



The use of Molecular Dynamics simulations for the interpretation of EXAFS spectra

Alexei Kuzmin

Dr.phys., Head of the Lab

Institute of Solid State Physics, University of Latvia

Kengaraga street 8, LV-1063 Riga, Latvia

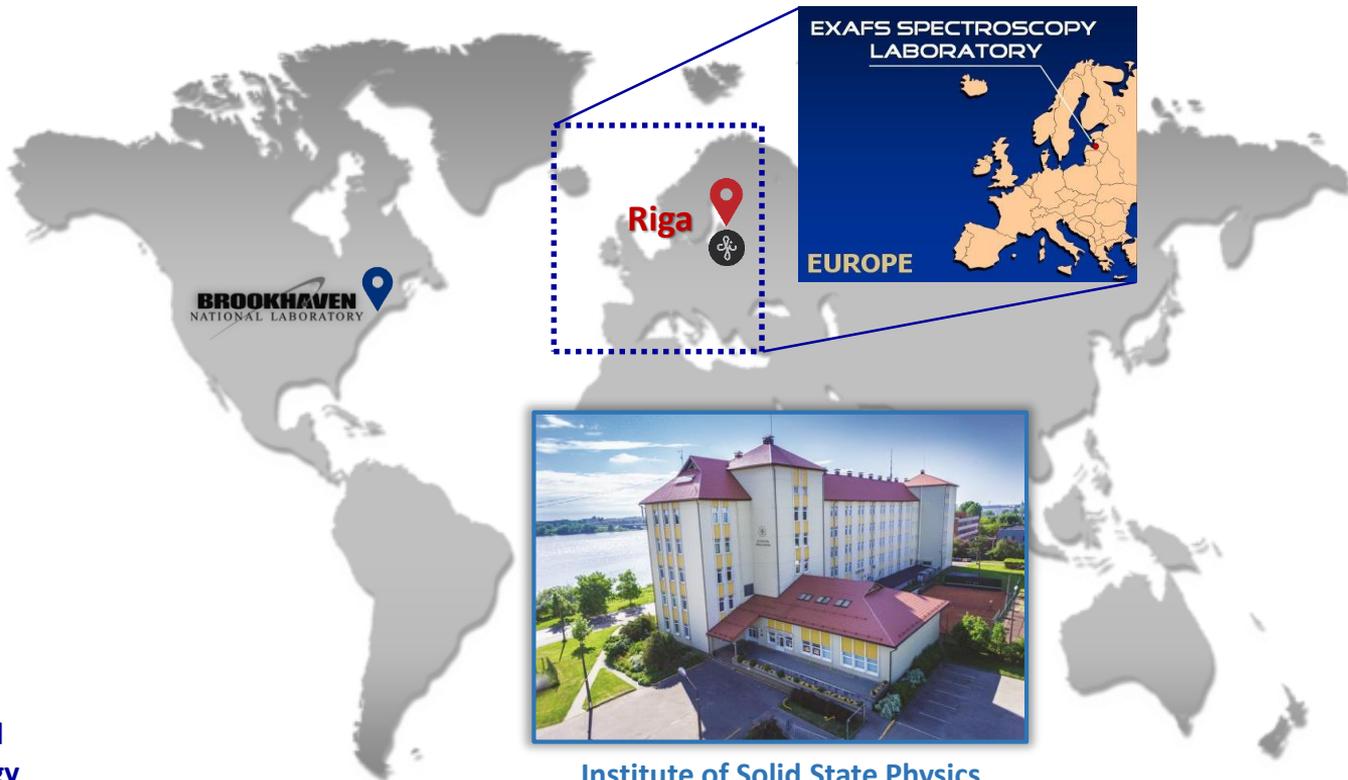
E-mail: a.kuzmin@cfi.lu.lv

Internet: www.cfi.lu.lv

Internet: www.dragon.lv/exafs



EXAFS Spectroscopy Laboratory



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Outline

- **Introduction**
 - Motivation & problems
- **MD-EXAFS approach**
 - Concept
 - State of the Art
 - Basics of Molecular Dynamics (MD)
 - EXAFS simulations
 - Interpretation of results
- **Examples of MD-EXAFS simulations**
 - ReO₃, ScF₃, BCC & FCC metals
- **Conclusions**

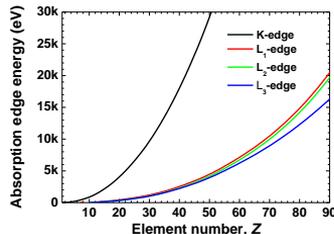


Introduction

- Characteristic times in X-ray absorption process vs thermal vibrations
- How local is X-ray absorption spectroscopy?
- EXAFS formalism beyond the first coordination shell: multiple-scattering effects
- Problems and possible solutions

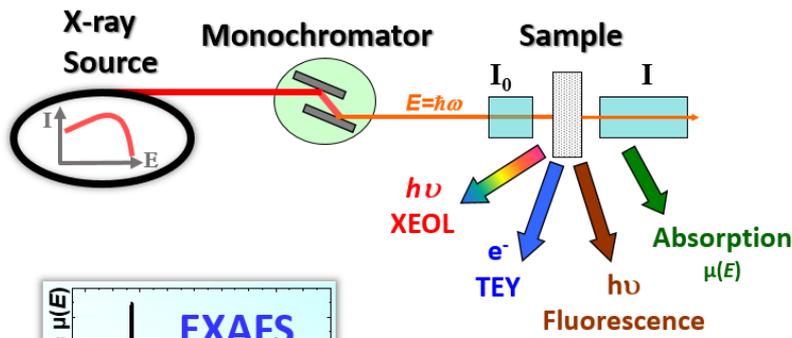
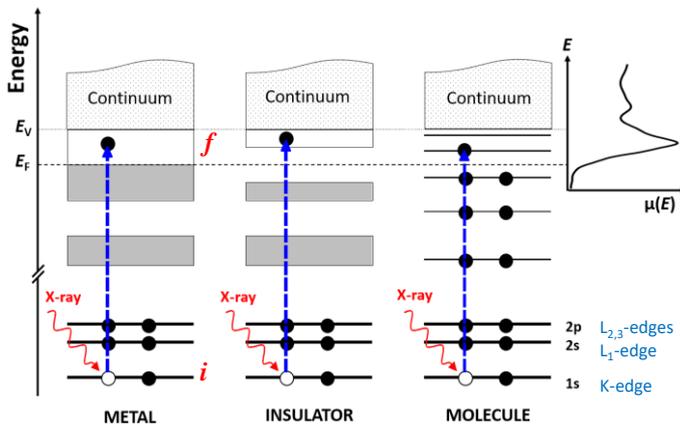
Basics of X-ray absorption spectroscopy (XAS)

X-rays:
soft (<5 keV)
hard (>5 keV)



Fermi's Golden Rule:

$$\mu(E) \propto \sum_f |\langle f | \hat{H} | i \rangle|^2 \delta(E_f - E_i - \hbar\omega)$$

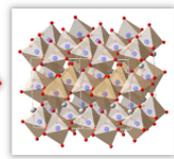
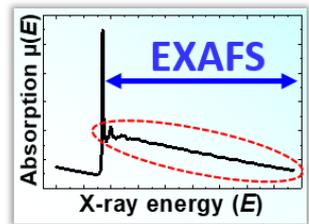


$$\mu(E) = \frac{1}{x} \ln \left(\frac{I_0(E)}{I(E)} \right)$$

$$\mu(E) \propto \frac{I_{\text{fluor}}(E)}{I_0(E)}$$

$$\mu(E) \propto \frac{I_{\text{TEY}}(E)}{I_0(E)}$$

$$\mu(E) \propto \frac{I_{\text{XEOL}}(E)}{I_0(E)}$$



Characteristic time of X-ray absorption process:

$$t_{\text{ab}} \sim 10^{-15} - 10^{-16} \text{ s}$$

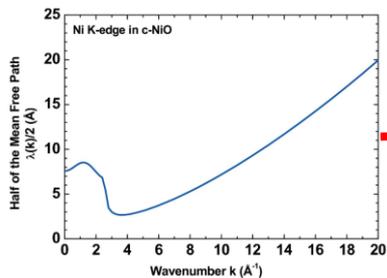
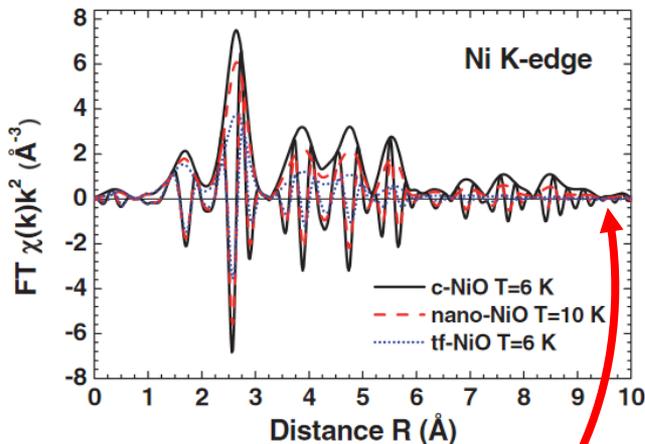
Characteristic time of thermal vibrations:

$$t_{\text{th}} \sim 10^{-13} - 10^{-14} \text{ s}$$

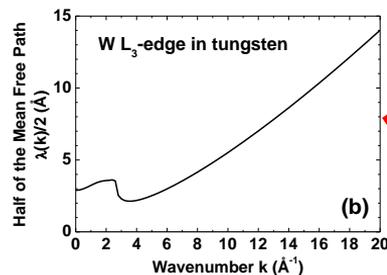
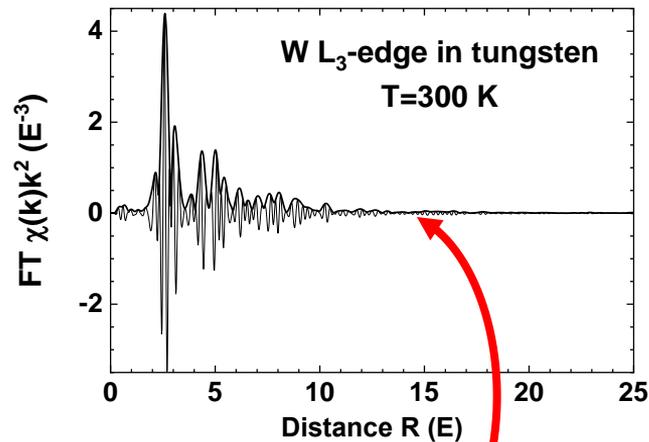
- Atoms are frozen during X-ray absorption
- Static & thermal disorder contribute similarly

How local is X-ray absorption spectroscopy?

