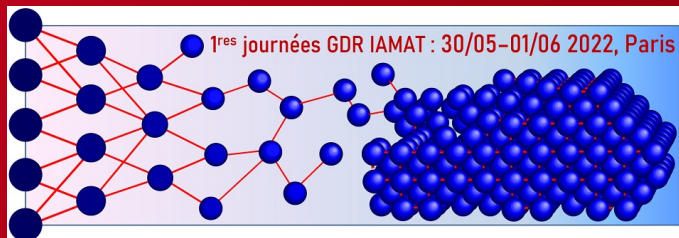


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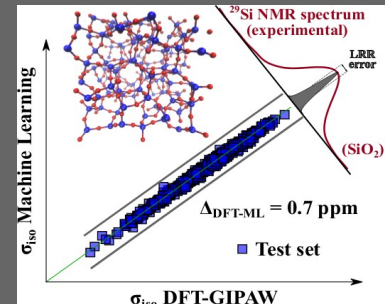
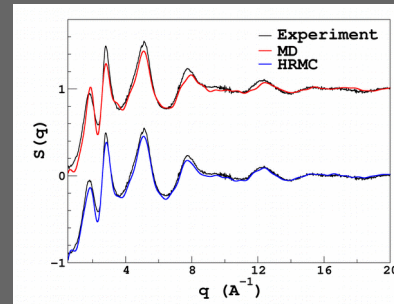
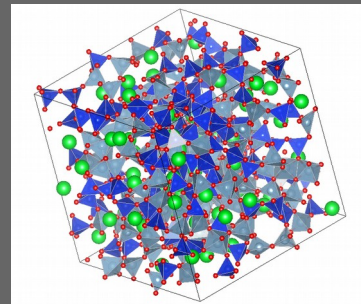
30 mai - 1 juin 2022
Sorbonne Université
Paris



1^{ères} journées GDR IAMAT : 30/05-01/06 2022, Paris

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Modelling NMR Spectroscopy of Oxide Glasses with Machine Learning



Thibault Charpentier
Université Paris-Saclay, CEA, CNRS,
NIMBE, CEA Saclay France
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IRAMIS

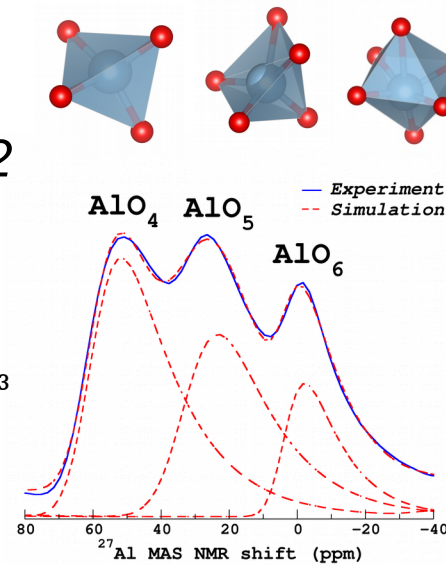
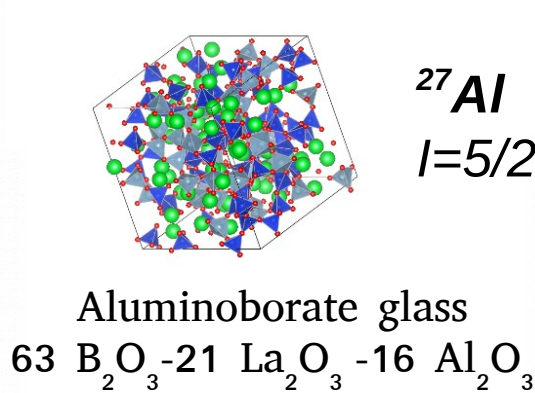
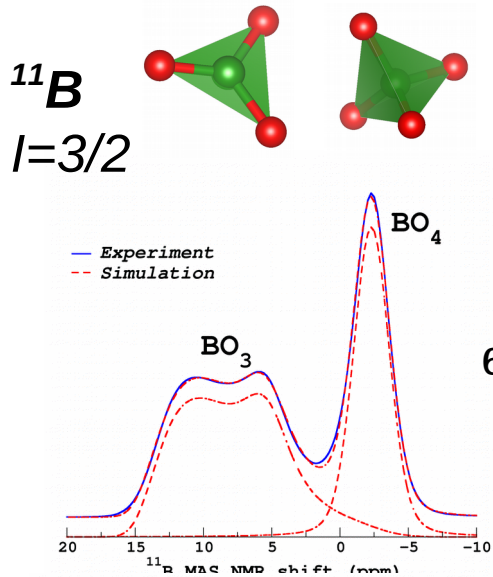


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CONTEXT: MAS NMR IN OXIDE GLASSES

MAS NMR of Quadrupolar Nuclei in Glasses

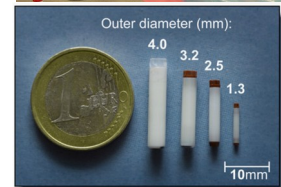
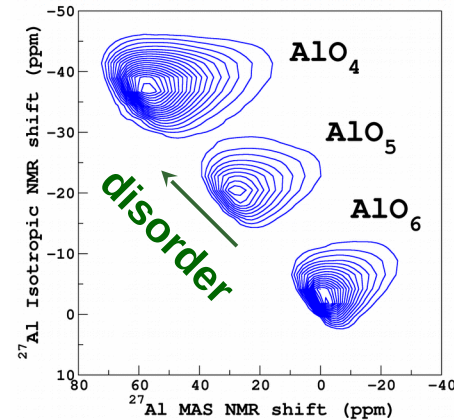
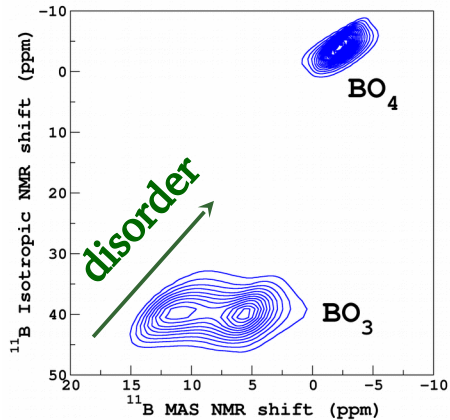
Deciphering the Short Range Order (SRO) in Glasses



1D MAS NMR

Gain in resolution with 2D NMR

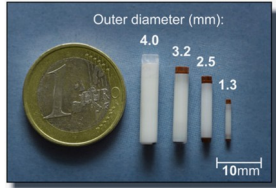
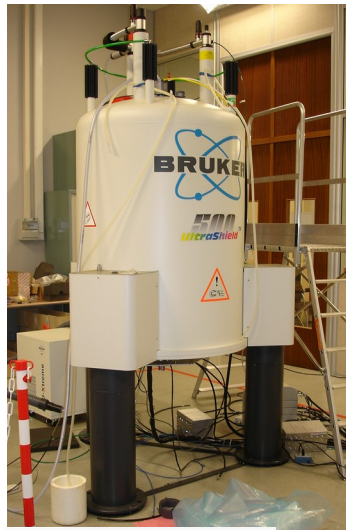
2D MQMAS NMR (Triple Quantum MAS)



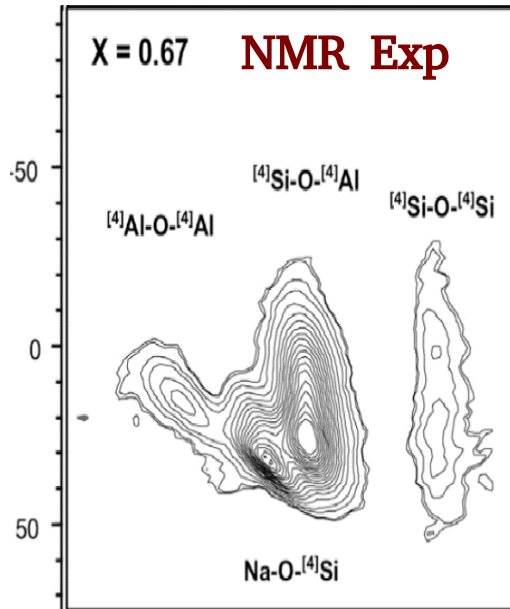
High-resolution MAS NMR reveals the structural motifs (network formers) building the glass network. NMR lineshapes are reflective of the local disorder and reveals the distribution of NMR parameters.

