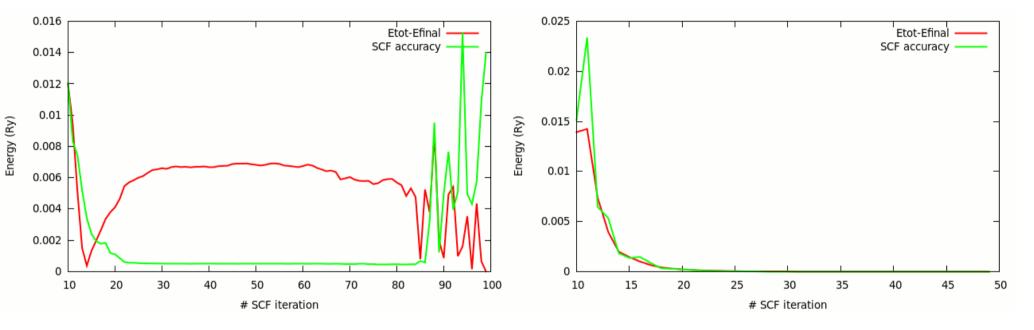
(in serial)
/path/to/pw.x <betacrist-scf.in >betacrist-scf.out &

(in paralllel, maybe in a job script) mpirun -np 8 /path/to/pw.x -npool 4 -ndiag 1 <betacrist-scf.in >betacrist-scf.out

(the most crazy line I've used, on Fermi) runjob --ranks-per-node 8 --np 2048 --envs OMP_NUM_THREADS=4 : pw.x -ndiag 16 -npool 4 -ntg 8 <AI-AI2O3-slab300atom-run09-vcmd.in >AI-AI2O3-slab300atom-run09-vcmd.out

3 - Ground state and relaxation

Provided that planewave cutoff and k-point mesh are sufficiently converged, the most important quantities to monitor during SCF or relaxation are: **total energy** and **SCF accuracy**



problematic convergence

fast, good convergence

In SCF convergence is problematic, maybe the system is a metal, or has a small bang gap. In that case, set: **occupations='smearing'**, **degauss=0.01** (in Ry), and reduce **mixing_beta** to 0.1 or smaller

- After having obtained the ground state, you can run GIPAW
- GIPAW input is much shorter and keywords have reasonable default values
- Here is an example to compute NMR shielding tensors

```
&inputgipaw
```

```
job = 'nmr'  ! for NMR shielding
prefix = 'betacrist'  ! same as in PW input
tmp_dir = './scratch/'  ! same as in PW input
isolve = 0  ! diagonalization, 0 or 1
iverbosity = 1  ! increase for more details
q_gipaw = 0.01  ! the small "q"
spline_ps = .true.  ! same as in PW input
use_nmr_macroscopic_shape = .true.  ! consider sample macroscopic shape
```

Finally, you can run GIPAW in a similar fashion to PW:

(serial) /path/to/gipaw.x <betacrist-nmr.in >betacrist-nmr.out

(parallel) mpirun -np 8 /path/to/gipaw.x -ndiag 1 -npool 4 <betacrist-nmr.in >betacrist-nmr.out

The most important quantities in output are: f-sum rule, susceptibility and total sigmas:

. . .

f-su	m rule (should H -30.7050 0.0000 0.0000	-0.0000	0.0000 0.0000 -30.7048	very importar fulfilled as m possible	uch as	
_	bare pGv (HH) in -44.2541 0.0000 0.0000	0.0000 -44.2541 0.0000	0.0000 0.0000 -44.5654	the true i susceptibili	ty is in th	is
CUT	bare vGv (VV) iı -42.8223 0.0000 -0.0000	-0.0000	0.0000 0.0000	ran	ige	
			or anisotropy and	asymmetry)		
Atom	3 0 pos: (223.7763 26.8036 32.1875	26.8036	31.1766	Total sigma: Mielding tensor	225.13	isotropic in ppm
0 0 0 0	3 sigma_11= 3 sigma_22=	196.9727 a 192.9773 a	xis=(0.707107 - xis=(0.434141	Simpson convention 0.707107 0.000000) 0.434141 -0.789331) 0.558142 0.613967)	n principal	axes

To calculate electric field gradient (EFG), you must provide the nuclear quadrupole in 10⁻³⁰ m²:

In the output, you can find Cq, eta and principal axes:

