Molecular bonding with the RPAX: from weak dispersion forces to strong correlation

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In a recent article [Phys. Rev. B 90, 125102 (2014)] we showed that the random phase approximation with exchange (RPAX) gives accurate total energies for a diverse set of systems including the high and low density regime of the homogeneous electron gas, the N\textsubscript{2} molecule, and the H\textsubscript{2} molecule at dissociation. In this work, we present results for the van der Waals bonded Ar\textsubscript{2} and Kr\textsubscript{2} dimers and demonstrate that the RPAX gives superior dispersion forces as compared to the RPA. We then show that this improved description is crucial for the bond formation of the Mg\textsubscript{2} molecule. In addition, the RPAX performs better for the Be\textsubscript{2} dissociation curve at large nuclear separation but, similar to the RPA, fails around equilibrium due to the build up of a large repulsion hump. For the strongly correlated LiH molecule at dissociation we have also calculated the RPAX potential and find that the correlation peak at the bond mid-point is overestimated as compared to the RPA and the exact result. The step feature is missing and hence the delocalisation error is comparable to the RPA. This is further illustrated by a smooth energy versus fractional charge curve and a poor description of the LiH dipole moment at dissociation.

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\section{I. INTRODUCTION}

In its formulation due to Kohn and Sham (KS)\textsuperscript{1} Density Functional Theory (DFT)\textsuperscript{2} allows the computation of the ground state density and total energy of a many-particle system by solving a set of single particle Schrödinger-like equations. Within current approximations to the unknown exchange-correlation (xc) energy, this can be done with a computational cost that grows as a power of the number of particles, thus making KS-DFT very attractive from a computational point of view and the ideal tool to study complex systems. The local spin density approximation (LSDA)\textsuperscript{13} and its generalized gradient corrections (GGAs)\textsuperscript{14-16} are the most widely used xc energy functionals. Notwithstanding their capability to successfully predict the properties of a wide class of electronic systems, there are still many situations in which they perform poorly or even fail qualitatively. One such fundamental problem is the description of dispersion interactions. Although one of the weakest, the dispersion interaction is ubiquitous and can add up to give a substantial contribution to the interaction energy in large assemblies of atoms and molecules thus becoming very important in systems like, for instance, biomolecules, adsorbates or nanomaterials. Different solutions have been proposed to deal with weakly interacting systems in DFT, ranging from semi-empirical approaches\textsuperscript{9,13} to truly non-local and parameter-free functionals.\textsuperscript{10,12}

Another class of materials for which standard Density Functional Approximations (DFAs) badly fail is represented by the strongly correlated systems. Strong correlation is usually meant to refer to situations where multiple determinants associated with degeneracy or near degeneracy are needed and a single-particle picture becomes inadequate. This limitation leads, for instance, to a dramatic failure of standard KS methods when studying the dissociation of systems into open shell fragments, whose simplest and paradigmatic example is the stretched hydrogen molecule. However, this is not a failure of DFT itself, but just a break-down of present DFAs. In this respect many different corrective approaches have been developed for \textit{ab initio} calculations on strongly correlated systems ranging from Self Interaction Correction\textsuperscript{13} to Exact-Exchange and hybrid functionals\textsuperscript{14} to DFT+U\textsuperscript{15}

However, rather than introducing \textit{ad hoc} corrections to describe selected problems, it would be highly desirable to find a new class of functionals that perform uniformly better than the present ones as already happened passing from LDA to GGAs and more recently with the development of hybrid functionals. In this respect it is illuminating to start with the formally exact way of constructing the exchange and correlation functional using the Adiabatic Connection Fluctuation-Dissipation (ACFD) theory\textsuperscript{16-17} In this framework an exact expression for the exchange-correlation energy can be derived in terms of the density-density response functions of a continuous set of fictitious systems defining a path that couples the non-interacting KS system with the real many-body interacting one. The formalism thus provides a powerful theoret-
ical framework for a systematic development of advanced functionals but also a practical way to compute accurate correlation energies. Moreover, all ACFD methods treat the exchange energy exactly thus cancelling out the spurious self-interaction error present in the Hartree energy, while the correlation energy is fully non-local and automatically includes van der Waals interactions.