MOTIVATIONS: NMR from first-principles

Coupling MD simulations with NMR Experiments through DFT/NMR computations (GIPAW)

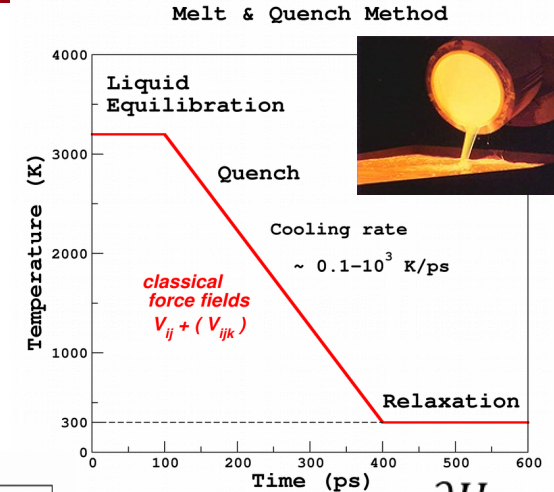


^{17}O MQMAS (11.7 T)
~1-4 days



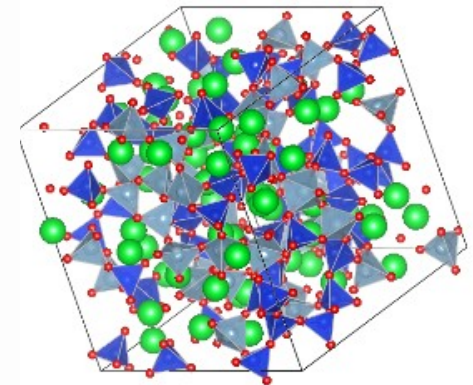
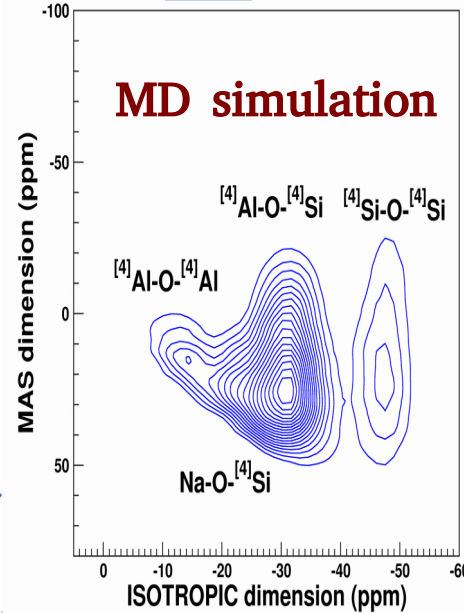
Experiment Interpretation

MD assessment



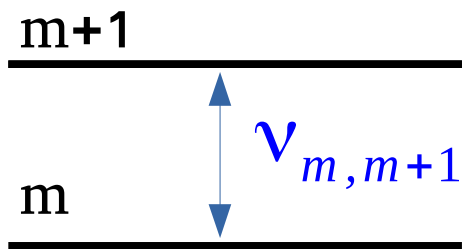
$$m_i \ddot{r}_i = F_i = - \frac{\partial U}{\partial r_i}$$

MD simulation



VASP-GIPAW, 800at.
~ 5h, 96 procs

NMR transitions



$$H_{NMR} = -\hbar \gamma_i \vec{I} \cdot \underbrace{(1 - \sigma)}_{\text{DFT}} \cdot \vec{B}_0 + \vec{I} \cdot \vec{V} \cdot \vec{I}$$

NMR

MAS NMR simulation :

$$\nu_{m,m+1} = \delta_{iso} + \nu_{Q,(m,m+1)}^{(2)}$$

Magnetic Shielding Tensor

$$\sigma = \frac{\delta^2 E_0}{\delta \vec{u} \delta \vec{B}_0}$$

Isotropic magnetic shielding : $\sigma_{iso} = Tr \{ \sigma \}$
(isotropic chemical shift) scalar

today

Electric Field Gradient Tensor (Traceless Symmetric Matrix)

$$V_{\alpha,\beta}(r) = \int du \frac{n_0(r)}{|r-u|} \left\{ \delta_{\alpha,\beta} - 3 \frac{(r_\alpha - u_\alpha)(r_\beta - u_\beta)}{|r-u|^2} \right\}$$

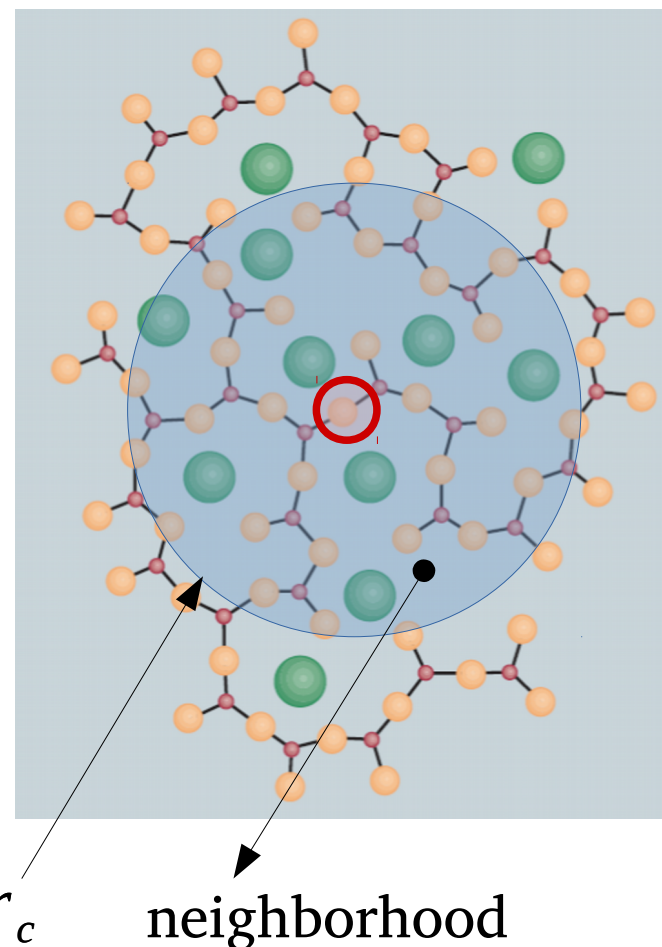
matrix

Atomic Centered Descriptors

Chemical and Geometrical Properties

It is assumed that the quantity of interest is **local** (e.g., NMR shift). The Cartesian coordinates of each atom cannot be used as such in ML. They must be transformed into a set of descriptors that fulfill some invariance/symmetry properties.

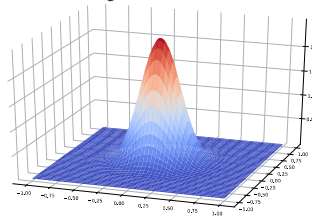
- Structural description (faithfulness)
- Translational invariance
- **Rotational invariance for a scalar quantity**
(Energy, **NMR isotropic chemical shift**, ...)
- Atomic permutation independent
- Zero value and derivative at cutoff radius
(locality, short range)
- Continuous in value and derivative (smoothness)
- Of course, the design of descriptors must be driven by the properties of interest...



Atom-Centered Descriptors Smooth Overlap of Atomic Positions (SOAP)

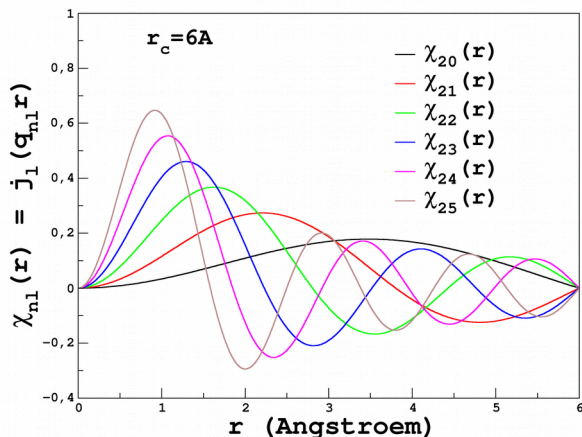
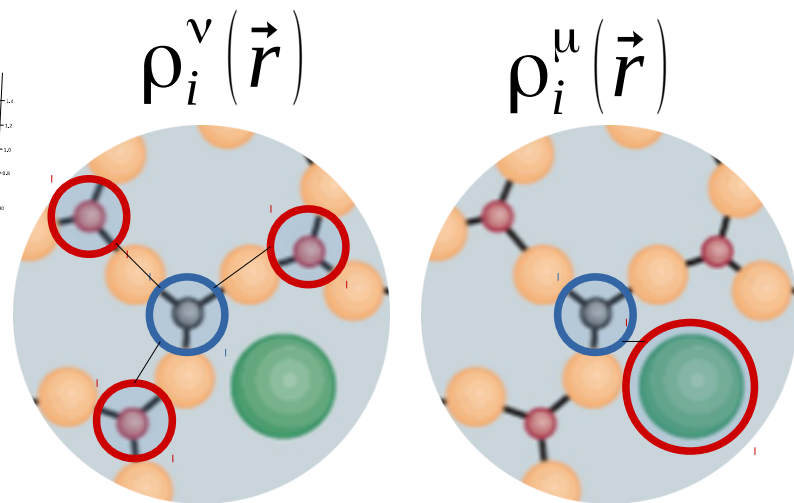
Smooth expansion of the atomic density with a radial cutoff function

$$\rho_i^v(\vec{r}) = \sum_{j \in N_i^v} G_\sigma(\vec{r} - \vec{r}_{ij}) f_c(r_{ij})$$

