

Documentation for the RTA code in 3D

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Abstract

This is a first draft for the CPC presenting the TDLDAs+RTA code to the public. The layout of presentation has yet to be discussed. Presently it mixes theory and algorithm with code. We may also consider collecting all theory & numerics together then followed by a huge block detailing the code.

Keywords: electronic dissipation, time-dependent density functional theory, ...

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1. Introduction

PGR2all: ... yet to be written ...
PGR2all: Two general points yet to be clarified:
The code is still capable of coupling to a dielectric environment. We should skip that for the CPC publication.
Another question to be discussed is whether we maintain the option to run finite differences instead of FFT. If we do so, then we have to test that branch carefully.

2. Formal background: electronic DFT coupled to ionic motion

2.1. Brief introduction to TDLDA-MD

PGR2all: The explanation of formal background is extremely compact and reduced to the basic formula. Here we should provide a brief overview of the history of DFT and of the physics behind.

2.2. The total energy of the model

The basic dynamical variables are the set of single-particle (s.p.) wavefunctions with their occupation probabilities at the side of electrons and classical coordinates and momenta for the ions, together

$$\begin{array}{ll} \text{s.p. wavefunctions:} & \varphi_\alpha, \alpha = 1 \dots \Omega, \\ \text{s.p. occupation probabilities:} & w_\alpha, \alpha = 1 \dots \Omega, \\ \text{ionic coordinates:} & \mathbf{R}_I, I = 1, N_{\text{ion}}, \\ \text{ionic momenta:} & \mathbf{P}_I, I = 1, N_{\text{ion}}. \end{array} \quad (1)$$

A key quantity in connection with energy-density functionals is the electronic local density

$$\varrho_{\uparrow}(\mathbf{r}) = \sum_{\alpha \in \uparrow} w_\alpha \varphi_\alpha^+(\mathbf{r}) \varphi_\alpha(\mathbf{r}), \quad \varrho_{\downarrow}(\mathbf{r}) = \sum_{\alpha \in \downarrow} w_\alpha \varphi_\alpha^+(\mathbf{r}) \varphi_\alpha(\mathbf{r}), \quad (2)$$

which covers, in fact, two separate densities for spin-up and spin-down. To simplify the presentation of the formalism we ignore this distinction in the following and use simply one density $\varrho(\mathbf{r})$.

Starting point is an expression for the total energy of the coupled electronic and ionic

system:

$$E_{\text{total}} = E_{\text{kin}} + E_C + E_{\text{xc}} + E_{\text{ext}} + E_{\text{el,ion}} + E_{\text{kin,ion}} + E_{\text{pot,ion}}, \quad (3a)$$

$$E_{\text{kin}}[\{\varphi_\alpha\}] = \int d\mathbf{r} \sum_\alpha w_\alpha \varphi_\alpha^+ \frac{\hat{p}^2}{2m} \varphi_\alpha, \quad (3b)$$

$$E_C[\varrho] = \frac{e^2}{2} \int d^3r d^3r' \frac{\varrho(\mathbf{r})\varrho(\mathbf{r}')}{|\mathbf{r} - \mathbf{r}'|} \quad (3c)$$

$$E_{\text{xc}}[\varrho] = \int d\mathbf{r} \varrho(\mathbf{r}) \epsilon_{\text{xc}}(\varrho(\mathbf{r})) , \quad (3d)$$

$$E_{\text{ext}}[\varrho, t] = \int d\mathbf{r} \varrho(\mathbf{r}) V_{\text{ext}}(\mathbf{r}, t) , \quad (3e)$$

$$E_{\text{el,ion}}[\{\varphi_\alpha\}, \{\mathbf{R}_I\}] = \sum_I \int d\mathbf{r} \sum_\alpha w_\alpha \varphi_\alpha^+ \hat{V}_{\text{PsP}}(\mathbf{r} - \mathbf{R}_I) \varphi_\alpha , \quad (3f)$$

$$E_{\text{kin,ion}}(\mathbf{P}_I) = \sum_I \frac{\mathbf{P}_I^2}{2M_I} , \quad (3g)$$

$$E_{\text{pot,ion}}(\mathbf{R}_I) = \frac{1}{2} \sum_{J \neq I} \frac{e^2}{|\mathbf{R}_I - \mathbf{R}_J|} + V_{\text{ext,ion}}(\mathbf{R}_I, t) , \quad (3h)$$

where functionals of density are wavefunctions are indicated by square brackets and functions of coordinates by round brackets. The E_C is the direct part of the electronic Coulomb energy. The E_{xc} is the energy-density functional for electronic exchange and correlations for which we use in the code two options: the functional of [1] or the older from from [2]. The E_{ext} stands for the excitation mechanisms by external sources, either from a photon pulse or from the Coulomb field of a fast bypassing ion, see section 2.2.5. This part is, of course, absent in static calculations of the ground state. The $E_{\text{el,ion}}$ carries the interaction with the ions which is usually described by a pseudo-potential \hat{V}_{PsP} , see section 2.2.2, or may be simplified in terms of the jellium model, see section 2.2.3. Ionic kinetic and potential energy are described by the obvious classical expressions. The $V_{\text{ext,ion}}$ in the ionic potential energy describes the action of an external field on the ions (which is usually negligible as compared to the effect on the electrons).

2.2.1. Self-interaction correction (SIC)

PGR2all: We have yet to decide which levels of SIC we keep in the code and then explain them here properly.

2.2.2. The pseudo-potentials

There is a great variety of pseudo-potentials available. The code employs two variants. Particularly efficient and simple to use are local pseudo-potentials integrated from a pseudo-density which is represented as sum of Gaussians. The corresponding potential is then a sum of error functions

$$V_{\text{PsP}}(\mathbf{r}) = \sum_{i=1}^2 c_i \frac{\text{erf}(\mathbf{r}/\sigma_i)}{|(\mathbf{r})|} , \quad (4a)$$

$$\text{erf}(x) = \sqrt{\frac{2}{\pi}} \int_0^x dy e^{-y^2} , \quad (4b)$$

where the σ_i are widths and the strength parameters c_i have to line up to the total charge of the ionic core $c_1 + c_2 = Z_{\text{ion}}$. This pseudo-potential is well suited for alkaline atoms for which it was originally developed [3].

More involved, but also more versatile in the applicability, is the pseudo-potential from [4] which is composed from a local and a non-local part as

$$\hat{V}_{\text{PsP}} = V_{\text{loc}}(\mathbf{r} - \mathbf{R}) + \hat{V}_{\text{nloc}} , \quad (5a)$$

$$V_{\text{loc}}(\mathbf{r} - \mathbf{R}) = -\frac{Z_{\text{ion}}}{x} \text{erf}(x) + e^{-x^2} \sum_{n=0}^3 C_{n+1} 2^n x^{2n} , \quad x = \frac{|\mathbf{r} - \mathbf{R}|}{r_{\text{loc}}} , \quad (5b)$$

$$\hat{V}_{\text{nloc}}(\mathbf{r}, \mathbf{r}') = \sum_{jl\mu} \mathcal{G}(\mathbf{r} - \mathbf{R}) \mathcal{H}_{jl} \mathcal{G}(\mathbf{r}' - \mathbf{R}) , \quad (5c)$$

where \mathbf{R} is the position of the ionic core with respect to which the pseudo-potential is defined. The \mathcal{G} in the non-local part (5c) serve to project out the electronic states which are occupied in the ionic core. These projector functions for the first few quantum numbers read

$$\mathcal{G}_{000}(\mathbf{x}) = r_{\text{nloc}}^{-3} \pi^{-3/2} \exp\left(-\frac{x^2}{2r_{\text{nloc}}^2}\right) , \quad (5d)$$

$$\mathcal{G}_{01\mu}(\mathbf{x}) = \sqrt{\frac{2}{3}} r_{\text{nloc}} \nabla_\mu \mathcal{G}_{000}(\mathbf{x}) , \quad (5e)$$

$$\mathcal{G}_{100}(\mathbf{x}) = \frac{r_{\text{nloc}}^2}{\sqrt{6}} \left(2\Delta - \frac{3}{r_{\text{nloc}}^2}\right) \mathcal{G}_{000}(\mathbf{x}) . \quad (5f)$$

The projection looks expensive at first glance. However, one can exploit the fact that the Gaussians cover only a small region space of order of a few multiples of the non-local radius r_{nloc} . Thus one needs to evaluate the projector only on a small sub-grid which reduces expense dramatically.

2.2.3. The soft jellium model for the ionic background

The electronic wavefunctions of metals and metal clusters are spread softly over the whole system and hardly resolve the spatial structure the ionic cores. This allow to replace the detailed ionic background by a smooth, positive background density. Such jellium approximation is a standard in the theory of bulk metals [5] and the adaptation to a finite cluster is straightforward. One carves from bulk jellium a finite element of constant positive charge corresponding to the average bulk density. The volume is chosen such that its total charge coincides with the given ionic charge. For finite clusters, it is advantageous to use jellium with a soft surface profile. This is more suited for numerical handling and it improves the quality of the model, e.g., by producing a correct peak energy for the optical response of metal clusters [6]. Versatile and easy to handle in this

respect is a Woods-Saxon profile for the jellium density

$$\varrho_{\text{jel}}(\mathbf{r}) = \frac{3}{4\pi r_s^3} \left[1 + \exp \left(\frac{|\mathbf{r}| - R(\vartheta, \phi)}{\sigma_{\text{jel}}} \right) \right]^{-1}, \quad (6a)$$

$$\text{with } R(\vartheta, \phi) = R_{\text{jel}} \left(1 + \sum_{lm} \alpha_{lm} Y_{lm}(\vartheta, \phi) \right), \quad (6b)$$

$$\int d\mathbf{r} \varrho_{\text{jel}} = N_{\text{ion}}. \quad (6c)$$

The jellium radius R_{jel} is determined by eq. (6c), the normalization to desired number of ions. The central density is determined by the bulk density $\rho_0 = 3/(4\pi r_s^3)$ and the Wigner-Seitz radius r_s is a genuine material parameter [5]. The σ_{jel} parametrizes the surface width and the transition from 90% to 10% bulk density is achieved within $4\sigma_{\text{jel}}$. The model does also allow to describe deformed clusters by angular dependence $R(\vartheta, \phi)$ of the extension. This is achieved through the deformation coefficients α_{lm} weighting the impact of the spherical harmonics. For example, axially symmetric ellipsoids are tuned by α_{20} , positive values producing prolate and negative values oblate shapes. The α_{30} generate octupole (pear-like) shapes which have considerable influence in metal cluster spectroscopy and α_{40} generates hexadecapole which play a role for fine-tuning the shape [7]. Triaxial shapes are produced by moments with $m \neq 0$. The cluster radius R_{jel} is fixed by the others parameters through eq. (6c) which is to be solved numerically by a root finding procedure. After all, the leading parameters of the soft jellium model Eq. (6) are the Wigner-Seitz radius r_s and the surface thickness σ_{jel} . They are universal for a given material. Typical values are $r_s \sim 4 a_0$ with $\sigma_{\text{jel}} \sim 0.9 a_0$ for Na clusters, $r_s \sim 2.66 a_0$ and $\sigma_{\text{jel}} \sim 0.76 a_0$ for Mg clusters, or $r_s \sim 3 a_0$ and $\sigma_{\text{jel}} \sim 0.78 a_0$ for Ag clusters. The deformation parameters α_{lm} depend on the actual cluster and strongly vary with system and size.

The jellium approximation then consists in discarding the ionic contribution to the total energy (3), i.e. the terms $E_{\text{kin,ion}}$ and $E_{\text{pot,ion}}$, and to replace the pseudo-potential background in $E_{\text{el,ion}}$ by the Coulomb potential of the jellium density (6). There is no dynamics associated with the jellium. The model thus applies to situations where ionic motion can be ignored.

2.2.4. Phenomenological electronic shell models

Experience shows that KS potentials for metal clusters have a typical profile characterized by a flat bottom and a smooth transition to zero. For neutral clusters, one can approximate that very well by a Woods-Saxon profile

$$V_{\text{WS}}(\mathbf{r}) = -V_0 \left[1 + \exp \left(\frac{|\mathbf{r}| - R(\vartheta, \phi)}{\sigma_{\text{WS}}} \right) \right]^{-1}, \quad (7)$$

where, again, a possible deformation can be parametrized by an angular dependence of the radius $R(\vartheta, \phi) = R_0 (1 + \sum_{lm} \alpha_{lm} Y_{lm}(\vartheta, \phi))$, similar to the jellium model of Eq. (6) for the density. In Eq. (7) the potential depth V_0 is taken as the average binding potential in bulk matter. Computations in a fixed potential are somewhat simpler than with the self-consistent scheme. The Woods-Saxon model has thus been very often used

for describing the electronic shell structure of clusters, particularly in early studies, for a review see [8]. It is meanwhile somewhat out of fashion. But we provide it in the code for quick tests and comparison.

A substantial further simplification is achieved by realizing that shell effects are determined by the states near the Fermi surface and that these states practically see a harmonic potential. This suggests to use for first estimates a simple harmonic oscillator shell model. To mimic surface profile of the Woods-Saxon potential and so reproduce the sequence of shell closures in metal clusters a phenomenologically tuned \hat{l}^2 term is added to the oscillator model. This yields the Clemenger-Nilsson model for the mean field potential

$$\hat{V}_{\text{CN}} = \frac{m}{2} (\omega_x^2 x^2 + \omega_y^2 y^2 + \omega_z^2 z^2) - V_{l2} \hbar \bar{\omega} \left(\hat{l}^2 - n(n+3)/6 \right) \quad (8)$$

where n is the global shell number ($n = n_x + n_y + n_z$). The three separate curvatures ω_i allow one to accommodate deformed situations including triaxiality. Volume conservation restricts their choice to $\omega_x \omega_y \omega_z = \omega_0^3 = \text{constant}$. The separate values can also be expressed in terms of the quadrupole deformation α_{lm} as introduced above and in Eq. (6), e.g. for axially symmetric shapes by $\omega_x = \omega_y = \omega_0 \exp(\alpha_{20} \sqrt{5/4\pi})$, $\omega_z = \omega_0 \exp(-2\alpha_{20} \sqrt{5/4\pi})$. The parameter V_{l2} serves to tune the downshift of high angular momentum orbits and thus of the shell sequence. The Clemenger-Nilsson model was introduced into clusters physics in [9] taking up a much similar nuclear oscillator model [10]. An extensive review of early applications is given in [11]. The oscillator model is kept in the code mainly for testing because one has analytical the solutions for $V_{l2} = 0$.