Although much more expensive than the simple approximations to it, a direct evaluation of the exact formula is also possible when the interacting density-density response function is given. A Dyson-like equation linking the latter to its non-interacting counterpart can be derived from time dependent density functional theory (TDDFT), but still requires a suitable approximation for the unknown exchange-correlation kernel, $f_{xc}$. Nevertheless it has been shown that using the Random Phase Approximation (RPA), i.e., by completely neglecting the xc kernel, plus a local-density correction for short range correlation, leads to a qualitatively correct description of van der Waals interactions and systems with mixed bonds.

On the other hand, the RPA total energy is far too negative because of the well known overestimation of the correlation energy; moreover, the binding energy of van der Waals systems is systematically underestimated and an erroneous repulsion hump appears in the dissociation curve of covalently bonded systems at intermediate distances. For an exhaustive review on the RPA see for instance Refs. 32 and citations therein.

Staying within the ACFD framework, a rigorous possibility to address the shortcomings of RPA is to include a xc contribution to the kernel that appears in the Dyson-like equation defining the interacting density-density response function. For a consistent description to first order in the electron-electron interaction, not only the Coulomb kernel, defining RPA, but also an exchange contribution has to be taken into account. The frequency-dependent exact-exchange kernel, $f_{EXX}$, has been derived by Görling from the time-dependent optimized effective potential (TDOEP) method, by Hellgren and von Barth from a variational formulation of many-body perturbation theory (MBPT) and more recently by us from a perturbative approach along the adiabatic-connection path. The corresponding approximation for the density-density response function, named RPAn, is obtained by solving the Dyson equation approximating the xc kernel, $f_{xc}$, with its exact-exchange contribution, $f_{EXX}$, and has been successfully used in the ACFD formula to compute correlation energies of atoms and molecules. In a recent work by us we showed that with a slight redefinition of the re-summation procedure one can overcome a severe deficiency of the RPAx when it is applied to the low density regime of the electron gas and the stretched Ne dimer.

In this work, we investigate the RPAx for molecular dissociation of diatomic molecules with different types of chemical bonds. The binding energies as well as the entire potential energy surfaces are obtained. We study the weakly interacting Kr$_2$ and Ar$_2$ dimers which allows us to test how well the van der Waals forces are described. We also study the Be$_2$ and Mg$_2$ dimers which are challenging due to the complex interplay between van der Waals and covalent bonding forces. Finally, we address the question of how well the RPAn can describe strong correlation by analyzing the LiH molecule.

The paper is organized as follows. Section II briefly summarizes the basic equations of the ACFD theory, the RPAn approximation and the computational details of our calculations. Results for dissociation curves and structural parameters of different kind of molecules are reported and discussed in Sec. III. In Sec. IV, we give our conclusions.

II. THEORY

In this section we recall some basic ideas concerning the ACFD theory as well as some technical aspects of the plane-wave and pseudopotential implementation of the RPA(x) total energy.

A. Exchange and correlation energy within ACFD theory

Within the ACFD framework a formally exact expression for the exchange-correlation energy $E_{xc}$ of an electronic system can be derived:

$$E_{xc} = - \frac{1}{2} \int d\lambda \int dr dr' \frac{e^2}{|r - r'|} \times \left\{ \frac{\hbar}{\pi} \int_{0}^{\infty} \chi_{\lambda}(r, r'; iu) du + \delta(r - r')n(r) \right\}$$

where $\chi_{\lambda}(r, r'; iu)$ is the density-density response function at imaginary frequency, $iu$, of a system whose electrons interact with a scaled Coulomb interaction, $\lambda e^2/|r - r'|$, and move in a local potential chosen in such a way to keep the electronic density fixed to the ground state density of the fully interacting system ($\lambda = 1$). At $\lambda = 1$, the local potential is equal to the external potential (usually the nuclear potential) of the fully interacting system and $H_{\lambda=1}$ coincides with the fully interacting Hamiltonian while at $\lambda = 0$ the local potential coincides with the KS potential and $H_{\lambda=0}$ is the KS Hamiltonian.

