

Pulsed EPR spectroscopy with a focus on DEER (PELDOR)

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Why pulsed EPR?



How does it work? FID & echoes



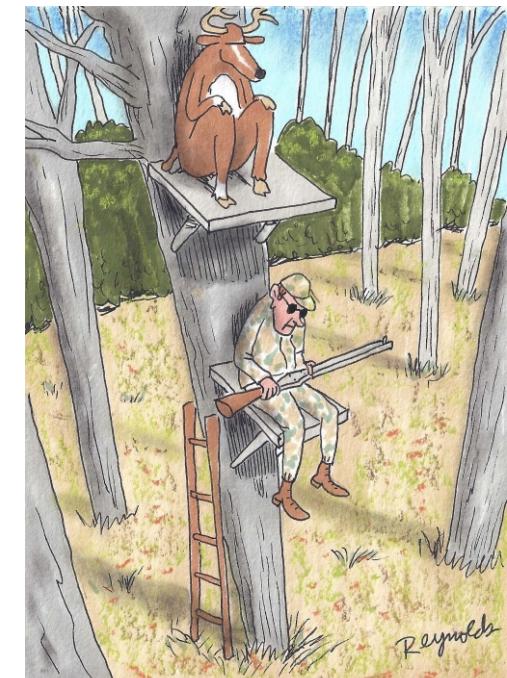
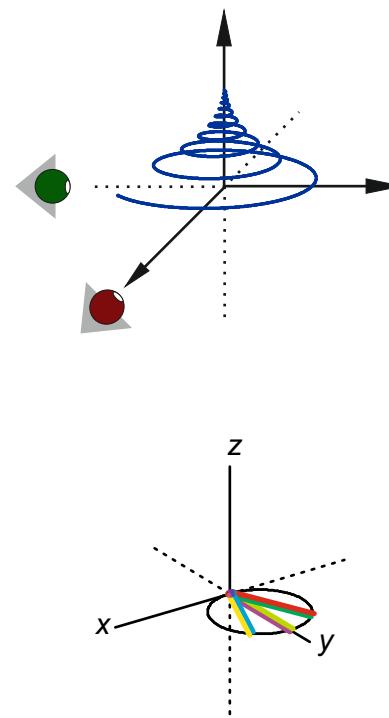
A few concepts and pulse sequences



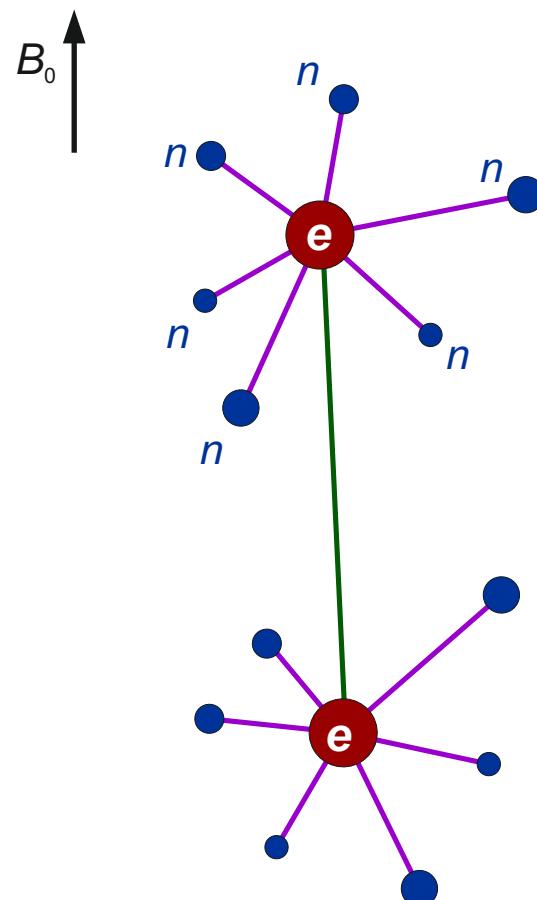
DEER: Basics and implementation



DEER: Data analysis

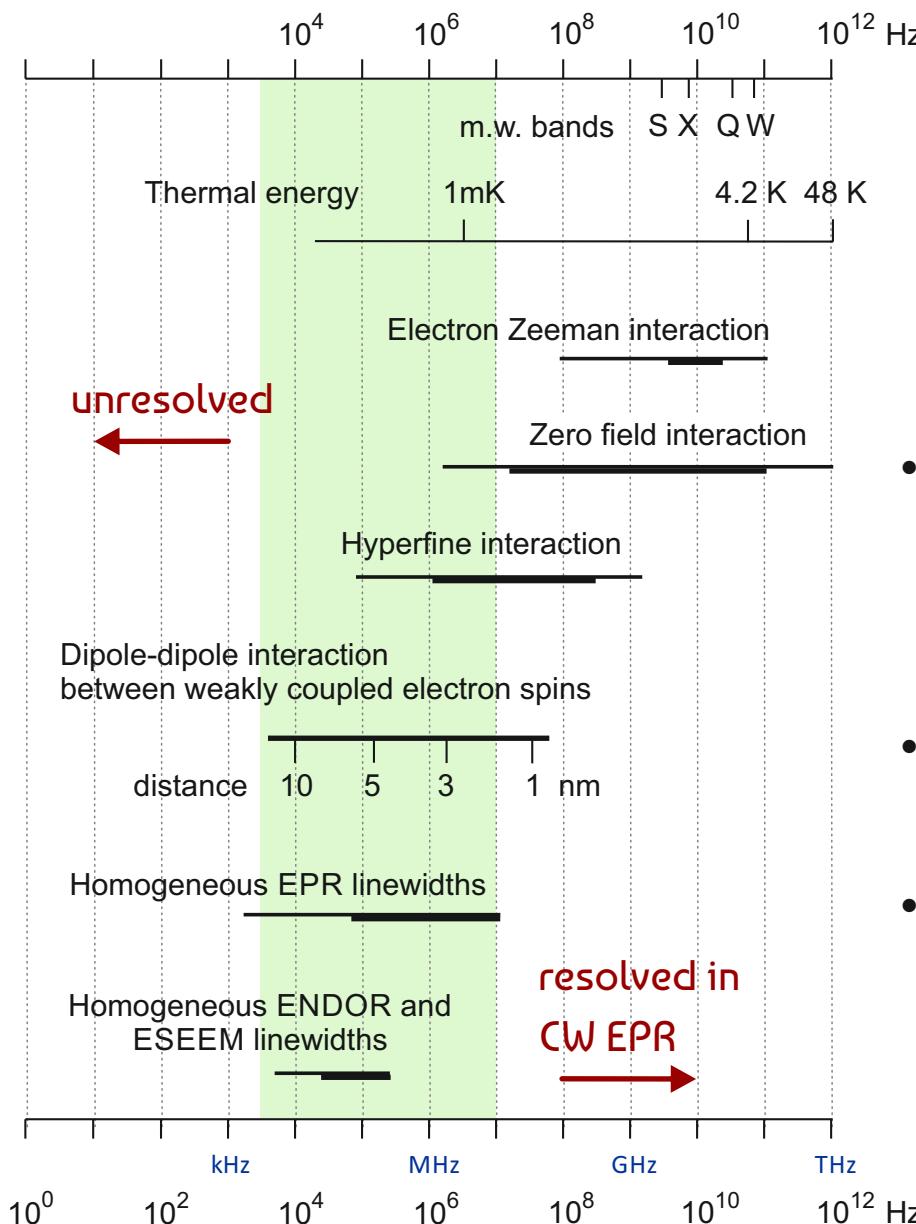


Interactions and the information that they provide



Name	Information
electron Zeeman	fingerprinting of radical type or metal coordination
hyperfine	distribution of the SOMO (reactivity) distance of protons from the center of spin density
nuclear Zeeman	identification of nuclei that give rise to hfi
nuclear quadrupole	binding situation of the nucleus for $I > 1/2$ (chemical shift is not available)
zero-field	fingerprinting of triplet type or metal coordination spin state for metal ions (low or high spin)
exchange	orbital overlap (important for electron transfer)
dipole-dipole	distances in the nanometer range (15 - 100 Å)

An overview of microwave bands and interactions



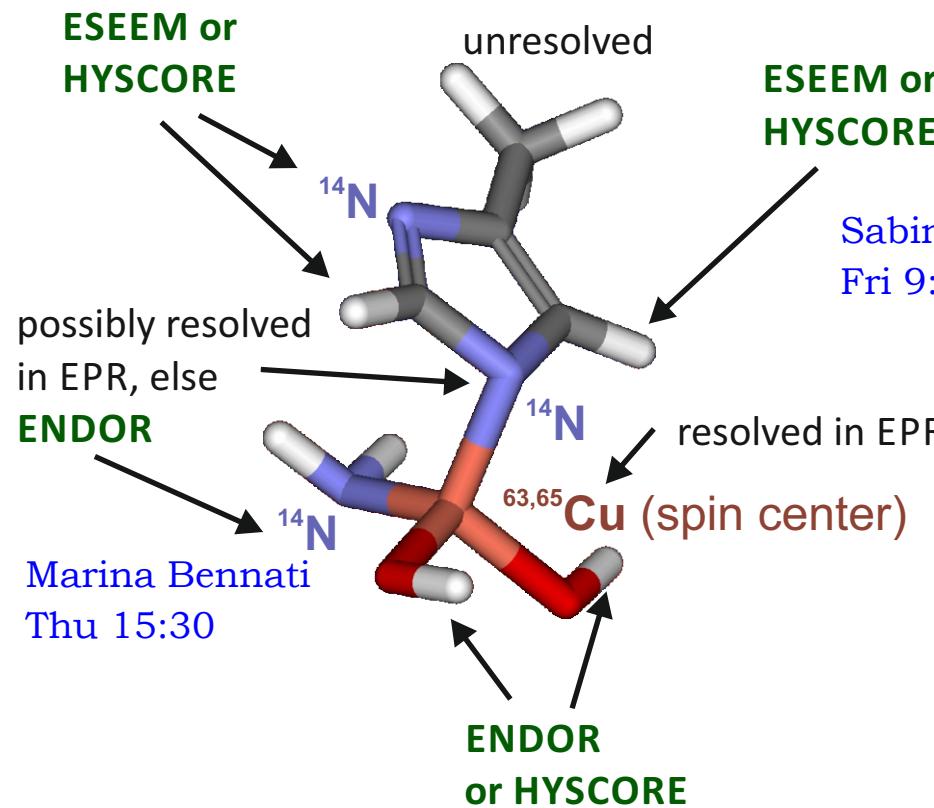
By separating interactions,
pulsed EPR provides information
lost by line broadening in CW EPR

- for electron group spin > 1/2 (more than one unpaired electron)
- most valuable source of EPR restraints on structure
- resolution limit depends on sample preparation

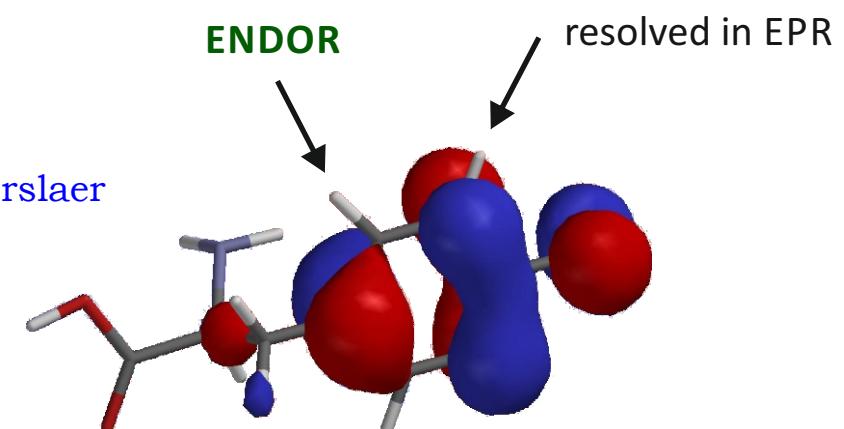
1 mT corresponds to 28 MHz

Example: Measuring hyperfine couplings

Histidine-coordinated Cu(II)

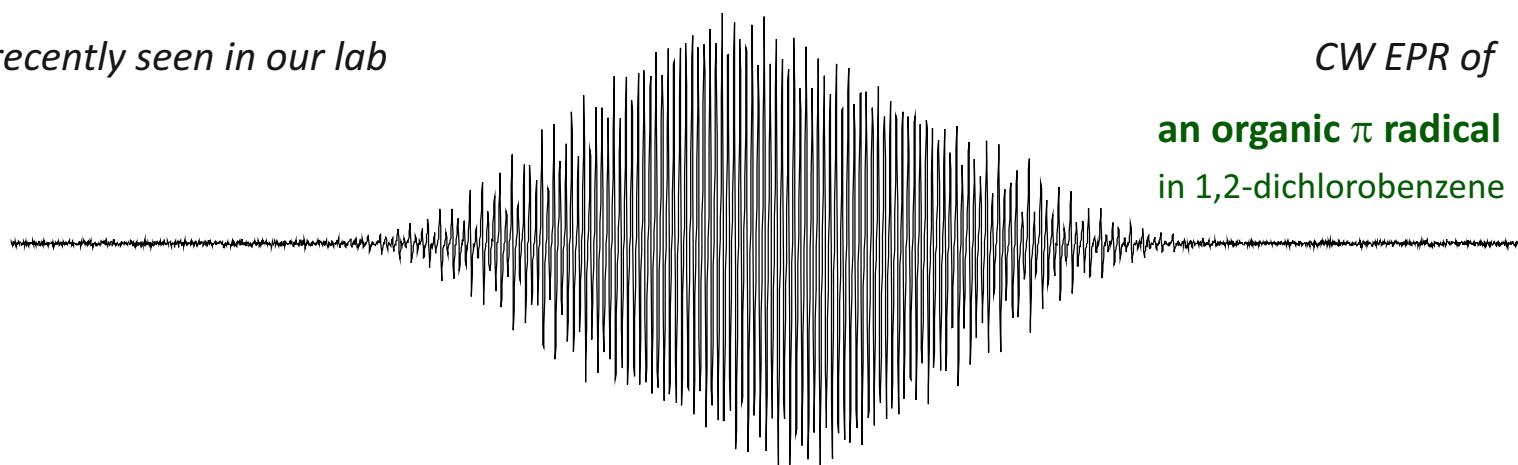


Tyrosyl radical

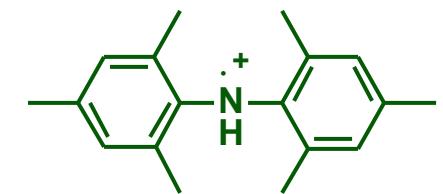


Even if it's resolved, ...