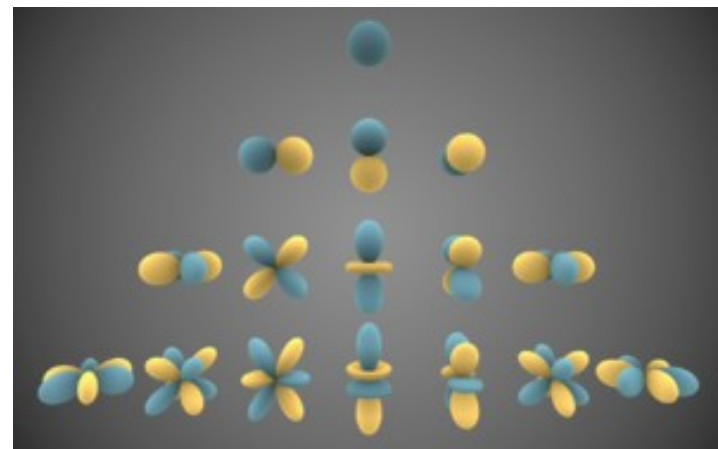


Spherical Harmonics & radial basis expansion

$$\rho_i^v(\vec{r}) = \sum_{nlm} C_{nlm}^v \chi_{nl}(r) Y_{lm}(\hat{r})$$



We found the spherical Bessel functions to be superior to other functions for the radial Basis Expansion. They form a natural orthonormal basis on the segment $[0, r_c]$.



A.P. Bartok et al. (2013) On representing chemical environments, Phys. Rev. B

R. Jinnouchi et al. (2019) On-the-fly machine learning force field generation: Application to melting points, Phys. Rev. B

E. Kocer et al. (2020) Continuous and optimally complete description of chemical environments using Spherical Bessel descriptors, AIP Advances 10, 015021

Smooth Overlap of Atomic Positions (SOAP)

$$\rho_i^v(\vec{r}) = \sum_{nlm} c_{nlm}^v \chi_{nl}(r) Y_{lm}(\theta, \phi)$$

The problem is that the only **rotational invariant** descriptors are obtained for $l=m=0$, which limits severely the number of descriptors (typically $l_{\max}=4$, $n_{\max}=8$)

$$c_{n00}^v$$

2-body terms

One option is to couple channels in such a way that they form rotational invariants:

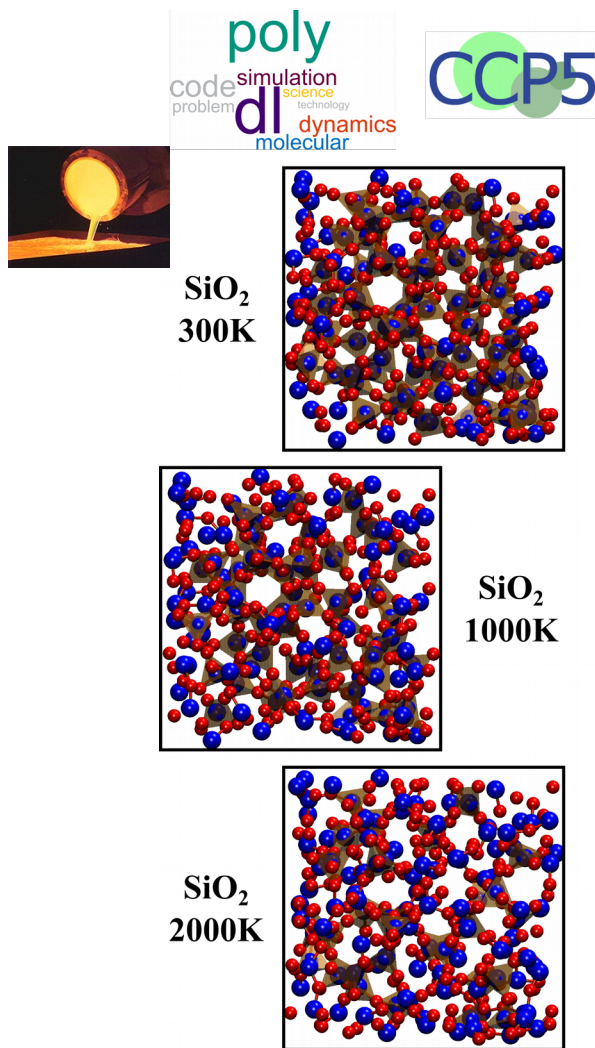
$$c_{nl+m}^v \times c_{n'l-m}^u$$

This is the so-called **SOAP** power spectrum

$$p_{nn'l}^{v\mu} = \sum_m c_{nl+m}^v \times c_{n'l-m}^u$$

3-body terms

Methodology : DATABASE Generation



DFT-GIPAW

QM/MM



ab initio MD

Scalar (Nmr shifts, charges)
Vector (Forces, dipole moments)
Tensors (Polarizability, Stress tensor)

Database for Machine Learning (ML)

Machine learning engine

Invariant descriptors construction
(Atom-centered functions)

ML algorithms
(Regression, classification)



Cross-Validation
(Optimize hyperparameters)

10-20 MD models
(400 at.) for each T

OUTPUT (DFT-NMR)
10³-10⁴ NMR shift values

ML Box

Kernel Ridge Regression builds a linear model with a **non-linear** similarity measurement (Kernel) :

$$\delta(\chi) = \sum_{(j \in \text{train})} \alpha_j K_G(\chi, \chi_j)$$

$$\alpha = \left(\mathbf{K}_G^{\text{train}} + \lambda \mathbf{I} \right)^{-1} \delta^{\text{train}}$$

χ : local environment (descriptors)

χ_j, δ_j : database (train)

α_j : regression parameters

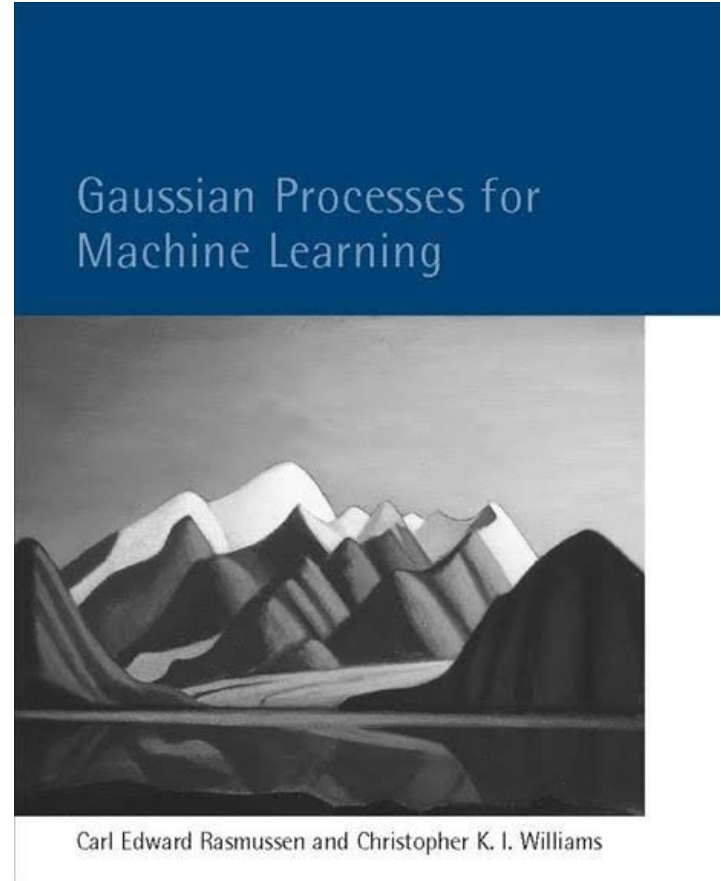
$$\text{Kernel: } K_G(\chi, \chi_j) = e^{-\kappa \|\chi - \chi_j\|^2}$$