The exchange-correlation energy can be further separated into the KS exact-exchange energy, $E_{x}^{\text{EXX}}$, and the correlation energy $E_{c}$. The former has the same expression as the Hartree-Fock exchange energy but is evaluated with the KS orbitals $\phi_i(r)$,

$$E_{x}^{\text{EXX}} = - \frac{e^2}{2} \int dr dr' \frac{\sum_i \phi_i^*(r) \phi_i(r') |r - r'|^2}{|r - r'|}.$$

It is easy to verify that it can be derived from Eq. [1] replacing $\chi_{\lambda}$ with the non-interacting density response
function

\[ \chi_0(r, r'; iu) = \sum_{ij} (f_i - f_j) \phi_i^*(r) \phi_j(r) \phi_i^*(r') \phi_j(r') \frac{\epsilon_i - \epsilon_j + i\hbar u}{\epsilon_i - \epsilon_j + i\hbar u} \]  

(3)

where \( \epsilon_i, \phi_i(r) \) and \( f_i \) are the KS eigenvalues, KS orbitals and occupation numbers, respectively. Subtracting the KS exchange energy from Eq. (1) the correlation energy \( E_c \) is obtained in terms of linear density-density responses:

\[ E_c = -\frac{\hbar}{2\pi} \int_0^1 d\lambda \int_0^\infty du \text{Tr} \left\{ \nu_c [\chi_\lambda(iu) - \chi_0(iu)] \right\} \]  

(4)

where \( \nu_c = e^2/|r - r'| \) is the Coulomb kernel and Tr indicates the integrals over spatial coordinates.

For \( \lambda > 0 \) the interacting density response function \( \chi_\lambda(iu) \) can exactly be related to the KS non-interacting one via a Dyson equation obtained from time-dependent density functional theory (TDDFT):

\[ \chi_\lambda(iu) = \chi_0(iu) + \chi_0(iu)[\nu_c + f^{xc}_{\lambda}(iu)]\chi_\lambda(iu) \]  

(5)

where \( f^{xc}_{\lambda}(iu) \) is the scaled frequency-dependent xc kernel. Spatial coordinate dependence is implicit in the matrix notation. When the xc kernel is specified one can thus determine a corresponding correlation energy via Eq. (4). Alternatively, the interacting density response function can be obtained from many-body perturbation theory which allows truncation at arbitrary order and various re-summations of the higher order terms. For example, by truncating to first order we generate an approximation similar to the MP2 (Moller-Plesset second order perturbation theory) approximation.\(^{18-22}\) A different RPAx re-summation based on a first order expansion of the irreducible polarisability was used in our previous work to eliminate a pathology in the original formulation of RPAx.\(^{18}\)

### B. RPAx energy functional

The RPA and RPAx correlation energies \( E^{\text{RPA}(x)} \) are obtained by solving the Dyson equation setting \( f^{xc}_{\lambda} = 0 \) and \( f^{\lambda}_{xc} = \lambda f_x \), respectively, and inserting the result in the ACFD formula. The RPA(x) total energy is then defined as

\[ E^{\text{RPA}(x)} = E_s + E_{\text{ext}} + E_H + E^{\text{EXX}} + E_c \]  

(6)

where \( E_s, E_{\text{ext}} \) and \( E_H \) are the KS kinetic, external and Hartree energy, respectively.

In this work we have also exploited alternative re-summations based on the RPAx defined as

\[ \chi_\lambda^{\text{RPAx}} = P^{(1)}_\lambda + \chi_\lambda P^{(1)}_\lambda \nu_c \chi_\lambda^{\text{RPAx}} \]  

(7)

with \( P^{(1)}_\lambda = \chi_0 + \lambda \chi_0 f_x \chi_0 \) and

\[ \chi_\lambda^{\text{tRPAx}} = \chi_\lambda^{\text{RPA}} + \lambda \chi_\lambda^{\text{RPA}} f_x \chi_\lambda^{\text{RPA}}. \]  

(8)