NQR/	NQR/NMR SPECTROSCOPIC PARAMETERS:						
Са	1	Vxx=	-0.0229	axis=(0.788473	0.612926	0.051306)
Са	1	Vyy=	-0.1861	axis=(-0.609218	0.789729	-0.071989)
Са	1	Vzz=	0.2090	axis=(-0.084641	0.025505	0.996085)
Са	1	Q=-4.4	4 1e-30 m^2	Cq=	-2.1806 M	Hz eta=	0.78044

To calculate the **EPR g-tensor**, your system must be spin polarized. The input file is similar to that for NMR shieldings:

```
&inputgipaw
    job = 'g_tensor'
    prefix = 'h2cn'
    tmp_dir = './scratch/'
    isolve = 0
    iverbosity = 1
    q_gipaw = 0.01
    spline_ps = .true.
    use_nmr_macroscopic_shape = .false.
/
```

In output, the deviation Δg in ppm from 2.002319... is (there are two different ways to compute the spin-other-orbit contributions):

Delta_g total	(SOO a la Parate	ec):	
3104.68	0.00	0.00	
0.00	-82.23	0.00	
0.00	0.00	-1925.19	
Delta_g total	(SOO as in Eq.(7	7)):	
3195.44	0.00	0.00	
0.00	-66.90	0.00	
0.00	0.00	-1935.92	

Likewise, to compute EPR hyperfine couplings you must provide the nuclear g-factor for each atomic specie:

```
&inputgipaw
    job = 'hyperfine'
    prefix = 'h2cn'
    tmp_dir = './scratch/'
    spline_ps = .true.
    iverbosity = 11
    hfi_output_unit = 'MHz' ! 'MHz' or 'G'
    hfi_nuclear_g_factor(1) = 1.404844 ! 13C
    hfi_nuclear_g_factor(2) = 0.40376087 ! 15N
    hfi_nuclear_g_factor(3) = 5.585694 ! 1H
```

In output you can find the dipolar and Femi contact in the chosen units:

	to	tal dipolar (symme	trized)	
С	1	-18.889103	0.00000	-0.000001
С	1	0.000000	2.676450	0.00001
С	1	-0.000001	0.000001	16.212653

...

IS0	ISOTROPIC (FERMI-CONTACT) COUPLINGS WITHOUT ZORA:						
War	Warning: core-relaxation is an experimental feature						
	<	bare	GIPAW	core-relax	total		
С	1	-1.471958	-71.844171	-0.494637	-73.810766		
Ν	2	1.158516	68.877881	-33.425295	36.611101		
Н	3	133.194574	96.213231	0.00000	229.407804		
Н	4	133.194577	96.213234	0.00000	229.407811		

Troubleshooting

E: Error reading pseudopotential

S: open the UPF files with a text editor, locate the PP_PAW_VERSION and PP_PAW_GIPAW_VERSION sections, and change the number therein (0.1 or 1.0) to 1

E: Very strange NMR shielding or EPR g-tensor, not respecting crystal symmetry, especially for hexagonal systems

S: GIPAW is incompatible with symmetry operation that do not map cartesian axes into cartesian axes. Orient the hexagonal cell like this, or disable symmetry in SCF (nosym=.true. in &system)

E: Cell parameters relaxation stops with error: non orthogonal operation S: The symmetry of you system is changing, disable it (nosym=.true.)

E: Difficult SCF convergence, crazy results, atoms too close

S: Check the crystal structure with Xcrysden: xcrysden --pwi betacrist-scf.in

Troubleshooting

E: Error computing Cholesky...

S: Did you compile with SCALAPACK? if not the default CAVAPACK(*) routines are very sensitive to numerical noise. Re-configure espresso with: --with-scalapack Still not working? some versions of MKL are reported to be broken. Disable parallel diagonalization with '-ndiag 1' on the command line

E: compute_u_kq, some eigenvectors not converged

S: try to change isolve (from 0 to 1 or viceversa) in GIPAW

E: error in davcio, if you use a different number of CPUs between PW and GIPAW S: add wf collect=.true. in PW &control namelist

E: task groups not supported

S: this will be fixed in the next GIPAW release

E: GIPAW says: cgsolve_all: root not converged

S: maybe your system is metallic, or with a small band gap. Metallic systems, including the Knight shift term will be supported in the next GIPAW release