recently seen in our lab



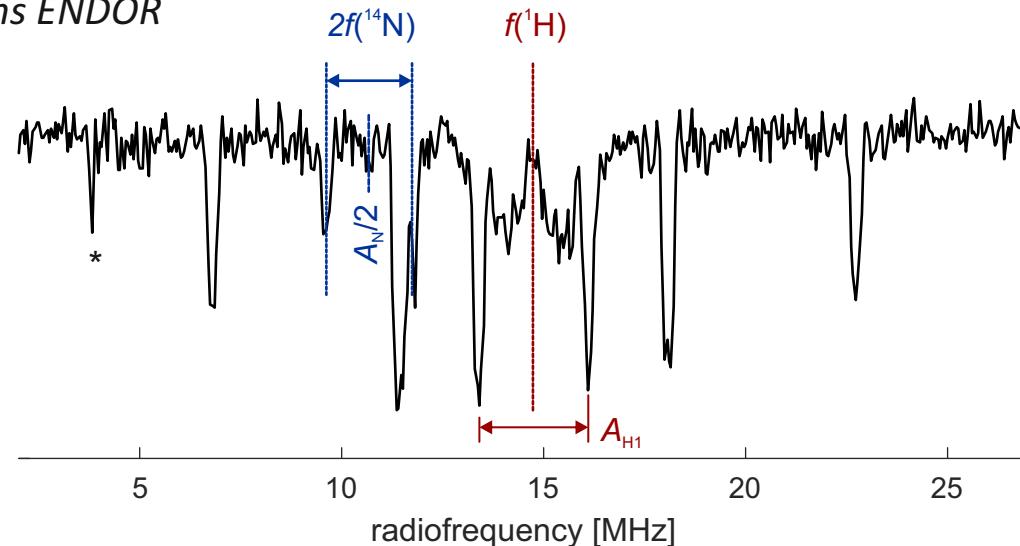
*CW EPR of
an organic π radical
in 1,2-dichlorobenzene*



sample courtesy
Agnes Kütt
University Tallin

a more complicated experiment can make it simpler

Mims ENDOR



⊕ identity of the nuclei (^1H , ^{14}N)

⊕ direct read-off of hyperfine couplings

Pulsed EPR is a zoo of techniques

EPR spectra

- Fourier-transform EPR
- field-swept echo-detected EPR

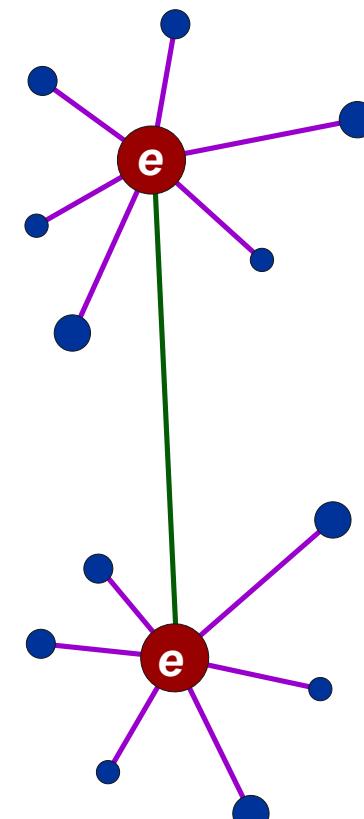
Electron spin relaxation

- Hahn echo decay (T_2/T_m)
- Inversion recovery (T_1)

Hyperfine couplings

nuclear Zeeman & nuclear quadrupole interactions

- ESEEM
 - HYSCORE
 - Davies ENDOR
 - Mims ENDOR
 - ELDOR-detected NMR
- Sabine, Fri 9:00
 Marina, Thu 15:30
 Daniella, Fri 11:00

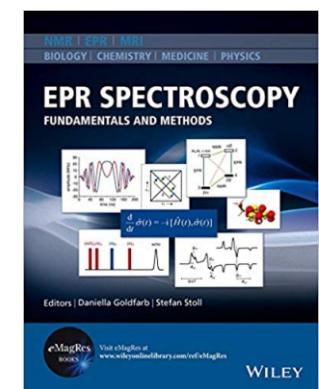
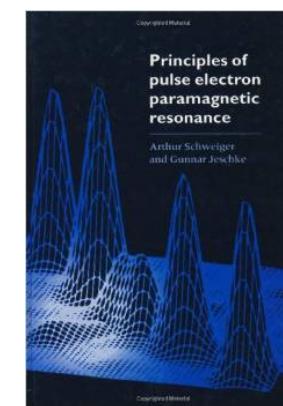


Electron-electron couplings

- DEER/PELDOR
- RIDME
- SIFTER
- DQC-EPR

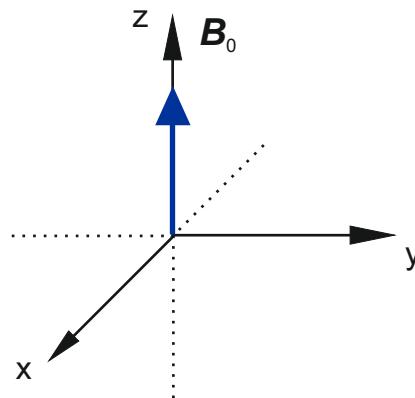
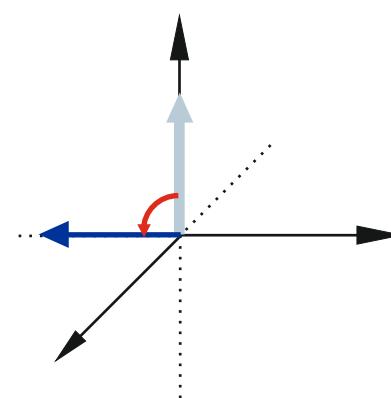
Fine print

- more techniques exist
- new ones are still coming up
- almost all of them fit into one of the four boxes

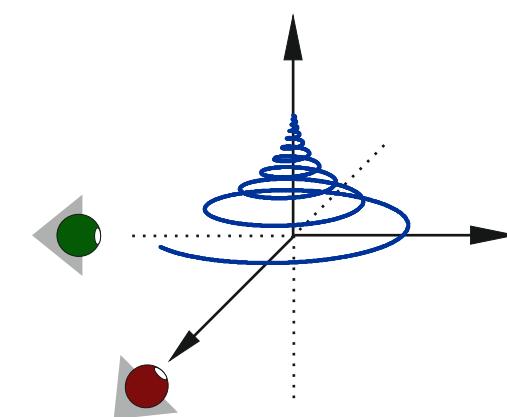


Free induction decay (FID) and Fourier transform EPR

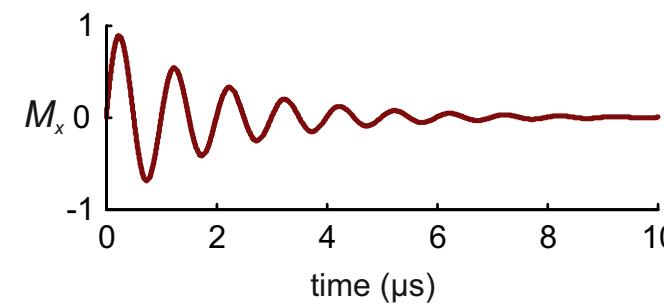
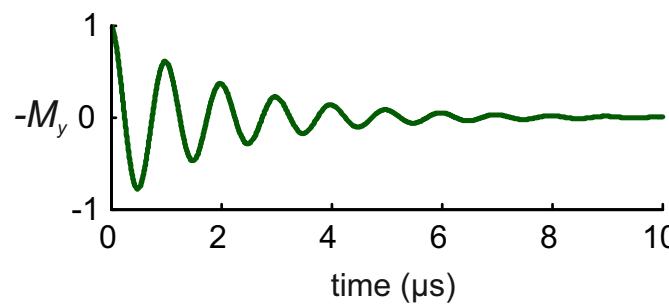
Thermal equilibrium

 $\pi/2$ (90°) pulse

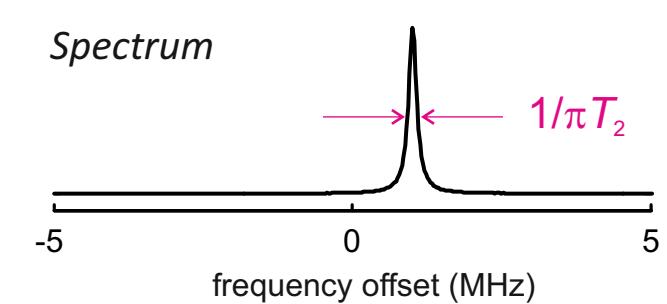
Free evolution (FID)



Complex (quadrature) signal: $-M_y + i M_x$



Fourier transform



Motion of the magnetization vector without relaxation

Classical equation of motion

spin has, both, angular momentum J and a magnetic moment μ

$$\frac{d\vec{J}}{dt} = \underbrace{\vec{\mu} \times \vec{B}(t)}_{\text{torque}}$$

\vec{J} total angular momentum vector, $\vec{B}(t)$ includes static and oscillatory magnetic fields

as $\gamma \vec{J} = \vec{\mu}$ and $\vec{M} = \vec{\mu}/V$, we have

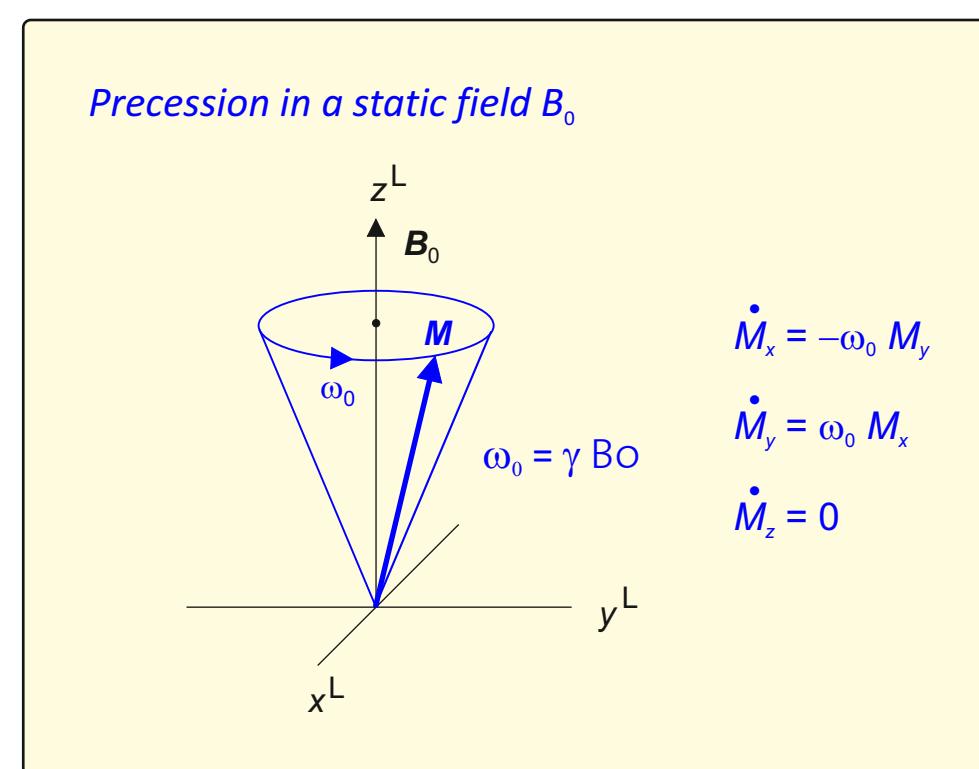
$$\frac{d\vec{M}}{dt} = \vec{M} \times \gamma \vec{B}(t)$$

and by expanding the cross product

$$\dot{M}_x = -\gamma B_z(t) M_y + \gamma B_y(t) M_z$$

$$\dot{M}_y = \gamma B_z(t) M_x - \gamma B_x(t) M_z$$

$$\dot{M}_z = \gamma B_x(t) M_y - \gamma B_y(t) M_x$$



Relaxation in the magnetization vector model

Relaxation is due to flips of individual spins

⇒ only phenomenological description possible in a classical picture

⇒ length of \vec{M} is not a constant of motion

Two types of relaxation just a reminder, Thomas Prisner talked about this

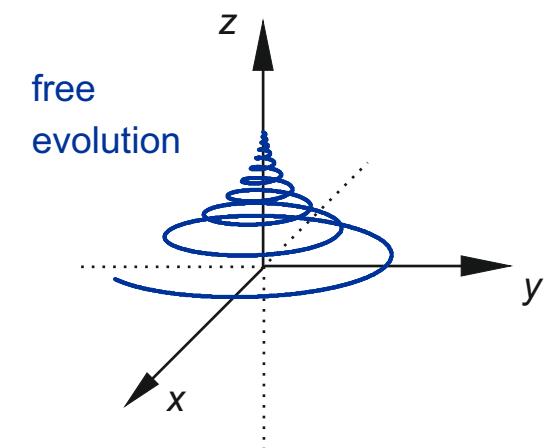
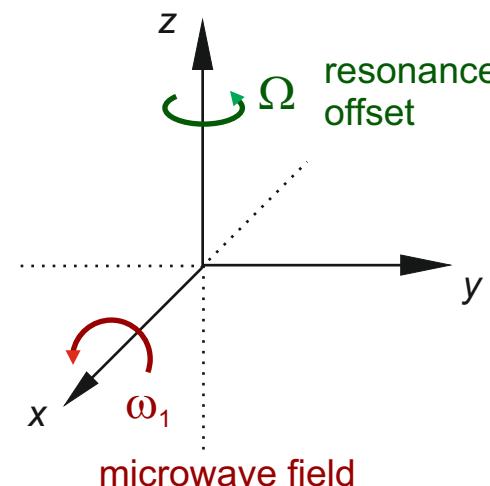
- longitudinal relaxation requires energy exchange with the environment, enthalpic, time constant T_1
- transverse relaxation reduces phase coherence of the ensemble, entropic, time constant T_2