The case of NiO.



The case of W foil.



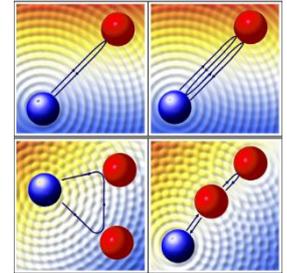
Contributions from outer shells are often available and contain useful structural information.

EXAFS simulations beyond the first coordination shell

1. Multiple-scattering (MS) expansion (FEFF code):

$$\chi^l(k) = \sum_{n=2}^{\infty} \chi_n^l(k) \quad \chi_n^l(k) = \sum_i A_n^l(k, R_i) \sin(2kR_i + \Phi_n^l(k, R_i) + 2\delta_c^l(k))$$

A.L. Ankudinov, B. Ravel, J.J. Rehr, S.D. Conradson, *Phys. Rev. B* 58 (1998) 7565-7576.



2. N-body expansion (GNXAS code):

$$\begin{aligned} \chi(k) = & \int 4\pi R^2 \rho_0 g_2(R) (\chi_2^{oio}(k) + \chi_4^{oioio}(k) + \dots) dR \\ & + \iiint 8\pi^2 R_1^2 R_2^2 \sin(\theta) \rho_0^2 g_3(R_1, R_2, \theta) \\ & \times (2\chi_3^{oijjo}(k) + 2\chi_4^{oiojo}(k) + \chi_4^{oijio}(k) + \chi_4^{ojijjo}(k) + \dots) dR_1 dR_2 d\theta \\ & + \iiint \iiint 8\pi^2 R_1^2 R_2^2 R_3^2 \sin(\theta) \rho_0^3 g_4(R_1, R_2, \theta, R_3, \Omega) \\ & \times (2\chi_4^{oijkjo}(k) + 2\chi_4^{oikjo}(k) + 2\chi_4^{ojikjo}(k) + \dots) dR_1 dR_2 d\theta dR_3 d\Omega \\ & + \dots \end{aligned}$$

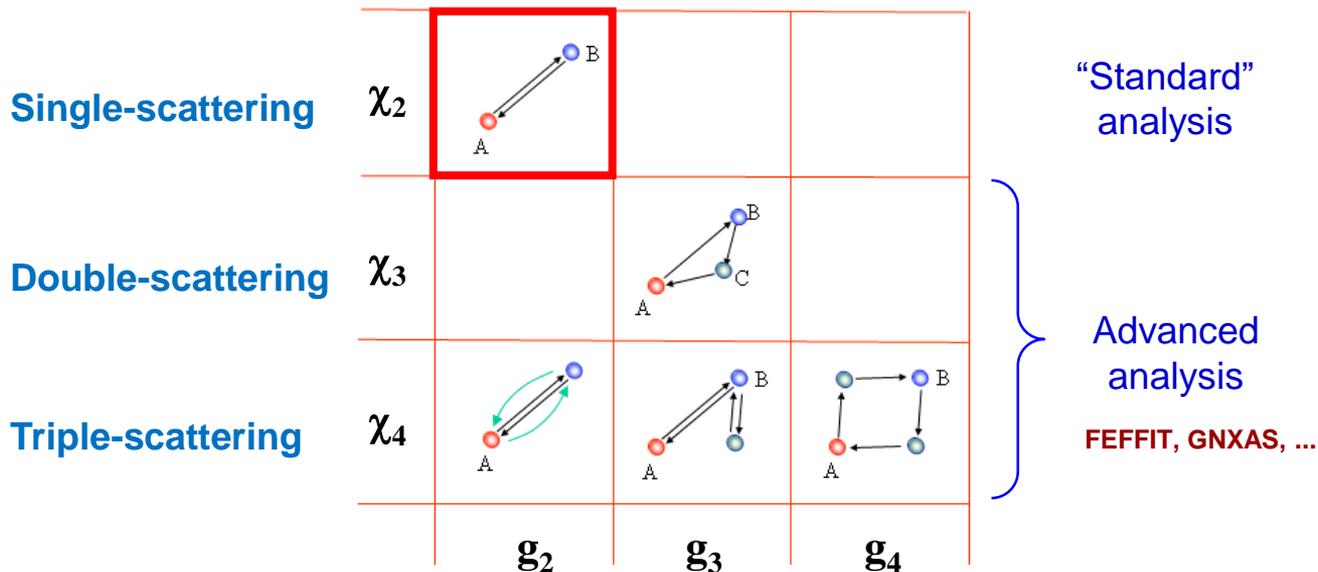
A. Filipponi, A. Di Cicco, C. R. Natoli, *Phys. Rev. B* 52 (1995) 15122-15134.

**EXAFS depends on
atomic distribution
functions!**

Multiple-scattering terms and many body correlation functions

$$\mu(k) = \mu_0(k)[1 + \chi_2(k) + \chi_3(k) + \chi_4(k) + \dots]$$

$$\mu(k) = \mu_0(k)[1 + g_2(k) + g_3(k) + g_4(k) + \dots]$$

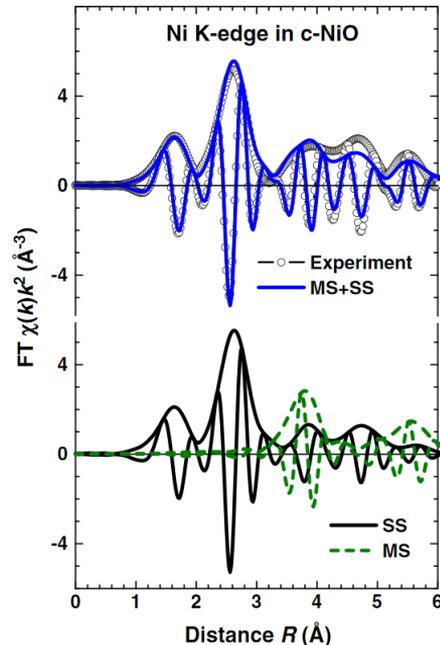
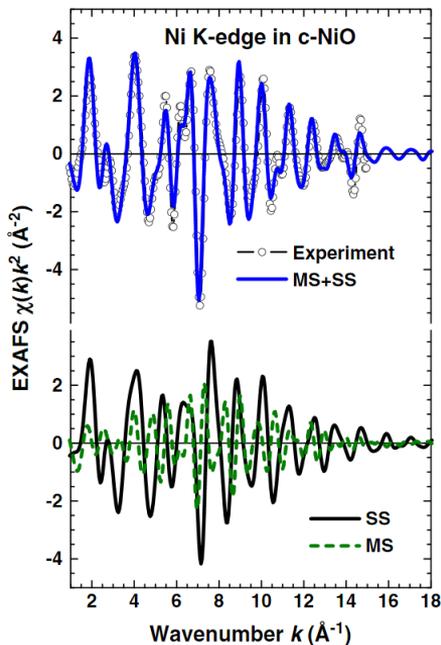
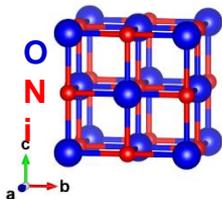


The rows group the different possible paths which give the same order contributions to the absorption coefficient, χ_2 , χ_3 or χ_4 . The columns group the paths corresponding to two, three, four body correlation functions g_i .

EXAFS problems (1)

1) One needs to account for many-body correlations through the so-called multiple-scattering (MS) effects.

NiO
Space group Fm-3m
 $a_0 = 4.1773 \text{ \AA}$



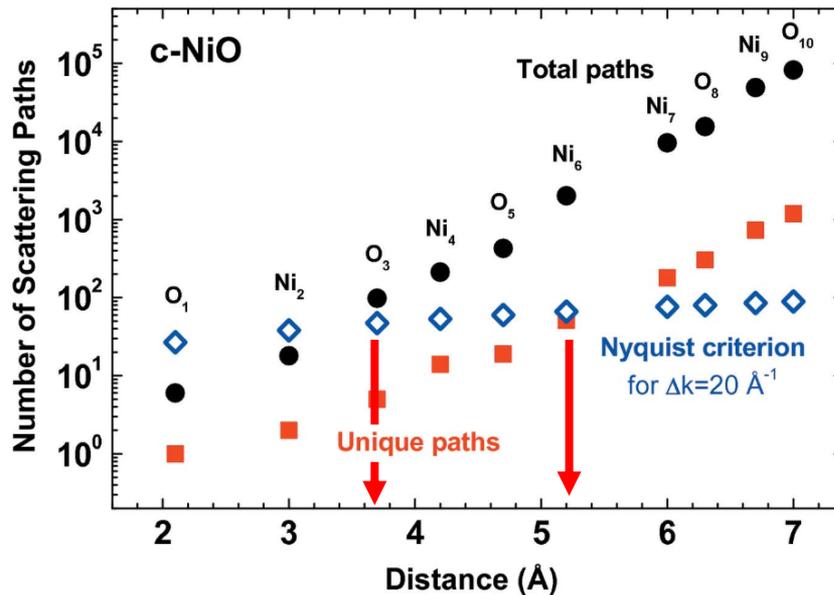
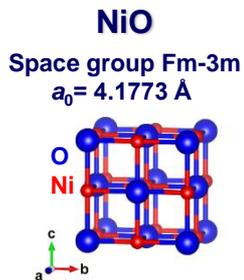
A. Anspoks and A. Kuzmin,
J. Non-Cryst. Solids 357 (2011) 2604.

EXAFS problems (2)

2) A number of fitting parameters grows rapidly upon an increase of analyzed region size around the absorbing atom. This problem is especially relevant for disordered and nanocrystalline materials.