Both shell models, Woods-Saxon or harmonic oscillator, overrule the total energy (3). What remains is only the kinetic energy (3b) together with the given model potentials. Also this approach is, as the jellium model, applicable only in circumstances where ionic motion plays little role.

2.2.5. External excitation fields

Lasers are the most important and very flexible means for a dedicated, well tuned excitation of electronic systems. They produce a strong coherent electromagnetic field which can be well approximated by a classical time-dependent electromagnetic field. Typical wavelengths are in the range of several hundredths of nm. This is a huge distance as compared to the spatial extension of atoms, molecules, and (most) clusters. One can thus treat the laser field in the limit of long wavelengths ($k \rightarrow 0$) which means to deal with a spatially homogeneous electrical field \mathbf{E} at the cluster site and we can neglect the effect of the magnetic field. The coupling Hamiltonian leaves of gauge transformation [12]. The external-field operator in velocity gauge reads¹

$$V_{\text{ext}} = e \mathbf{E}_0 F(t) \hat{\mathbf{p}} \quad , \quad F(t) = \int dt' f(t') \exp(-i\omega_{\text{las}} t') \quad , \quad (9)$$

where $f(t')$ is the envelope of the photon pulse. The same pulse in space gauge becomes

$$V_{\text{ext}} = e \mathbf{E}_0 f(t) \cdot \hat{\mathbf{r}} \exp(-i\omega_{\text{las}} t) \quad (10)$$

¹I am not sure that we should show velocity gauge. It may be useful when explaining the computation of PES later on.

which is simpler to handle because the laser field acts here simply as a time-dependent local operator. The code thus uses the external field in space gauge. Transformation to velocity gauge, if needed, can be performed a posteriori by standard rules of gauge transformation [13]². The photon pulse is characterized by frequency ω_{las} , peak field strength $E_0 = |\mathbf{E}_0|$, polarization \mathbf{E}_0/E_0 , and time profile $f(t)$. The field strength is usually parametrized in terms of the laser intensity I as

$$E_0 = c_{EI} I^{1/2} \quad , \quad c_{EI} = 1.07 * 10^{-8} \frac{\text{eV}}{\text{a}_0} \left(\frac{\text{W}}{\text{cm}^2} \right)^{-1/2} . \quad (11)$$

The profile is a matter of debate. The experimental pulse profiles are not precisely known either. It is assumed to be a well peaked function with a certain full-width at half maximum (FWHM), in practice well approximated by a Gaussian. In order to have a finite support for the pulse, we use the \sin^2 pulse

$$f(t) = \begin{cases} \sin^2 \left(\pi \frac{t}{2T_{\text{pulse}}} \right) & \text{for } t \in \{0, 2T_{\text{pulse}}\} \\ 0 & \text{else} \end{cases} . \quad (12)$$

It is close to a Gaussian pulse in the vicinity of peak field strength and combines high spectral selectivity with finite bounds. Note that the form (12) is scaled such that the pulse parameter T_{pulse} is identical with the full-width at half-maximum (FWHM).

Probing electronic systems by beams of charged particles is a standard tool in atomic and molecular physics [14] and is also used in cluster physics. Highly charged ions, protons and to some extend also electrons can be considered as being structureless. What counts is only their Coulomb field. Charged ions are heavy and can be treated with classical trajectories $\mathbf{R}_{\text{ext}}(t)$. For sufficiently heavy and fast projectiles, one can approximate these trajectories by straight lines and that is what is done in the code. In any case, the effect of the ion (charge Z_{ext}) on the system can again be described by a time dependent external field

$$V_{\text{ext}}(\mathbf{r}, t) = \frac{Z_{\text{ext}} e^2}{|\mathbf{r} - \mathbf{R}_{\text{ext}}(t)|} . \quad (13)$$

Magnetic effects are neglected. They may play a role only for extremely fast ions in the relativistic domain. The ionic trajectories are characterized by the ion velocity v_{ion} and the impact parameter b_{ion} which is the distance of closest approach. The velocity is more or less well defined by the experimental setup. But the impinging ion beam will cover a broad range of impact parameters. From the theoretical side, one has then to run several calculations with systematically varied impact parameters. Reaction cross sections are then computed by integration of the reaction probability over impact parameters, for examples see [15].

2.3. Equations of motion

2.3.1. Electronic Kohn-Sham equations

Static and dynamical equations determining structure and dynamics of the electron cloud are determined by variation of the total energy with respect to the s.p. wavefunctions φ_α . This yields what is called the Kohn-Sham (KS) equations [16]. We use

²Cross ref to phase correction in PES in section 2.5.

energy-density functionals [1, 2] i.e. functionals in local density approximation (LDA). The dynamical application is called time-dependent LDA (TDLDA) and we use this acronym also in cases where we add SIC (see section 2.2.1) to the treatment.

The time-dependent KS equations determining the propagation of electronic ground wavefunctions are

$$\hat{h}_{\text{KS}}\varphi_{\alpha} = i\partial_t\varphi_{\alpha} \quad \text{for } \alpha \in \{1, \dots, N_{\text{el}}\} , \quad (14a)$$

$$\hat{h}_{\text{KS}} = \frac{\hat{p}^2}{2m} + \underbrace{V_C(\mathbf{r}) + \hat{V}_{\text{xc}} + \hat{V}_{\text{back}} + V_{\text{ext}}(\mathbf{r}, t)}_{V_{\text{KS}}} , \quad (14b)$$

$$V_C(\mathbf{r}) = e^2 \int d^3 r' \frac{\varrho(\mathbf{r}')}{|\mathbf{r} - \mathbf{r}'|} , \quad (14c)$$

$$\hat{V}_{\text{xc}}(\mathbf{r}) = \epsilon(\varrho(\mathbf{r})) + \varrho(\mathbf{r}) \frac{\partial \epsilon_{\text{xc}}(\varrho(\mathbf{r}))}{\partial \varrho} . \quad (14d)$$

The exchange-correlation potential \hat{V}_{xc} involves a construction $\partial \epsilon_{\text{xc}} / \partial \varrho$ which is the formal derivative of the exchange correlation energy per particle ϵ_{xc} with respect to the density ϱ considered as an independent variable. The time-dependent KS equations analogously read

$$\hat{h}_{\text{KS}}\varphi_{\alpha} = \varepsilon_{\alpha}\varphi_{\alpha} \quad \text{for } \alpha \in \{1, \dots, N_{\text{el}}\} , \quad (14e)$$

where \hat{h}_{KS} is composed in the same manner as in the time-dependent case (instantaneous approximation for V_{xc}). The Kohn-Sham equations (14e) pose a stationary eigenvalue problem. They provide the electronic ground state of a system. The time-dependent Kohn-Sham equations (14a) pose an initial value problem. The natural starting point is the ground state as obtained from the stationary Kohn-Sham equation. The numerical solution of the KS equations is explained in section 2.4. The time evolution delivered by the dynamical KS Eqs. (14a) can be expressed formally by the unitary one-body time-evolution operator

$$\hat{U}(t, t') = \hat{\mathcal{T}} \exp \left(-i \int_t^{t'} \hat{h}(t'') dt'' \right) \quad (15a)$$

where $\hat{\mathcal{T}}$ is the time-ordering operator. This yields a closed expression for the time-evolution of s.p. states

$$|\phi_{\alpha}(t)\rangle = \hat{U}(t, t')|\phi_{\alpha}(t')\rangle . \quad (15b)$$

This compact form will be used later on in connection with the extension of the propagation by dissipation, see section 4.

2.3.2. LDA and TDLDA with mixed states

So far, we dealt with the equations for pure states. In case of stationary states at finite temperature and in case of dissipative we encounter mixed states. This is described by associating an occupation weight w_{α} with each s.p. state φ_{α} as provided already in the initial setup in section 2.2. The weights w_{α} becomes crucial in the dissipative dynamics, see section 4, they are kept frozen during TDLDA propagation, and they matter again for stationary states at finite temperature in which case they are determined by

$$w_{\alpha} = \frac{1}{1 + \exp((\varepsilon_{\alpha} - \mu)/T)} \quad (16)$$

where the ε_α are the s.p. energies from the stationary KS Eqs. (14e) and μ is the chemical potential tuned such that the total electron number is reproduced by $\sum_\alpha w_\alpha = N$.

In case of mixed states, it is advantageous to express the entity $\{\varphi_\alpha, w_\alpha\}$ in compact manner by the one-body density operator

$$\hat{\rho} = \sum_{\alpha=1}^{\infty} |\phi_\alpha\rangle w_\alpha \langle\phi_\alpha| . \quad (17)$$

The solution of the thermal KS Eqs., i.e. Eqs. (14e) together with (16), provides the density operator immediately in this form which is called natural orbital representation (diagonal in the s.p. states). In general, the one-body density operator can be non-diagonal with respect to given s.p. states. This will play a role later on.

As said above, the occupation weights w_α are kept frozen during TDLDA. The KS equations (14a) then can be written compactly as³

$$i\partial_t \hat{\rho} = [\hat{h}[\varrho], \hat{\rho}] \quad (18)$$

where $\hat{h}[\varrho]$ is the KS Hamiltonian as above. This pure mean-field propagation (18) leaves the occupation weights W_α unchanged and propagates only the s.p. states. The mean-field propagation of an initial state (17) then reads

$$\hat{\rho}(t) = \sum_{\alpha=1}^{\infty} |\phi_\alpha(t)\rangle w_\alpha \langle\phi_\alpha(t)| = \hat{U}(t,0)\hat{\rho}(0)\hat{U}^{-1}(t,0) \quad (19)$$

where \hat{U} is the mean-field evolution operator (15a).

2.3.3. Ionic dynamics

We finally add the ionic dynamics to complete the picture. Ions are described by classical Molecular Dynamics (MD), i.e. classical equations of motion, under the influence of their mutual Coulomb force, the forces experienced from the electrons, and possibly external forces. We start from the total energy (3) which serves as classical Hamiltonian in terms of \mathbf{R}_I and \mathbf{P}_I . Standard variation yields the classical equations-of-motion for the ions of positions \mathbf{R}_I and momenta \mathbf{P}_I

$$\partial_t \mathbf{P}_I = -\nabla_{\mathbf{R}_I} \left[E_{\text{pot,ion}}(\mathbf{R}_I) + \sum_{\alpha=1}^{N_{\text{el}}} (\varphi_\alpha | V_{\text{PsP}}(\mathbf{r} - \mathbf{R}_I) | \varphi_\alpha) \right] , \quad (20a)$$

$$\partial_t \mathbf{R}_I = \frac{\mathbf{P}_I}{M_I} . \quad (20b)$$

They are to be propagated simultaneously with TDLDA for the electrons, here represented by wavefunctions φ_α . The ionic motion is coined molecular dynamics (MD). The simultaneous propagation scheme is called TDLDA-MD. It applies to all dynamical situations including those that are far from the adiabatic limit and embraces truly dia-batic scenarios. The practical realization adds the (simple) classical propagation to the evolution of the electronic states. For solution schemes, see section 2.4.⁴

³I wonder whether we should shift this introduction of the one-body density matrix to the RTA section.

⁴Should we add a word on the limitations of TDLDA-MD?

2.4. Numerical schemes

2.4.1. Grid representation and derivatives

All wave functions and fields are defined on a three-dimensional (3D) equidistant Cartesian grid of nx by ny by nz grid points. The spacing between the points is given as dx , dy , and dz in units of a_0 . For reasons of equilibrated accuracy, it is highly recommended to give the same value to all of them. Typical values depend on the atoms/ions involved. The pseudo-potentials (section 2.2.2) set the pace. The dx , dy , dz must not be larger than $\sqrt{2 \log 2}$ times the smaller radius in the pseudopotential. In case of the jellium model (section refsec:jell), the spacing should be of order of the surface width σ_{jel} .

The grid is automatically arranged in such a way that in each direction the same number of grid points are located on both sides of the origin. The coordinate values for e. g., the x -direction are thus⁵:

$$\begin{aligned} -\frac{\text{nx}-1}{2}\text{dx}, \quad &-\frac{\text{nx}-1}{2}\text{dx} + \text{dx}, \quad \dots \quad -\frac{\text{dx}}{2}, \\ &+\frac{\text{dx}}{2}, \quad \dots \quad \frac{\text{nx}-1}{2}\text{dx}. \end{aligned} \quad (21)$$

The corresponding values are available in the arrays $x(\text{nx})$, $y(\text{ny})$, and $z(\text{nz})$.

The computation of currents and the action of the kinetic energy operator first and second derivatives. We have two options for that in the code. Standard is the definition of derivatives via Fourier transformation⁶. This delivers high precision at affordable expense space. For simplicity, we explain here the Fourier strategy for one dimension.

Given are nx discrete grid points x_ν in coordinate space. They are mapped to the same number of grid points k_n in Fourier space (physically equivalent to momentum space) as

$$x_\nu = \left(-\frac{\text{nx}-1}{2} + \nu \right) \text{dx} , \nu = 1, \dots, \text{nx} , \quad (22a)$$

$$k_n = (n-1)\text{dk}, \quad n = 1, \dots, \text{nx}/2 , \quad (22b)$$

$$k_n = (n - \text{nx} - 1)\text{dk}, \quad n = \text{nx}/2 + 1, \dots, \text{nx} , \quad (22c)$$

$$\text{dk} = \frac{2\pi}{\text{nx} \cdot \text{dx}} .$$

Note the particular indexing for the k -values. In principle, the values $k_n = (n-1)\text{dk}$ for all n are equivalent for the Fourier transform, but for the second half of this range the negative k -values should be chosen because of their smaller magnitude. For the Fourier expansion, $k = -\text{dk}$ and $k = (\text{nx}-1)\text{dk}$ are equivalent because of periodicity in k -space.

A function $f(x_\nu)$ in coordinate space is mapped to a function $\tilde{f}(k_n)$ in Fourier space by

$$\tilde{f}(k_n) = \sum_{\nu=1}^{\text{nx}} \exp(-ik_n x_\nu) f(x_\nu) , \quad (22d)$$

$$f(x_\nu) = \frac{1}{\text{nx}} \sum_{n=1}^{\text{nx}} \exp(ik_n x_\nu) \tilde{f}(k_n) \quad (22e)$$

⁵PGRtoFC: please check and correct if necessary.