Bloch equations with relaxation rotating frame

$$\dot{M}_x = -\Omega M_y - \frac{M_x}{T_2}$$

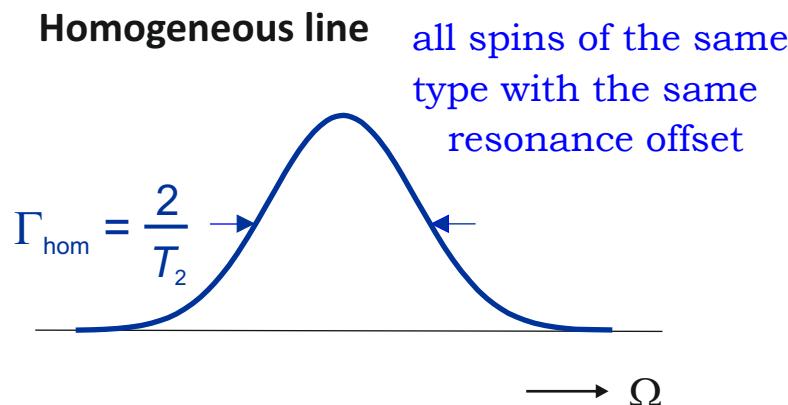
$$\dot{M}_y = \Omega M_x - \omega_1 M_z - \frac{M_y}{T_2}$$

$$\dot{M}_z = \omega_1 M_y - \frac{M_z - M_0}{T_1}$$



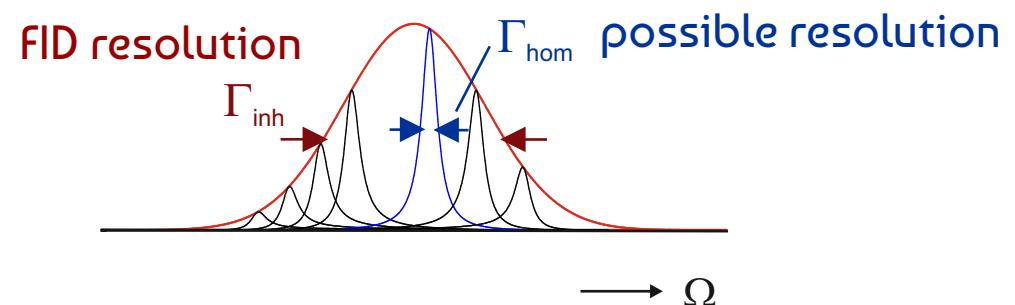
Homogeneous and inhomogeneous lines

Homogeneous line

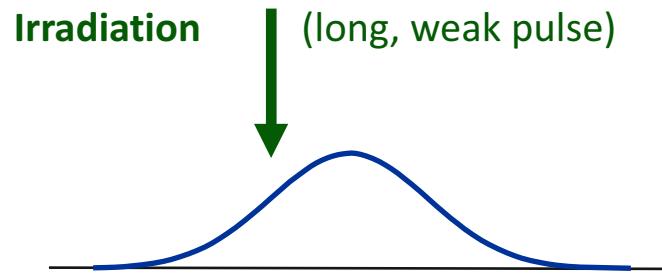


all spins of the same type with the same resonance offset

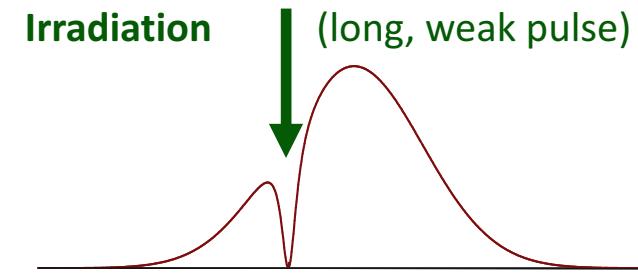
Inhomogeneous line



possible resolution



- saturation



only a spin packet is saturated

- spectral hole burning

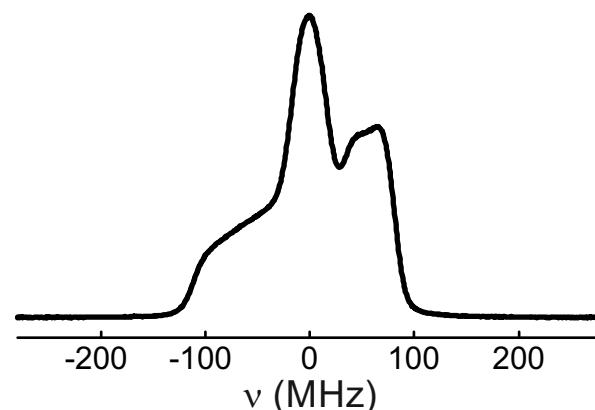
(used in Davies ENDOR, ELDOR-detected EPR)



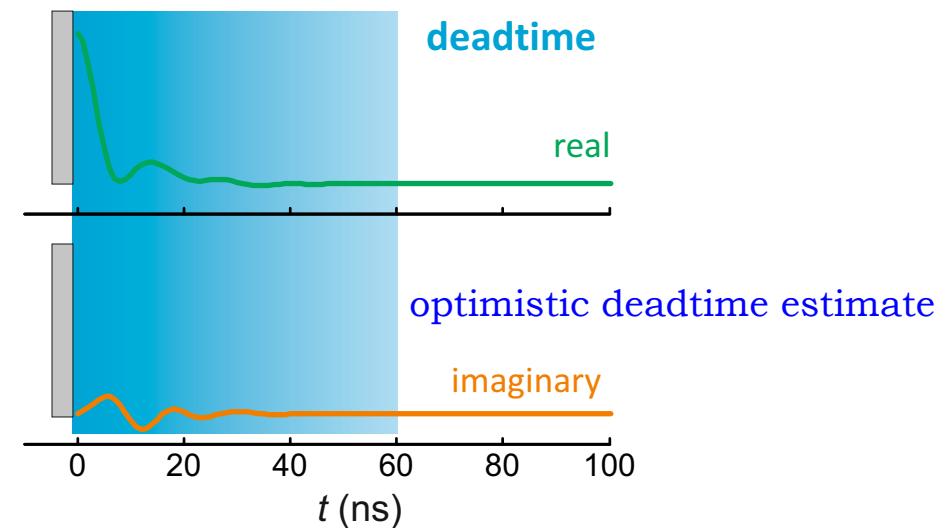
! in the solid state, inhomogeneous broadening by anisotropic interactions is often dominant

Why a single pulse is not sufficient for solids

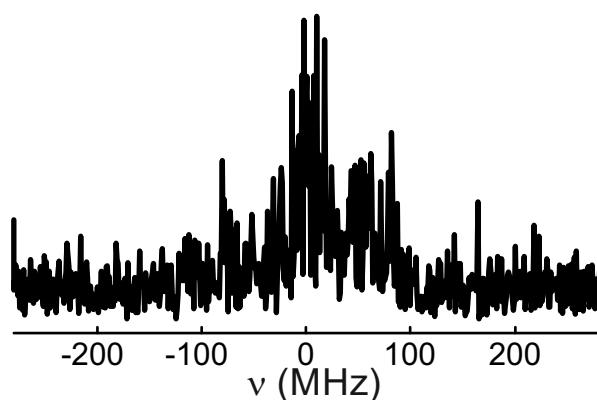
EPR absorption spectrum of a nitroxide
(X band)



... and the corresponding FID



FT of the accessible part of the FID



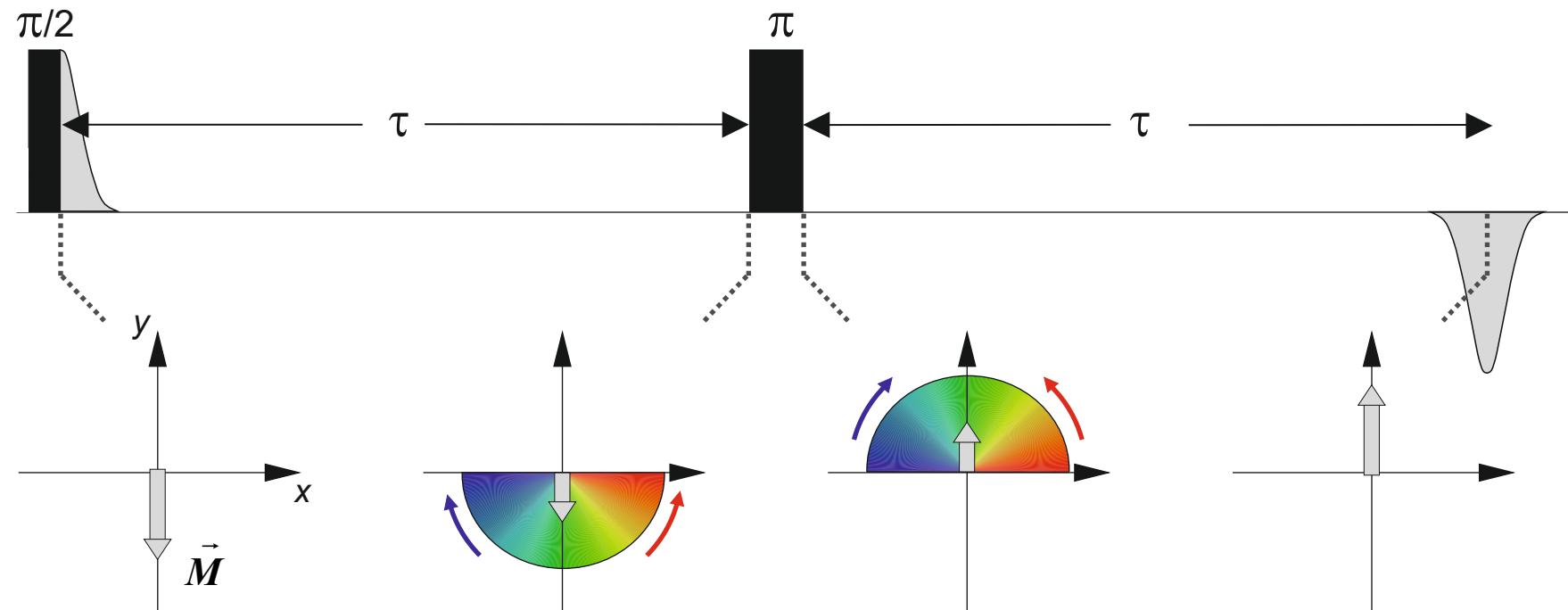
- destructive interference of spin packets with different resonance offset dampens FID within dead time

FT EPR by FID detection possible only for:

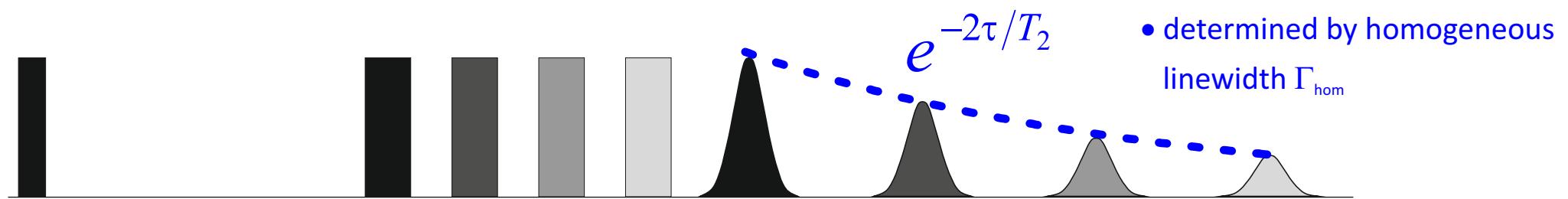
- spectral widths < 100 MHz*
- line widths ≈ 3 MHz

*with modern arbitrary wavefunction generators: up to 800 MHz

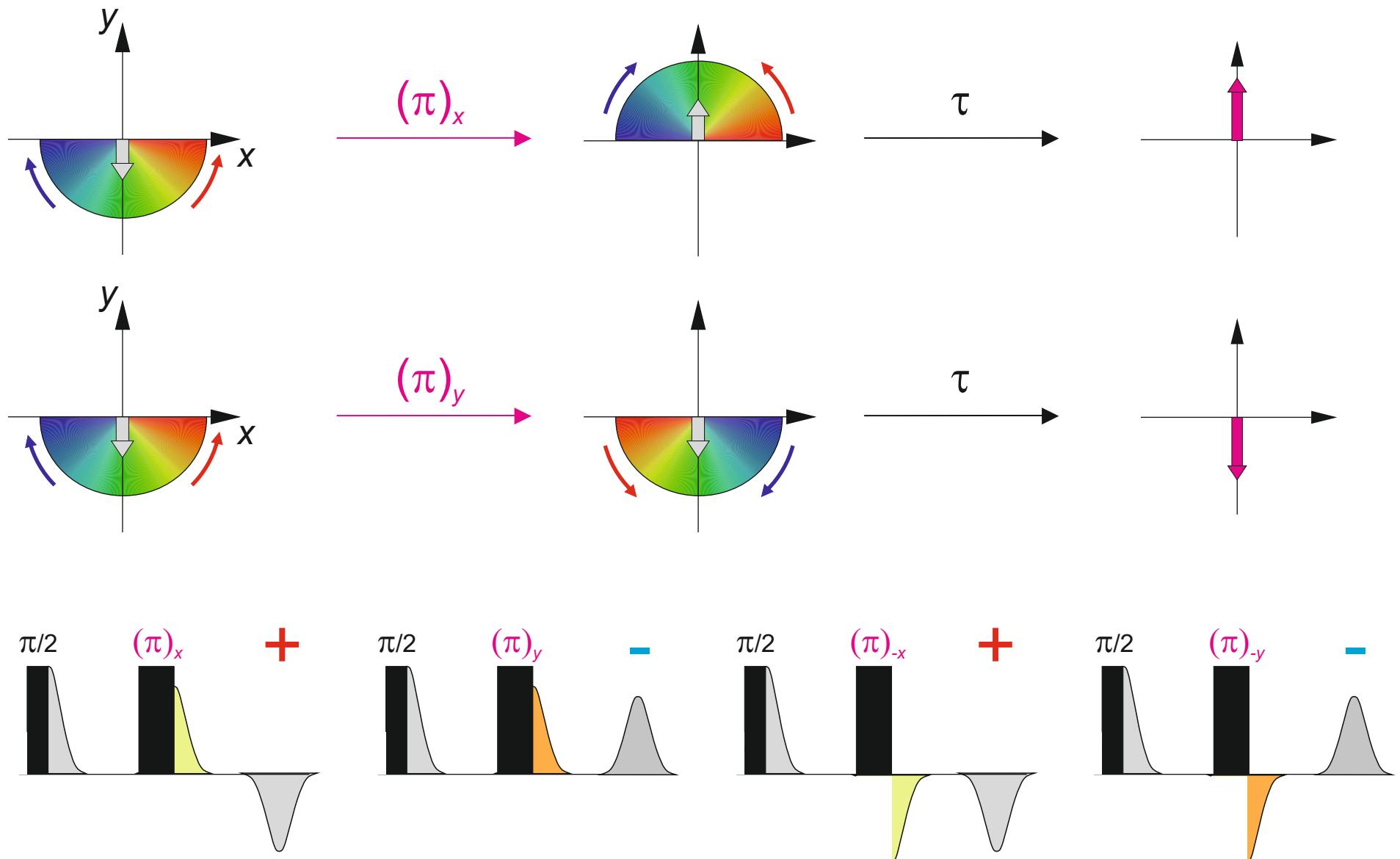
The primary echo (Hahn echo)