Number of scattering paths
vs
independent parameters

$$N_{\text{par}} = \frac{2\Delta k \Delta R}{\pi}$$



A. Kuzmin and J. Chaboy, IUCrJ 1 (2014) 571-589.

Possible solutions

The problem of disorder in EXAFS can be addressed using

- **Molecular Dynamics (MD),**
- **Monte Carlo (MC),**
- **Reverse Monte Carlo (RMC)**

Configuration-averaged EXAFS

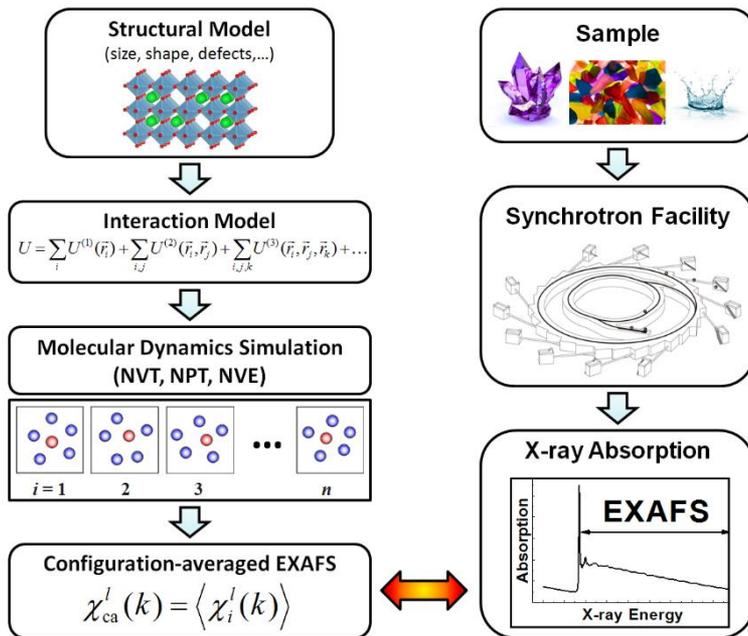
$$\chi_{ca}^l(k) = \langle \chi^l(k) \rangle$$

simulations.

Advantages of MD-EXAFS approach:

- ✓ has no structural fitting parameters
- ✓ allows interpretation of EXAFS spectra far beyond the first coordination shell
- ✓ allows a validation of the interatomic potential models

MD-EXAFS approach: Concept



EDACA code:

- A. Kuzmin and R.A. Evarestov, *J. Phys.: Condens. Matter* **21** (2009) 055401.
- A. Kuzmin and J. Chaboy, *IUCr* **1** (2014) 571-589.
- A. Kuzmin, A. Anspoks, A. Kalinko, J. Timoshenko, *Z. Phys. Chem.* **230** (2016) 537-549.



Author: Dr. phys. Alexei Kuzmin
Institute of Solid State Physics
University of Latvia
Kengaraga street 8, LV-1063 Riga, Latvia
Internet: <http://www.dragon.lv/edaca>
E-mail: a.kuzmin@csl.lu.lv

Preface

X-ray absorption spectroscopy (XAS) at synchrotron radiation sources is a structural tool providing information on the local atomic and electronic structure around an atom of a particular type. Today XAS is successfully applied to a study of crystalline, nanocrystalline and disordered solids, liquids and gases in a wide range of external conditions defined by temperature, pressure, etc. The size of the region, probed by XAS, depends on the degree of thermal and static disorder present in a material and is limited by the mean-free path of the excited photoelectron. Typically the information reach region extended up to 3-10 Å around the absorbing atom.

An advantage of the XAS method is its sensitivity to many-atom distribution functions, giving rise to multiple-scattering (MS) contributions, and to correlation effects in atom dynamics. Note that accurate account of both effects is still challenging.

The time-scale (about $10^{-15}, 10^{-16}$ s) of the X-ray absorption process is much shorter than the characteristic time (about $10^{-13}, 10^{-14}$ s) of thermal vibrations. Therefore, the atoms may be considered as frozen at their instantaneous positions during a single photoabsorption process, and the total experimentally measured X-ray absorption spectrum corresponds to the configurational average of all atomic positions over the time of the experiment. This situation can be straightforwardly modelled combining the molecular dynamics (MD) simulation with the extended X-ray absorption fine structure (EXAFS) calculations, known as the MD-EXAFS approach.

Finally, the agreement between the experimental and configuration-averaged EXAFS spectra can be used to validate the accuracy of the interatomic potential (force-field) models employed in the MD simulations.

<http://www.dragon.lv/edaca/>

XAFS 2023 - Short Course on X-ray Absorption Fine Structure: Advanced topics in data analysis and modeling, BNL, November 1–3, 2023

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First Molecular Dynamics Simulation

Phase Transition for a Hard Sphere System

B. J. ALDER AND T. E. WAINWRIGHT

University of California Radiation Laboratory, Livermore, California

(Received August 12, 1957)

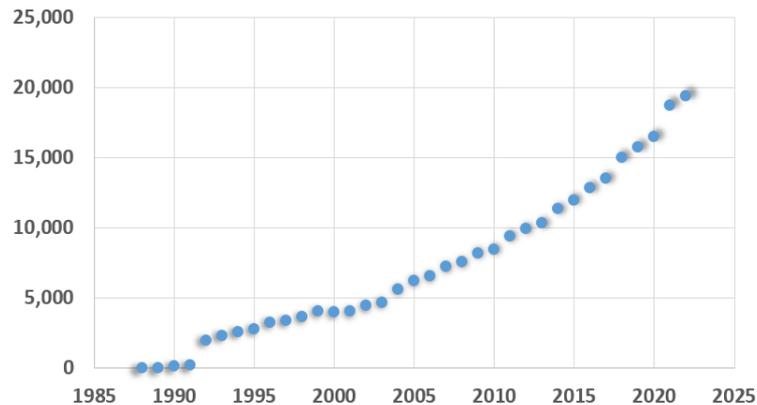
J. Chem. Phys. **27**, 1208 (1957); doi: <http://dx.doi.org/10.1063/1.1743957>

J. Chem. Phys. **31**, 459 (1959); doi: <http://dx.doi.org/10.1063/1.1730376>

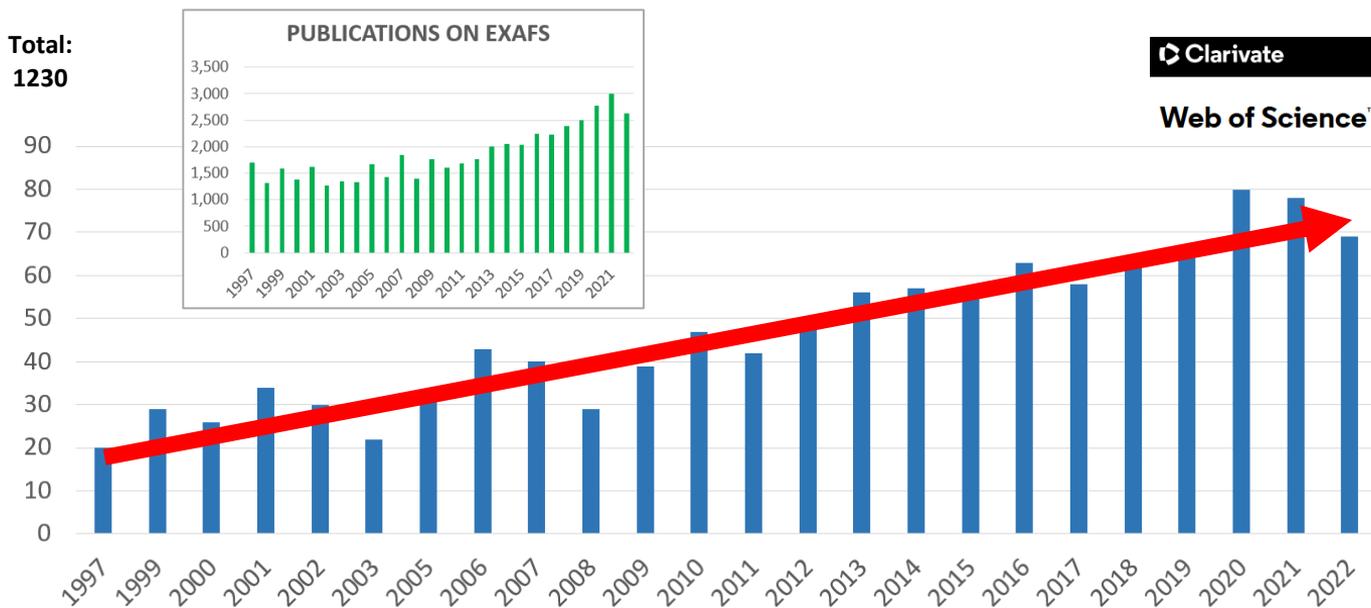


IBM 704

Publications on Molecular Dynamics simulations



MD-EXAFS approach: State of the Art

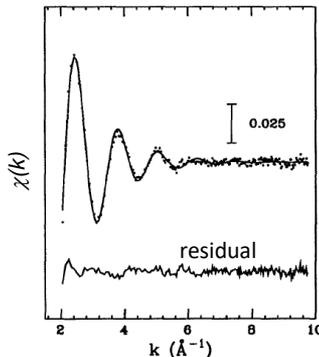


- Comparison of MD and EXAFS results
- **Using MD results for EXAFS calculation** ←

MD-EXAFS approach: History

An extended x-ray absorption fine structure study of aqueous solutions by employing molecular dynamics simulations

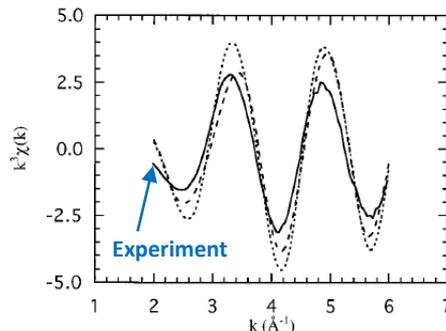
P. D'Angelo and A. Di Nola
Dipartimento di Chimica, Università degli Studi di Roma "La Sapienza," P.le Aldo Moro 5, 00185 Roma, Italy
A. Filipponi
Dipartimento di Fisica, Università degli Studi dell'Aquila, Via Vento, 67100 Coppito-L'Aquila, Italy
N. V. Pavel and D. Roccatano
Dipartimento di Chimica, Università degli Studi di Roma "La Sapienza," P.le Aldo Moro 5, 00185 Roma, Italy



**Br K-edge in 0.15M
RbBr aqueous
solution**

- [1] P. D'Angelo, A.D. Nola, A. Filipponi, N.V. Pavel, D. Roccatano, *J. Chem. Phys.* **100** (1994) 985-994.
- [2] P. D'Angelo, A.D. Nola, M. Mangoni, N.V. Pavel, *J. Chem. Phys.* **104** (1996) 1779-1790.
- [3] B.J. Palmer, D.M. Pfund, J.L. Fulton, *J. Phys. Chem.* **100** (1996) 13393.
- [4] A. Kuzmin, S. Obst, J. Purans, *J. Phys.: Condensed Matter* **9** (1997) 10069-10078.
- [5] M. I. McCarthy, G. K. Schenter, M. R. Chacon-Taylor, J. J. Rehr, G. E. Brown, Jr., *Phys. Rev. B* **56** (1997) 9925-9936.