⁶Do we keep 3-point derivatives? Then we have to explain them as well.

This complex Fourier representation implies that the function f is periodic with $f(x + \mathbf{d}\mathbf{x} \cdot \mathbf{n}\mathbf{x}) = f(x)$. The appropriate integration scheme is the trapezoidal rule which complies with the above summations adding up all terms with equal weight. The derivatives of the exponential basis functions are

$$\frac{d^m}{dx^m} \exp(i k_n x) = (ik_n)^m \exp(i k_n x) . \quad (23)$$

Computation of the m -th derivative thus becomes a trivial multiplication by $(ik_n)^m$ in Fourier space. Time critical derivatives are best evaluated in Fourier space using the fast Fourier transformation (FFT). To that end a forward transform (22d) is performed, then the values $\tilde{f}(k_n)$ are multiplied by $(ik_n)^m$ as given in Eq. (23) and finally transformed $(ik_n)^m \tilde{f}(k_n)$ back to coordinate space by the transformation (22e).

A word is in order about the first derivative. The upper point in the k -grid, $\mathbf{d}\mathbf{k} \mathbf{n}\mathbf{x}/2$, is ambiguous. Exploiting periodicity, it could be equally well $-\mathbf{d}\mathbf{k} \mathbf{n}\mathbf{x}/2$. Both choices introduce an unwanted bias. We circumvent the problem by setting $k_{\mathbf{n}\mathbf{x}/2} = -k_{\mathbf{n}\mathbf{x}/2} = 0$.

2.4.2. Electronic ground state

The static KS equations (14e), optionally combined with thermal occupation (16) of s.p. states are solved iteratively. The wave functions are iterated with a gradient step which is accelerated by pre-conditioning with kinetic-energy damping [17, 18]

$$\varphi_\alpha^{(n+1)} = \mathcal{O} \left\{ \varphi_\alpha^{(n)} - \frac{\delta}{\hat{T} + E_0} \left(\hat{h}^{(n)} - \langle \varphi_\alpha^{(n)} | \hat{h}^{(n)} | \varphi_\alpha^{(n)} \rangle \right) \varphi_\alpha^{(n)} \right\} \quad (24)$$

where $\hat{T} = \hat{p}^2/(2m)$ is the operator of kinetic energy, \mathcal{O} means ortho-normalization of the whole set of new s.p. wavefunctions, and the upper index indicates the iteration number. Note that this sort of kinetic-energy damping is particularly suited for the fast Fourier techniques that we use in the present code. The damped gradient step has two numerical parameters, the step size δ and the damping regulator E_0 . The latter should be chosen typically of the order of the depth of the local potential V_{KS} . Typical values are $E_0 = 1...5$ Ry depending on the material under consideration. After proper choice of E_0 , the step size is of order of $\delta = 0.1...0.5$. Larger values yield faster iteration, but can run more easily into pathological conditions.

In case of calculations at finite temperature (switched by setting `temp > 0`), the Fermi distribution (16) is determined using the actual s.p. energies $\varepsilon_\alpha = \langle \varphi_\alpha | \hat{h} | \varphi_\alpha \rangle$ and adjusting the chemical potential μ such that the correct total electron number is reproduced, $\sum_\alpha w_\alpha = N$, which is done by bisection⁷.

Having the new s.p. wavefunctions and occupations, the new local electron densities (40b) are computed and then the new KS Hamiltonian (14). This then provides the starting point for the next iteration. The process is continued until sufficient convergence is achieved. We consider as the convergence criterion the averaged energy variance of the

⁷Check.

single particle states

$$\overline{\Delta\varepsilon} = \sqrt{\frac{\sum_{\alpha} w_{\alpha} \Delta\varepsilon_{\alpha}^2}{N_{\text{el}}}}, \quad (25a)$$

$$\Delta\varepsilon_{\alpha}^2 = \langle \psi_{\alpha} | \hat{h}^2 | \psi_{\alpha} \rangle - \varepsilon_{\alpha}^2, \quad (25b)$$

$$\varepsilon_{\alpha} = \langle \psi_{\alpha} | \hat{h} | \psi_{\alpha} \rangle. \quad (25c)$$

Vanishing total variance $\overline{\Delta\varepsilon}$ signals that we have reached minimum energy, i.e. a solution of the KS equations. However, this may be only a local minimum (isomeric state). It requires experience to judge whether one has found the absolute energy minimum. In case of doubt, one should redo a couple of static iterations from very different initial configurations.

Initialization of the s.p. wavefunctions can be done in two ways. one is to take the wave functions of the deformed harmonic oscillator, for details see e.g. [19]. These are characterized by $\vec{n} = (n_x, n_y, n_z)$, the number of nodes in each direction. We sort the states in order of increasing oscillator energy $\epsilon_{\alpha}^{(0)} = \hbar\omega_x n_x + \hbar\omega_y n_y + \hbar\omega_z n_z$ and stop if the desired number of states is reached. The deformation of the initializing oscillator influences the initial state in two ways: first, through the deformation of the basis wave functions as such, and second, through the energy ordering of the $\epsilon_{\alpha}^{(0)}$ and corresponding sequence of levels built. Variation of initial conditions means basically a variation of the oscillator radius and deformation. In particular, the initial deformation decides in which local minimum the KS iteration will terminate. This initialization by harmonic oscillators is well suited for metallic bonding where the wavefunctions spread over the whole system. Covalent bonding produces localized states and here it is more appropriate to start also from localized states. In that case, we place Gaussians at each ionic site and, of more is needed, higher harmonic oscillator wavefunctions. Bookkeeping is more involved then and will be explained in connection with the input to the code.

2.4.3. Ionic ground state

Three strategies are implemented for the optimizing the ionic configuration: steepest descend, dynamical cooling, and simulated annealing. The simplest method is steepest descent. For given electronic configuration, one computes the Hellmann-Feynman forces on the ions and follows their direction for a short step. One re-iterates to the new electronic ground state and repeats the steps until convergence. This is the fastest method but prone to get stuck in side-minima. one should use it only if one knows a reliable starting configuration.

Dynamical cooling allows to explore more of the energy landscape and thus to avoid distraction by unimportant side-minima. From a given starting configuration, one runs full TDLDA-MD and keeps a protocol of ionic kinetic energy. Starting from a non-minimal configuration, this kinetic energy will first increase which is accompanied by a corresponding decrease of potential energy. As soon as kinetic energy turns to decrease, we stop propagation and reset all ionic velocities to zero. This defines the new starting point for the next round. The procedure is stopped if gain in kinetic energy falls below a given level of precision. This method is more forgiving than steepest descent. Still, it is also often kept in isomeric minima. One has to rerun it from different initial configurations to explore the landscape of minima.

The most elaborate, expensive, and reliable method is simulated annealing, for a detailed description see [20]. Simulated annealing explores in Monte-Carlo fashion the energy landscape with a supposed thermal distribution of configurations thereby reducing the temperature successively which leads at the end to a ground state minimum for temperature zero. The method has several parameters which need to be tuned carefully to a given situation. It requires some experience to use it efficiently.

2.4.4. Electronic propagation

Electronic dynamics at the level of TDLDA is governed by the time-dependent KS equation (14a). formally resolved as. The code offers two different ways of determining the solution. Both methods starts directly from the formulation as propagator (15a). The first method, called exponential evolution, employs simply a Taylor expansion of the propagator

$$\begin{aligned} |\psi_\alpha(t+\Delta t)\rangle &= \hat{\mathcal{T}} \exp\left(-\frac{i}{\hbar} \int_t^{t+\Delta t} dt' \hat{h}(t')\right) |\psi_\alpha(t)\rangle \\ &\approx \sum_{n=0}^m \frac{(-i\Delta t)^n}{\hbar^n n!} \hat{h}^n \psi, \end{aligned} \quad (26)$$

where \hat{h} in the expansion is taken at fixed time. At time t , only $\hat{h}(t)$ is known. However, using that in the approximate propagator (26) generates undue bias on t with disastrous consequences for energy conservation. The solution is to use a predictor-corrector scheme. For the predictor, we perform a half time step, i.e. using Eq. (26) with $\hat{h}(t)$ and $\Delta t/2$. This produces an intermediate wavefunction $|\psi_\alpha(t+\Delta t/2)\rangle$ with subsequent densities and KS Hamiltonian $\hat{h}(t+\Delta t/2)$. For the corrector step, we perform a full time step (26) with using the intermediate KS Hamiltonian $\hat{h}(t+\Delta t/2)$ in the expansion. Appropriate values for the order of expansion are $m = 4\dots 6$. Below that, conservation laws are at stake, and above that, we will not gain much because the time step is also limited by the speed of change of $\hat{h}_{\text{KS}}(t)$. Altogether, this exponential evolution provides a reliable propagation with satisfying norm and energy conservation.

An alternative propagation scheme is T - V -splitting [21] we we extend here for use in connection with non-local pseudo-potentials 2.2.2. The KS Hamiltonian is split into three pieces

$$\hat{h}_{\text{KS}} = \hat{T}_{\text{kin}} + \hat{V}_{\text{nonl}} + V_{\text{KS,loc}}(\mathbf{r}, t) \quad (27a)$$

where \hat{V}_{nonl} is the piece stemming from the non-local part of the pseudo-potentials while $V_{\text{KS,loc}}(\mathbf{r}, t)$ collects all parts which form together a local potential. The latter is strongly time dependent due to the self-consistent electronic contributions. The non-local part depends only on ionic motion which is snail-slow at electronic scale such that we can ignore time dependence during an electronic step. We define a propagator for each one

of the three parts and factorize the full propagator as

$$|\varphi_\alpha(t_1)\rangle = \hat{U}_{\text{KS}}(t_1, t_0)|\varphi_\alpha(t)\rangle \approx e^{-\frac{i}{\hbar} \frac{\Delta t}{2} V_{\text{KS,loc}}(\mathbf{r}, t_1)} \hat{U}_{\text{nonl}} e^{-\frac{i}{\hbar} \Delta t \hat{T}_{\text{kin}}} \hat{U}_{\text{nonl}} e^{-\frac{i}{\hbar} \frac{\Delta t}{2} V_{\text{KS,loc}}(\mathbf{r}, t_0)} |\varphi_\alpha(t_0)\rangle , \quad (27\text{b})$$

$$\hat{U}_{\text{nonl}} = \sum_{n=0}^m \frac{(-i\Delta t)^n}{2\hbar^n n!} \hat{V}_{\text{nonl}}^n , \quad (27\text{c})$$

$$t_0 = t , \quad t_1 = t + \Delta t . \quad (27\text{d})$$

The sequence (27b) is built in symmetric manner to minimize the separation error. The quality of the splitting method depends on the size of the commutators between the three pieces of \hat{h}_{KS} . It is the better the smaller the commutators are. The advantage of the method is that the two crucial propagators, kinetic and potential energy, are performed exactly. The local potential energy operator can be easily exponentiated in coordinate space while the kinetic energy operator is exponentiated in Fourier space for which we employ forward and backward FFT as for the evaluation of derivatives (see section 2.4.1). The propagator of the non-local potential could also be resolved exactly, however with considerable book-keeping expense. It is thus handled by Taylor expansion. The T - V -splitting has a particular advantage concerning evaluations of the KS potential. Note that the separation (27b) employs the potential at t_0 in the first potential step and at t_1 in the second. This avoid any bias by giving same weight to both times. The key saving now comes with the fact that the local propagator $e^{-\frac{i}{\hbar} \frac{\Delta t}{2} V_{\text{KS,loc}}(\mathbf{r}, t_1)}$ changes only the complex phase of the wavefunction and does not change the local density. Thus we can evaluate the final new density for t_1 already at the stage before applying this local propagator and no predictor-corrector strategy is needed any more.

2.4.5. Ionic propagation

Time stepping for ionic molecular dynamics (MD) is done with the robust and reliable Verlet algorithm [22, 20], also known as leap-frog algorithm. We present it here summarizing all ionic coordinates by one vector \mathbf{R} and calling the ionic time step Δt , not to be mixed with the electronic time step from the previous section. Propagating the ionic equations of motion (20) raises the same problem with update of the forces as we had experienced in quantum mechanical propagation. The Verlet algorithm solves this by placing coordinates \mathbf{R} and momenta \mathbf{P} different time grid shifted by half a time step $\Delta t/2$. This then proceeds as

$$\mathbf{R}(t + \Delta t) = \mathbf{R}(t) + \Delta t \mathbf{P}(t + \frac{1}{2} \Delta t) , \quad (28\text{a})$$

$$\mathbf{P}(t + \frac{3}{2} \Delta t) = \mathbf{P}(t + \frac{1}{2} \Delta t) + \Delta t \vec{F}(\mathbf{R}(t + \Delta t), t + \Delta t) . \quad (28\text{b})$$

The advantage is that the force is needed only for propagating \mathbf{P} and it is placed exactly at a half \mathbf{P} -step in the shifted-grid setup, readily available at the moment when it is needed. The disadvantage of the method appears at initialization and for computation of observables. In both cases one has to move one of the variables \mathbf{R} or \mathbf{P} by half a time step to have both at the same time. This problem is avoided by slight modification of the Verlet algorithm, called velocity Verlet [20]. Here, both variables are handled at the same time grid. The momentum $\mathbf{P}(t + \frac{1}{2} \Delta t)$ at intermediate time is estimated by a

predictor and the force at intermediate time is taken as average of force at t and $t+\Delta t$. This amounts to

$$\begin{aligned}\mathbf{R}(t+\Delta t) &= \mathbf{R}(t) + \Delta t \mathbf{P}(t+\Delta t) + \frac{1}{2} \Delta t^2 \vec{F}(\mathbf{R}(t), t) \\ \mathbf{P}(t+\Delta t) &= \mathbf{P}(t) + \Delta t \frac{1}{2} \left(\vec{F}(\mathbf{R}(t), t) + \vec{F}(\mathbf{R}(t+\Delta t), t+\Delta t) \right) .\end{aligned}\quad (29a)$$

Note that the force at the new time $\vec{F}(\mathbf{R}(t+\Delta t), t+\Delta t)$ is available for the \mathbf{P} -step because the \mathbf{R} -step has been completed before. Both steps are equivalent what precision, energy conservation, and stability is concerned.⁸

2.4.6. Absorbing boundary conditions

A grid representation naturally leads to reflecting or periodic boundary conditions. Reflection emerges for finite difference schemes. A representation of the kinetic energy by complex Fourier transformation is associated with periodic boundary conditions where flow leaving the box at one side is re-fed at the opposite side. Both can lead to artifacts if a sizable fraction of electronic flow hits the boundaries. There are several ways to solve the problem [23]. The present 3D code employs the simple strategy of applying a mask function during time evolution [24]. This technique is particularly easy to implement and has been widely used in the past. Its robustness and efficiency allow one to develop advanced analyzing techniques on the grid as, e.g., the computation of PES and PAD [25]. We give here a short explanation. A detailed description and discussion of this approach and its proper choice of numerical parameters is found in [26].