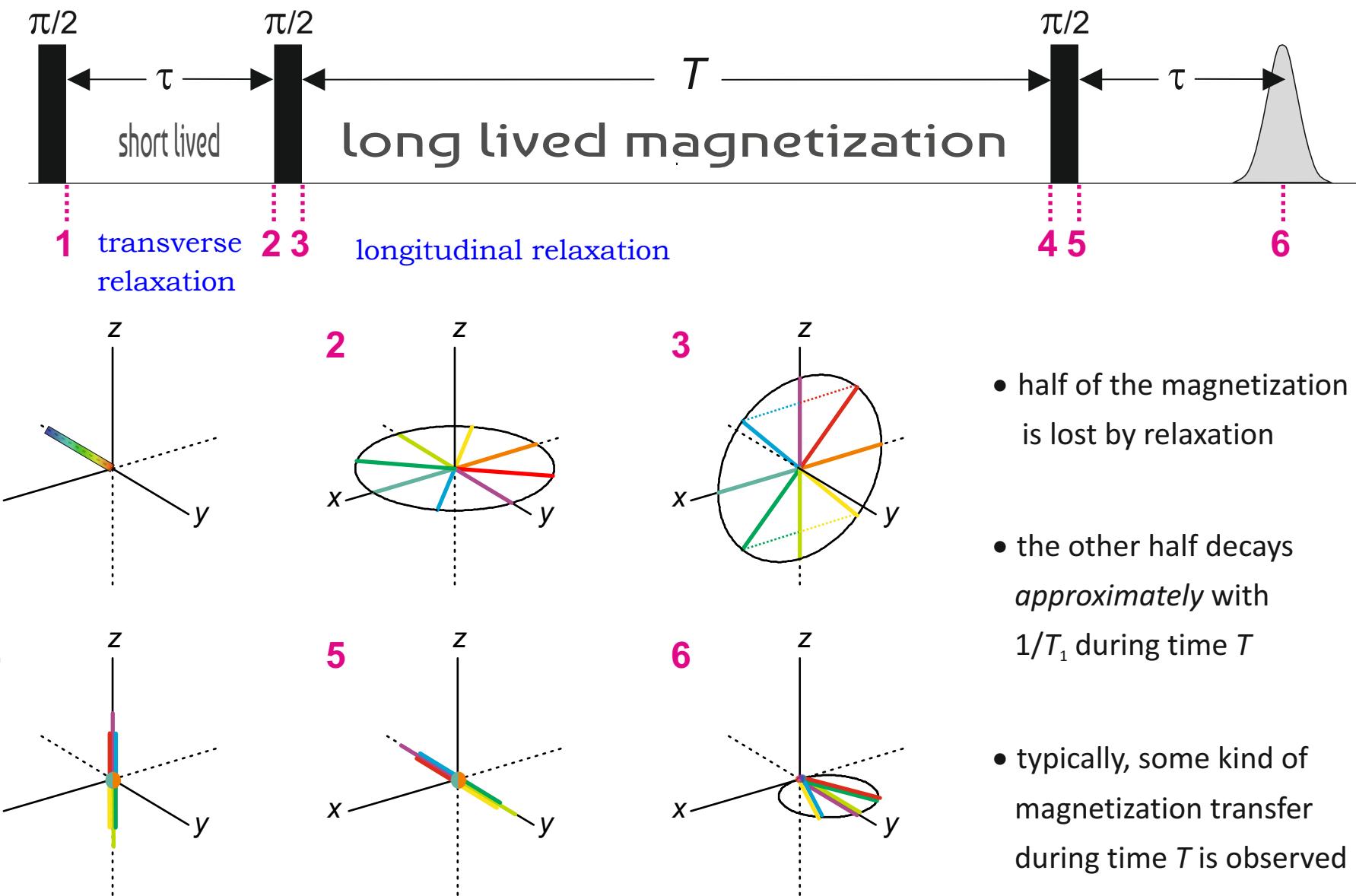
Echo decay



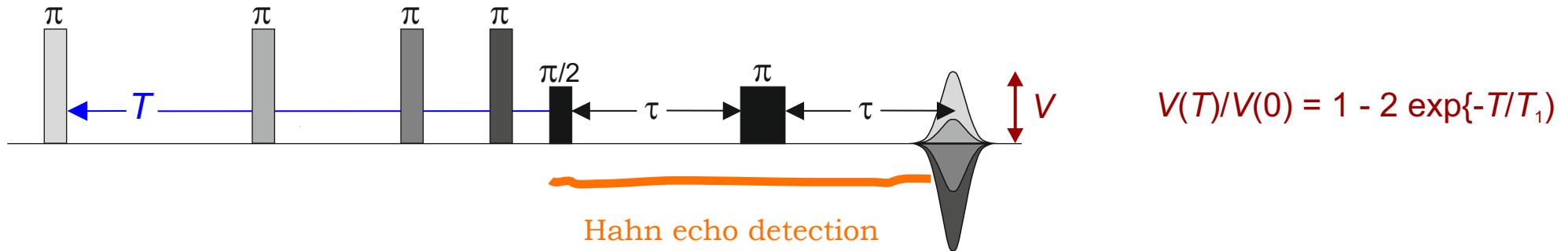
Basics of phase cycling



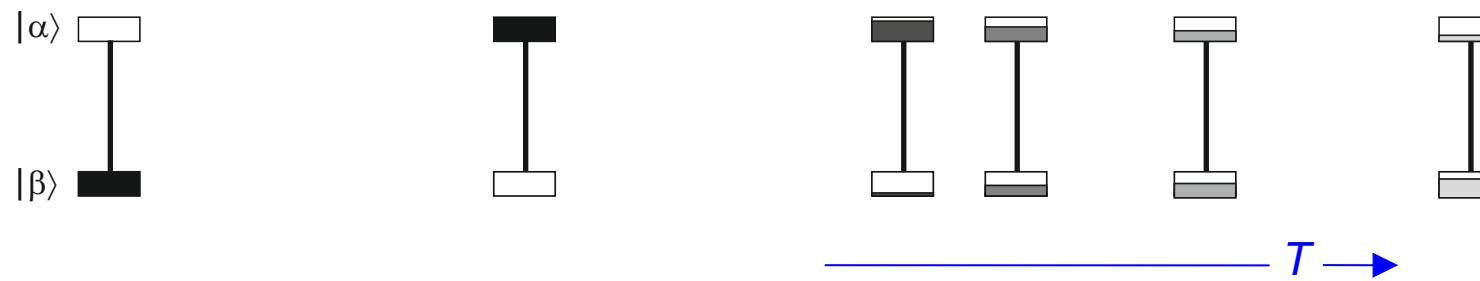
The stimulated echo experiment



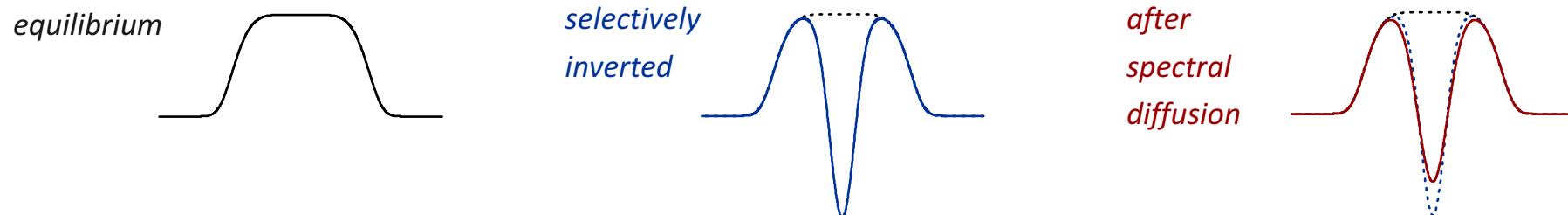
The inversion recovery pulse sequence



Thermal equilibrium After inversion Longitudinal relaxation



Beware of spectral diffusion spins change resonance frequency or exchange magnetization



Nutation and excitation bandwidth

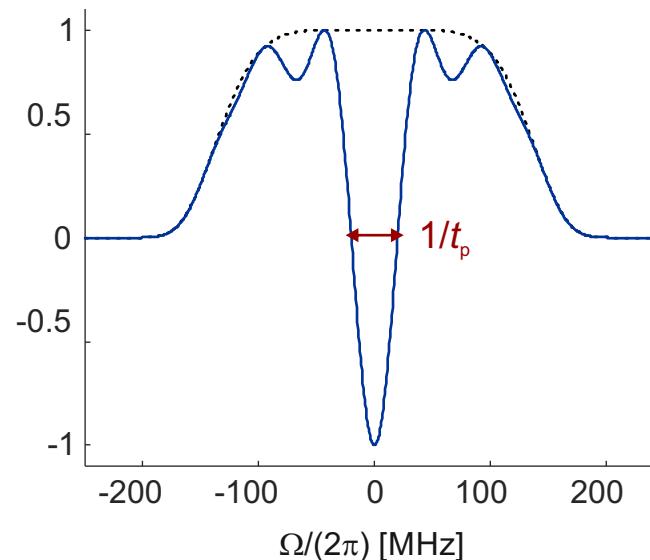
Rotating-frame Hamiltonian during irradiation

$$\mathcal{H} = \Omega S_z + \omega_1 S_x$$

The flip angle is defined on resonance

$$\beta = \omega_1 t_p$$

Excitation profile



Nutation

$\omega_{\text{eff}} = \sqrt{\Omega^2 + \omega_1^2}$

$\theta = \text{atan}(\omega_1/\Omega)$

precession about the effective field

- the Fourier transform of a boxcar function approximates the excitation profile for a *small* flip angle:

$$P(\Omega) \approx \frac{\sin(\Omega t_p/2)}{\Omega t_p/2}$$

Overcoming the microwave power limit to bandwidth

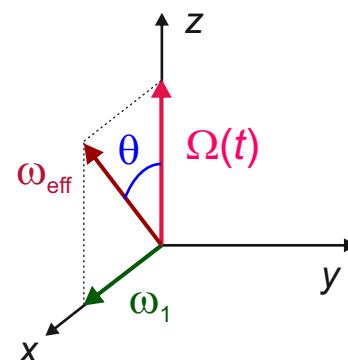
The power-bandwidth relation for rectangular monochromatic pulses

$$\text{BW} \approx 1/t_p \quad t_p = \beta/\omega_1 \quad \Rightarrow \text{BW} \approx \omega_1/\beta = g_x \mu_B B_1/(\hbar\beta) \quad \text{with } B_1 \propto \sqrt{P_{\text{mw}}}$$

! the time available for spin excitation is of the order of T_2 (microseconds) or T_1 (milliseconds),
 whereas spectral width is of the order of 100 MHz ... 10 GHz, corresponding to $t_p = 10 \dots 0.1 \text{ ns}$

Frequency-swept pulses

Accelerated frame picture

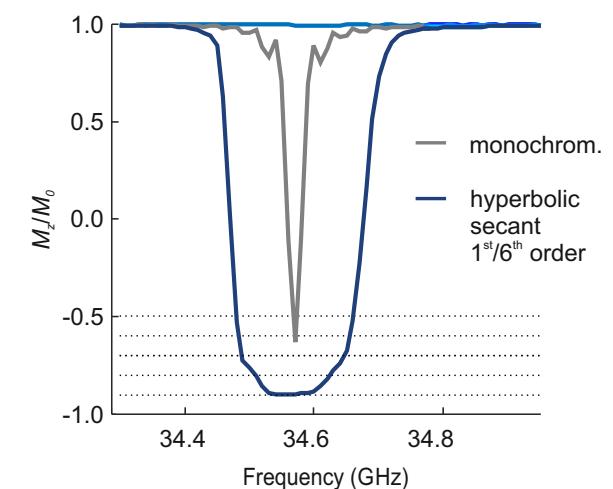


Adiabaticity

$$Q = \omega_{\text{eff}} / |\text{d}\theta/\text{dt}|$$

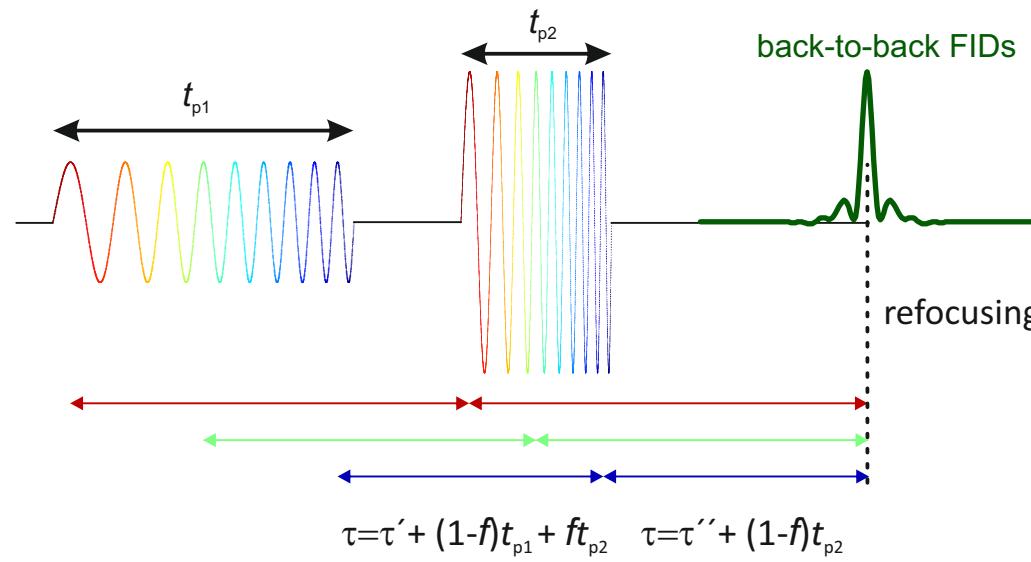
- for $Q \gg 1$ the magnetization vector follows the effective field from $+z$ to $-z$
 \Rightarrow **perfect** wideband inversion