Sr K-edge in strontium nitrate aqueous solution

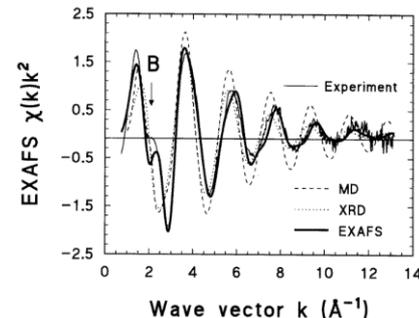


Direct Modeling of EXAFS Spectra from Molecular Dynamics Simulations

Bruce J. Palmer,* David M. Pfund, and John L. Fulton
Energy and Environmental Sciences Division, Pacific Northwest National Laboratory, Richland, Washington 99352
Received: January 17, 1996; In Final Form: April 29, 1996

X-ray absorption spectroscopy and molecular dynamics studies of Zn²⁺ hydration in aqueous solutions

A. Kuzmin], S. Obst] and J. Purans]
] Institute of Solid State Physics, Kengaraga Str. 8, LV-1063 Riga, Latvia
] Institut für Kristallographie, Freie Universität Berlin, Takustrasse 6, 14195 Berlin, Germany



**Zn K-edge in the 0.125M
aqueous solution of ZnSO₄**

} aqueous solutions
Na⁺(H₂O)_n-MgO (100) interface



Basics of classical Molecular Dynamics (I)

Molecular dynamics (MD) simulation is a technique to simulate the motion of atoms under predefined conditions, such as **temperature (T)**, **pressure (P)**, etc.

The **NVT** ensemble is also called the *canonical* ensemble. In the **NVT** ensemble, the number (N) of particles (atoms) and the supercell size (V) are fixed, whereas the temperature (T) is controlled using a surrounding virtual heat bath (**thermostat**).

The **NPT** ensemble is also called *isothermal-isobaric* ensemble. In the **NPT** ensemble, the number (N) of particles (atoms) is fixed, whereas the temperature (T) and pressure (P) are controlled using a **thermostat** and a **barostat**.

- J. C. Berendsen, J. P. M. Postma, W. F. van Gunsteren, A. DiNola, J. R. Haak, Molecular-Dynamics with Coupling to an External Bath, *J. Chem. Phys.* **81** (1982) 3684.
- G. J. Martyna, M. L. Klein, M. Tuckerman, Nosé-Hoover chains: The canonical ensemble via continuous dynamics, *J. Chem. Phys.* **97** (1992) 2635.
- G. J. Martyna, D. J. Tobias, M. L. Klein, Constant pressure molecular dynamics algorithms, *J. Chem. Phys.* **101** (1994) 4177.

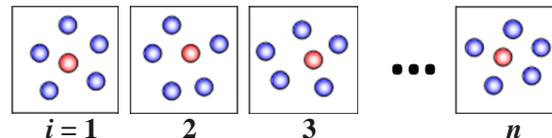
Basics of classical Molecular Dynamics (II)

During classical MD simulation the **Newton's equations** of motion are numerically integrated for a set of atoms starting from a given **initial configuration**. This is performed via numerical integration by discretizing time into small intervals called the **time step (Δt)**. The time step must be smaller than the fastest vibrational frequency in the system, typically in the order of 1 femtosecond ($\approx 10^{-15}$ s).

The interactions between the atoms, i.e. the interatomic forces, can be calculated based on **empirical potentials** or **first-principles theory (*ab initio*)**.

Machine Learning (ML) potentials become popular nowadays.

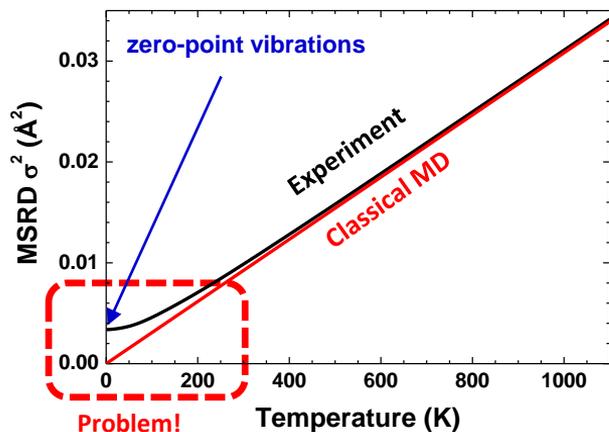
These interatomic forces determine the acceleration of the atoms, allowing the positions and velocities propagate towards the next time step. Repeating this procedure many times yields a **series of snapshots**, describing the trajectory of the system in phase space, which can be analyzed to extract the desired properties, including EXAFS/XANES.



Temperature in the classical MD simulation

$$\langle E_k \rangle = \left\langle \frac{1}{2} \sum_{i=1}^N (m_i \vec{v}_i)^2 \right\rangle = \frac{3}{2} N k_B T$$

$T \rightarrow 0\text{K}, v_i \rightarrow 0$



The Debye temperature (θ_D) and characteristic temperature (T_0) of zero-point vibration.

The unit for temperature is K.

| Element | θ_D | T_0 |
|---------|------------|--------|
| Be | 1440 | 481.06 |
| Al | 428 | 142.98 |
| Ti | 420 | 140.31 |
| Ta | 240 | 80.18 |
| Pb | 105 | 35.08 |

Y. Yang and Y. Kawazoe, EPL 98 (2012) 66007

Measurements are often performed using cryostats!

N.B. The magnitude of zero-point vibrations at T_0 equals that of the excited vibrations.

Quantum effects

Quantum effects are expected to become significant when the **thermal de Broglie wavelength**

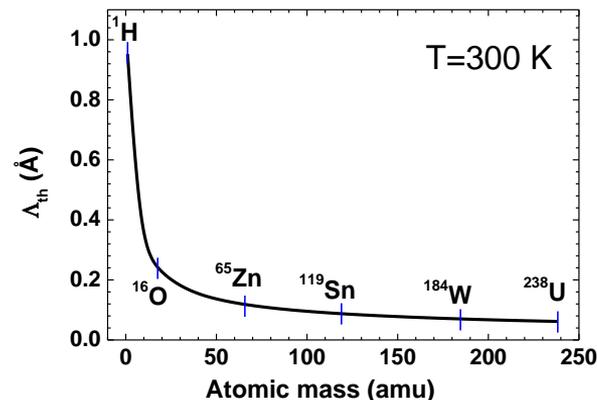
$$\Lambda_{\text{th}} = \frac{h}{\sqrt{2\pi m k_B T}}$$

is much larger than the interatomic distance d .

Typical interatomic distance in solids is about $d \approx 1\text{-}3 \text{ \AA}$.

For example, at $T = 300 \text{ K}$:

- $\Lambda_{\text{th}} = 1 \text{ \AA}$ for a H atom ($m_{\text{H}} = 1 \text{ amu}$)
- $\Lambda_{\text{th}} = 0.24 \text{ \AA}$ for a O atom ($m_{\text{O}} = 16 \text{ amu}$)
- $\Lambda_{\text{th}} = 0.12 \text{ \AA}$ for a Zn atom ($m_{\text{Zn}} = 65 \text{ amu}$)
- $\Lambda_{\text{th}} = 0.09 \text{ \AA}$ for a Sn atom ($m_{\text{Sn}} = 119 \text{ amu}$)
- $\Lambda_{\text{th}} = 0.06 \text{ \AA}$ for a U atom ($m_{\text{U}} = 238 \text{ amu}$)



All atoms, except for the lightest ones such as H, He, can be considered as “point” particles at **sufficiently high temperature** (or $d \gg \Lambda_{\text{th}}$), and classical mechanics can be used to describe their motion.

Interatomic potentials

Empirical potentials

- Depend on a set of parameters which must be determined in advance by fitting to some physical properties (structure, phonons, elastic constants, etc).
- Must be validated using experiment or more accurate simulation.
- MD simulations are fast and can be done using desktop computer or a server.

Machine Learning potentials

- Must be trained using ab initio (DFT) simulations.
- Must be validated using experiment.
- MD simulations are slower than with empirical potentials but much faster than doing ab initio simulations. Server or cluster are required.

Ab initio (DFT) potentials

- Calculated from first principles.
- Must be validated using experiment.
- MD simulations are slow. Supercomputer is required.

Interatomic potentials or force-fields

as a consequence of the Born-Oppenheimer separation of electrons and nuclei motion

$$V_{TOT} = \sum_i^N V_1(\vec{r}_i) + \sum_{i,j}^N V_2(\vec{r}_i, \vec{r}_j) + \sum_{i,j,k}^N V_3(\vec{r}_i, \vec{r}_j, \vec{r}_k) + \dots$$

External field

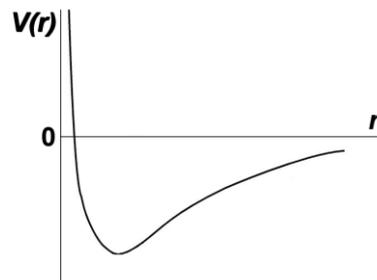
Two-body

Three-body

Example:

The *Born-Mayer* potential

$$V(r) = Ae^{-\alpha r} - \frac{C}{r^6}$$



Databases of interatomic potentials:

<https://gulp.curtin.edu.au/gulp/models.cfm>

<https://www.ucl.ac.uk/klmc/Potentials/>

<https://www.ctcms.nist.gov/potentials/>

<https://openkim.org/>

When more accurate description is needed, electronic behavior can be obtained from first-principles calculations using *ab initio* Molecular Dynamics (AIMD).