λ : ridge parameter

κ : Kernel width

hyperparameters

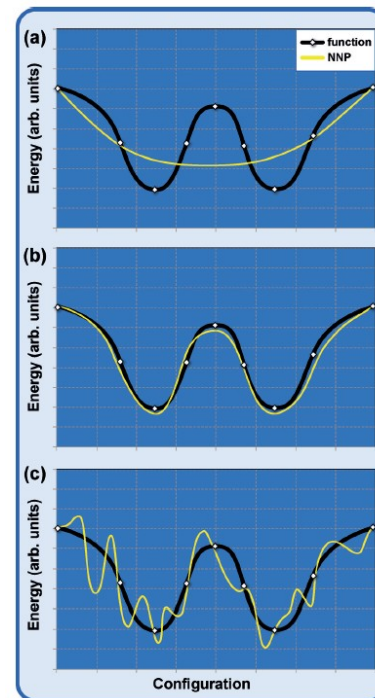
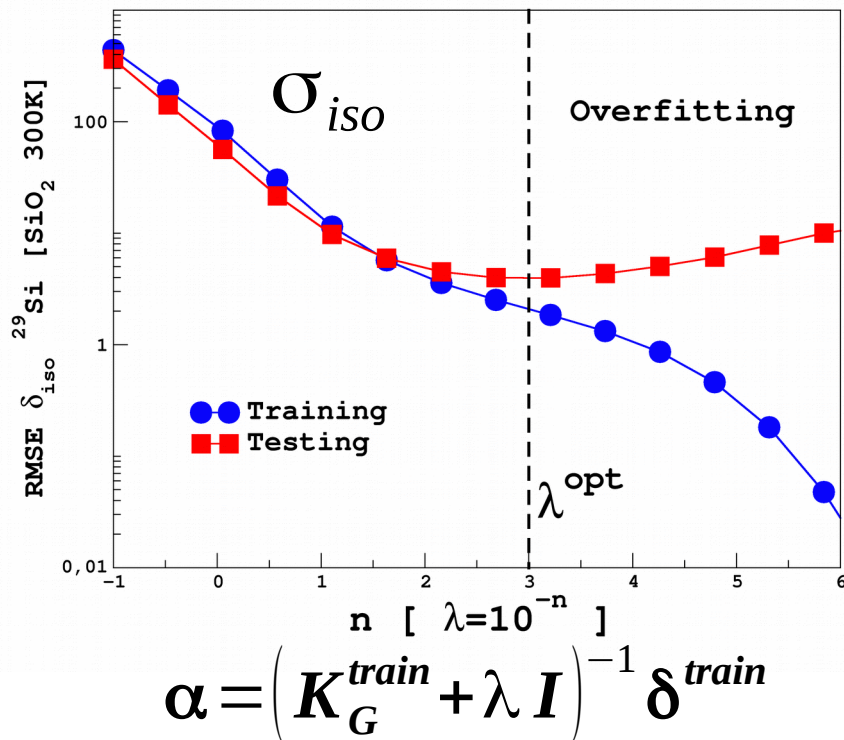
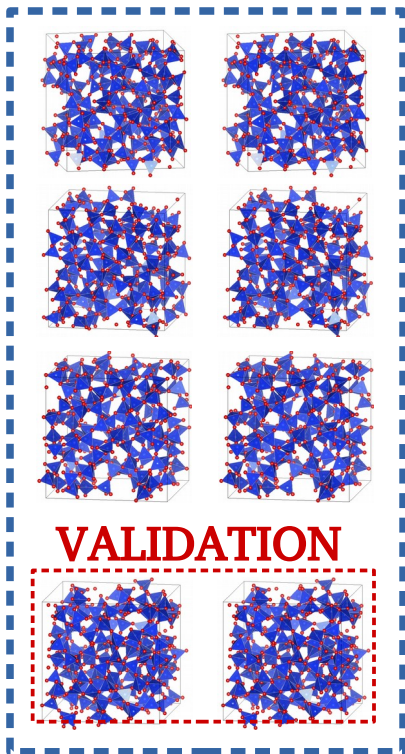
The ridge parameter prevents from overfitting and ensures a good transferability.



Kernel Ridge Regression : Hyperparameters optimization

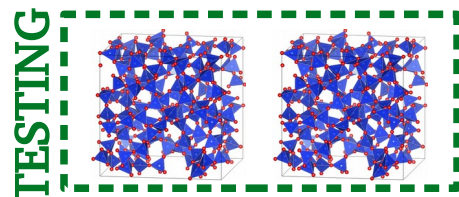
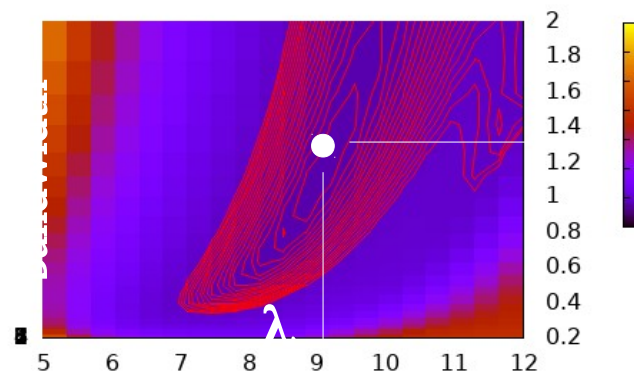
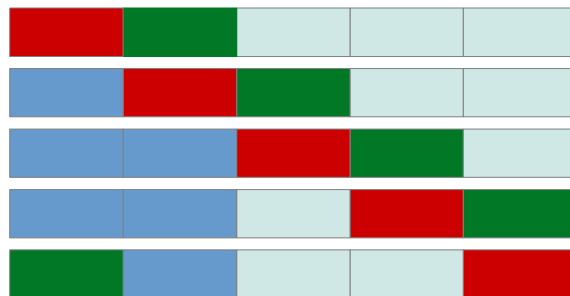
J. Behler et al. Angewandte Chemie 2017