Well, nobody is perfect



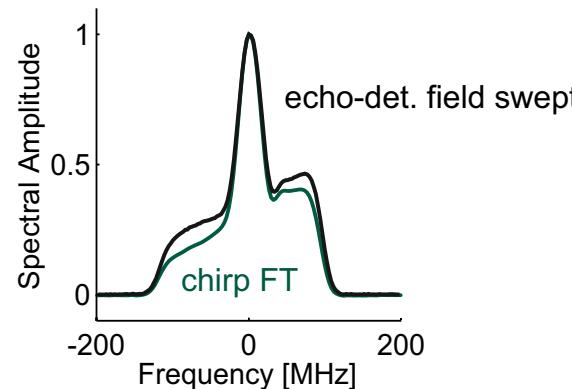
Chirp echoes

Primary chirp echo

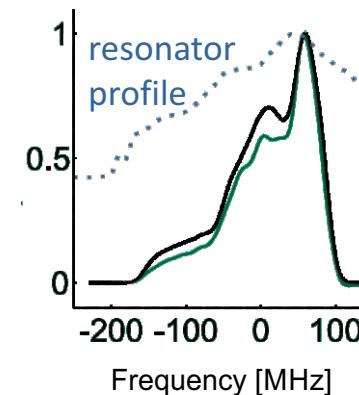


- a spin packet Ω is excited after time fraction $f = (\Omega - \omega_{\text{start}})/B$
- for all packets to refocus at the same time, we must have $\tau' + (1-f)t_{p1} + ft_{p2} = \tau'' + (1-f)t_{p2}$ for all f , which gives $t_{p1} = 2t_{p2}$
- Fourier transformation of the echo gives the frequency-domain EPR spectrum

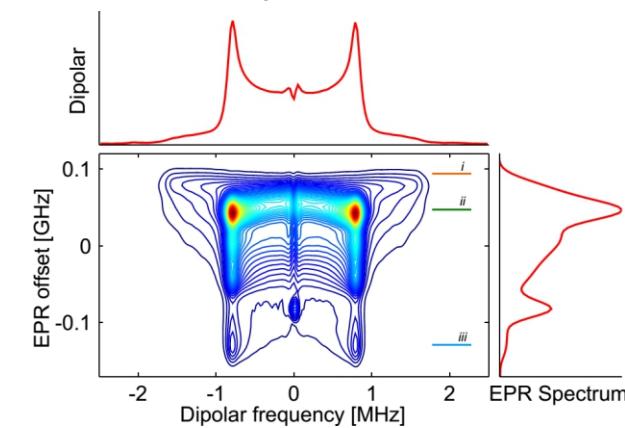
X-band chirp FT EPR



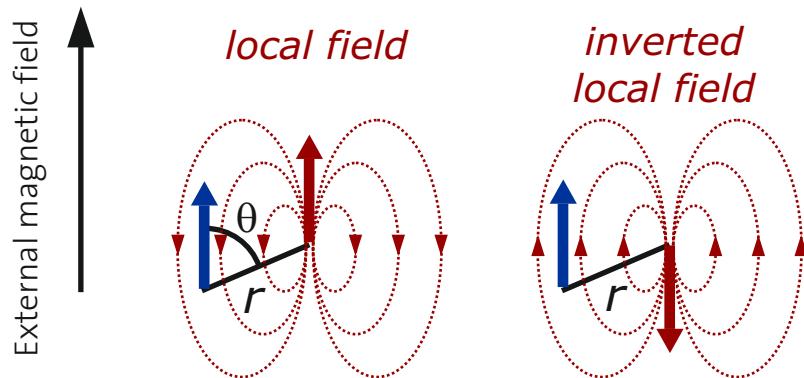
Q-band chirp FT EPR



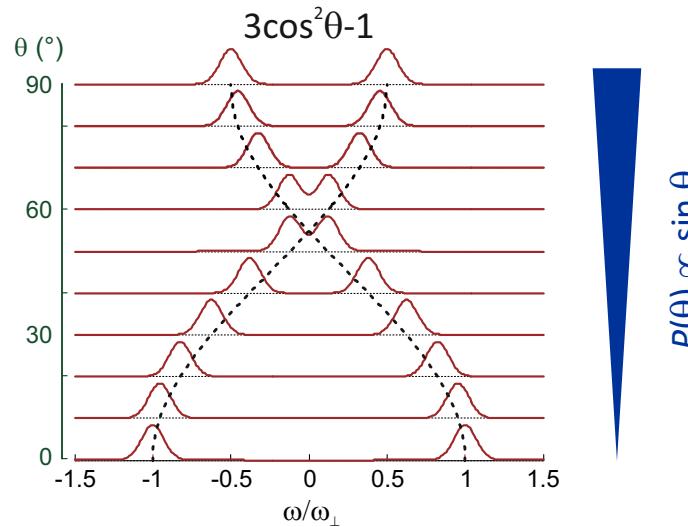
Q-band SIFTER/FT EPR correlation



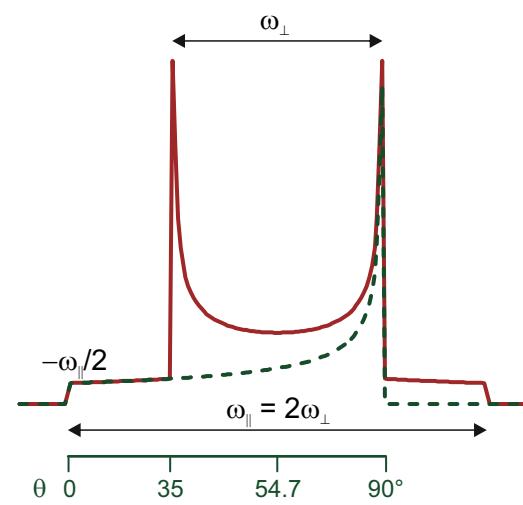
Dipole-dipole coupling



Orientation dependence



Pake pattern



spin vector
operators

$$\hat{H}_{dd} = \frac{1}{r_{SI}^3} \frac{\mu_0 \hbar}{4\pi} \gamma_S \gamma_I \left[\hat{\mathbf{S}} \cdot \hat{\mathbf{I}} - 3 \frac{1}{r_{SI}^2} (\hat{\mathbf{S}} \cdot \mathbf{r}_{SI}) (\hat{\mathbf{I}} \cdot \mathbf{r}_{SI}) \right]$$

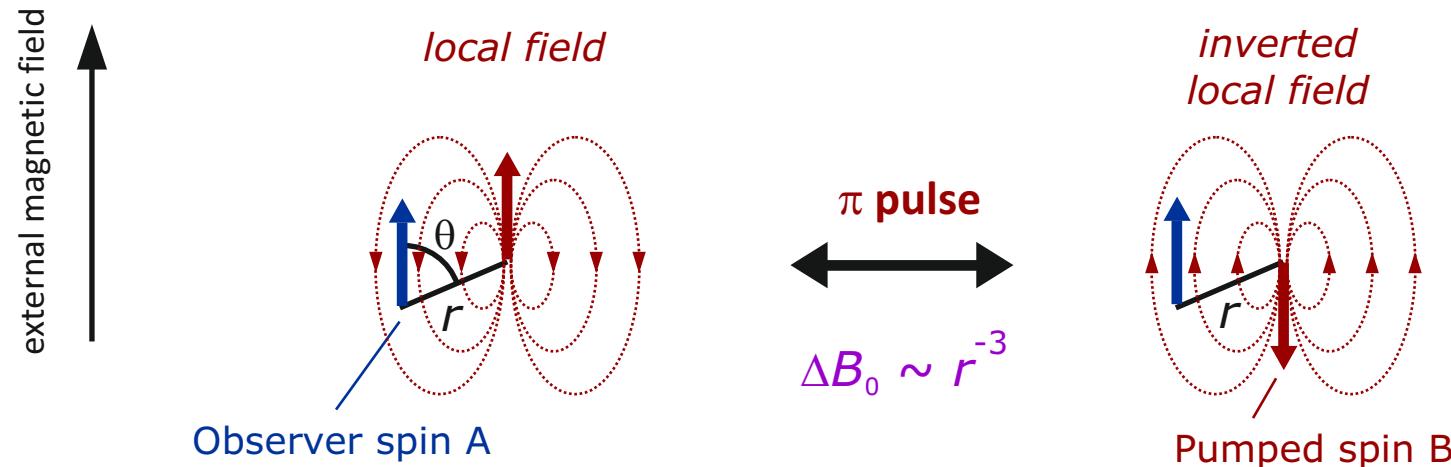
Relevant terms

$$\frac{1}{r_{SI}^3} \frac{\mu_0 \hbar}{4\pi} \gamma_S \gamma_I [3\cos^2\theta - 1] [\hat{S}_z \hat{I}_z - \frac{1}{2}(\hat{S}^+ \hat{I}^- + \hat{S}^- \hat{I}^+)]$$

To infer the distance r_{SI} , we must

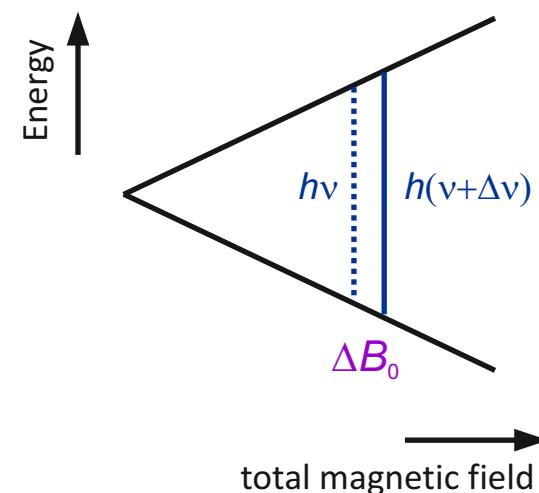
- separate dipole-dipole coupling from stronger interactions
- get rid of the orientation dependence

Separation of the electron-electron coupling from other interactions



1. Signal is labeled with resonance frequency $v = g\mu_B B_o / h$

2. Pumped spin is inverted, field shifts



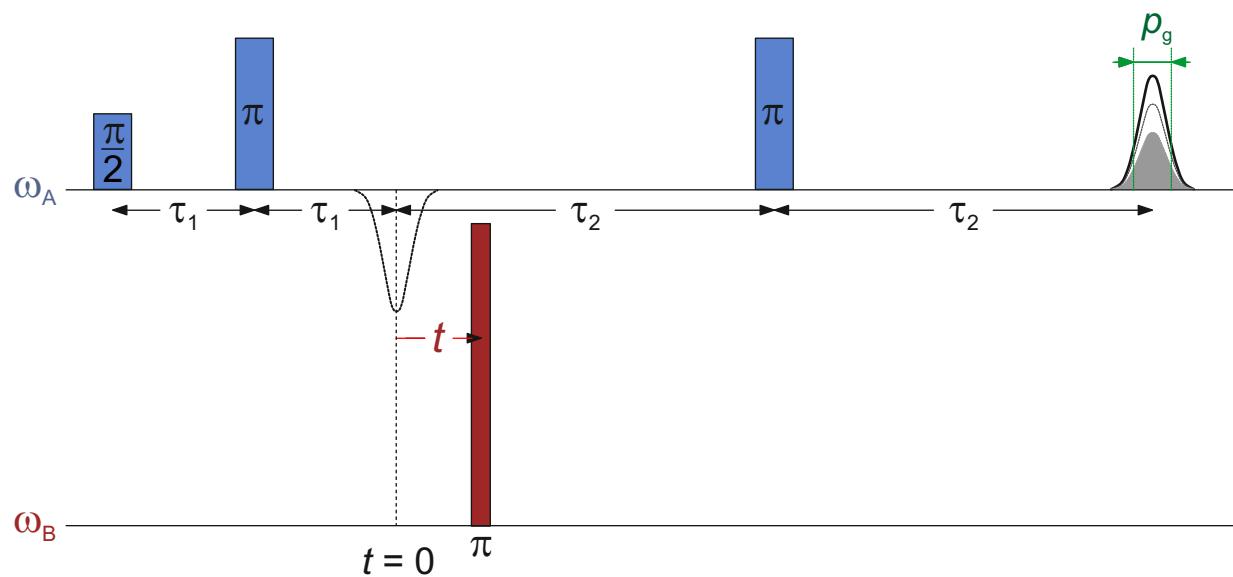
3. Field shift ΔB_0 is measured via change Δv of resonance frequency

4. Distance r is computed from Δv

$$r^3 = (52.04 \text{ MHz}/\Delta v) \text{ nm}$$

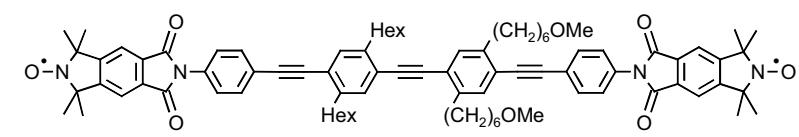
What is DEER?