Fig. 1⁹ sketches the implementation of absorbing boundary conditions with computation of PES and PAD on a coordinate space grid. Proper handling of electron emission requires absorbing boundary conditions. These are indicated by the ring area in the figure covering here 3 grid points in each direction (actual calculations typically use 6 and more points.) The absorption is performed in each time step as :

$$\varphi(\mathbf{r}, t) \longrightarrow \tilde{\varphi}(\mathbf{r}, t+\delta t) = \hat{\mathcal{U}}_{\text{KS}}(t+\delta t, t) \varphi(\mathbf{r}, t) , \quad (30a)$$

$$\varphi(\mathbf{r}, t+\delta t) = \mathcal{M}(\mathbf{r}) \tilde{\varphi}(\mathbf{r}, t+\delta t) , \quad (30b)$$

$$\mathcal{M}(\mathbf{r}) = \begin{cases} 1 & \text{for } |\mathbf{r}| < R_{\text{in}} , \\ \cos \left(\frac{|\mathbf{r}| - R_{\text{in}}}{R_{\text{out}} - R_{\text{in}}} \frac{\pi}{2} \right)^{\gamma_{\mathcal{M}}} & \text{for } R_{\text{in}} < |\mathbf{r}| < R_{\text{out}} , \\ 0 & \text{for } R_{\text{out}} < |\mathbf{r}| . \end{cases} \quad (30c)$$

First comes one standard KS step ¹⁰ expressed here in terms of the TDLDA propagator $\hat{\mathcal{V}}_{\text{KS}}$, which yields the intermediate wave function $\tilde{\varphi}(\mathbf{r}, t+\delta t)$. This is followed by the action (30b) of the mask function \mathcal{M} defined in Eq.(30c), which removes gradually any amplitude towards the bounds. We use here a spherically symmetric mask. The spherical profile is helpful to minimize gridding artifacts when computing angular distributions [27] (simpler rectangular masks may be used if PAD are not of interest). The absorbing bounds steadily reduce the norm of the wave functions from the inner mask radius R_{in} to

⁸The question is whether we want to keep both ionic steps in the code.

⁹Figure needs to be redrawn for copyright reasons.

¹⁰Crossref to numerical section.

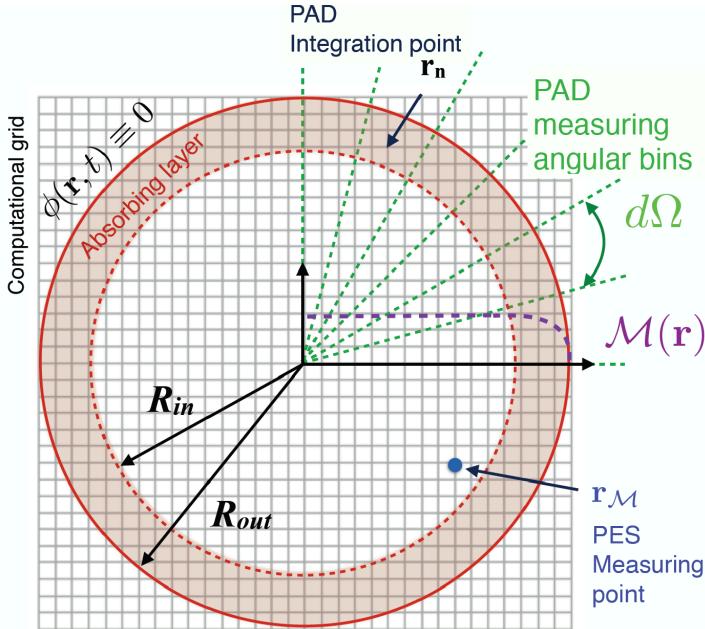


Figure 1: Schematic view of a coordinate-space grid with absorbing bounds (ring zone), a sampling direction for accumulating PAD, and measuring points \mathbf{r}_M for the PES.

the outer one R_{out} . This looks simple and straightforward. However, the mask technique is not perfect. One will always encounter a small amount of reflected flow, particularly for electrons with low kinetic energy. One can minimize the back-flow by proper choice of the exponent γ_M entering the mask profile, see Eq. (30c). This depends, however, on the actual numerics (number of absorbing points, size of time step), for a detailed discussion see [26]. Typical values of γ_M are of order 1/8 or lower.

2.5. Observables

2.5.1. Energies

The total binding energy $E(N_{el}, \mathbf{R}^{(N_{ion})})$, depending in ionic configuration and electron number, is a most prominent observable and it is naturally result of any calculations with energy-density functionals. Comparison with measurements is usually done in terms of differences of energies, e.g., the monomer separation energy as the adiabatic energy difference $E_{mon} = E(N_{el}, \mathbf{R}^{(N_{ion})}) - E(N_{el}-1, \mathbf{R}^{(N_{ion}-1)})$ where the both energies are to be taken from fully relaxed ionic configurations, or the vertical ionization potential (IP) $E_{IP} = E(N_{el}, \mathbf{R}^{(N_{ion})}) - E(N_{el}-1, \mathbf{R}^{(N_{ion})})$ where the new electronic state in the $N_{el}-1$ system has relaxed but the ions are kept in their original configuration.

As a byproduct of mean field calculations, one obtains also the series of s.p. energies ε_α . But it is known that ε_α from (TD)LDA are spoiled by the self-interaction error. This defect can be cured by a SIC, see section 2.2.1, after which the set of ε_α provides a fair map of electron separation energies, particularly of the IP [28, 29]. Experimental data on s.p. energies are mainly the IP and the sequence of peaks in PES from one-photon

processes with weak laser pulses. Note that the latter can also be computed directly from TDLDA and analysis of emitted electrons, see section ??, which is anyway compulsory above the regime of weak pulses. Besides comparison with data, the s.p. energies are very instructive observables for theoretical interpretation as, e.g., analyzing electronic shell structure [11, 8].

Other “theoretical” observables are the separate energy contributions, particularly potential and kinetic energy. The latter can be used to define an “intrinsic kinetic energy” $E_{\text{kin,intr}}$ which characterizes the amount of internal, so to say thermal, kinetic excitation of the electron cloud. This is a quantity which will play a key role in the model for dissipation outlined in section 4.

2.5.2. Densities and shapes

Energy-density functional produce also the electronic local density $\varrho(\mathbf{r})$ as natural outcome. This together with the ionic configuration $\{\mathbf{R}_I\}$ constitutes the shape of a system and they do that in all detail which is in most cases overwhelmingly much information. The gross structure of a shape is well sorted in terms of its multipole moments. Leading quantity is the root-mean-square (r.m.s.) radius which reads for the electrons

$$r_{\text{rms}} = \sqrt{\frac{\int d^3r \varrho(\mathbf{r}) r^2}{N}} . \quad (31\text{a})$$

The further multipoles are best quantified in terms of dimensionless moments

$$\alpha_{lm} = \frac{4\pi}{5} \frac{\int d^3r \varrho(\mathbf{r}) r^l Y_{lm}}{N_{\text{el}} r_{\text{rms}}^l} . \quad (31\text{b})$$

Axially symmetric systems are distinguished by $\alpha_{lm \neq 0} = 0$. The appearance of $\alpha_{lm \neq 0} \neq 0$ can have two causes: first, the system is not aligned along its principal axes, and second, there is a true breaking of axial symmetry. The first action is then to rotate the system such that the z -axis is identical with one principal axis of the cluster. Axial symmetry is truly broken if we then still find some $\alpha_{lm \neq 0} \neq 0$. As far as the quadrupole is concerned, the only remaining case is $\alpha_{22} = \alpha_{2-2} \neq 0$ signaling triaxial shapes. One often regroups the quadrupole deformation parameters into total deformation β_2 and triaxiality γ as

$$\beta_2 = \sqrt{\sum_m \alpha_{2m}^2} , \quad \gamma = \tan \left(\frac{\sqrt{2}\alpha_{22}}{\alpha_{20}} \right) . \quad (32)$$

This convention has been originally introduced to characterize shapes of nuclei [30] and has been taken over for clusters at several places, e.g. [31, 32, 33]. It is to be noted that $\gamma = 0^\circ$ as well as $\gamma = 60^\circ$ represent axially symmetric shapes. The case $\gamma = 0^\circ$ corresponds to prolate shapes and $\gamma = 60^\circ$ to oblate ones.

The same definitions apply to ionic shapes if we replace $\int d^3r \varrho(\mathbf{r}) r^l Y_{lm}(\Omega_r) \rightarrow \sum_I R_I^l Y_{lm}(\Omega_R)$ in Eqs. (31). In case of ions, the quadrupole shape is often alternatively characterized by the moments of inertia

$$I_{ii} = M \sum_I (R^2 - R_i^2) \quad \text{for } i \in \{x, y, z\} , \quad (33)$$

to be evaluated in the frame of principal axes of the cluster ions. The mass M is here the ion mass. The relation between these moments of inertia and the above dimensionless quadrupole moments is

$$r_{\text{rms}} = \sqrt{\frac{I_{xx} + I_{yy} + I_{zz}}{2MN}} , \quad (34\text{a})$$

$$\alpha_{20} = \frac{4\pi}{5} \sqrt{\frac{5}{16\pi} \frac{\langle I_{xx} + I_{yy} - 2I_{zz} \rangle}{NMr^2}} , \quad (34\text{b})$$

$$\alpha_{22} + \alpha_{2-2} = \frac{4\pi}{5} \sqrt{\frac{15}{8\pi} \frac{\langle I_{yy} - I_{xx} \rangle}{NMr^2}} . \quad (34\text{c})$$

Radius and multipole moments are at first glance static observables used to characterize cluster structure. But they are also important ingredients to evaluate response properties as will become apparent in the next two sections.

2.5.3. Polarizability

The static polarizability is a key observable of atoms, molecules, and clusters. Its computation is straightforward. We discuss it here the most important case of dipole polarizability α_D . One applies a static external dipole field $V_{\text{ext}}(\mathbf{r}) = e\mathbf{E}_0 \cdot \mathbf{r}$ and performs static calculations for a couple of \mathbf{E}_0 . This delivers a dipole momentum $\bar{\mathbf{D}} = \langle e\mathbf{r} \rangle$ as a function of \mathbf{E}_0 . The tensor of static dipole polarizability is then

$$(\alpha_D)_{ij} = \left. \frac{\partial \bar{D}_i}{\partial E_{0,j}} \right|_{\mathbf{E}_0=0} . \quad (35)$$

The polarizability matrix simplifies if the system has spatial symmetries. For example, an axially symmetric system aligned with the z -axis has $(\alpha_D)_{xz} = (\alpha_D)_{yz} = (\alpha_D)_{xy} = 0$ and $(\alpha_D)_{xx} = (\alpha_D)_{yy}$ and a spherically symmetric system has additionally $(\alpha_D)_{xx} = (\alpha_D)_{zz}$.

2.5.4. Optical response

Optical response is a key observable in cluster physics. In the limit of long wavelengths, the photon field at the system site is related the dipole momentum $\mathbf{D} = \mathbf{r}$ where \mathbf{r} is taken with respect to the center-of-mass of the electron cloud. With the given code, the dipole excitation strength can be computed by spectral analysis of the dipole response to an instantaneous dipole excitation of the system. To that end, one starts from a well relaxed ground state and applies the instantaneous initial excitation by a small dipole boost

$$\varphi_\alpha(\mathbf{r}, t=0) = e^{i\mathbf{p}_0 \cdot \mathbf{r}} \varphi_{\alpha, \text{g.s.}}(\mathbf{r}) \quad (36\text{a})$$

where \mathbf{p}_0 is a (small) boost momentum. One then propagates electrons with TDLDA and samples a protocol of the dipole momentum

$$\mathbf{D}(t) = \int d\mathbf{r} \mathbf{r} \rho(\mathbf{r}, t) . \quad (36\text{b})$$

After a sufficient time (T_{\max}), one Fourier transforms the dipole signal with an appropriate window function $\mathcal{W}(t)$ and finally obtains the spectral strength $S_{D_i}(\omega)$ in x , y , or

z directions and corresponding spectral power \mathcal{P}_{D_i} as¹¹

$$S_{D_i}(\omega) = \Im\{\tilde{D}_i(\omega)\} , \quad \tilde{D}_i(\omega) = \int dt \mathcal{W}(t) e^{i\omega t} D_i(t) , \quad (36c)$$

$$\mathcal{P}_{D_i}(\omega) = |\tilde{D}_i(\omega)|^2 . \quad (36d)$$

The maximum possible spectral resolution is given by $\delta\omega = 2\pi/T_{\max}$. The window function $\mathcal{W}(t)$ serves to attenuate the dipole signal toward the end point T_{\max} in order to avoid artifacts from non-zero $\mathbf{D}(T_{\max})$ [20]. Useful windows are $\mathcal{W}(t) = \cos^{2n}(t\pi/(2T_{\max}))$ where n is an integer number. The choice $n = 1$ produces often still a bit rough $\Im\{\tilde{D}_i(\omega)\}$ while $n = 2$ performs usually satisfactorily well. It is interesting to note that this treatment in connection with absorbing boundary conditions (see section 2.4.6) allows to compute correctly the escape width of spectral states lying in the electron continuum. For details of spectral analysis and variants thereof see [34].

Spectral analysis can equally well be performed with other observables, as e.g. higher multipoles, spin modes etc. The full TDLDA furthermore allows one to go beyond the linear regime. The more appropriate observable is then the power spectrum.