DATABASE

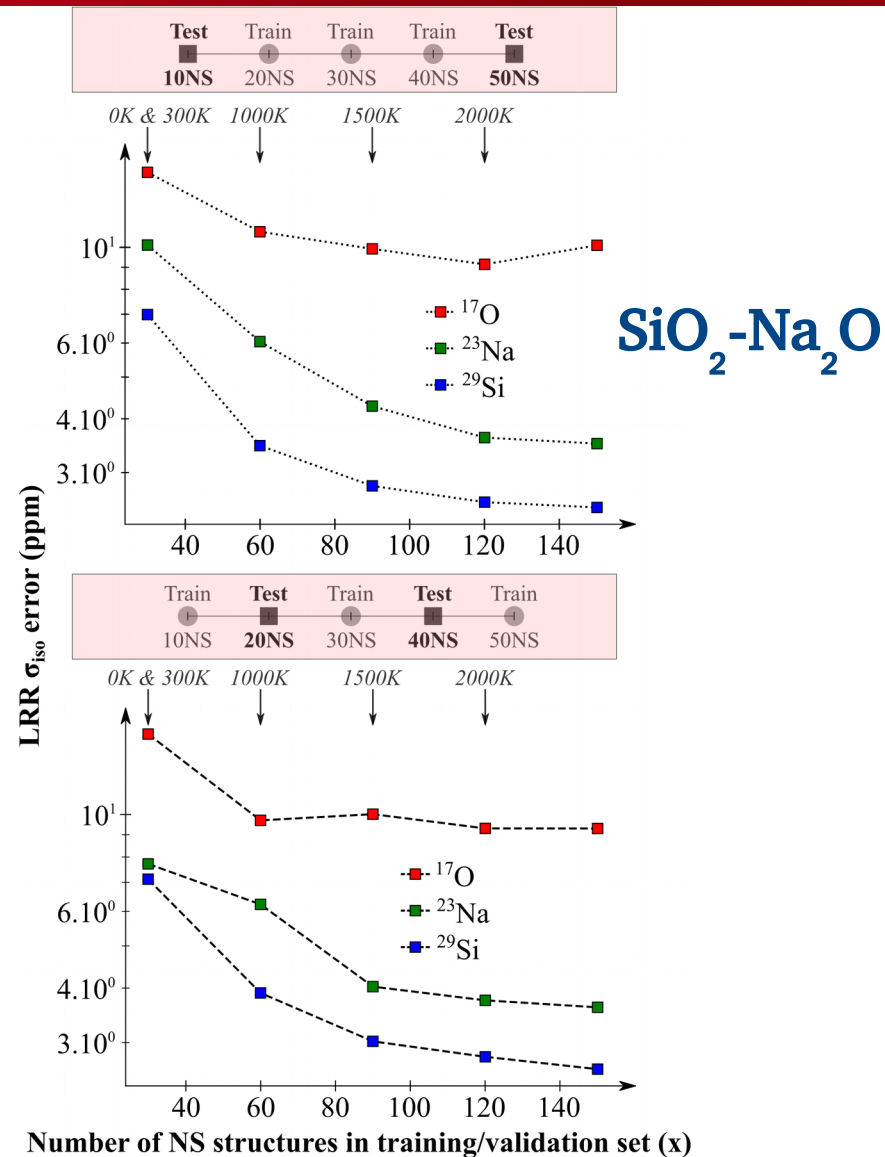
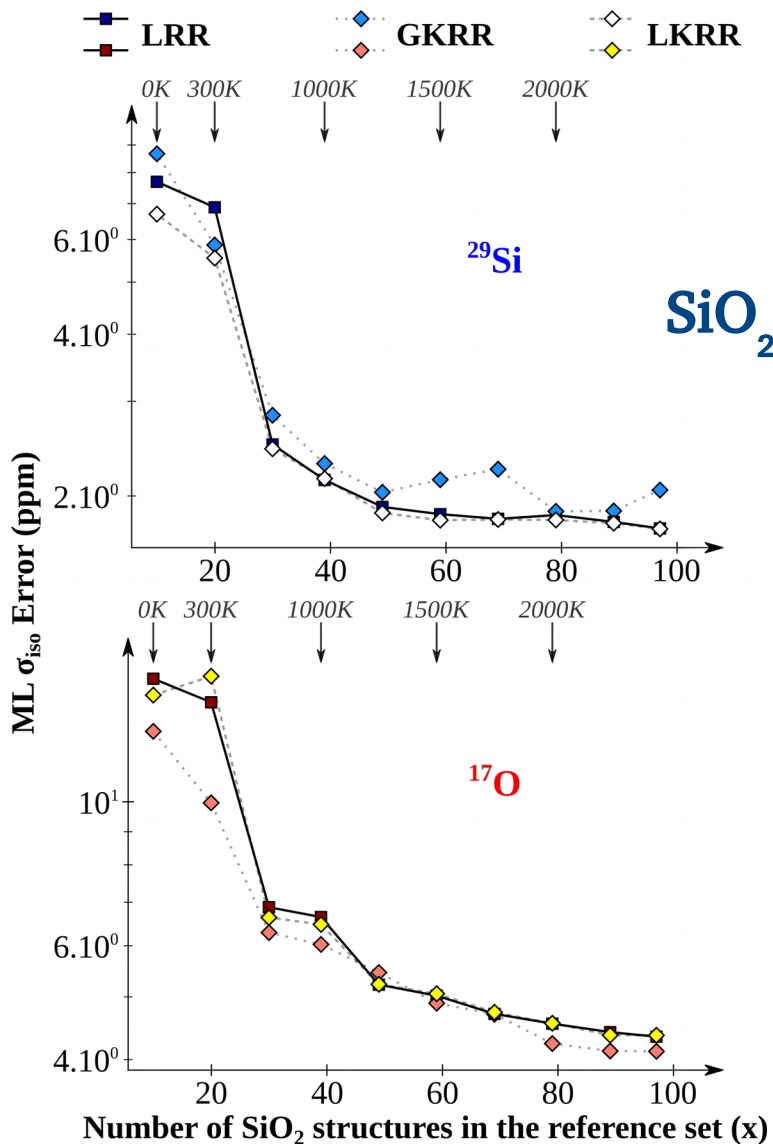


λ and κ must be optimized on a validation set before testing.

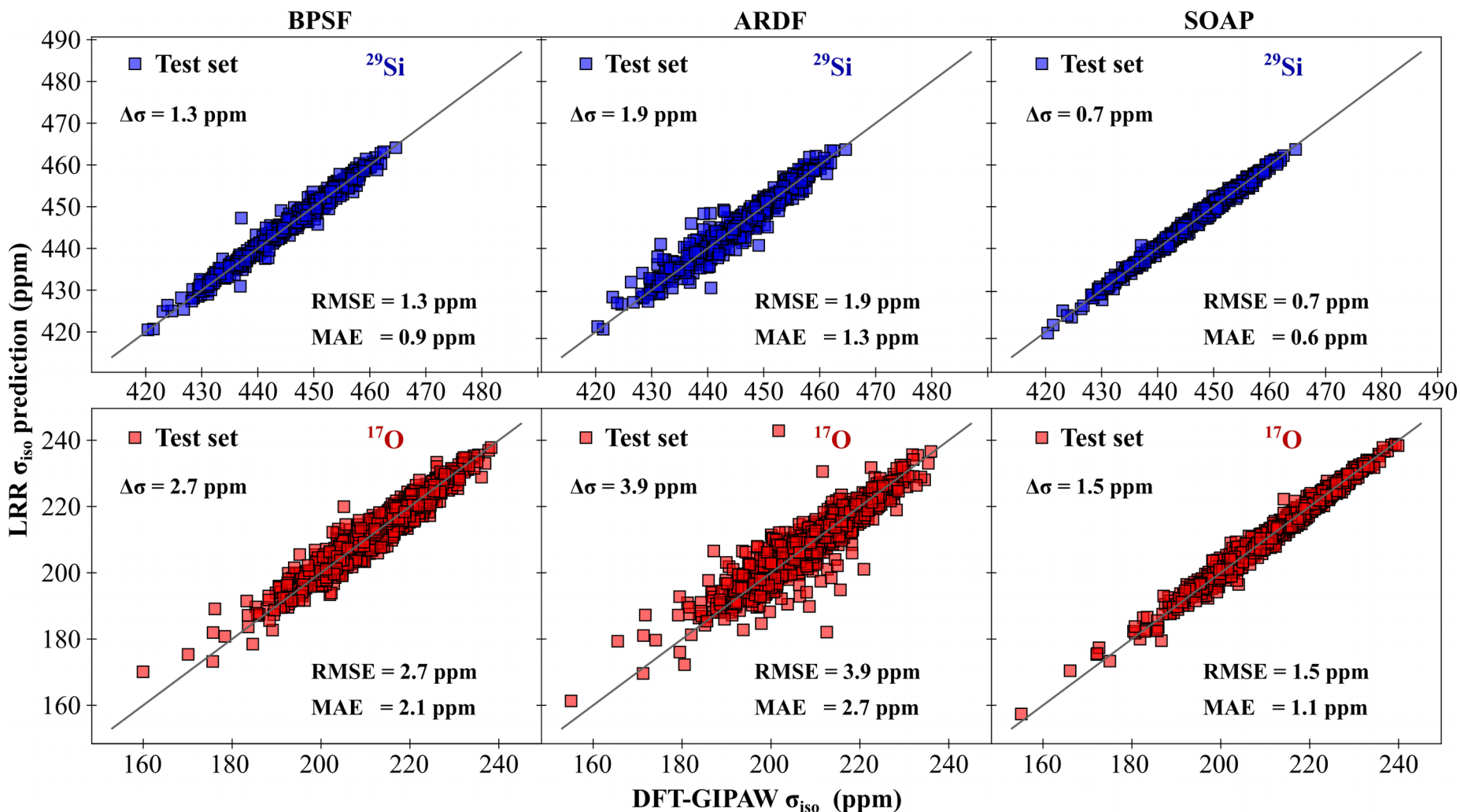
k-fold Cross Validation



Machine Learning NMR shifts : Database Generation MD-generated structures at various Temperatures