Both tRPAx and tRPAx have shown to give results either very similar to the original RPAx or results which qualitatively improve upon the RPAx. In particular, we observed that the break-down of the RPAx approximation in the low density regime of the homogeneous electron gas and in the stretched nitrogen dimer is lifted by using the tRPAx and tRPAx resummations.\(^{18}\) The difference between the RPAx and the alternative resummations for \( \chi \) lies in the treatment of the terms beyond first order in the Coulomb interaction. Crucial to the original RPAx is the so-called exchange kernel which is defined as the functional derivative of the exchange OEP potential. When inserted into the linear response equation it can be shown to exactly reproduce first order particle-hole and self-energy effects as defined within linear response time-dependent Hartree-Fock (TDHF) theory.\(^{22}\) However, higher orders cannot be represented in terms of higher order particle-hole and self-energy diagrams and therefore a direct comparison between the RPAX- and the TDHF response function cannot be made. On the other hand, it is well known that TDHF produces a particle-hole interaction in solids that is too strong.\(^{10}\) This is to some extent compensated by the HF band structure which overestimates the quasi-particle gap. It can therefore be argued that the instabilities of the RPAx response function that are not present in the TDHF response function are due to the fact that to higher orders the exchange kernel is not able to correctly balance the particle-hole attraction with a DFT band structure that has an energy gap much smaller than the HF band structure. The use of tRPAx and tRPAx removes this type of instability by simply neglecting higher order particle-hole terms in the response function. In the cases where RPAx is well-behaved we found that this procedure has only a minor effect on the total energies.

In this work we focus on systems with a rather large gap (or on two-electron systems for which the exact-exchange kernel simply reduces to a half of the Hartree kernel) and we do not expect any critical behavior of the original RPAx. Nevertheless we tested all the alternative resummations for all the systems considered in this work finding additional cases (Be\(_2\) and Mg\(_2\)) which have a relatively small gap around equilibrium where the alternative resummations perform quantitatively better than the original RPAx.

In principle a fully self consistent field (scf) evaluation of the RPA(x) energy can be done by relying on the optimized effective potential (OEP) method which provides a way to compute the local multiplicative potential that minimizes a given orbital dependent energy functional. The first approximate RPA-potential calculations were performed on bulk materials.\(^{37-31}\) Fully scf-RPA calculations have been performed for closed shell atoms\(^{30,52}\) and simple molecules.\(^{33-55}\) For what concerns plane wave implementations, an efficient way to compute the scf-RPA energy and potential has been recently proposed by Nguyen \textit{et al.}\(^{56}\) stemming from the non-scf implementation of Nguyen and de Gironcoli\(^{11}\) paving the
way for extensive calculations also for extended systems. The functional derivative of the RPAx correlation energy was first calculated for spherical atoms [25] and recently for molecules [26]. However, despite the efforts in trying to reduce the computational workload of RPA and RPAx calculations, they still remain computationally very demanding. For this reason, most of the RPA/RPAx calculations are limited to a post self-consistent correction where the xc energy is computed from the charge density obtained from a self-consistent calculation performed with a more traditional xc functional.

The calculations of the RPA and RPAx correlation energies presented in this work are based on an eigenvalue decomposition of the non-interacting response function and of its first order correction in the limit of vanishing electron-electron interaction, respectively. Only a small number of eigenvalues contributes significantly to the decomposition meaning that accurate results for the RPA and RPAx correlation energy can be obtained considering only the most relevant ones. A detailed and general (base-independent) description of the schemes can be found in Ref. [57] and Ref. [18]. In the plane-wave pseudopotential approach any system is treated as periodic and the response functions are block diagonal and can be classified by a vector \( \mathbf{q} \) in the first Brillouin zone. The final expression of the RPAx correlation energy (Eq. (12) in Ref. [18]) in a plane-wave basis set becomes:

\[
E_{c}^{\text{RPAx}} = -\frac{\hbar}{2\pi} \int_{0}^{\infty} du \sum_{\mathbf{q}=0}^{N_{q}} \sum_{\alpha} s_{\alpha}(\mathbf{q};iu) \times \left\{ a_{\alpha}(\mathbf{q};iu) + \ln[1-a_{\alpha}(\mathbf{q};iu)] \right\}
\]

where

\[
s_{\alpha}(\mathbf{q};iu) = \langle \omega_{\alpha}^{0}\rangle \langle \chi_{0}^{0}(iu) \chi_{0}^{0}(iu) \omega_{\alpha}^{0} \rangle
\]

\[
\{a_{\alpha}(\mathbf{q};iu),|\omega_{\alpha}^{0}\rangle\}
\]

are the eigenpairs solution of the generalized eigenvalue problem

\[
-h_{\text{H}}^{0}(iu)|\omega_{\alpha}^{0}\rangle = a_{\alpha}(\mathbf{q};iu)|\omega_{\alpha}^{0}\rangle,
\]

where

\[
h_{\text{H}}^{0}(iu) = \chi_{0}^{0}(iu)[v(\mathbf{q}) + f_{x}(\mathbf{q};iu)]\chi_{0}^{0}(iu).
\]

When the exact exchange kernel is neglected the RPA energy is recovered [25].