The same principles of spectral analysis apply, of course, also for computing the vibrational spectra of clusters. Here we concentrate on the MD part of TDLDA-MD. The analyzing times are, of course, to be taken much longer to supply sufficient spectral resolution for the excitations in the meV range, characteristic of ionic vibrational states, and the ionic multipole momenta ought to be used for the spectral analysis of ionic motion [35].

2.5.5. Ionization

Absorbing boundary conditions as explained in section 2.4.6 provide a pertinent picture of electron emission. There are several observables associated with emission from (photo-)excited systems: net ionization, photo-electron angular distribution (PAD), and photo-electron spectra (PES). We discuss them in the next three sections.

The first observable is the total ionization, i.e. the number of escaped electrons N_{esc} . This can be computed simply from the, now decreasing, single-particle norms as :

$$N_{\text{esc}}(t) = \sum_{i=1}^N N_{\text{esc},i}(t) , \quad N_{\text{esc},i}(t) = 1 - \sum_i \langle \varphi_i(t) | \varphi_i(t) \rangle . \quad (37)$$

This shows that we have access to even more than the mere net ionization. Indeed each $1 - N_{\text{esc},i}$ yields the depletion of s.p. state i separately. Both, total ionization and detailed level depletion are very instructive observables [36].

2.5.6. Photo-electron angular distribution (PAD)

The angular distributions $d\sigma/d\Omega(\vartheta, \phi)$ are evaluated in angular segments labeled by the azimuthal angle ϑ and the polar angle ϕ , see Fig. 1. The reference frame for these two angles is usually the z axis identical with the laser polarization axis. We collect

¹¹The dipole strength function, also known as dynamic polarizability, is in principle a tensor as α_D is. Should we address that or better restrict discussion to one coordinate only?

all probability which was removed by the absorption step (30b) and accumulate it. A straightforward collection of grid points in a segment tends to produce noisy results because the number of grid points per segment fluctuates. We therefore associate with each grid point a smoothing function $\mathcal{S}(\mathbf{r})$ which distributes the strength over a vicinity of order of grid spacing. This suffices to produce acceptable smooth distributions. The PAD is thus computed as :

$$\mathcal{A}(\vartheta, \phi) = \sum_{i=1}^N \mathcal{A}^{(i)}(\vartheta, \phi) , \quad (38a)$$

$$\mathcal{A}^{(i)}(\vartheta, \phi) = \sum_{\mathbf{n} \in \text{abs.b.c.}} \int d\mathbf{r} r^2 \mathcal{S}(r\mathbf{e}_r - \mathbf{r}_n) n_{\text{esc},i}(\mathbf{r}_n) , \quad (38b)$$

$$\mathcal{W}(\mathbf{r}) = \frac{\max(\Delta x - |x|, 0)}{\Delta x} \frac{\max(\Delta y - |y|, 0)}{\Delta y} \frac{\max(\Delta z - |z|, 0)}{\Delta z} , \quad (38c)$$

$$n_{\text{esc},i}(\mathbf{r}_n) = \int dt |\varphi_i(\mathbf{r}_n, t)|^2 [1 - \mathcal{M}(\mathbf{r}_n, t)] , \quad (38d)$$

where $\mathbf{e}_r = (\sin \vartheta \cos \phi, \sin \vartheta \sin \phi, \cos \vartheta)$ is the unit vector in the direction of the wanted angles. The smoothing is done by simple tent functions which comply with the integration rule used in the normalization. The angular segments in Fig. 1 try to symbolize this smoothing which collects (weighted) information in the vicinity of a ray. The above recipe applies to state specific PAD \mathcal{A}_i as well as the total PAD $\mathcal{A} \equiv d\sigma/d\Omega(\vartheta, \phi)$.

2.5.7. Photo-emission spectra (PES)

The PES can be deduced from the temporal phase oscillations of the wave functions $\varphi(\mathbf{r}_{\mathcal{M}}^{(v)}, t)$ at measuring points $\mathbf{r}_{\mathcal{M}}$ close to the absorbing bounds [37]. The result for the PES sampled at $\mathbf{r}_{\mathcal{M}}$ is

$$\mathcal{Y}(E_{\text{kin}}, \Omega_{\mathcal{M}}) = \sum_{\alpha} w_{\alpha} \mathcal{Y}_{\alpha}(E_{\text{kin}}, \Omega_{\mathcal{M}}) , \quad (39a)$$

$$\mathcal{Y}_{\alpha}(E_{\text{kin}}, \Omega_{\mathcal{M}}) = \left| \int \frac{dt}{\sqrt{2\pi}} e^{iE_{\text{kin}}t - i\delta q \sqrt{2E_{\text{kin}}} + i\delta\Omega + iE_0 F(t) \mathbf{e}_E \cdot \mathbf{r}_{\mathcal{M}}} \varphi(\mathbf{r}_{\mathcal{M}}, t) \right|^2 , \quad (39b)$$

$$\delta q(t) = E_0 \int_0^t dt' F(t') , \quad (39c)$$

$$\delta\Omega(t) = \frac{E_0^2}{2} \int_0^t dt' F(t')^2 , \quad (39d)$$

where \mathbf{e}_E is the direction of (linear) polarization of the electrical field, $\Omega_{\mathcal{M}}$ the space angle associated with $\mathbf{r}_{\mathcal{M}}$, and $F(t) = \int dt' f(t') \exp(-i\omega_{\text{las}} t')$ the time integrated laser pulse envelope introduced in Eq. (9). This form applies for the wavefunction $\varphi(\mathbf{r}_{\mathcal{M}}, t)$ in space gauge as computed in the code. Note that the formula does not only give the total PES, but also the PES \mathcal{Y}_{α} for emission specifically from state α .

A detailed derivation of Eqs. (39) is found in [38]. We summarize here the ideas behind that compact formula. It is deduced under the assumption that the measuring point $\mathbf{r}_{\mathcal{M}}$ is sufficiently far away from the system (placed around $\mathbf{r} = 0$) such that that outgoing electron wave has direction $\mathbf{e}_k = \mathbf{e}_{\mathcal{M}} = \mathbf{r}_{\mathcal{M}}/r_{\mathcal{M}}$. We expand the wavefunction

at \mathbf{r}_M into outgoing waves with direction \mathbf{e}_k and momenta $k > 0$. These are plane waves $e^{ik\mathbf{e}_k \cdot \mathbf{r}_M}$ for weak fields and for stronger fields the corresponding electron waves in the time-dependent photon field (Volkov states). The fact that we have only outgoing waves in one direction allows to identify uniquely energy and momentum as $\mathbf{k} = \mathbf{e}_k \sqrt{2E_{\text{kin}}}$. The energy is read off from the phase oscillations of $\varphi(\mathbf{r}_M^{(v)}, t)$ and the direction $\mathbf{e}_k \Rightarrow \Omega_M$ from the position of the measuring point.

The formula (39) applies for weak and for strong field, but still fails for extremely strong fields. The limits of validity depend on system and time structure of the pulse. To give an order of magnitude, a intensity for long photon pulses impinging on Na clusters is $I \approx \times 10^{15} \text{W/cm}^2$. For details see [38].

3. The structure of the TDLDA package

3.1. Storage of wavefunctions, densities and potentials

Naturally, wavefunctions and other fields in spatial 3D representation are would be handled as three-dimensional arrays, e.g. `chpcoul3D(1:kxbox, 1:kybox, 1:kzbox)` for the Coulomb potential. However, in order to simplify loops, all 3D fields are addressed as linear arrays, e.g., for the Coulomb field `chpcoul(1:kxbox*kybox*kzbox)` where `kxbox`, `kybox`, `kzbox` are the number of grid points in x , y , and z direction. The parameters defining and handling the grid are:

<code>kxbox</code>	:	total number of grid points in x -direction
<code>kybox</code>	:	total number of grid points in y -direction
<code>kzbox</code>	:	total number of grid points in z -direction
<code>nx2</code>	:	equivalent to <code>kxbox</code>
<code>ny2</code>	:	equivalent to <code>kybox</code>
<code>nz2</code>	:	equivalent to <code>kzbox</code>
<code>nx</code>	:	= <code>nx2/2</code> , number of grid points with $x > 0$, see eq. (??)
<code>ny</code>	:	= <code>ny2/2</code> , number of grid points with $y > 0$, see eq. (??)
<code>nz</code>	:	= <code>nz2/2</code> , number of grid points with $z > 0$, see eq. (??)
<code>kdfull2</code>	:	= <code>kxbox*kybox*kzbox</code> , total number of grid points
<code>nxyz</code>	:	equivalent to <code>kdfull2</code>
<code>ix</code> or <code>i1</code>	:	running index for x values
<code>iy</code> or <code>i2</code>	:	running index for y values
<code>iz</code> or <code>i3</code>	:	running index for z values
<code>ind</code>	:	= <code>ix+nx2*(iy-1)+nx2*ny2*(iz-1)</code> , running index for linear storage

The basic arrays concern densities, potentials, and s.p. wavefunctions. Densities and potentials distinguish spin which is denoted by \uparrow = spin-up and \downarrow = spin-down. Each s.p. wavefunction is associated with one unique spin carried in the array `ispin(1:kstate)`. The fields are arranged the following way:

```

rho(2*kdfull2) = density, linearly stored in two blocks of length kdfull2:  

rho(1:kdfull2) = total density  $\rho(\mathbf{r}) = \rho_\uparrow(\mathbf{r}) + \rho_\downarrow(\mathbf{r})$   

rho(kdfull2+1:2*kdfull2) = difference  $\rho(\mathbf{r}) = \rho_\uparrow(\mathbf{r}) - \rho_\downarrow(\mathbf{r})$ 

```

```

aloc(2*kdfull2) local Kohn-Sham potential, linearly stored in two blocks of length  

kdfull2:

```

```

aloc(1:kdfull2) = local KS potential for spin-up =  $U_{\uparrow}(\mathbf{r})$ 
aloc(kdfull2+1:2*kdfull2) = local KS potential for spin-down =  $U_{\downarrow}(\mathbf{r})$ 

chpcoul(kdfull2) = Coulomb potential, linearly stored

psi(kdfull2,kstate) = set of complex s.p. wavefunctions
1. index for spatial distribution, 2. index counts the states
complex array psi for dynamics, real array psir for static case
each s.p. state has unique spin given in the array ispsin(1:kstate)

```

3.2. The TDLDA calling tree

The TDLDA packages is a rather complex collection of routines. Thus the tree structure of the code is sketched only at the major level of callings and is presented in three separate diagrams: the main routine with all initializations and two calls to the major drivers for static and dynamic calculations in diagram 2, the static driver in diagram 3, and the dynamic driver in 4.

3.3. The TDLDA subroutines in detail

PGR2all: Here is to come a detailed description of the subroutines similar as for RTA in 5.3, but here for the LDA part. Only one example is given as appetizer. The problem is that it cost enormous amounts of space to present all routines and functions at that level of detail. We should shift that to the supplementary material. It will comes there anyway if we decide to transfer all these details to doxygen.

```

SUBROUTINE coul_mfield(rho)
  rho(1:2*kdfull2)           in/out
  density for which Coulomb field is computed

```

Computes Coulomb potential for given density by invoking Poisson solver. Th emerging coulomb potential is communicated as chpcoul via module **params**. In case of dielectric external media, adds pseudo-density for image charge.

*PGR2all: Why is rho also INTENT OUT? Is all 1:2*kdfull2 used or only the first block 1:kdfull2?*

4. Relaxation-time approximation (RTA)

4.1. The formal background of RTA

The quantum Boltzmann equation is the quantum mechanical counterpart of the semi-classical Vlasov-Uehling-Uhlenbeck equation [39, 40]. It complements the self-consistent TDLDA propagation of the one-body density matrix $\hat{\rho}$ by dynamical correlations through a collision term. It reads in general [41, 42] $i\partial_t \hat{\rho} - [\hat{h}, \hat{\rho}] = \hat{I}[\hat{\rho}]$ where the left hand side contains the mean-field propagation. the \hat{I} at the right-hand side consists stands for the quantum-mechanical collision term which, however, is extremely hard to handle for finite Fermion systems. A great simplification can be achieved by the relaxation-time approximation (RTA) which was used successfully in a wide variety of homogeneous systems

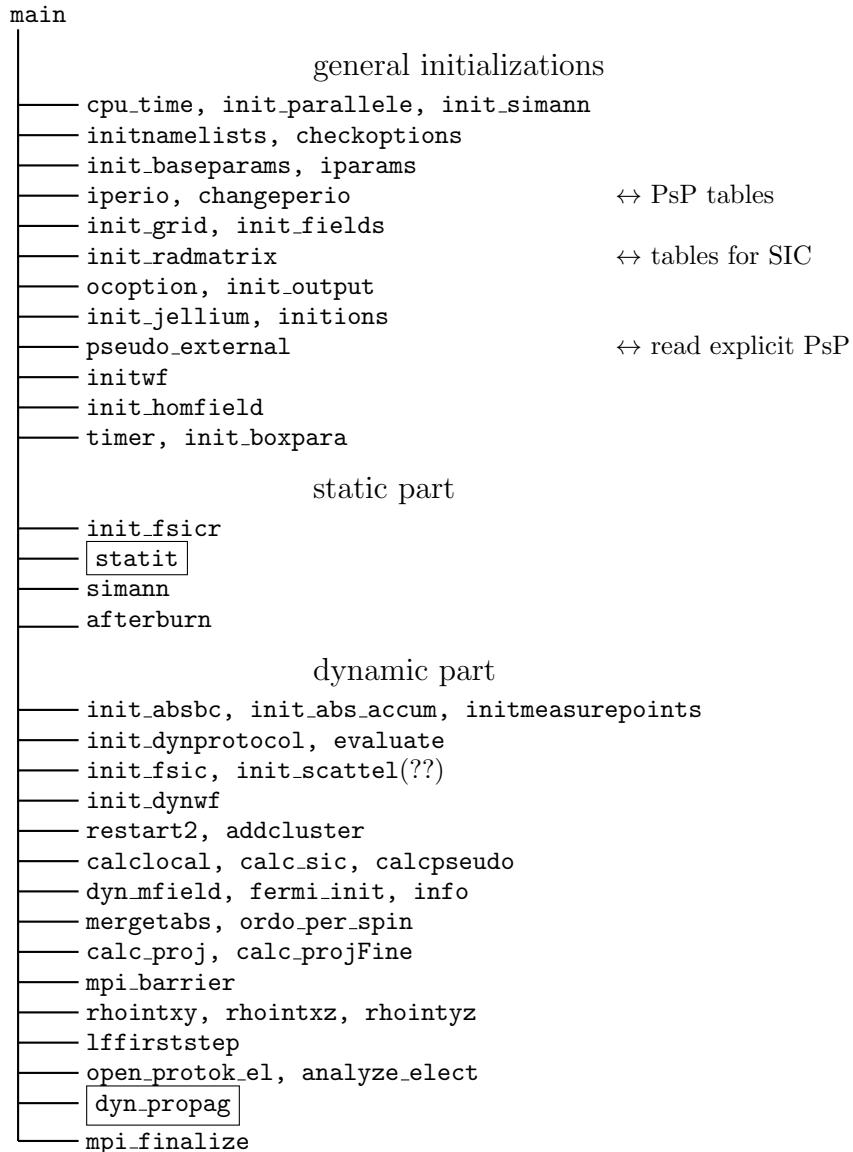


Figure 2: Schematic calling tree for the main routine in `main.F90`. The calling trees for the two major subroutines in framed boxes are explained in subsequent figures 3 and 4.

```

static
├── calcrhor
├── falr, solv_posson
└── calcpseudo, static_mfield
  ├── pricm, infor, prifld
  └── sstep
    ├── cpu_time, system_clock
    ├── exchgr, subtr_sicpot
    ├── nonlocalr
    ├── rfft, rfftback, rkin3D
    ├── project, givens
    └── schmidt, reocc
  ├── prifldz, pri_pstat, printfield
  ├── localizer, mtv_field
  ├── resume, rsave
  └── infor_sic, diag_lagr