Machine Learning NMR shifts : Descriptor's Performances



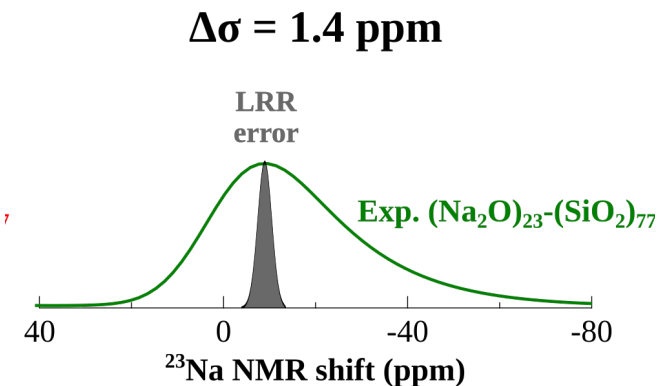
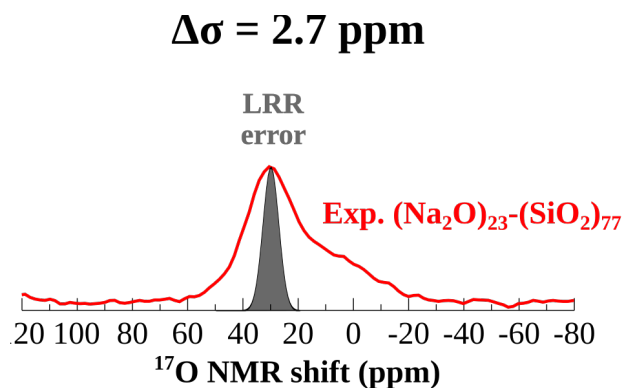
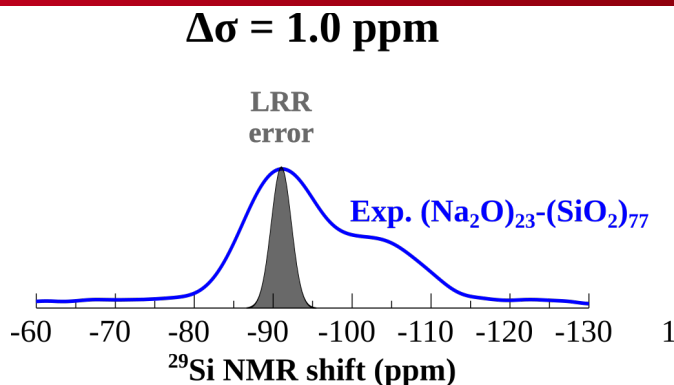
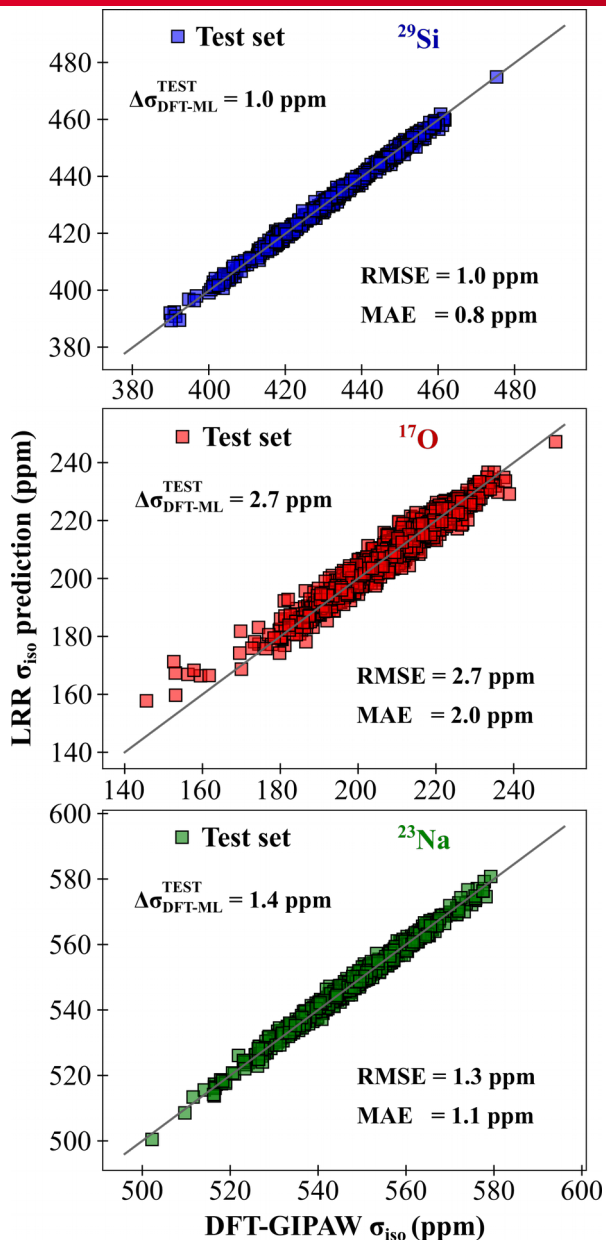
Symmetry Functions (**BPSF**) : J. Behler and M. Parrinello, Phys. Rev. Lett. 98, 146401 (2007)

Angular-Radial Distribution Functions (**ARDF**) : N. Artrith, A. Urban, and G. Ceder, PHYSICAL REVIEW B 96, 014112 (2017)

Smooth Overlap of Atomic Positions (**SOAP**) : A. P. Bartók, R. Kondor, and G. Csányi, Phys. Rev. B 87, 184115 (2013)

Machine Learning NMR shifts

Accuracy versus experimental data



NMR shifts prediction : sparcification Least-Square Support Vector Regression

The Nyström approximation of the Kernel uses a small set of landmark/inducing points (support vectors) :

$$K(\chi, \chi) \approx K(\chi, \xi) K^{-1}(\xi, \xi) K(\xi, \chi)$$

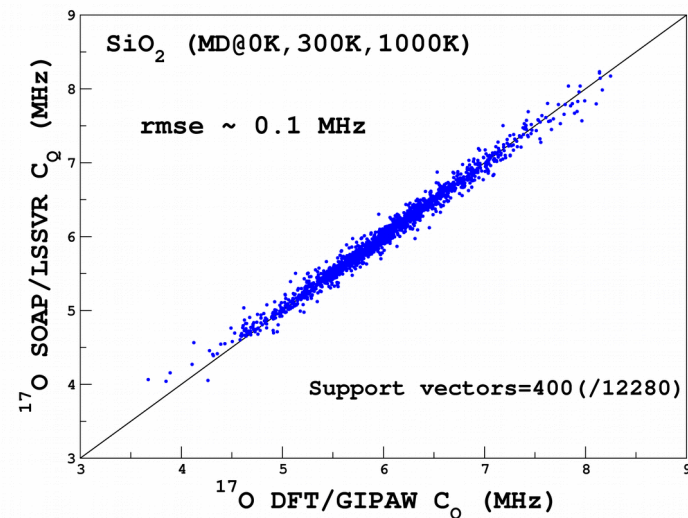
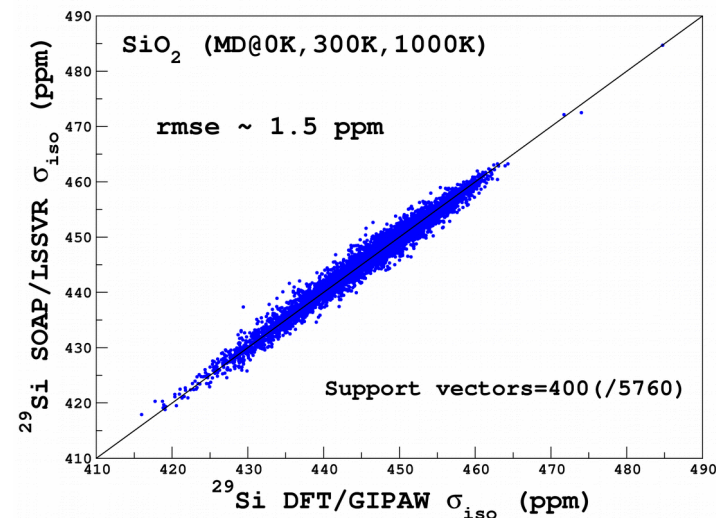
In the RKHS, the Kernel can be defined as a scalar product of non-linear feature map (Mercer Th.) :

$$K(\chi_i, \chi_j) = \langle \phi(\chi_i) \phi(\chi_j) \rangle$$

Then, according to the Reproducing Kernel Hilbert Space (RKHS) formalism, the KRR prediction can be expressed as linear combination of the non-linear feature maps approximated on the support vectors :