a proud animal that also accepts to be named PELDOR

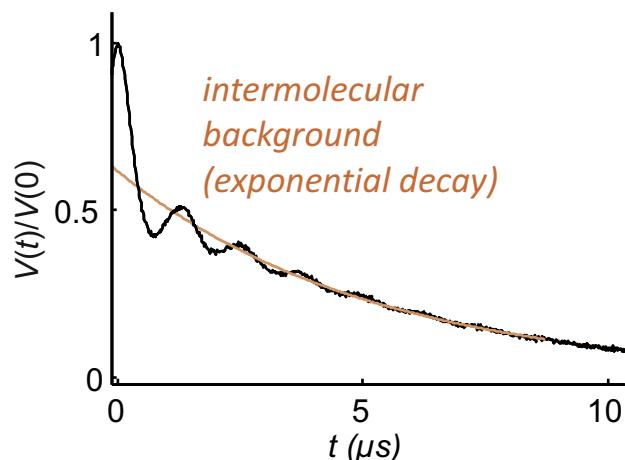


The **echo amplitude** is observed as a function of **time t**

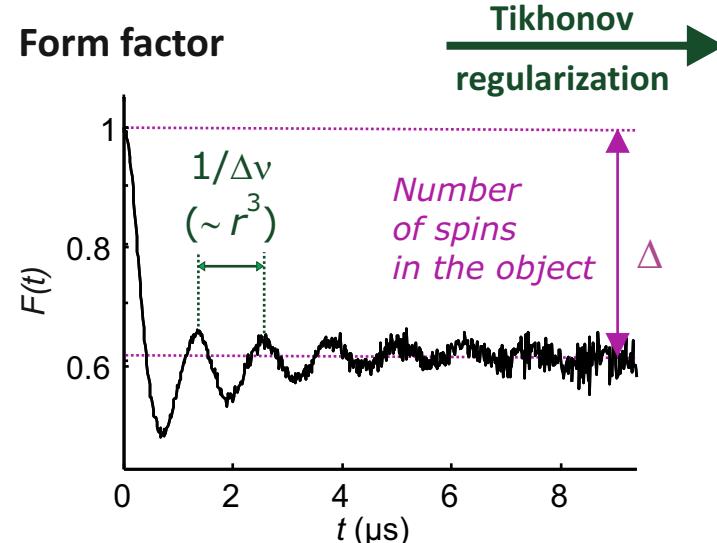
Model compound



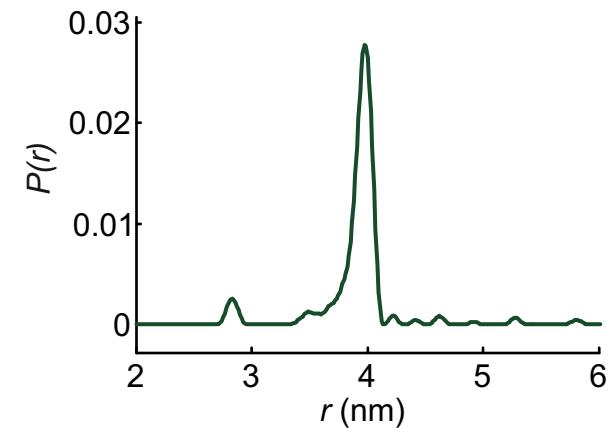
Primary data



Form factor



Distance distribution



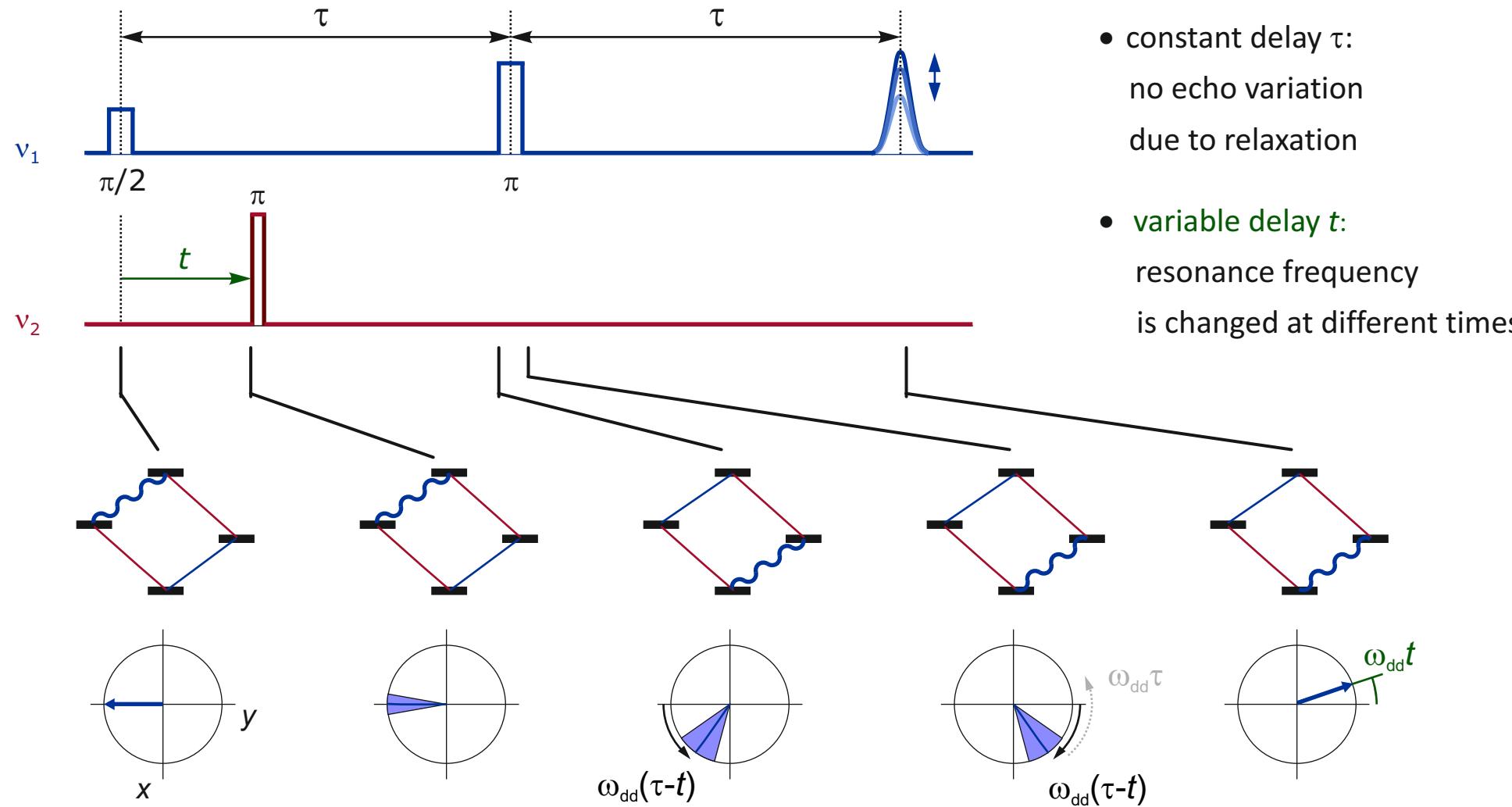
MARTIN RE et al., *Angew. Chem. Int. Ed.* **1998**, 37, 2834

PANNIER M, VEIT S, GODT A, JESCHKE G, SPIESS HW, *J. Magn. Reson.* **2000**, 142, 331

JESCHKE G et al. *J. Magn. Reson.* **2002**, 155, 72

JESCHKE G et al. *Appl. Magn. Reson.* **2006**, 30, 473

Separating the dipole-dipole interaction by a spin flip

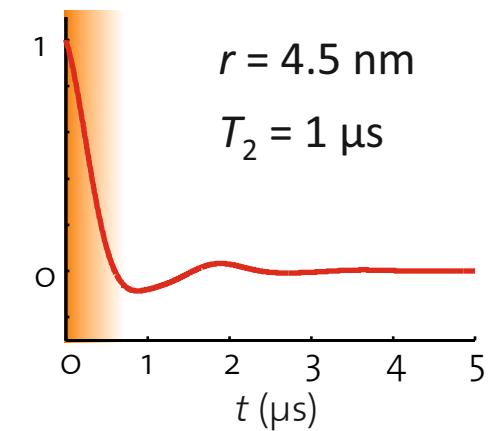
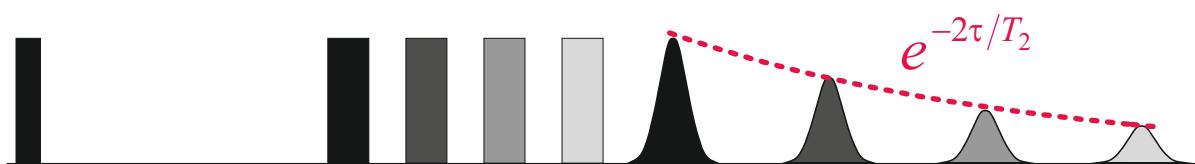


- constant delay τ : no echo variation due to relaxation
- variable delay t : resonance frequency is changed at different times

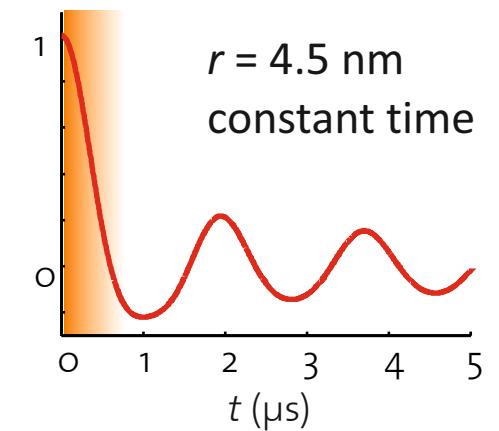
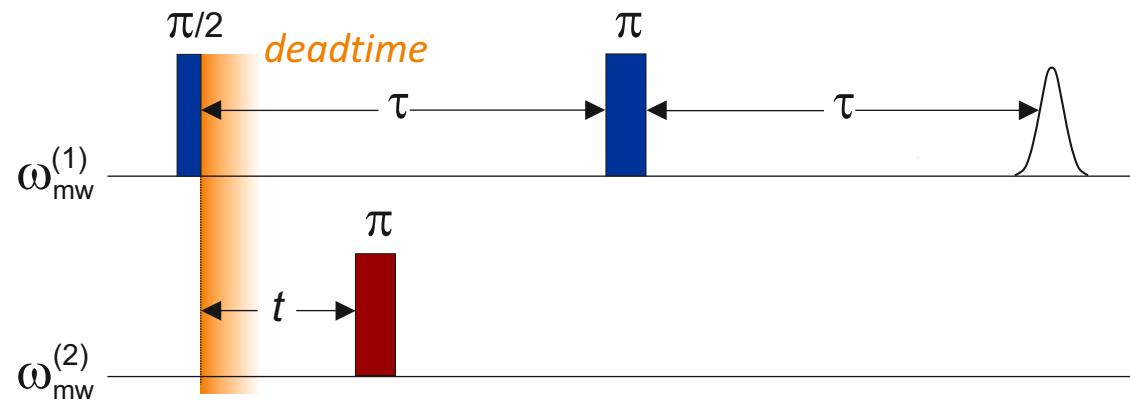
original experiment by Salikhov, Milov, Shirov, and Tsvetkov

Ultimate resolution by a constant-time experiment

- if the total length of the pulse sequence changes, dipolar evolution is damped by relaxation
- for distances larger than $\sim 3\text{-}4 \text{ nm}$, dipolar modulation is overdamped

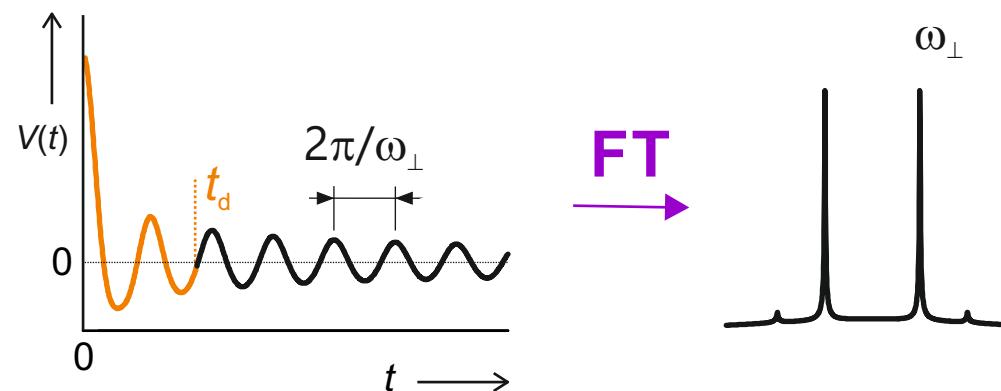


Keep τ fixed, vary only t , pulse sequence has constant length



We trade sensitivity for resolution

Eliminating dead-time by the four-pulse DEER experiment



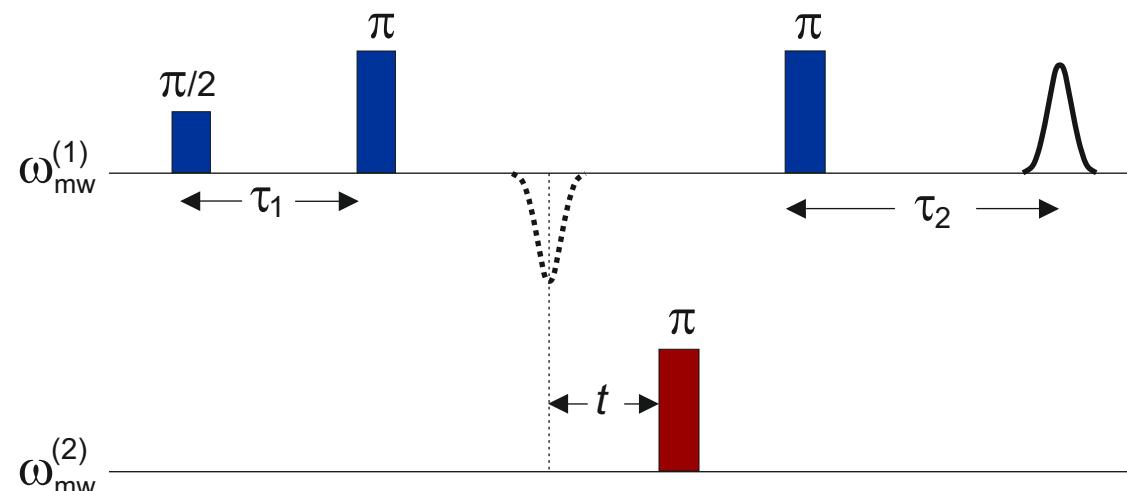
unreliable results due to deadtime for

- distributions with $r < 2.2$ nm
- direct extraction of distance distributions

sensitivity loss for small r

Remedy: Refocusing of the echo

\Rightarrow accessible $t: -\tau_1 + t_d < t < \tau_2 - t_d$



Typical values (nitroxides):