The dependence of the RPA correlation energy on the Brillouin-zone sampling and super-cell size has been already carefully analyzed by Harl et al. [25] and Nguyen and de Gironcoli [57] with particular emphasis on how to treat the singularity of the Coulomb kernel in the \( \mathbf{q} \to 0 \) limit. In the special case of finite systems the singularity problem can be easily avoided using a modified coulomb interaction. The long range tail of the coulomb interaction in real space is truncated after a distance of the order of half of the super-cell size making its Fourier transform remaining finite for \( \mathbf{q} \to 0 \) thus leading to a well defined problem even exactly at \( \mathbf{q} = 0 \). The approximation becomes exact in the limit of infinite super-cell meaning that a careful check of the convergence with respect to the super-cell size is needed. In Fig. 1 we report the result for the RPA and RPAx correlation energies of Be dimer at equilibrium distance as a function of the volume of the simulation box calculated within different approximations. Simply setting to zero the \( |\mathbf{q} + \mathbf{G}| = 0 \) component (\( G_{0} = 0 \) in the figure) leads to a slowly convergent behavior for the correlation energies that eventually goes to a limiting value (extrapolated with a linear fit of the \( G_{0} = 0 \) data) represented by the dashed black lines in the figure. For the RPA correlation energy (top panel) we also report the result from a calculation with a
III. BOND DISSOCIATION OF DIMERS

The correct description of chemical bonds of different nature is a fundamental prerequisite for any accurate electronic structure method. In the following we will consider diatomic molecules with different types of chemical bonds ranging from weak van der Waals bonds (Ar$_2$ and Kr$_2$) to mixed covalent-van der Waals bonds (Be$_2$ and Mg$_2$) to an ionic/covalent bond (LiH). We study the full potential energy curves which allows us to assess the general accuracy and applicability of the RPAX functional.

A. van der Waals dimers

Although one of the weakest, the dispersion (or van der Waals) interaction is of primary importance for an accurate understanding of, for instance, biological processes as well as adsorption on surfaces and chemical reactions. Dispersion forces originate from the response of an electron in one region to instantaneous charge-density fluctuations in another one. The leading contribution arises from the dipole–induced-dipole interaction and leads to an attractive energy with the well-known $-1/R^6$ decay with the inter-atomic separation $R$. Standard xc functionals give a contribution only when there is an overlap of density charges of the individual components of the system. Because the overlap decay exponentially with the inter-atomic separation so does the binding energy.

Correlation energy functionals derived within the ACFD formalism are fully non-local and therefore include automatically and seamlessly dispersion interactions (see, for instance, Ref. [60] for a detailed proof). They perfectly combine with the exact-exchange energy, thus removing the ambiguity, present in almost any other approach to van der Waals systems, in the choice of how

shifted $q$ mesh (in particular we set $q = (0,0,0.01)$) and notice that a much faster convergence toward the same limit of the $G_0 = 0$ calculation is obtained. We then tested two different modified Coulomb interactions implemented in the QUANTUM ESPRESSO distribution.$^{55}$ The first one, referred to as “Spherical cut” in the figure, is an abrupt truncation of the coulomb interaction for distances greater than half of the super-cell size. In the second one, “WS cut”, the coulomb interaction is unchanged inside the Wigner-Seitz cell and periodically repeated outside. Both these two approximations converge fast and to the correct limit increasing the simulation box volume. In particular the Wigner-Seitz truncation is the most effective and already for a super-cell lattice size of 20 bohr gives very well converged results for both RPA and RPAX correlation energies.