Molecular Dynamics Programs

https://en.wikipedia.org/wiki/Comparison_of_software_for_molecular_mechanics_modeling

LAMMPS

- a Large-scale Atomic/Molecular Massively Parallel Simulator.

<https://www.lammps.org/>

GULP

- the General Utility Lattice Program.

<https://gulp.curtin.edu.au/gulp/>

DL_POLY

- a general purpose serial and parallel molecular dynamics simulation package.

https://www.scd.stfc.ac.uk/Pages/DL_POLY.aspx

CP2K

- a program to perform atomistic and molecular simulations of solid state, liquid, molecular, and biological systems. (also *ab initio*)

<https://www.cp2k.org/>

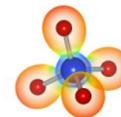
Important: MD trajectory must be saved as *.XYZ file with a specific structure.

Interpretation of results

VESTA

a 3D visualization program for structural models, volumetric data such as electron/nuclear densities, and crystal morphologies.

<http://jp-minerals.org/vesta/en/>



OVITO

a scientific visualization and data analysis solution for atomistic and other particle-based models.

<https://www.ovito.org/>





MD-EXAFS simulation related parameters

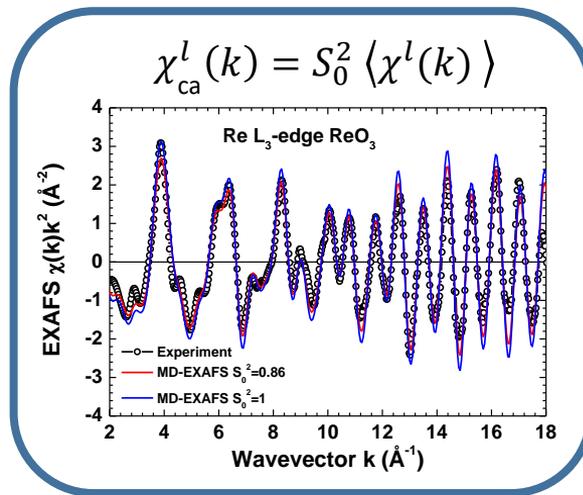
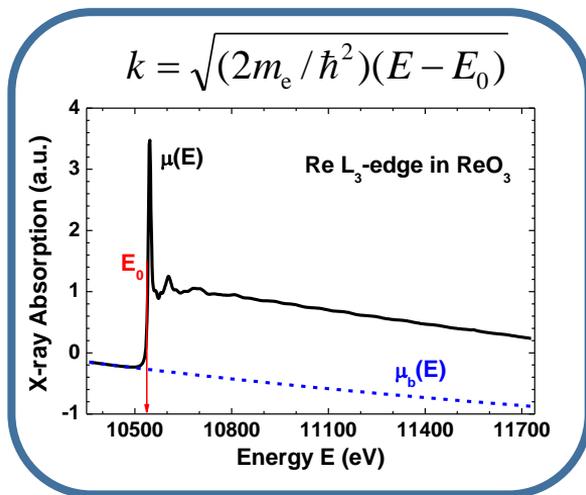
- Equilibration time
- Proper averaging (number of configurations & time step)
- The simulation box size must be large enough to avoid boundary condition artifacts such as, for example, artificial correlations (simulation box size $> 2R_{\max}$)
- Multiple-scattering series truncation problem in FEFF8.5L:
 - NLEG** = 8 default 8 order * (bond length 2-3 Å) / 2 = 8-12 Å
 - CRITERIA** 4.0 2.5 * default critcw=4.0% critpw=2.5%
 - CRITERIA** 0 0 * use all paths, if possible (cw and pw criteria turned off)
- A “configuration” average over the spectra of multiple absorbing atoms (for nanoparticles)
 - CFAVERAGE** iphabs nabs rclabs
 - iphabs** potential index for the type of absorbing atoms over which to make the configuration average
 - nabs** the configuration average is made over the first nabs absorbers in the `feff.inp` file of type iphabs.
 - rclabs** radius to make a small atom list from a bigger one allowed in `feff.inp`.

Non-structural parameters in MD-EXAFS simulation

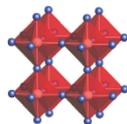
The configuration-averaged EXAFS spectrum $\chi_{ca}(k)$ is calculated **uniquely** from the results of the MD simulation.

Comparison with the experiment requires two more parameters:

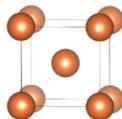
$$E_0 \text{ and } S_0^2$$



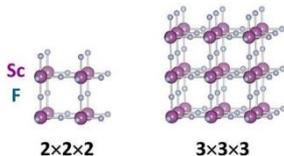
Examples of MD-EXAFS simulations



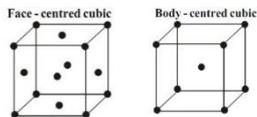
Cubic perovskite-type ReO_3 - force-field development



Bcc tungsten - validation of interatomic potentials

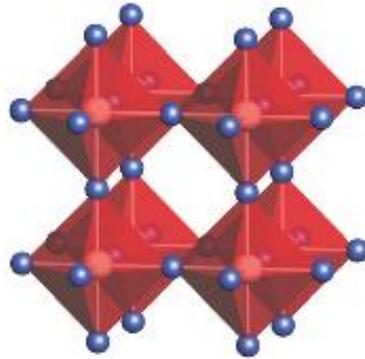


Cubic perovskite-type ScF_3 - the effect of the supercell size



FCC and BCC metals - validation of Machine Learning potentials

Cubic perovskite-type ReO_3



A. Kalinko, R.A. Evarestov, A. Kuzmin, J. Purans, *J. Phys.: Conf. Ser.* **190** (2009) 012080.

Force-field model for ReO_3 used in the MD simulations

| | Exper.* | Calc. |
|-------------------------|---------|-------|
| Lattice constant (Å) | 3.75 | 3.75 |
| Bulk Modulus (GPa) | 211 | 198 |
| Elastic constants (Gpa) | | |
| C_{11} | 572 | 503 |
| C_{12} | 7 | 46 |
| C_{44} | 68 | 46 |

} for central pairwise forces

* T. Chatterji, P. F. Henry, R. Mittal, S. L. Chaplot, *Phys. Rev. B* 78 (2008) 134105;
N. Tsuda, Y. Sumino, I. Ohno, T. Akahane, *J. Phys. Soc. Jpn.* 41 (1976) 1153–1158.

Buckingham Potential

$$U_{ij} = Ae^{-\frac{r_{ij}}{\rho}} + \frac{C}{r_{ij}^6}$$

Covalent Exponential Potential

$$U_{ij} = -De^{-\frac{n(r_{ij}-r_0)^2}{2r_{ij}}}$$

The parameters of the force-field model for cubic ReO_3

| Coulomb potential | | | |
|-------------------|-------|---|--------------------------|
| Atom | Z | e | |
| Re | 3.36 | | |
| O | -1.12 | | ← from Quantum Chemistry |

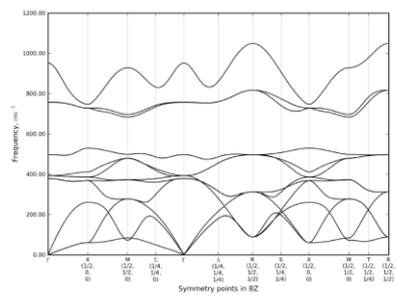
| Buckingham potential, cutoff 20 Å | | | |
|-----------------------------------|---------|------------|---------------------|
| Pair of atoms | A, eV | ρ , Å | C, eVÅ ² |
| Re-O | 1194.24 | 0.3426 | 98.10 |
| O-O | 3224.54 | 0.2652 | 47.40 |

| Covalent exponential potential, cutoff 2 Å | | | |
|--|--------|--------|-----------|
| Pair of atoms | D | n | r_0 , Å |
| Re-O | 4.8283 | 6.2107 | 1.875 |

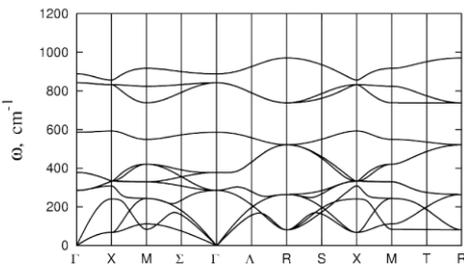


Force-field parameters were optimized to reproduce:

- structural parameters,
- elastic constants,
- ab initio phonon frequencies.



Phonon dispersion curves calculated using frozen phonon (LCAO, CRYSTAL06) method.

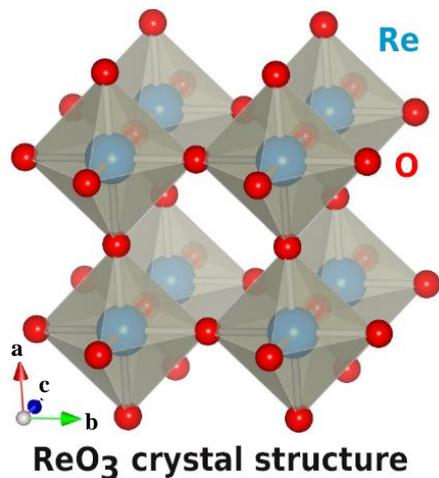


Phonon dispersion curves calculated by the force-field (GULP 3.1) method.

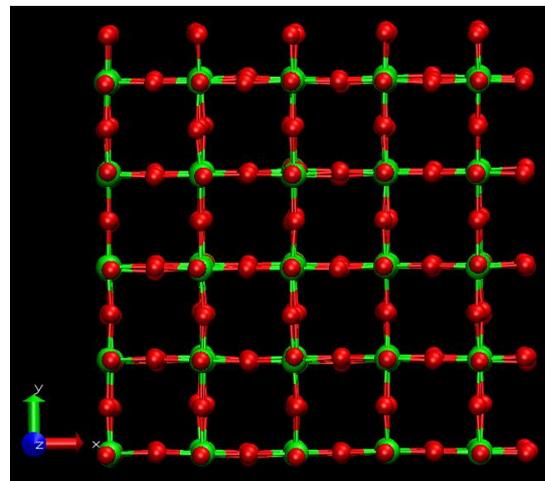
R. A. Evarestov, A. Kalinko, A. Kuzmin, M. Losev, J. Purans, *Integr. Ferroelectrics* 108 (2009) 1-10.