```

Figure 3: Schematic calling tree for the static driver routine in `static.F90`.

```

dyn_propag
├── stimer, print_densdiff, savings
  └── rta                                ↔ see section 4
  ├── init_occ_target, init_psitarget
  ├── tstep_exp, CrankNicholson_exp
  └── tstep
    ├── cpu_time, system_clock
    ├── nonlocstep
    ├── kinprop, fftf, fftback
    ├── eval_unitrot
    └── dyn_meanfield, escmask, checkzeroforce
  ├── rhointxy, rhointxz, rhointyz, testcurrent
  ├── itstep, itstepv, enerkin_ions, reset_ions
  ├── calc_pseudo, calclocal, calc_sic
  ├── calc_proj, calc_projFine
  ├── analyze_elect, analyze_ions, energ_ions
  └── mpi_barrier

```

Figure 4: Schematic calling tree for the static driver routine in `dynamic.F90`. The routine in the framed box is explained in great detail in section 4.

[43, 5]. The RTA equations for the present case of finite Fermion systems read [44]

$$\partial_t \hat{\rho} = -i[\hat{h}, \hat{\rho}] - \frac{1}{\tau_{\text{relax}}} (\hat{\rho} - \hat{\rho}_{\text{eq}}[\varrho, \mathbf{j}, E_{\text{sp}}]) , \quad (40a)$$

$$\varrho(\mathbf{r}, t) = \sum_{\alpha} |\phi_{\alpha}(\mathbf{r}, t)|^2 W_{\alpha} , \quad (40b)$$

$$\mathbf{j}(\mathbf{r}, t) = \sum_{\alpha} W_{\alpha} \phi_{\alpha}^*(\mathbf{r}, t) \frac{\vec{\nabla} - \vec{\nabla}}{2i} \phi_{\alpha}(\mathbf{r}) , \quad (40c)$$

$$E_{\text{sp}} = \sum_{\alpha} W_{\alpha} \varepsilon_{\alpha} , \quad (40d)$$

$$\frac{\hbar}{\tau_{\text{relax}}} = 0.40 \frac{\sigma_{ee}}{r_s^2} \frac{E_{\text{intr}}^*}{N} , \quad r_s = \left(\frac{3\bar{\varrho}}{4\pi} \right)^{-1/3} , \quad \sigma_{ee} = \sigma_{ee}(\bar{\varrho}) , \quad (40e)$$

where $\hat{\rho}_{\text{eq}}$ is the density operator of the thermal equilibrium for local density $\varrho(\mathbf{r}, t)$, current distribution $\mathbf{j}(\mathbf{r}, t)$. The reference energy should be, in fact, the total energy $E(t)$ and computed from the actual state $\hat{\rho}(t)$. We replace that by the simpler total s.p. energy $E_{\text{sp}}(t)$ which is legitimate because the difference to E is a functional of $\varrho(\mathbf{r})$ only, a quantity which is kept frozen by construction. The forms (40b,40c) hold for the diagonal representation. A crucial parameter is the relaxation time τ_{relax} which is taken over from semi-classical Fermi liquid theory, for details see [44]. Key entries are: the intrinsic (thermal) energy of the system E_{intr}^* (see appendix AppendixA), the actual number of particles N , the in-medium electron-electron cross section σ_{ee} , the effective Wigner-Seitz radius r_s of the electron cloud, and the average electron density $\bar{\varrho}$. Note that r_s and σ_{ee} depend on an average density $\bar{\varrho}$ because a spatially varying τ_{relax} would be very cumbersome to implement in a quantum mechanical expression. The average density is deduced from the r.m.s. radius r of the actual electron cloud as $\bar{\varrho} = (3N/(4\pi r^3))^{12}$.

This RTA equation (40) is rather involved because its entries depend in various ways on the actual state $\hat{\rho}(t)$. The most expensive piece is the instantaneous equilibrium density operator

$$\hat{\rho}_{\text{eq}}[\varrho, \mathbf{j}, E] = |\phi_{\alpha}^{(\text{eq})}\rangle W_{\alpha}^{(\text{eq})} \langle \phi_{\alpha}^{(\text{eq})}| \quad (41a)$$

which minimizes LDA energy with constraint on the actual $\varrho(\mathbf{r})$, $\mathbf{j}(\mathbf{r})$ and energy E_{sp} . It is determined by the density-constrained mean-field (DCMF) equation

$$\hat{h}_{\text{DCMF}}[\varrho, \mathbf{j}, E] \phi_{\alpha}^{(\text{eq})} = \varepsilon_{\alpha}^{(\text{eq})} \phi_{\alpha}^{(\text{eq})} \quad (41b)$$

$$\begin{aligned} \hat{h}_{\text{DCMF}}[\varrho, \mathbf{j}] &= \hat{h} - \int d^3 r \lambda(\mathbf{r}) \hat{\varrho}(\mathbf{r}) - \int d^3 r \boldsymbol{\lambda}_j(\mathbf{r}) \hat{\mathbf{j}}(\mathbf{r}) \\ &\quad - \mu \int d^3 r (\hat{\varrho}(\mathbf{r}) - \varrho(\mathbf{r}, t))^2 - \mu_j \int d^3 r (\hat{\mathbf{j}}(\mathbf{r}) - \mathbf{j}(\mathbf{r}, t))^2 \end{aligned} \quad (41c)$$

in combination with adjustment of particle number N and s.p. energy $E_{\text{sp}}(t)$ through a

¹²Should we add that as displayed equation?

Fermi distribution

$$W_\alpha^{(\text{eq})} = \frac{1}{1 + \exp((\langle \phi_\alpha^{(\text{eq})} | \hat{h} | \phi_\alpha^{(\text{eq})} \rangle - \mu^{(\text{eq})})/T^{(\text{eq})})} \quad (41\text{d})$$

$$\mu^{(\text{eq})} \leftrightarrow \sum_\alpha W_\alpha^{(\text{eq})} = N(t), \quad T^{(\text{eq})} \leftrightarrow \sum_\alpha W_\alpha^{(\text{eq})} \langle \phi_\alpha^{(\text{eq})} | \hat{h} | \phi_\alpha^{(\text{eq})} \rangle = E_{\text{sp}}(t). \quad (41\text{e})$$

Although cumbersome to evaluate, it is important to use exactly this local, instantaneous equilibrium in the relaxation term. This guarantees that the dissipative step conserves local density, current, and energy as it is mandatory for a good collision term [45].

4.2. Observables specific to relaxation

Most of the observables computed with RTA are exactly the same as for TDLDA, e.g., energy, density, excitation spectra, ionization, PES, or PAD. New are observables related to the mixed character of the one-body operator which is characterized by the occupation numbers W_α . A specific quantity in that respect is the entropy which is computed in diagonal representation (17) by the standard expression [46]

$$S = - \sum_\alpha [W_\alpha \log W_\alpha + (1-W_\alpha) \log(1-W_\alpha)] \quad (42)$$

in units of Boltzmann constant. It serves as a direct indicator of thermalization and allows to read off the typical time scale of relaxation processes.

4.3. Summary of the RTA procedure

The solution of the RTA equations (40) with (41) is rather involved. We briefly summarize the solution scheme for one step from $t \equiv t_0$ to $t + \Delta t \equiv t_1$, for more details see [44]¹³. The TDLDA propagation runs at a much faster pace than relaxation. We resolve it by standard techniques [47, 48] on a time step δt which is much smaller (factor 10–100) than the RTA step Δt . We summarize this TDLDA propagation in the evolution operator \hat{U} from Eq. (15a) and discuss only one RTA step:

1. We first propagate $\hat{\rho}$ by pure TDLDA. The s.p. states in diagonal representation (17) evolve as $|\phi_\alpha(t)\rangle \rightarrow |\phi_\alpha^{(\text{mf})}\rangle = \hat{U}(t_1, t_0)|\phi_\alpha(t)\rangle$, while the occupation weights $W_\alpha(t_1) = W_\alpha(t_0)$ are kept frozen (pure mean-field propagation).
2. Absorbing bounds may have removed parts from the s.p. wavefunctions and so destroyed ortho-normalization. We transform the propagated density operator to a representation in terms of natural orbitals which is diagonal representation (17) with an ortho-normal set of s.p. wavefunctions together with corresponding occupation weights $\{\phi_\alpha^{(\text{nat})}, W_\alpha^{(\text{nat})}\}$. This step can be overridden if reflecting (or periodic) boundaries are used in which case TDLDA preserves ortho-normalization.
3. We compute density $\varrho(\mathbf{r}, t_1)$, current $\mathbf{j}(\mathbf{r}, t_1)$, and total energy E_{mf} associated to the TDLDA-propagated density matrix $\hat{\rho}_{\text{mf}}$.

¹³PGR2all: can we outsource all details to reference [44]?

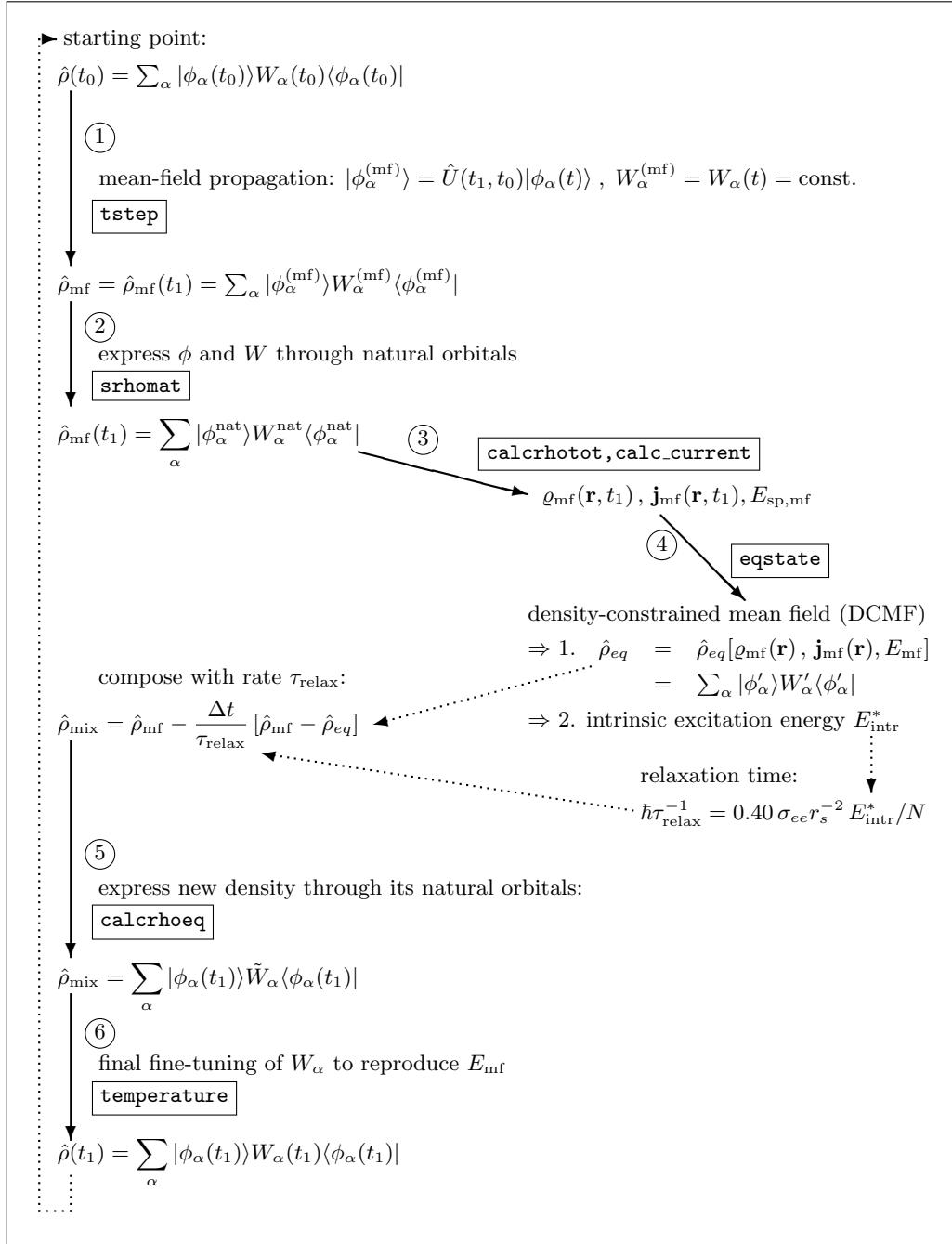


Figure 5: Sketch of the scheme for performing one large time step $t_0 \rightarrow t_1 = t_0 + \Delta t$ in solving the RTA equations. The numbers in open circles indicate the steps as outlined in the text. The names in **typewriter** font refer to subroutines in the code as detailed in section 5.