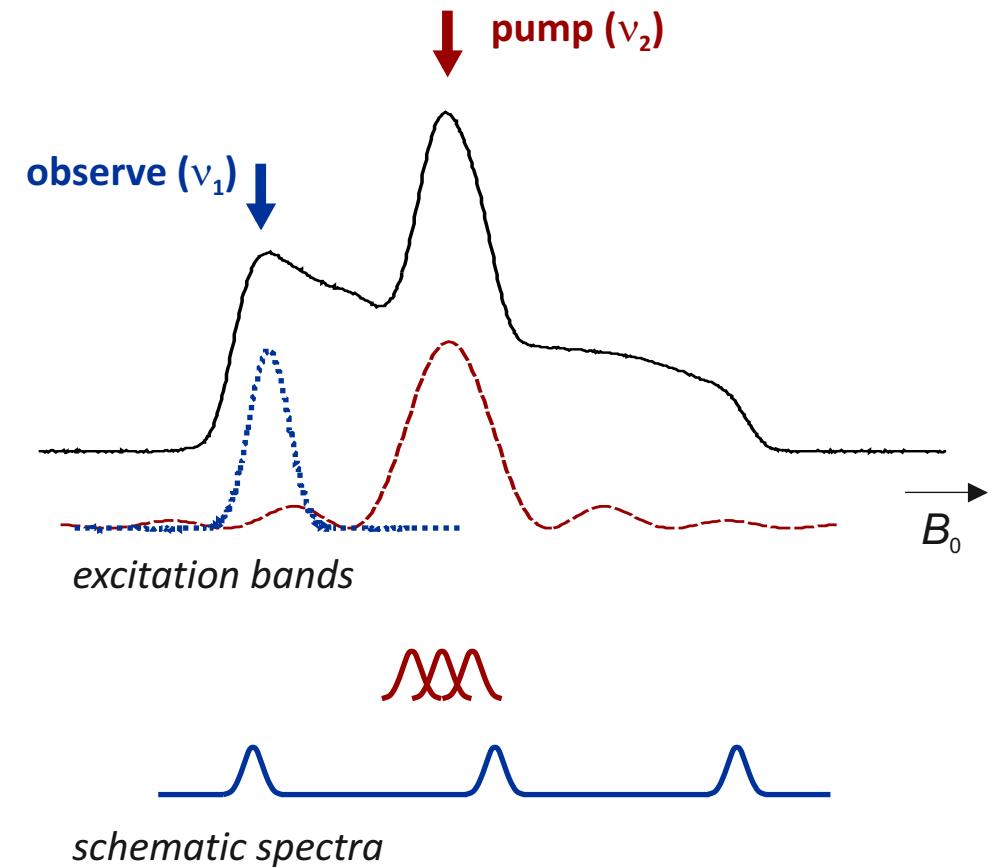
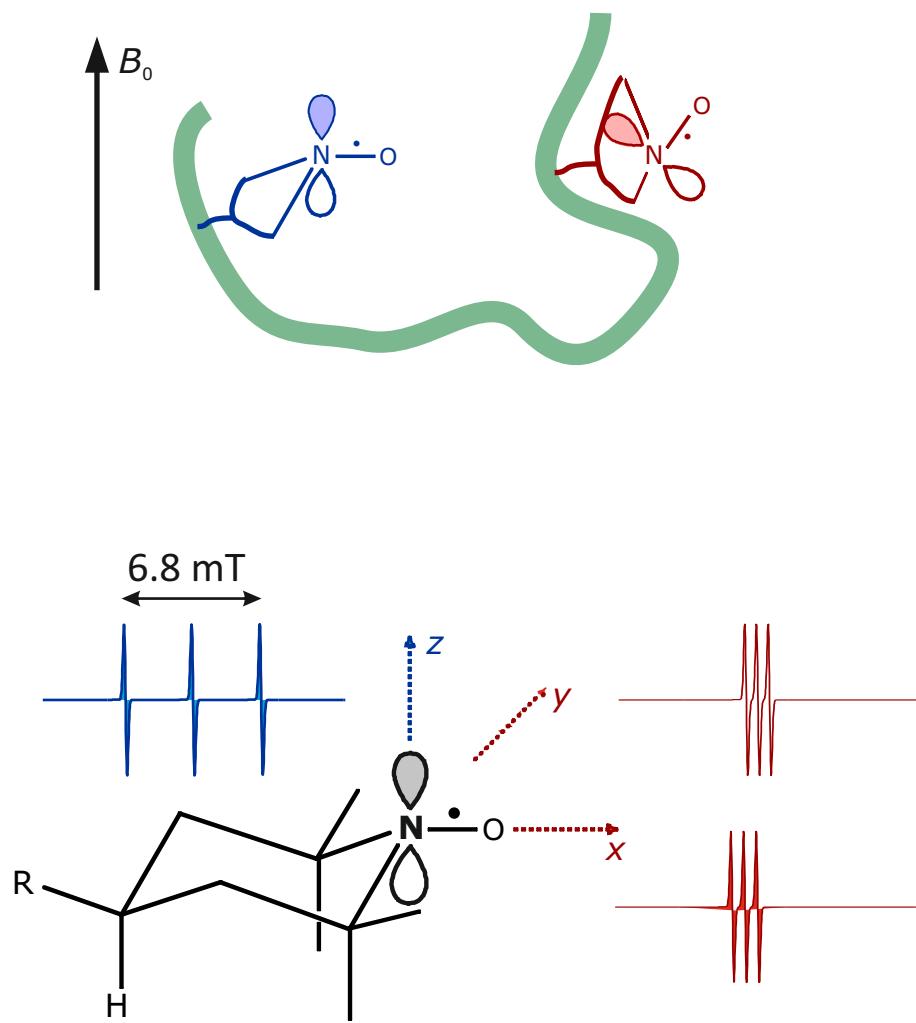
$\tau_1 \approx 120 \dots 200$ ns, 400 ns with deuterated matrix

$\tau_2 \approx 1 \mu\text{s}$ (2 nm) ... 6 μs (5-8 nm)

$t_p(1) = 32$ ns

$t_p(2) = 12$ ns (opt. sensitivity)

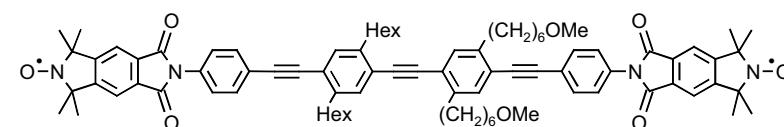
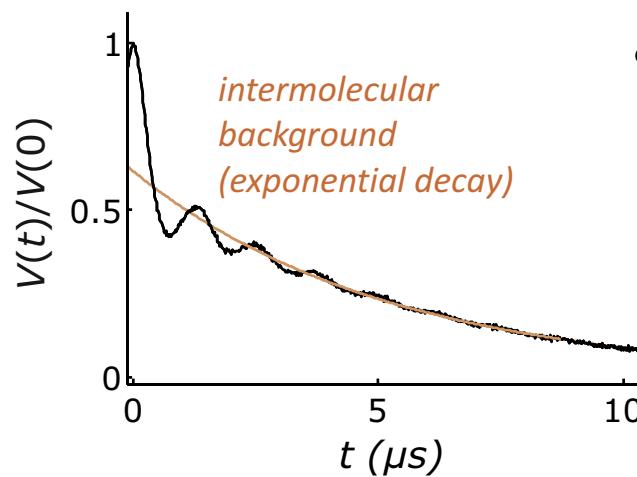
Why can we excite two nitroxide labels with two frequencies?



- only for a fraction λ of all spin pairs of the **observed spin A** the **coupled spin B** is indeed excited (pumped)

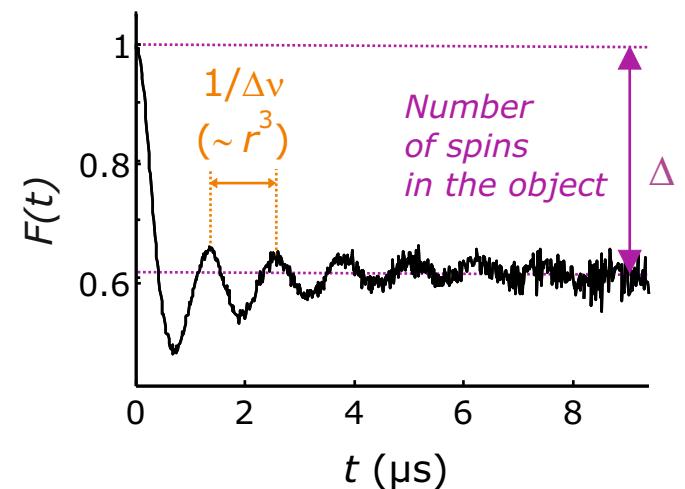
Information content of the DEER signal

Primary experimental data



background
correction

Form factor (intramolecular)



- decay envelope depends on distance distribution: fast decay of modulation \Rightarrow broad distribution

Dipole-dipole interaction $\omega_{dd} = \frac{1}{r^3} \frac{\mu_0}{4\pi h} g_1 g_2 \mu_B^2$

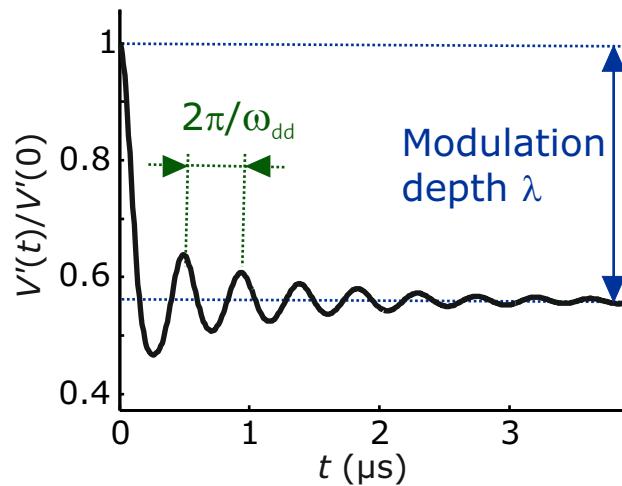
Dipolar frequency $d(\theta) = (1 - 3 \cos^2 \theta) \omega_{dd}$

Form factor
of an isolated spin pair

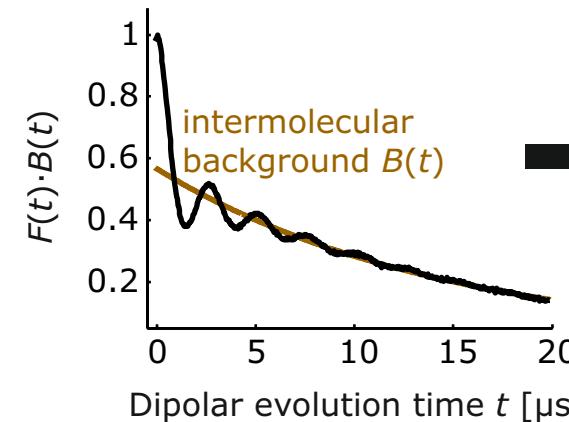
$$F(r, t) = \frac{V(t)}{V(0)} = 1 - \int_0^{\pi/2} \lambda(\theta) \{1 - \cos[2\pi v_{dd}(\theta)]\} \sin \theta d\theta$$

Data analysis by Fourier transformation

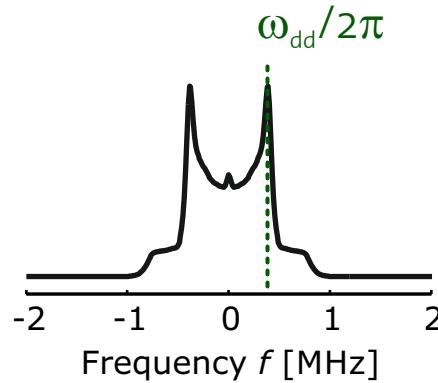
Dipolar evolution function (form factor)



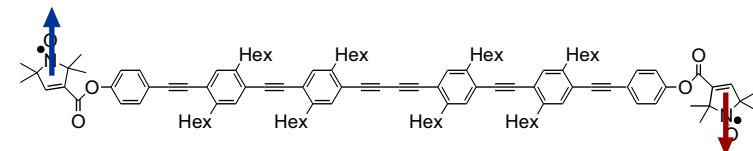
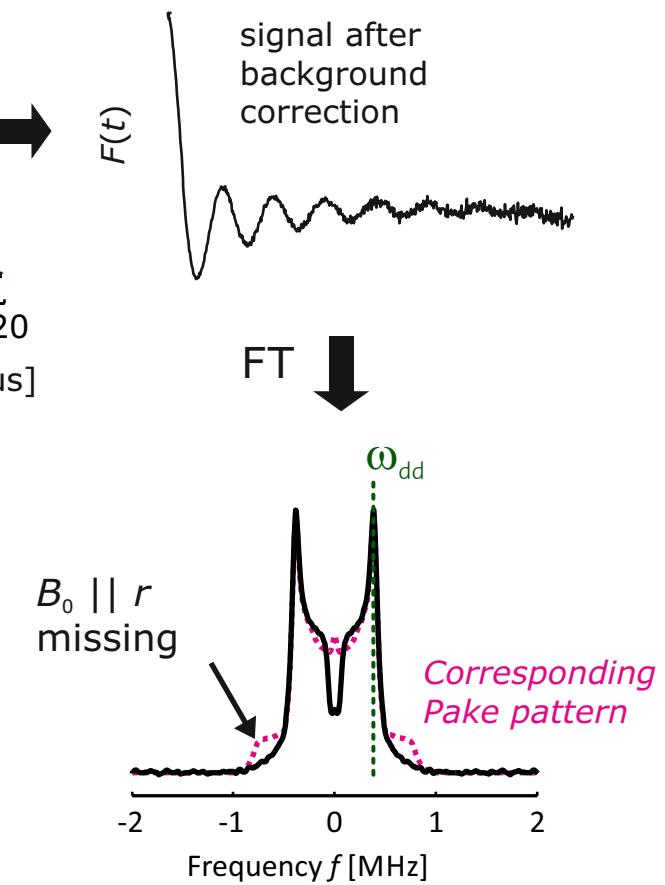
Normalized echo modulation



Fourier transformation

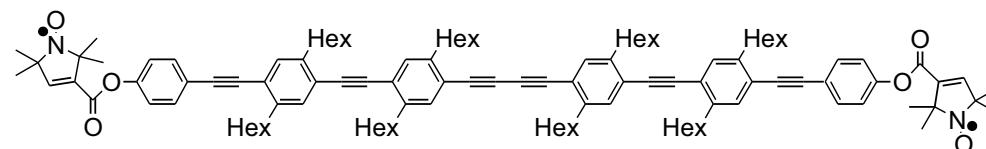


$$\begin{aligned}\omega_{dd}/2\pi &= 380 \text{ kHz} \\ \Rightarrow r &= 5.05 \text{ nm}\end{aligned}$$

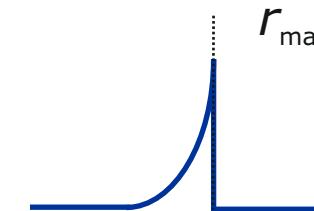


Average distance is not the full information

Molecular dynamics causes distribution of r

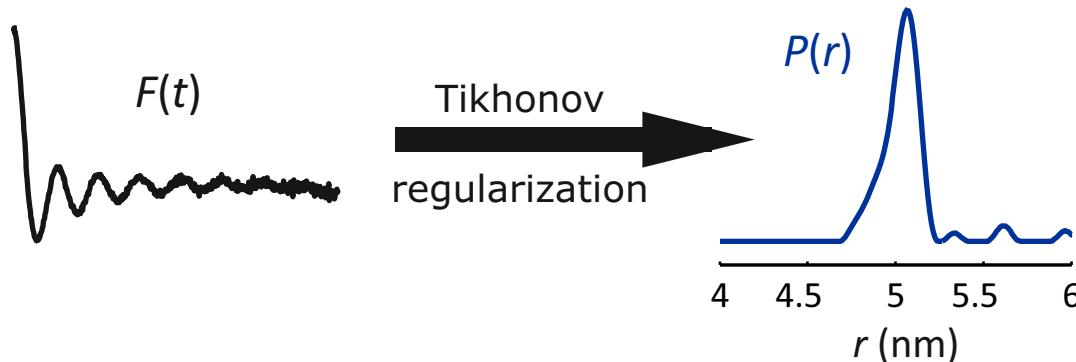


Expectation:



bent, higher energy, lower population, smaller r
stretched, min. energy, max. population, max. r