Molecular Dynamics simulations of ReO_3

using the rigid-ion model as implemented in the GULP3.1 code*



Cubic perovskite ReO_3
Space group No. 221, $Pm-3m$
Lattice parameter $a_0 = 3.7477 \text{ \AA}$

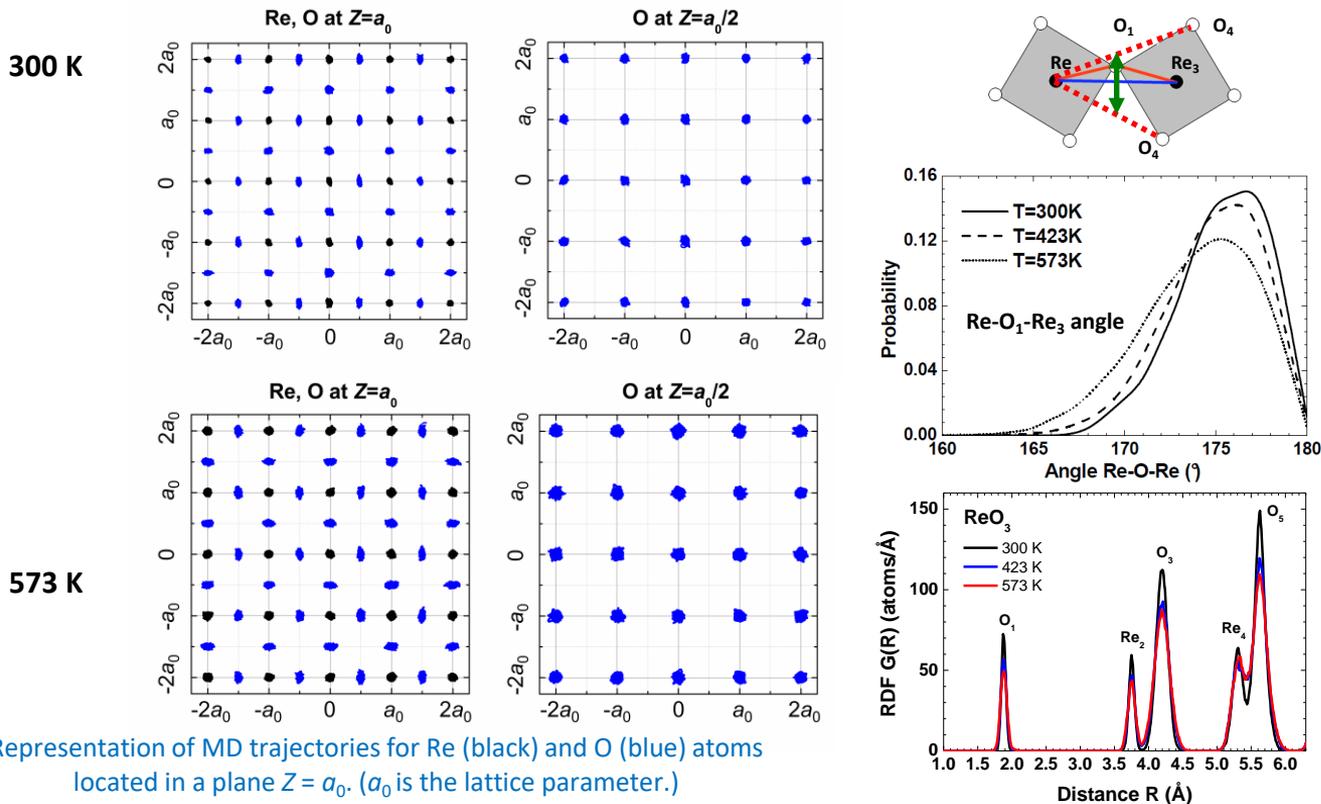


NVT/NPT ensembles

Leap-frog Verlet integration
Supercell **5x5x5** (500 atoms)
T = 300 K, 423 K, 573 K
Simulation time: 20 ps equilibration + 20 ps production
Time step: 0.5 fs
4000 atomic configurations

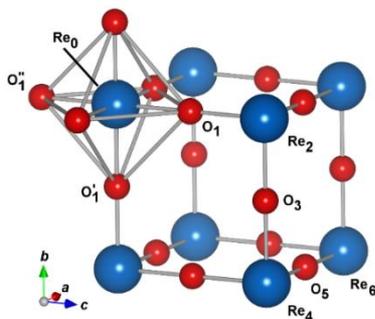
* J.D. Gale and A.L. Rohl, *Mol. Simul.* 9 (2003) 291.

Thermal disorder from MD simulations in ReO_3

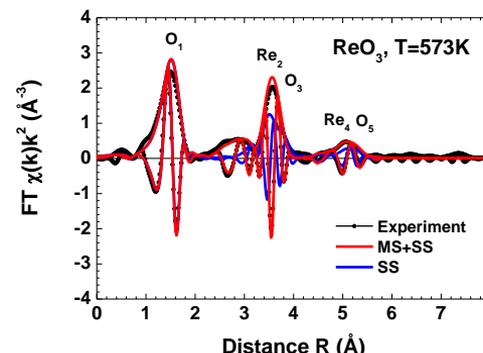
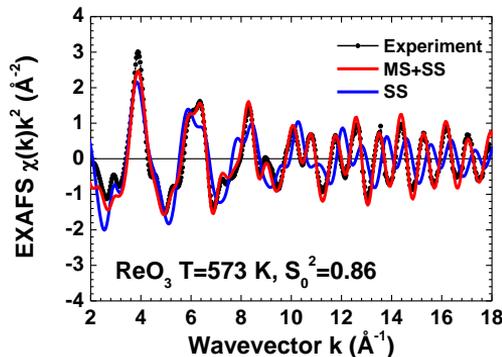
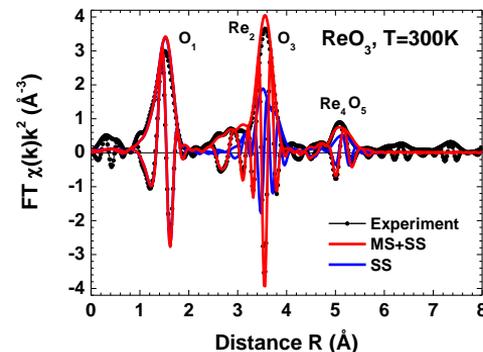
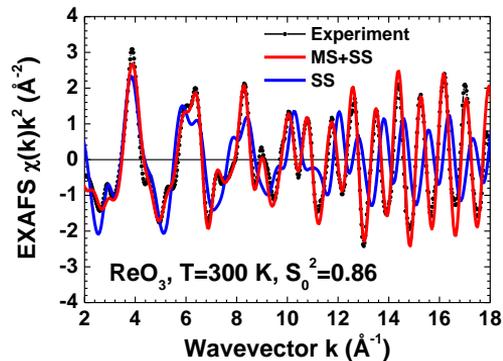


A. Kalinko, R.A. Evarestov, A. Kuzmin, J. Purans, J. Phys.: Conf. Ser. 190 (2009) 012080.

Comparison between experimental and configuration-averaged Re L_3 -edge EXAFS signals and their Fourier transforms in ReO_3



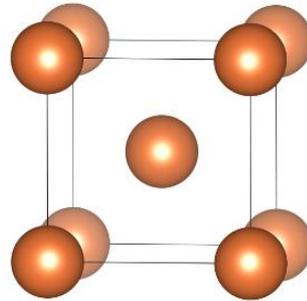
| Index | Path | Path type | Path degener. | Half-path length (Å) |
|-------|--|-----------|---------------|----------------------|
| 1 | Re ₀ -O ₁ -Re ₀ | SS | 6 | 1.875 |
| 2 | Re ₀ -O ₁ -O ₁ '-Re ₀ | DS | 24 | 3.201 |
| 3 | Re ₀ -Re ₂ -Re ₀ | SS | 6 | 3.750 |
| 4 | Re ₀ -O ₁ -O ₁ '-Re ₀ | DS | 6 | 3.750 |
| 5 | Re ₀ -O ₁ -Re ₂ -Re ₀ | DS | 12 | 3.750 |
| 6 | Re ₀ -O ₁ -Re ₀ -O ₁ '-Re ₀ | TS | 6 | 3.750 |
| 7 | Re ₀ -O ₁ -Re ₀ -O ₁ -Re ₀ | TS | 6 | 3.750 |
| 8 | Re ₀ -O ₁ -Re ₂ -O ₁ -Re ₀ | TS | 6 | 3.750 |
| 9 | Re ₀ -O ₁ -Re ₀ -O ₁ '-Re ₀ | TS | 24 | 3.750 |
| 10 | Re ₀ -O ₃ -Re ₀ | SS | 24 | 4.193 |
| 11 | Re ₀ -O ₁ -O ₃ -Re ₀ | DS | 48 | 4.360 |
| 12 | Re ₀ -O ₁ -O ₁ '-O ₁ -Re ₀ | TS | 24 | 4.527 |
| 13 | Re ₀ -O ₁ -O ₃ -O ₁ -Re ₀ | TS | 24 | 4.527 |
| 14 | Re ₀ -Re ₄ -Re ₀ | SS | 12 | 5.303 |
| 15 | Re ₀ -O ₅ -Re ₀ | SS | 30 | 5.625 |



A. Kalinko, R.A. Evarestov, A. Kuzmin, J. Purans, *J. Phys.: Conf. Ser.* **190** (2009) 012080.

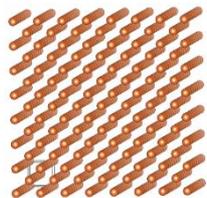
Validation of interatomic potentials for bcc tungsten using the W L₃-edge EXAFS

$$\sigma_{\text{th}}^2 = \langle u_A^2 \rangle + \langle u_B^2 \rangle - 2\langle u_A u_B \rangle$$



I. Jonane, A. Anspoks, A. Kuzmin, *Modelling Simul. Mater. Sci. Eng.* 26 (2018) 025004.

Validation of Molecular Dynamics simulations: How accurate are force-fields? EAM vs 2NN-MEAM