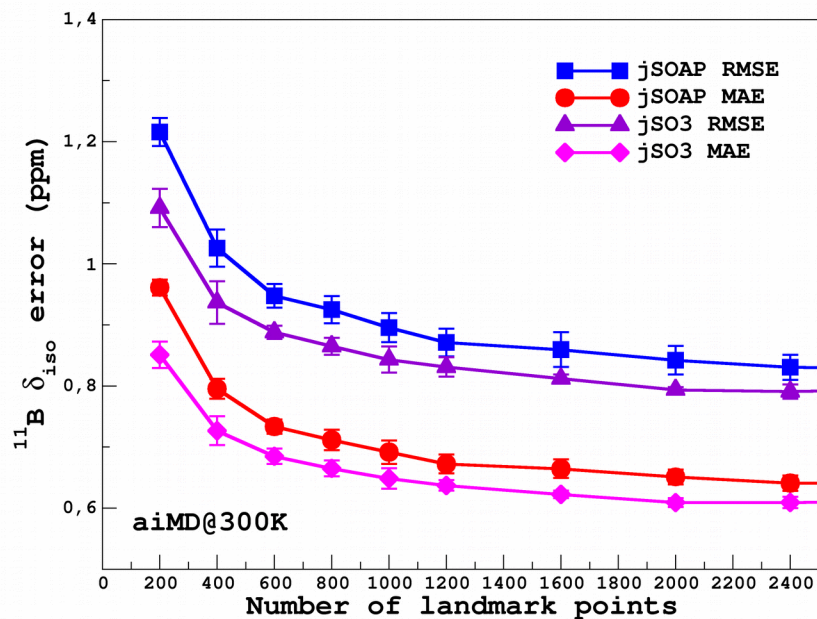
$$\delta(\chi) = \sum_{(\xi)} w_{\xi} \phi_{\xi}(\chi)$$

The dimensionality of the problem is reduced (sparcification). Several options exist for the choice of the support vectors (Kmeans, Max. Entropy). We found the **Incomplete Cholesky Decomposition** to be the most efficient.



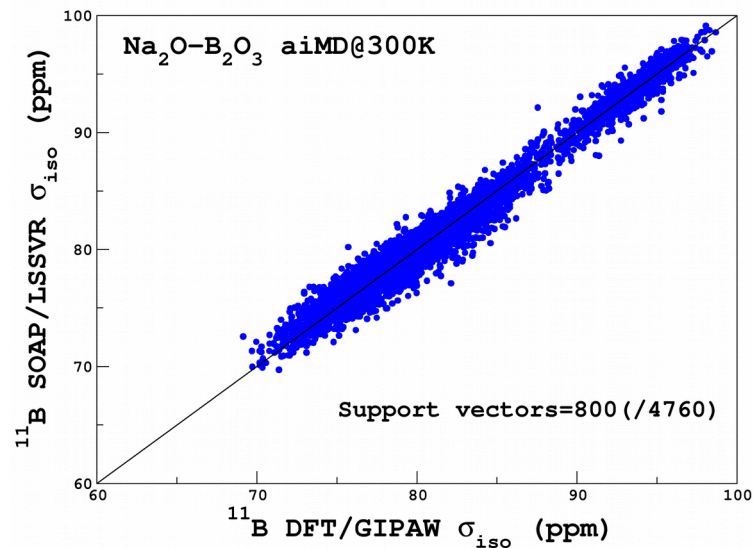
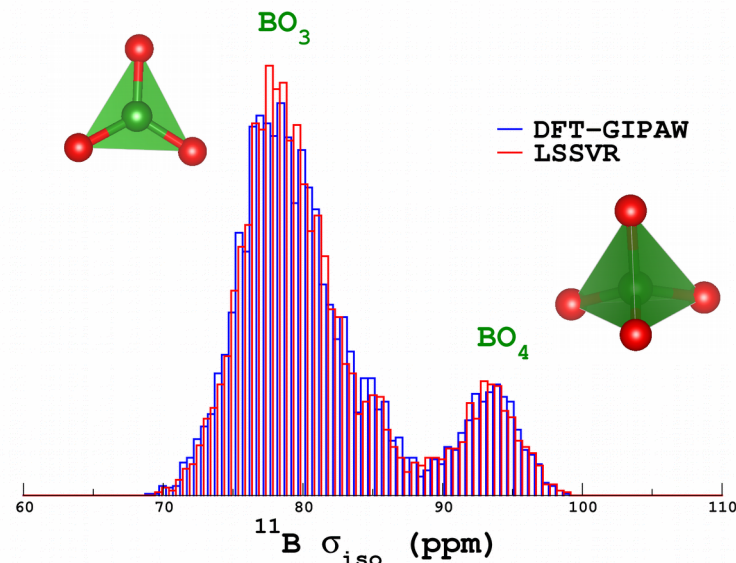
NMR Shifts in sodium borate glasses

Least Square Support Vector Regression (LSSVR)



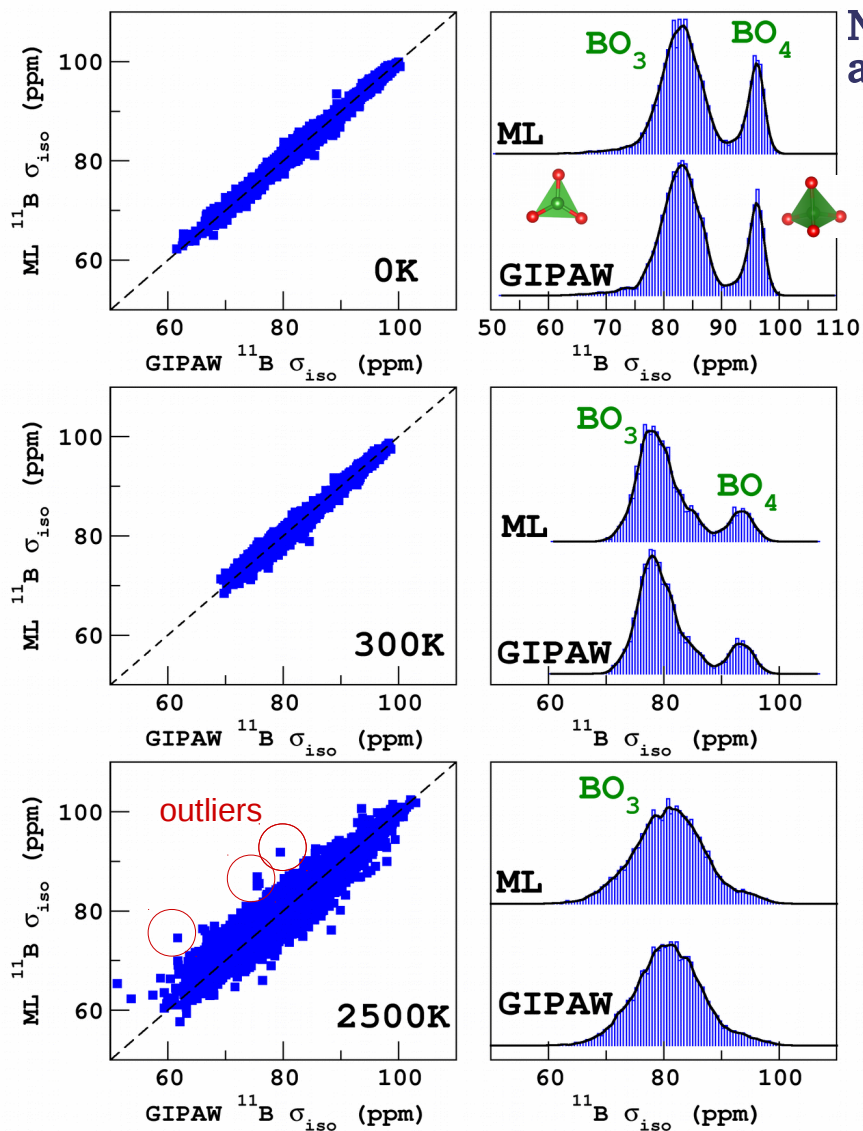
(Top) Comparison of convergence of the ^{11}B NMR shifts error in sodium borate glasses (aiMD at 300K) for two descriptors.

Right panel shows the comparison between DFT calculated and ML predicted distribution of isotropic ^{11}B NMR shifts.