Experimental distance distribution



Basic mathematics

fit simulated dipolar evolution function $S(t)=K P(r)$

Minimize

$$G_\alpha(P) = \| \mathcal{K}P(r) - F(t) \|^2 + \alpha \left\| \frac{d^2}{dr^2} P(r) \right\|^2$$

mean square deviation

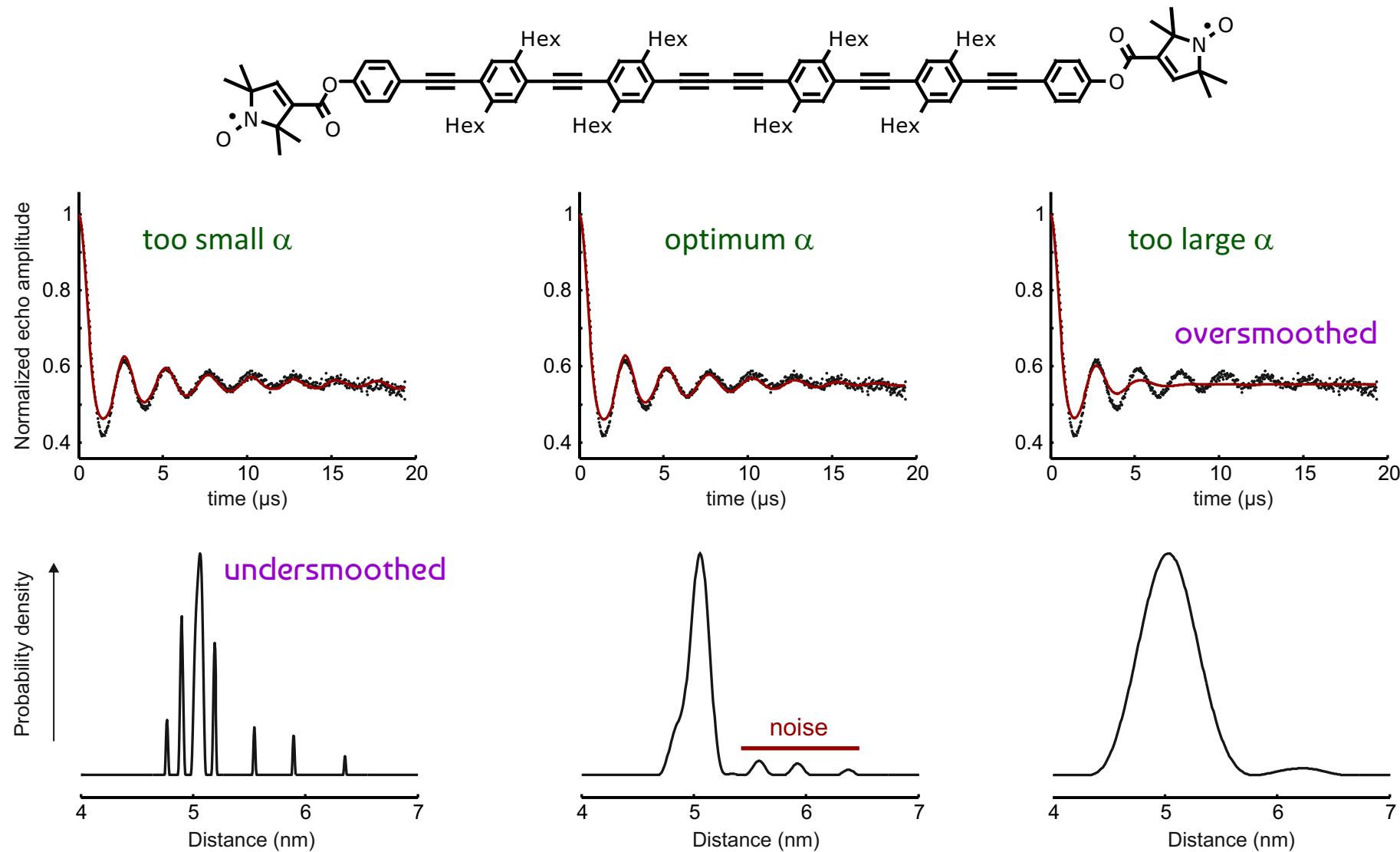
roughness of 2nd derivative

regularization parameter α

G. JESCHKE, A. KOCH, U. JONAS, A. GODT A, *J. Magn. Reson.* 155, 72-82 (2002)

G. JESCHKE et al. *Appl. Magn. Reson.* 30, 473-498 (2006)

Why regularization is required and how to do it

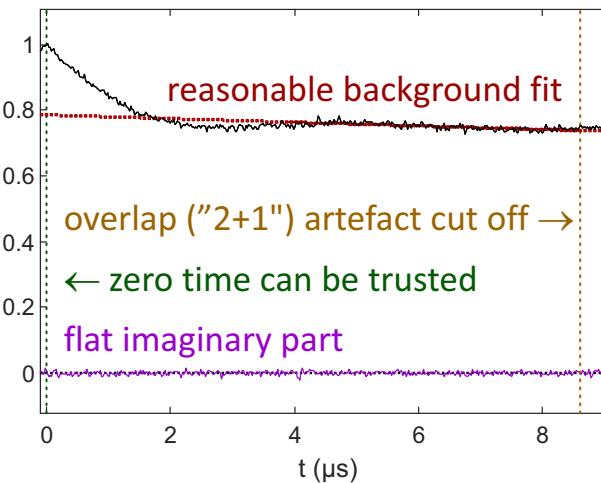


Don't forget to validate your data

Don't forget to validate your data

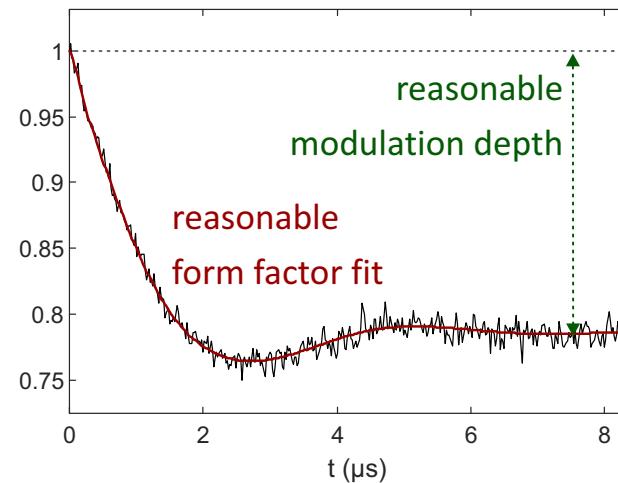
PTBP1/EMCV-IRES(D-F) 85 kDa protein-RNA complex, protein double mutant 235R1/388R1

Primary data

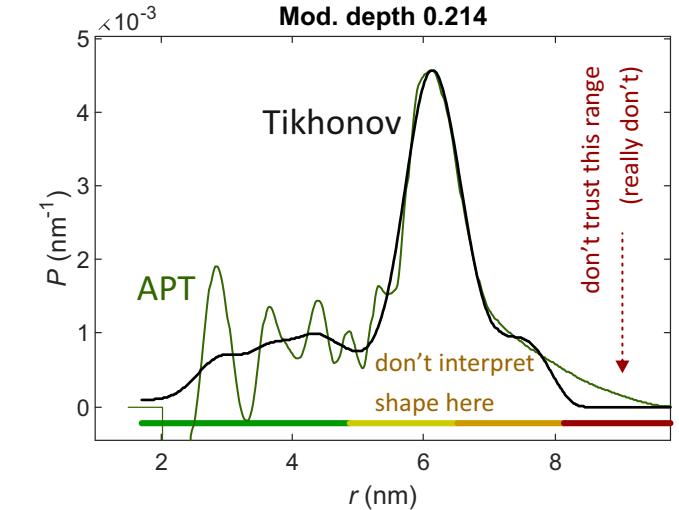


Form factor

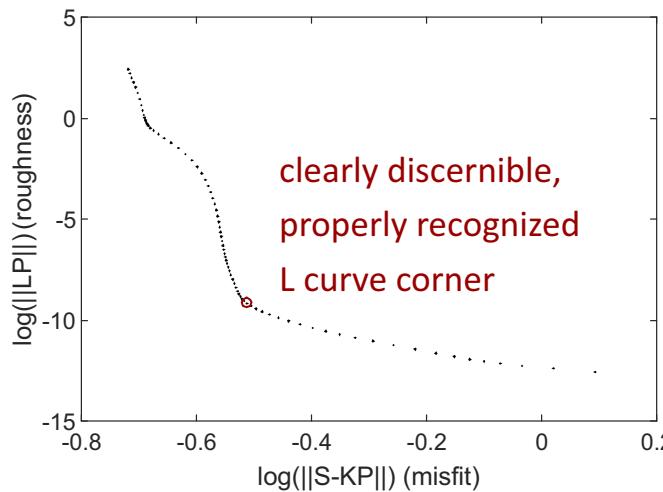
(with reasonable signal-to-noise ratio)



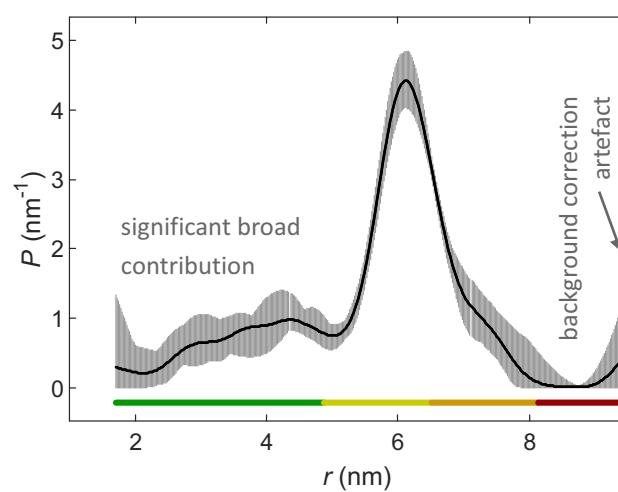
Distance distribution



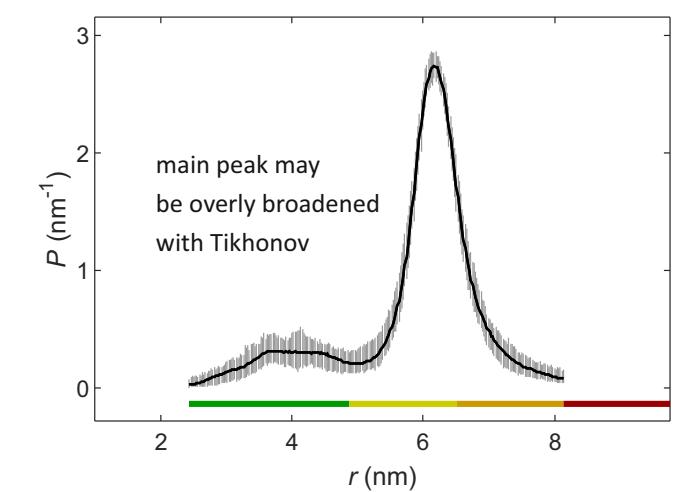
L curve



Validation



Second opinion (DEERNet)



Further reading & useful stuff on pulsed EPR

Product operator formalism and phase cycling

R.R. ERNST, G. BODENHAUSEN, A. WOKAUN, *Principles of Nuclear Magnetic Resonance in One and Two Dimensions*, Oxford University Press, Oxford, 1987.

O. W. SØRENSEN, *Progr. Nucl. Magn. Reson. Spectrosc.* 21, 503-570 (1989).

Topics specific to pulse EPR

A. SCHWEIGER, G. JESCHKE, *Principles of Pulse Electron Paramagnetic Resonance*, Oxford University Press, Oxford, 2001.

Adiabatic & fast passage

J. BAUM, R. TYCKO, A. PINES, *Phys. Rev. A* 32, 3435–3447 (1985).

A. DOLL, G. JESCHKE, *J. Magn. Reson.* 280, 46-62 (2017)

Signal processing

G. JESCHKE, *Lecture notes, ETH Zürich*

epr.ethz.ch/education/messtechnik.html

Numerical computations

EasySpin, Spinach

(Pulsed) EPR lecture notes

G. JESCHKE, *Lecture notes, ETH Zürich*

epr.ethz.ch/education.html

Product operator computations & beyond

SpinDynamica

Further reading & useful stuff on DEER & related

General overview on techniques for distance distribution measurements

P. P. BORBAT, J. H. FREED, eMagRes, Wiley, Vol. 5, 465–494 (2017). *Dipolar Spectroscopy – Single-Resonance Methods*
 G. JESCHKE, eMagRes, Wiley, Vol. 5, 1459-1475 (2016). *Dipolar Spectroscopy – Double-Resonance Methods*

Biological applications

G. JESCHKE, Annu. Rev. Phys. Chem. 63, 419-446 (2012), *DEER distance measurements on proteins*
 B. ENDEWARD, A. MARKO, V. P. DENYSENKO, S. T. SIGURDSSON, T. F. PRISNER, Methods in Enzymology
 564, 403-425 (2015) *Advanced EPR Methods for Studying Conformational Dynamics of Nucleic Acids*
 G. JESCHKE, Emerging Topics in Life Science, 2, 9-18, (2018) *The contribution of modern EPR to structural biology*

Structure modelling based on distance distributions

G. JESCHKE, Proteins 84, 544-560 (2016)
 G. HAGELUEKEN, D. ABDULLIN, O. SCHIEMANN, Methods in Enzymology, 563, 595-622 (2015) *mtsslSuite: Probing Biomolecular Conformation by Spin-Labeling Studies*
 G. JESCHKE, Protein. Sci. 27, 76-85 (2018)
MMM - A toolbox for integrative structure modelling

DeerAnalysis & MMM

epr.ethz.ch/software.html

deeranalysis.org