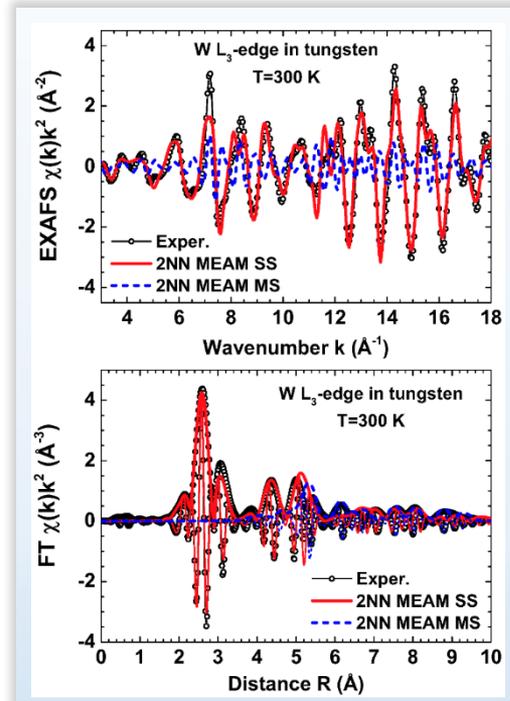
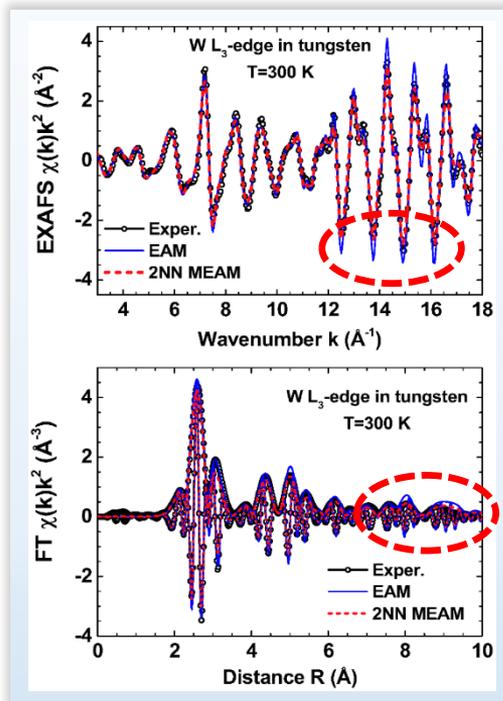


$bcc W a = 3.165 \text{ \AA}$

Supercell:
 $7a \times 7a \times 7a$
686 atoms

MD GULP code
NVT
 $T=300 \text{ K}$
 $\Delta t=0.5 \text{ fs}$
20 ps + 20 ps

EDACA + FEFF8.50L

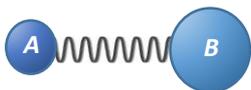


Embedded Atom Model (EAM): *M.W. Finnis and J.E. Sinclair, Phil. Mag. A 50 (1984) 45-55.*

2nd Nearest-Neighbor Modified EAM (2NN-MEAM): *B.J. Lee, M. Baskes, H. Kim, Y. Koo Cho, Phys. Rev. B 64 (2001) 184102.*

I. Jonane, A. Anspoks, A. Kuzmin, Modelling Simul. Mater. Sci. Eng. 26 (2018) 025004.

Simultaneous MSRD and MSD determination from EXAFS



$$\text{MSRD}_{AB} = \text{MSD}_A + \text{MSD}_B - 2\varphi\sqrt{\text{MSD}_A}\sqrt{\text{MSD}_B}$$

$$R \rightarrow \infty, \varphi \rightarrow 0 \ \& \ \text{MSRD}_{AB} \rightarrow \underbrace{\text{MSD}_A + \text{MSD}_B}$$

MD:

$$\text{MSD}(\text{EAM}) = 0.0023 \text{ \AA}^2$$

$$\text{MSD}(\text{MEAM}) = 0.0029 \text{ \AA}^2$$

RMC:

$$\text{MSD}(\text{RMC}) = 0.0039 \text{ \AA}^2$$

X-ray diffraction:

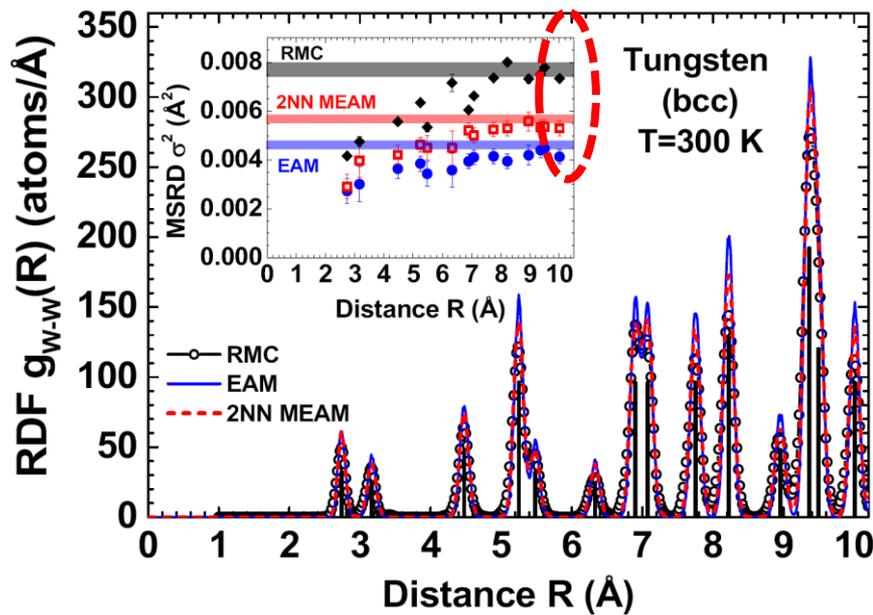
$$\text{MSD} = 0.0061 \text{ \AA}^2 \ [1]$$

$$\text{MSD} = 0.0022 \text{ \AA}^2 \ [2]$$

Lattice dynamics calculations:

$$\text{MSD} = 0.0023 \text{ \AA}^2 \ [3]$$

$$\text{MSD} = 0.0018 \text{ \AA}^2 \ [4]$$

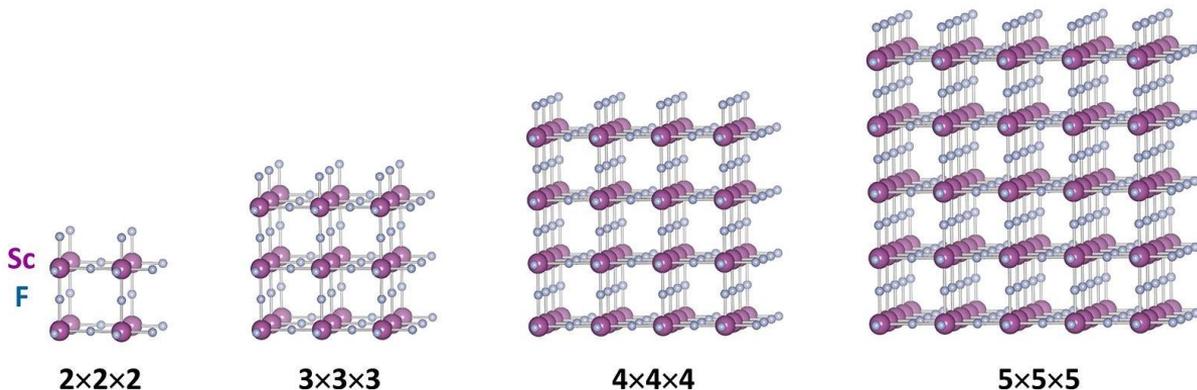


φ is a dimensionless correlation parameter

I. Jonane, A. Anspoks, A. Kuzmin, Modelling Simul. Mater. Sci. Eng. 26 (2018) 025004.

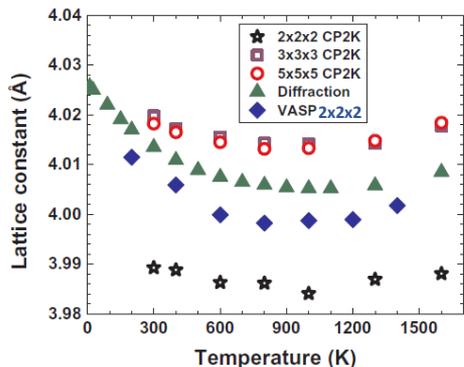
- [1] C.R. Houska, J. Phys. Chem. Solids 25 (1964) 359.
- [2] T. Paakkari, Acta Crystallogr. A 30 (1974) 83.
- [3] P.C. Fine, Phys. Rev. 56 (1939) 355.
- [4] L. Dobrzynski, P. Masri, J. Phys. Chem. Solids 33 (1972) 1603.

Cubic perovskite ScF_3 - the effect of the supercell size

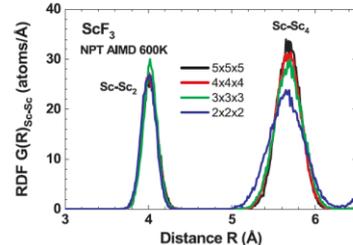
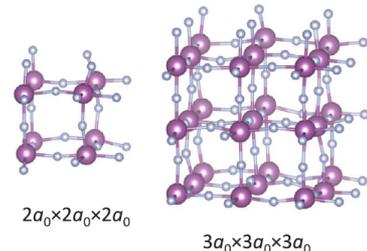
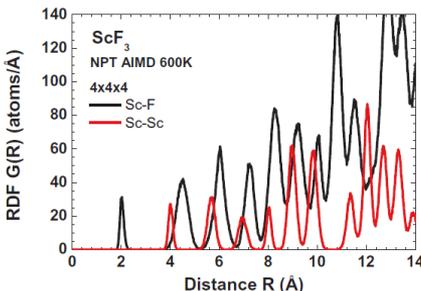
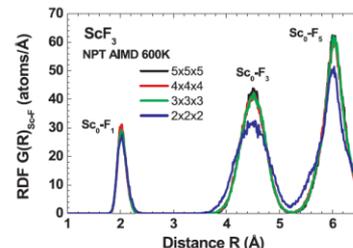
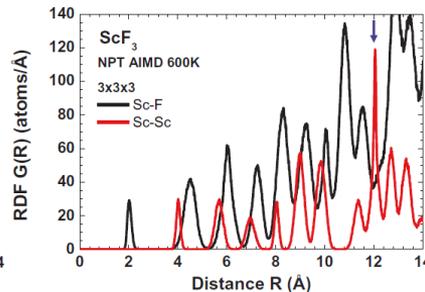
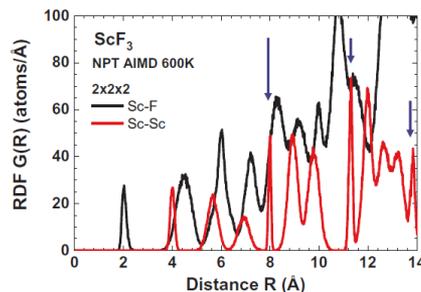
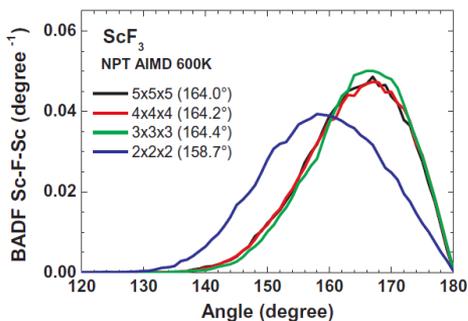


D. Bocharov, M. Krack, Yu. Rafalskij, A. Kuzmin, J. Purans, *Comp. Mater. Sci.* 171 (2020) 109198.

Ab initio molecular dynamics simulations of negative thermal expansion in ScF_3 : the effect of the supercell size



◆ P. Lazar, T. Bučko, J. Hafner, *Phys. Rev. B* 92 (2015) 224302.