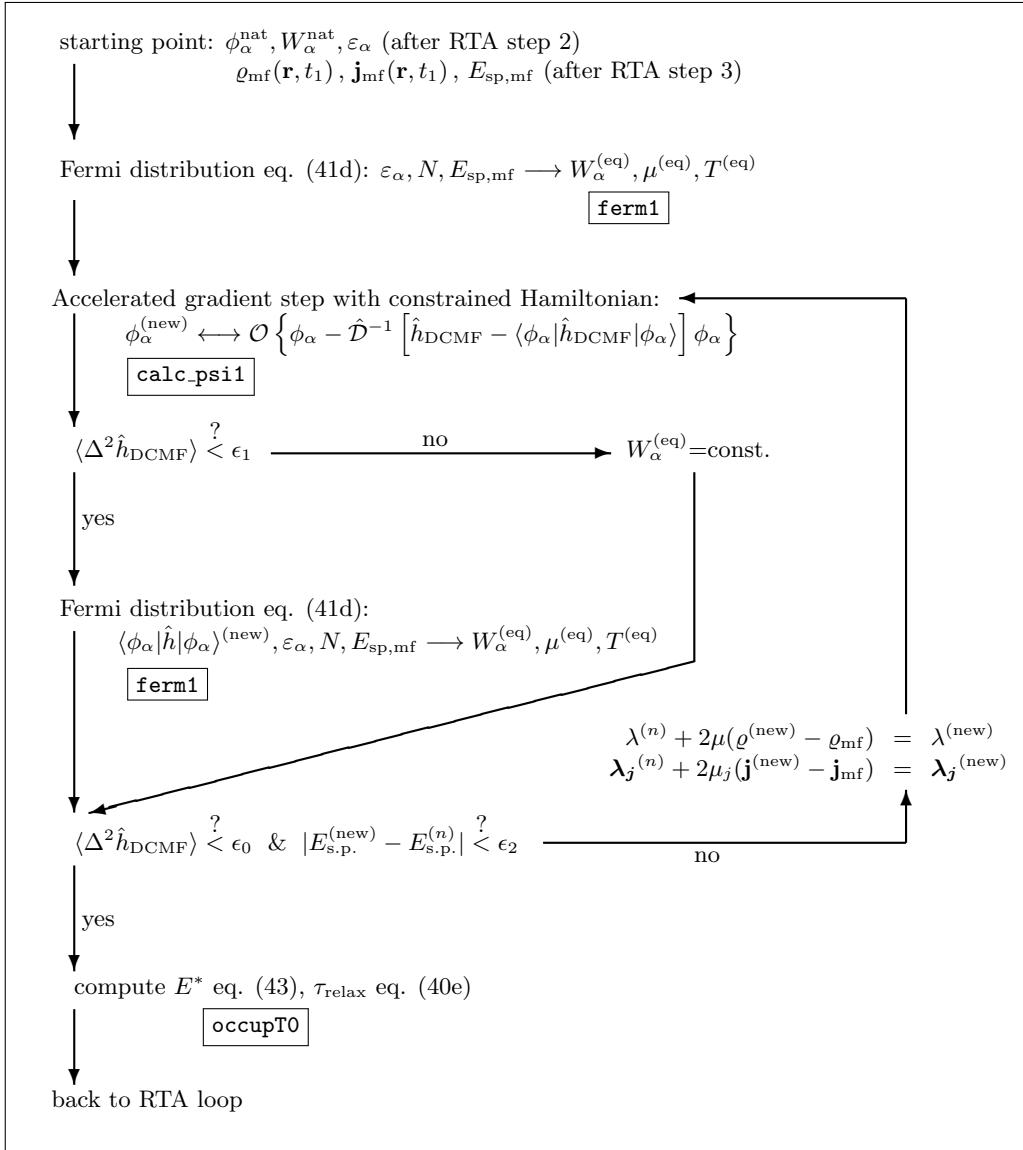


Figure 6: Sketch of the scheme for solving the DCMF Eqs. (41). This scheme expands SUBROUTINE `eqstate` in step 4 from the RTA scheme 5. The symbol \mathcal{O} stands for ortho-normalization of the new set of s.p. wavefunctions and \mathcal{D} for the damping operator in the accelerated gradient step ([PGR2all: Cross references to TDLDA section yet to be defined.](#)).

4. We determine the thermal mean-field equilibrium state $\hat{\rho}_{\text{eq}}$ constrained to the given ϱ , \mathbf{j} , and E_{mf} from step 3. This is achieved by the Density-Constrained Mean Eqs. (41) which is done iteratively as sketched in figure 6. The equilibrium state $\hat{\rho}_{\text{eq}}$ is represented by new s.p. states $\{|\phi'_{\alpha}\rangle\}$ and new occupation numbers W'_{α} in diagonal form (17). Having these, we determine finally the excitation energy as energy relative to the temperature zero state

$$E_{\text{intr}}^* = E_{\text{sp}} - \sum_{\alpha} W_{\alpha}^{(T=0)} \langle \phi_{\alpha}^{(\text{eq})} | \hat{h} | \phi_{\alpha}^{(\text{eq})} \rangle \quad (43)$$

where $W_{\alpha}^{(T=0)}$ are the occupation numbers determined with the s.p. energies $\langle \phi_{\alpha}^{(\text{eq})} | \hat{h} | \phi_{\alpha}^{(\text{eq})} \rangle$ at temperature $T = 0$. This E_{intr}^* thus measures the amount of thermal excitation energy in the system¹⁴

5. We compose the new density operator as mixture of TDLDA propagated state $\hat{\rho}_{\text{mf}}$ and equilibration driving term $\hat{\rho}_{\text{mf}} - \hat{\rho}_{\text{eq}}$ with weight $\Delta t / \tau_{\text{relax}}$ as

$$\hat{\rho}_{\text{mix}} = \hat{\rho}_{\text{mf}} - \frac{\Delta t}{\tau_{\text{relax}}} [\hat{\rho}_{\text{mf}} - \hat{\rho}_{\text{eq}}]$$

where the relaxation time τ_{relax} requires the actual intrinsic excitation energy E_{intr}^* which is also obtained from DCMF. While evaluating the mixing, the new state is expressed in natural-orbital representation Eq. (17). This yields the final s.p. states $\{|\phi_{\alpha}(t_1)\rangle\}$ for this step and preliminary new occupations \tilde{W}_{α} .

6. The mixing in step 5 may have slightly changed the energy such that we remain with a small energy mismatch as compared to the goal E_{mf} . We now apply a small iterative thermalization step to readjust the energy, as outlined in Appendix B. This then yields the final occupation weights $W_{\alpha}(t_1)$ which comply with energy conservation.

The above steps are sketched in Figure 5 whereby the step numbers here correspond to the ones in the Figure. The most involved part is step 4, the solution of the DCMF equations. It is expanded in details in figure 6. A word is in order about the termination criteria in DCMF iteration. The final check for total variance ϵ_0 is an input parameter `rta$sumvar2max` to the RTA part, see section 5.1. The two other switch criteria in figure 6 are hardwired in SUBROUTINE `eqstate` in file `rta.F90` with the values $\epsilon_1 = 10^{-1}$ and $\epsilon_2 = 10^{-4}$
¹⁴PGR2all: Switching DCMF iteration is more involved than that. There is also a parameter `err`. How is that defined?

We should turn ϵ_1 and ϵ_2 to input variables and also for `err`.

A similar problem exists for `ferm1`. The strategy for the choice of the search interval [`T0i`,`T1i`] requires explanation which should come at that place here .

As said above, the time step δt for propagation of TDLDA is very small because it is limited from above by the maximal energy on the grid representation. The stepping Δt for the relaxation term needs only to resolve the changes in the actual mean field which allows much larger values. Typically, we can do 50–200 TDLDA steps before calling one RTA step. For detailed values see the examples delivered with the code.¹⁵

¹⁴Is this definition sufficient? Then we could skip planned appendix AppendixA.

¹⁵PGR2all: A link to be set once we have the place for the benchmarks.

A word is in order about the system for which the present form of RTA can be used. The relaxation time τ_{relax} is allowed to depend on time which allows to accommodate changes of the dynamical state. But τ_{relax} is one global number chosen according to the average electron density $\bar{\rho}$, see eq. (40e). This requires systems which can be characterized by such an average density, i.e., systems having only small density variations in the bulk as it holds typically for metallic bonds. The RTA rate is insensitive to many details of the microscopic collision term as energy- and angle-dependent scattering cross sections [49] or a broad spectrum of relaxation rates. However, these details are usually resolved only (if at all) for fast and energetic processes which are anyway deep in the regime of semi-classical VUU. The grossly averaged treatment of RTA is acceptable for not too fast and not too energetic processes, preferably in compact systems.

5. The structure of the RTA package in `rta.F90`

5.1. Input and output related to RTA

PGR2all: This subsection needs yet to be worked out in detail.

The `NAMELIST dynamic` contains the following variables used in the RTA procedure:¹⁶

`jrtaint`: Modulus for calling the RTA subroutine, i.e., nr. of TDLDA steps per one RTA step. Course time step Δt for RTA and fine time step for TDLDA `dt1` are related as $\Delta t = jrtaint * dt1$.

`rtamu`: Parameter μ in front of the quadratic density constraint in the DCMF Hamiltonian (41c).

`rtamuj`: Parameter μ_j in front of the quadratic current constraint in the DCMF Hamiltonian (41c).

`rtasumvar2max`: Termination criterion ϵ_0 in the RTA step as used in figure 6.

`rtaeps`: Step size δ in the damping operator *PGR2all: (cross ref to be defined)* \mathcal{D} for the RTA step.

`rtae0dmp`: Energy offset E_0 in the damping operator *PGR2all: (cross ref to be defined)* \mathcal{D} for the RTA step.

`rtasigee`: In medium e^- - e^- cross section used for the relaxation time (40e).

`rtars`: Effective Wigner-Seitz radius r_s used for the relaxation time (40e).

`rtatempinit`: The value `rtatempinit`/10 is used as lower value for the search of temperature in SUBROUTINE `ferm1`.

`rtaforceTemperature`: *PGR2all: Seems to be obsolete?*

¹⁶We should define a new `NAMELIST` and shift the RTA variables to there.

Most observables were already defined at TDLDA stage and thus are returned in the standard output files as explained in the TDLDA section¹⁷. New output files specific to observables from RTA are.¹⁸

```
prta:  
peqstat:  
pspeed:  
prhov:
```

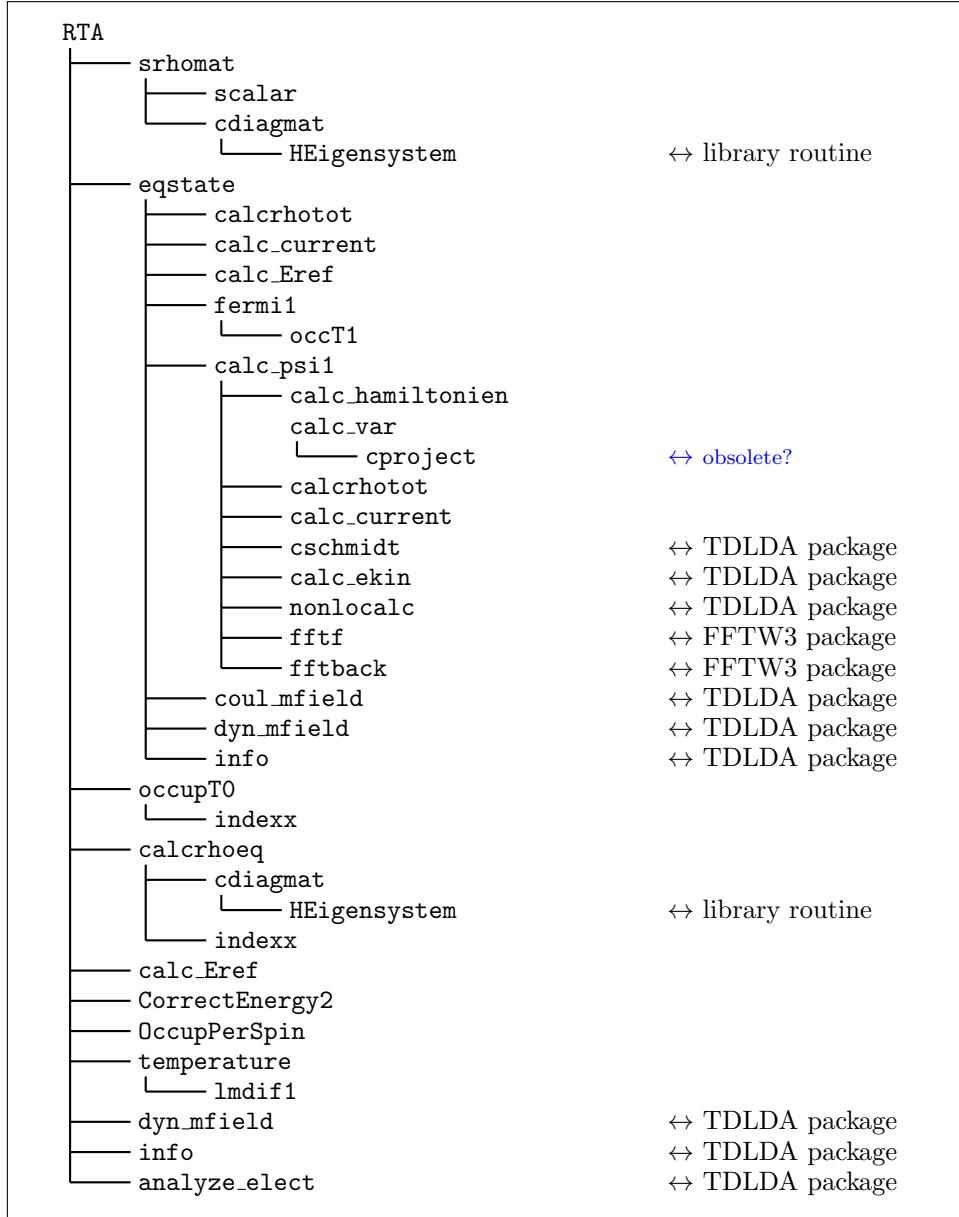
5.2. The calling tree

Here is an oversight over the tree structure of the RTA routines. Those subroutines contained in `rta.F90` are explained in detail in section 5.3. Subroutines coming from the TDLDA package or external sources are marked¹⁹

¹⁷Cross reference yet to be defined.