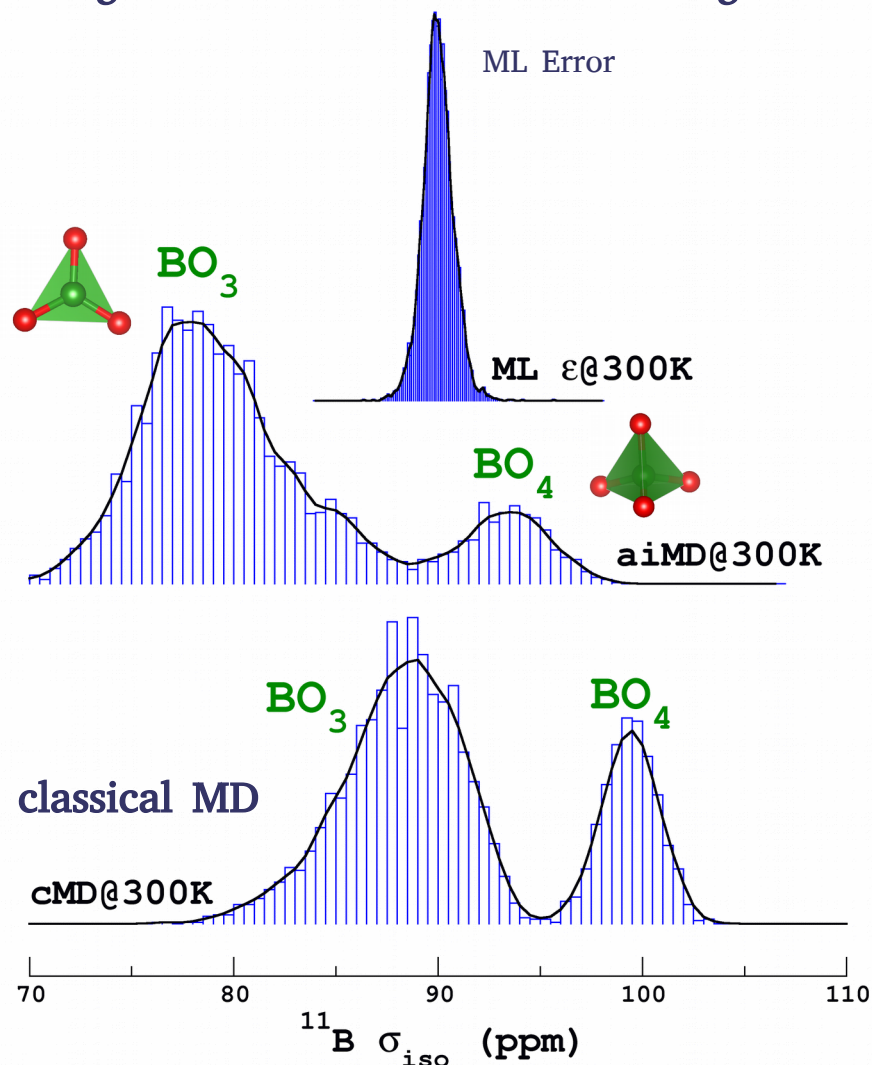


NMR Shifts in sodium borate glasses

Support Vector Regression (LSSVR)



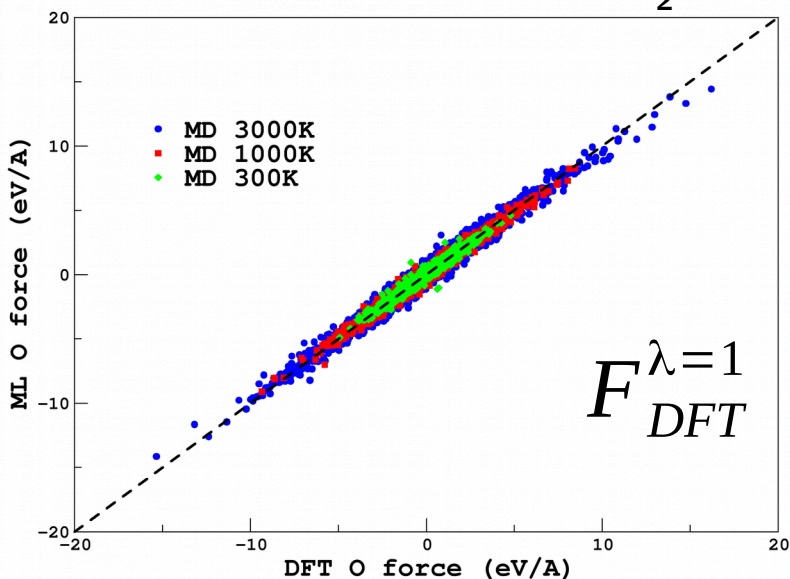
NMR shift :
a driving force for data driven modeling ?



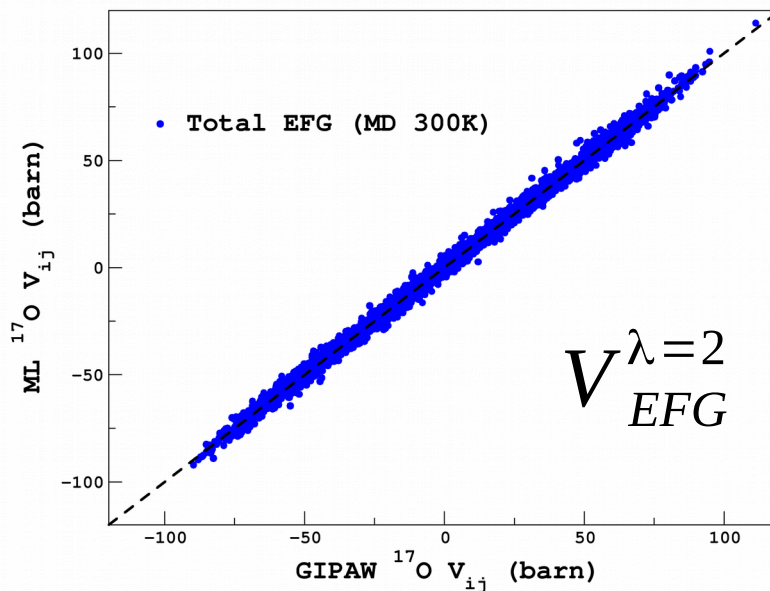
Each descriptor is a **Tensor** . With SOAP :

$$T_m^\lambda(\chi) = \sum_{n,v} \alpha_{n,v} c_{n\lambda m}^v(\chi) \quad \text{2-body terms}$$

Force prediction (SiO_2)



Matrix prediction (SiO_2)



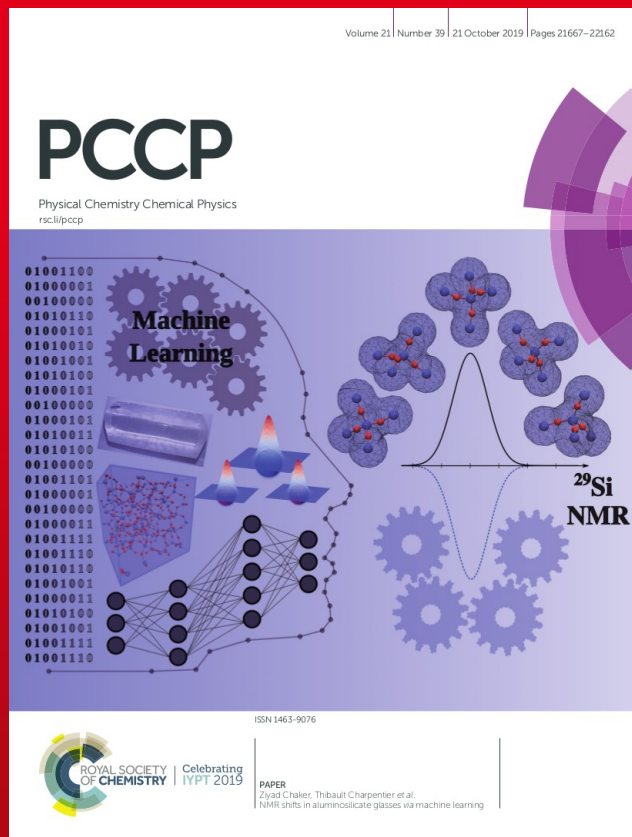
$$P_{n_1 n_2 l_1 l_2}^{\lambda, m}(\chi) = \sum_{m_1 m_2} c_{n_1 l_1 m_1}(\chi) c_{n_2 l_2 m_2}(\chi) C_{l_1, m_1, l_2, m_2}^{\lambda, m} \quad \text{3-body terms}$$



E(3) Equivariant Graph Neural Network (NequIP) [arXiv:2101.03164](https://arxiv.org/abs/2101.03164)

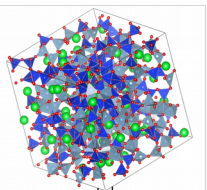


- **Z. CHAKER** (Machine Learning NMR)
- CEA Paris-Saclay, France, Post-Doc 2018-2020



Z. Chaker et al. Phys. Chem. Chem. Phys. **21**, 21667 (2019)

Perfect structure(s) ?



HRMC



MD, aiMD,
DFT-GIPAW

Experiments:
1D & 2D NMR, S(q),
Q(n) etc...



Machine Learning

We acknowledge the financial support of the Cross-Disciplinary Program on Numerical Simulation of CEA, the French Alternative Energies and Atomic Energy Commission

Thank you for your attention



FROM RESEARCH TO INDUSTRY



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