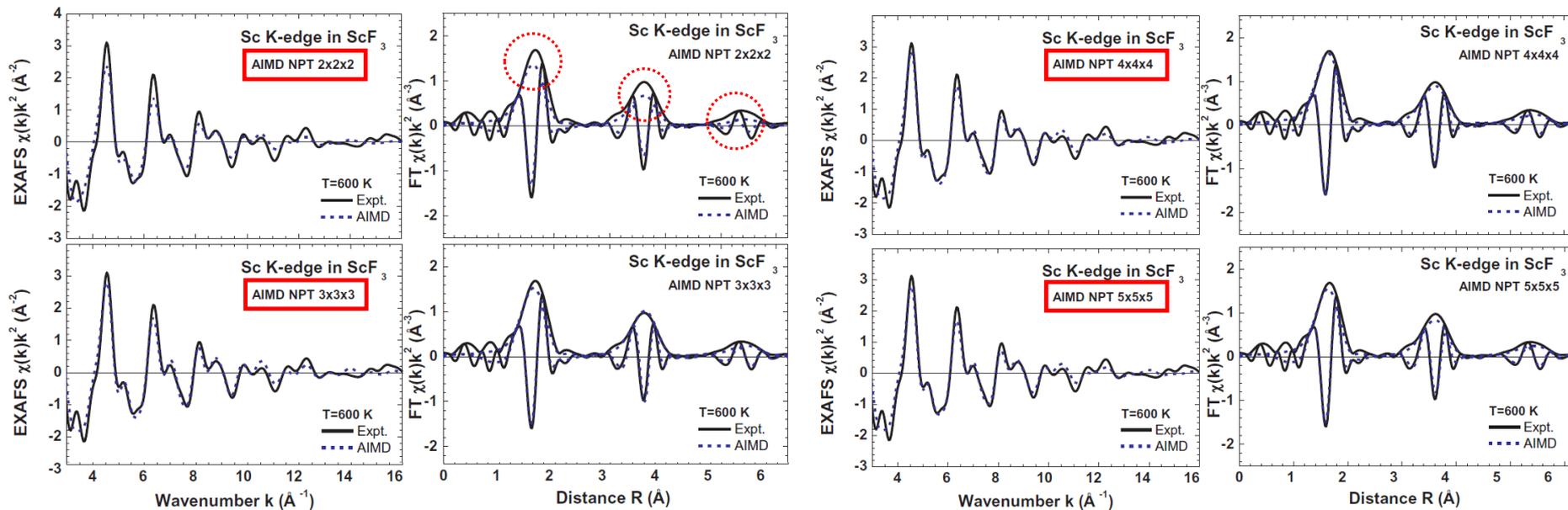
NpT AIMD simulations were performed using CP2K code.

D. Bocharov, M. Krack, Yu. Rafalskij, A. Kuzmin, J. Purans, *Comp. Mater. Sci.* 171 (2020) 109198.



<https://www.cp2k.org/>

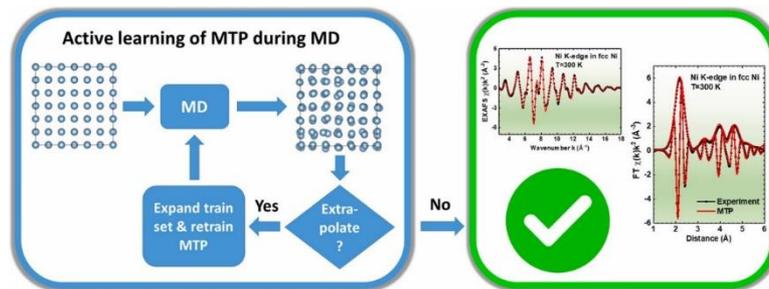
Ab initio molecular dynamics simulations of negative thermal expansion in ScF_3 : the effect of the supercell size



Small supercell size introduces artefacts into RDFs and results in inaccurate RDFs and BADFs.

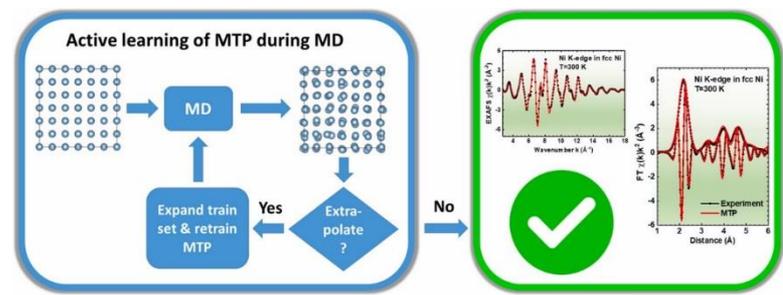
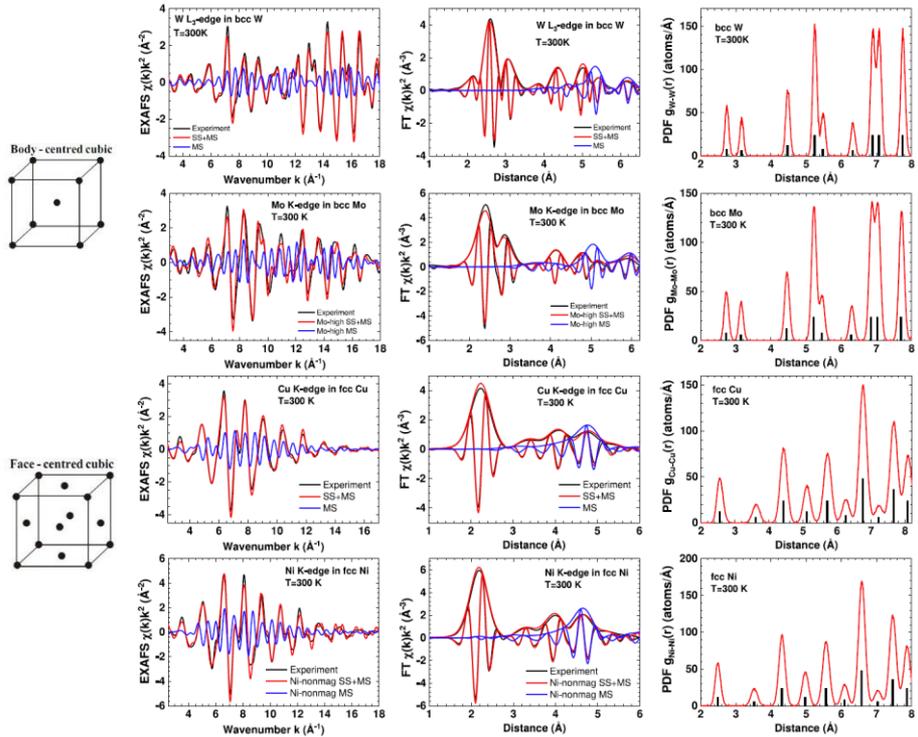
D. Bocharov, M. Krack, Yu. Rafalskij, A. Kuzmin, J. Purans, *Comp. Mater. Sci.* 171 (2020) 109198.

Validation of Machine Learning potentials



A. V. Shapeev, D. Bocharov, A. Kuzmin, *Comput. Mater. Sci.* 210 (2022) 111028.

Validation of moment tensor potentials for fcc and bcc metals using EXAFS spectra



NVT MD simulations were performed by LAMMPS using Moment Tensor Potentials (MTPs) [1,2] as implemented in the MLIP-2 software package [3].

MTP potentials were trained on-the-fly using the results of AIMD by VASP.

[1] A.V. Shapeev, *Multisc. Model. Simul.* 14 (2016) 1153–1173.
 [2] K. Gubaev, E.V. Podryabinkin, G.L. Hart, A.V. Shapeev, *Comput. Mater. Sci.* 156 (2019) 148–156.
 [3] I.S. Novikov, K. Gubaev, E.V. Podryabinkin, A.V. Shapeev, *Mach. Learn.: Sci. Technol.* 2 (2020) 025002.

<https://github.com/materialsvirtuallab/mllearn>

A. V. Shapeev, D. Bocharov, A. Kuzmin, *Comput. Mater. Sci.* 210 (2022) 111028.



Conclusions

- Molecular Dynamics is a natural way to include disorder (static and dynamic) into EXAFS simulations taking into account multiple-scattering effects.
- The MD-EXAFS approach allows one to interpret EXAFS spectra based on the multiple-scattering formalism far beyond the first coordination shell.

Good for EXAFS!

- No structural fitting parameters is used in the MD-EXAFS approach. The structural model is uniquely defined from the results of the MD simulation. But your EXAFS results will be as good as the MD ones.
- EXAFS spectra can be used as an additional physical property for interatomic potential/theory validation.

Good for Molecular Dynamics!

- Care should be taken regarding the temperature. At low temperature, special treatment is required, e.g. based on path-integral methodology*.

* D. Marx, M. Parrinello, *J. Chem. Phys.* **104** (1996) 4077-4082

Thank you for your attention!