¹⁸These files seem to carry only protocol for numerics. Where do we print energy balance (E_{intr}^* etc)?

¹⁹The `HEigensystem` seems copied from some library. This could cause copyright problems if we publish the code. Is it from BLAS/LINPACK? Then we could replace the Fortran source by a library call.



5.3. The subroutines in detail

```

SUBROUTINE rta(psi,aloc,rho,iterat)
  iterat          in     external iteration number (TDLDA time step)
  psi(1:kdfull2,1:kstate) in/out set of s.p. wavefunctions
  rho(1:2*kdfull2)   in/out local densities for spin up and down
  aloc(1:2*kdfull2)  in/out local potentials for spin up and down

```

Basic RTA routine performing density constrained mean-field (DCMF) iterations, energy

adjustment, admixing of local equilibrium states by calls to subroutines (see calling tree).

```
SUBROUTINE calcrhoeq(psiorthloc,psieqlc,psiloc,occuporthloc,occuploc,nstateloc)
  nstateloc                                in    number of s.p. states in spin block
  psiorthloc(1:kdfull2,1:nstateloc),       in    set of TDLDA wavefunctions (natural orbitals)
  occuporthloc(1:nstateloc)                in    occupations of TDLDA states
  psieqlc(1:kdfull2,1:nstateloc)          in    set of local-equilibrium wavefunctions
  psiloc(1:kdfull2,1:nstateloc)           out   set of final mixed wavefunctions
  occuploc(1:nstateloc)                  in/out
```

Encapsulated in **SUBROUTINE rta**. Performs the mixing of TDLDA states with local-equilibrium state according to relaxation rate for one spin block. The mixed density matrix is expanded in a representation by both sets of s.p. states.

```
SUBROUTINE calc_Eref(occup,ispin,Ei,Eref)
  occup(1:nstate)    in    occupation number for s.p. states.
  ispin(1:nstate)    in    spin assignment for s.p. states.
  Ei(1:nstate)      in    spin assignment for s.p. states.
  Eref(1:2)          out   sum of s.p. energies per spin.
```

Computes the weighted sum of s.p. energies as reference energy for DCMF. The sum is accumulated for each spin separately.

```
SUBROUTINE fermi1(ekmod,eref,occup,ispinact,T0i,T1i,T2,mu)
  ekmod(1:kstate)     in    given s.p. energies, spin up block first, then spin down
  efer                in    reference energy = wanted sum of s.p. energies
  ispinact            in    spin for which routine is run
  T0i, T1i            in    lower and upper temperature for search
  occup(1:kstate)    in/out occupation numbers, spin block-wise
  T2                  out   final temperature for which Fermi distribution matches efer
  mu                  out   final chemical potential
```

Determines thermal Fermi occupation such that given sum of s.p. energies **eref** and particle number is matched. Is done for each spin separately. Solution scheme is bracketing. Refers to **SUBROUTINE OccT1** while iterating temperature **T2**.

PGR2all: Nr. of spin-up/spin-down states comes through m_params. We should protocol all such entries. First step is to augment each USE by ONLY such that the explicitly communicated variables becomes visible. Important variables may then be listed explicitly.

PGR2all: Routine requires that arrays are sorted in continuous blocks of spin. Do we have an initial check for that? And we need to address that in the general part which explains the layout of arrays.

```

SUBROUTINE OccT1(occrefloc,enerloc,Etotloc,muloc,occtotloc,n,T,occuploc)
    enerloc(1:n)      in  s.p. energies for actual spin
    n                 in  number of s.p. states treated here
    T                 in  temperature
    occrefloc         in  wanted total number of particles
    occuploc(1:n)     out thermal occupation numbers for given T and s.p. energies
    muloc             out chemical potential (Fermi energy)
    occtotloc         out final total number of particles
    Etotloc           out sum of s.p. energies

```

Determines by bracketing chemical potential `muloc` for given array of s.p. energies, temperature `T`, and wanted number of particles `occrefloc` with precision 1D-12. Delivers with it thermal occupation numbers and corresponding total particle number and sum of s.p. energies.

PGR2all: This routine is specific to /tt SUBROUTINE ferm1. Could we encapsulate it by a /tt CONTAINS?

```

SUBROUTINE Calc_psi1(psi1,aloc,rhotot0,rhototloc,curr0,curr1,j,lambda,mu,
lambdaJ,muj,sumvar2,eal,ekmod)
combined with encapsulated SUBROUTINE calc_hamiltonien.
    j                  in  number of DCMF iteration, used here for print
    lambda(1:kdfull12,1:2)   in  Lagrange parameter for density for spin up&down
    lambdaJ(1:kdfull12,1:3)  in  Lagrange parameter for current
    mu, muj            in  driving parameter for augmented Lagrangian
    aloc(1:2*kdfull12)     in  local potentials for spin up and down
    rhotot0(1:kdfull12,1:2) in  initial density PGR2all: not used ??
    curr0(1:kdfull12,1:3)   in  wanted current
    psi1(1:kdfull12,1:kstate) in/out set of s.p. wavefunctions iterated
    rhototloc(1:kdfull12,1:2) out  actual density according to psi1
    curr1(1:kdfull12,1:3)   out  actual current from psi1
    ekmod(1:nstate)        out  final s.p. energies
    eal                 out  final sum of s.p. energies
    sumvar2             out  variance of s.p. energies

```

Performs one damped gradient step of with density & current constrained Hamiltonian.

PGR2all: The density array distinguishes spin up/down while the current array does not. Reason?

PGRcommThe IN & OUT assignments in this subroutine have to be updated.

```

SUBROUTINE eqstate(psi,aloc,rho,psi1,occuporth,iterat)
    iterat              in  actual iteration number (for printing)
    psi(1:kdfull12,1:kstate)  in  initial set of s.p. wavefunctions
    psi1(1:kdfull12,1:kstate) out  final set of s.p. wavefunctions
    aloc(1:2*kdfull12)       in/out local part of potential, spin up/down stacked in blocks
    rho(1:2*kdfull12)        in  initial density, spin up/down stacked in blocks
    occuporth(1:kstate)       in  occupation numbers for psi and still the same for psi1.

```

DCMF iterations by reapeatedly calling `Calc_psi1`, updating Lagrangian parameters for density & current constraints, and occassionally tuning temperature to achieve correct

energy. The latter is done by calling `fermi1`. The local potential is kept constant during DCMF iteration and updated only at the very end.

PGR2all: Fetches nr. of spin up/down from `m_params`.

PGR2all: Lagrange parameters are started from scratch. May it be faster to recycle the previous Lagrange parameters?

PGR2all: Density `rho` is entered via list and still recomputed as `rhotot0`. Unnecessary doubling?

```
SUBROUTINE OccupT0(occloc,esploc,Estar)
    esploc(1:nstate)    in   given s.p. energies
    occloc(1:nstate)   in   given occupation numbers
    Estar              out   excitation energy relative to T=0 distribution
```

Computes thermal excitation energy as difference of actual energy to the energy obtained by Fermi distribution for $T = 0$. The latter distributions is computed for the given s.p. energies which are the same as used for the thermal state.

```
SUBROUTINE calcrhotot(rho,q0)
    q0(1:kdfull2,1:kstate)   in   set of s.p. wavefunctions for which density is accumulated
    rho(kdfull2,2)           out   resulting density
```

Computes local density for set of wavefunctions `q0`. Note that two crucial information is communicated via module `params`, namely `occup`, the array of occupation numbers, `ispin` the array assigning spin top each s.p. state, and `nstate`, the number of s.p. states.

PGR2all: Exploiting the sorting of spin in blocks of s.p. states, we could rewrite the code with to `SUM` statements.

```
SUBROUTINE calc_var(hpsi,psi1,sumvar2)
    psi1(kdfull2,kstate)    in   set of s.p. states for which variance of s.p. energies of calculated
    hpsi(kdfull2,kstate)   in/out array  $H \rightarrow \psi_\alpha$ , on input in  $k$ -space, on output in  $r$ -space
    sumvar2                 out   summed variance of s.p. energies
```

Computes the sum of variances of the s.p. energies, $\langle \hat{h}^2 \rangle_{range}$.

PGRcommThe routine projects from each $hath\psi_\alpha$ all s.p. states ψ_β from the pool of states. That is too much. The s.p. variance should be $\sum_\alpha \langle |\psi_\alpha|(\hat{h} - \varepsilon_\alpha)^2|\psi_\alpha \rangle$ where $\varepsilon_\alpha = \langle |\psi_\alpha|\hat{h}|\psi_\alpha \rangle$.

```
SUBROUTINE forceTemp(amoy,occup,n,temp,mu)
    amoy(1:n)    in   given s.p. energies
    occup(1:n)   in   given thermal occupation
    n            in   number of s.p. states
    temp         in   temperature
    mu           out   emerging chemical potential
```

Determines chemical potential for given s.p. energies and temperature by call to `OccT1`.

PGR2all: Obsolete and never used.

```

SUBROUTINE fermi_init(ekmod,T,occup,ispinact)
    ekmod(1:nstate)      in     given s.p. energies
    T                    in     given temperature
    ispinact            in     actual spin
    occup(1:kstate)     in/out  initial occupation and resulting Fermi distribution for T.

```

Determines Fermi distribution for given s.p. energies and temperature. Searches appropriate chemical potential `mu` by bracketing. Use for repreated calls to FUNCTION `occ`.

PGR2all: This routine `fermi_init` and the related FUNCTION `occ` are never used, thus obsolete. May be removed.

```

SUBROUTINE srhomat(psi,aloc,psiorth,occuporth)
    psi(1:kdfull2,1:kstate)      in     set of s.p. wavefunctions, not orth-normalized
    psiorth(1:kdfull2,1:kstate)   out    ortho-normalized natural orbitals
    aloc(1:2*kdfull2)           in     actual local potential
    occuporth(1:kstate)         out    occupation numbers for ortho-normalized states

```

Computes the density matrix of initial state goiven by set of wavefunctions `psi` together with their occupations `occup`, the latter communicated through module `params`. Then diagonalizes the density matrix and computes on `psiorth` the new wavefunctions associated with diagonal representation of the density matrix.

Finally updates running transformation matrix `psitophi` which is communicated and stored through module `params`.

PGR2all: Usage and propagation of `psitophi` is somewhat hidden because it is handled through a module. Needs to be explained somewhere.

```

SUBROUTINE scalar(tab1,tab2,scal,ispin, mess)
    tab1(1:kdfull2,1:kstate)    in    1. set of s.p. wavefunctions
    tab2(1:kdfull2,1:kstate)    in    2. set of s.p. wavefunctions
    ispin(1:nstate)             in    spin of s.p. states
    mess                         in    message for print inside routine
    scal(nstate,nstate)         out   matrix of wavefunction overlaps

```

```

SUBROUTINE cdiagspin(mat, eigen, vect, N)
    mat(N,N)      in     complex Hermitean matrix to be diagonalized
    N              in     dimension of matrix
    eigen(N)       out    resulting eigenvalues
    Vect(N,N)     out    resulting eigenstates

```

Driver routine for diagonalization of a complex Hermitean matrix of dimension `N` which consists in a two blocks for separate spin. Refers for each single block to routine `cdiag` and subsequent library routines contained therein.

```

SUBROUTINE indexx (n,arrin,indx)
    n                  in     length of array
    arrin(1:n)        in     array to be sorted
    indx(1:n)         out    pointer array

```

Evaluates sorting of an array in ascending order.

```
SUBROUTINE occupPerSpin(mess,Occ)
    mess      in   character variable with comment printed inside routine
    Occ(1:2)  out   total number of particles in each spin
```

Computes number of particles in each spin block. Uses `nstate` and occupations `occup` from module `params`.

```
CorrectEnergy2(Wref,Eref,w,E,Wout,nloc)
    W(1:nloc)   in   initial occupations numbers
    E(1:nloc)   in   given s.p. energies
    Wref        in   reference particle number to be reached
    Eref        in   reference sum of s.p. energies to be reached
    nloc        in   actual number of states
    Wout(nloc)  out   readjusted occupation numbers
```

Final energy correction by one step along Fermi distribution (using Taylor expansion about actual distribution), see appendix AppendixB).

```
SUBROUTINE ordo_per_spin(psi)
    psi(1:kdfull2,1:kstate)  in/out  s.p. wavefunctions before and after reordering
Reorder states in two blocks of spin up and down. Applies that reshuffling to all relevant field of states, s.p. wavefunctions psi, spin per state ispin, and occupations occup.
PGR2all: Routine has been rendered obsolete by new initialization of states which produces immediately the correct sorting. But routine should be kept for possible later use (e.g., mixing states from different sources.
```

```
SUBROUTINE temperature(mu,T)
    mu   out   resulting chemical potential
    T    out   resulting temperature
```

Takes s.p. energies `amoy` and occupations `occup` from module `params` and fits a Fermi distribution to it. Temperature and chemical potentials of the fitted distribution are returned via list. Calls a fitting routine `lmdif1` using subroutine `ff` as argument.

```
SUBROUTINE ff(m,n,X,FVEC,IFLAG)
    X(1:n)      in   array handling chemical potential and temperature
    Fvec(1:m)   out   array of mismatches of distributions
    n           in   number of parameters of model, actually 2
    m           in   number of entries in array
    iflag       in   flaf possibly written (actually not used)
```

Mismatch of `occup` (via modules `params`) from Fermi distribution to given chemical potential and temperature. To be used in fitting routine `lmdif1`.

```
SUBROUTINE cproject(qin,qout,ispact,q0)
    qin(1:kdfull2)          in   s.p. wvaefunction to be projected
    q0(1:kdfull2,1:kstate)  in   set of s.p. wavefunctions which is projected out from qin
    ispact                  in   spin associated with qin
    qout(1:kdfull2)         out  projected s.p. wavefunction
```

Projects away from `qin` all contributions of the set `q0`.

PGR2all: This routine may become obsolete if we recode the the variance in routine calc_var to meet the standard definition.

AppendixA. The intrinsic excitation energy

PGR2all: Intrinsic excitation energy yet to be explained. May also become part of DCMF explanation.

AppendixB. Iterative correction of total energy

- PGR2all: Yet to be imported.*
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