To be consistent, the same strategy has been applied to the calculation of EXX energy. The integrable divergence appearing in the reciprocal-space expression of the EXX energy in Eq. [2] is a well known problem and can be dealt with, for example, using the scheme proposed by Gygi and Baldereschi$^{59}$ (GB). Using this scheme one gets a convergence behavior (black circle in Fig. [2]) that is proportional to the inverse of the simulation box volume and eventually goes to the infinite volume limit represented by the dashed black lines in the figure. An extrapolation scheme$^{60}$ can be used to mitigate this rather slow convergence with respect to the simulation box volume (“GB+extrapolation” in the figure). Also in this case, both modified coulomb interactions recover the correct limit and in particular the Wigner-Seitz renormalization gives converged results within few tenths of meV already for a super-cell lattice size of $a = 25$ bohr.
to combine exchange, short-range correlation and non-local correlation contributions.

Having binding energies (BEs) of the order of tens of meV, noble-gas dimers represent interesting test cases to investigate the accuracy of RPA and RPAX methods. Most of the RPA calculations for realistic systems in general have been performed in a non self-consistent-field (non-scf) fashion, namely, exact-exchange and RPA correlation energies were computed using single particle orbitals obtained from a local or semi-local self-consistent DFT calculation. Only recently Nguyen et al. have performed for the first time a fully scf calculation for \( \text{Ar}_2 \) and \( \text{Kr}_2 \) at the RPA level revealing a close agreement between the scf-RPA and the RPA@PBE dissociation curves. This indicates that the PBE density is rather close to the scf-RPA density and thus justifies the use of this density in non-scf calculations instead of performing a full scf-RPA one. According to these findings and in absence of a scf-RPAX method, we performed our RPAX correlation energy calculations starting from well converged PBE orbitals.

The dimers and the corresponding isolated atoms have been simulated using a simple-cubic super-cell with a linear size \( a = 25 \) bohr. The electron ion interactions have been described by conventional norm-conserving pseudo-potentials, a kinetic energy cut-off of 80 Ry and 50 Ry for \( \text{Ar} \) and \( \text{Kr} \), respectively, has been used. Finally we used up to 400 low-lying eigenvalues of the RPA/RPAX response function in order to calculate the corresponding correlation energies. Extensive tests have been conducted to ensure that these parameters give well converged binding energy with errors estimated to be less than 1 meV.

In Fig. 3 and Table I we report our results for the RPAX@PBE dissociation curve of \( \text{Ar} \) dimer (green squares) together with those of several DFA and compare them with an accurate model potential fitted on experimental data (black solid line). The RPAX and RPA binding energies have been calculated using well converged PBE orbitals while all the other density functional calculations are fully self-consistent. As expected LDA and PBE (and GGAs in general) give very poor results predicting either too large or too small binding energies and equilibrium distances. Including the exact-exchange kernel leads to an overall improvement of the RPA performance. This results is consistent with the improved van der Waals coefficients obtained in Ref. 39. Compared to the binding energy (12.4 meV), bond-length (3.76 Å) and vibrational frequency (31.2 cm\(^{-1}\)) obtained from a model potential fitted to experimental data, our RPAX results of 11.1 meV, 3.75 Å and 30.8 cm\(^{-1}\) show an impressive agreement. The RPAX dissociation curve turns out to be as good as the newly developed vdW functional which is a simple revision of the VV10 nonlocal density functional by Vydrov and Van Voorhis, specifically designed for van der Waals systems.

Similar results are also observed for Krypton dimer and reported in Fig. 4. Although comparison with dissociation energy curve obtained from an accurate model potential fitted to experimental data shows that RPAX scheme underestimate the binding energy by about 20%, the structural properties at the RPAX level show an improvement if compared to the RPA. Our RPAX results for the bond length and the vibrational frequency differ only by 1% and ~3% from the experimental values, respectively, and compare better than RPA (3%, ~15%) bond lengths and vibrational frequencies. Only the VV10 functional outshines the RPAX and gives result essentially identical to the experiments.

Despite the fact that \( \text{Ar}_2 \) and \( \text{Kr}_2 \) have very similar electronic structure, we notice a better agreement with the experiment for \( \text{Ar}_2 \) than for \( \text{Kr}_2 \). Fig. 5 shows that the discrepancy can be due in part to the different importance of the scf procedure in the two systems. The left panel shows that the dissociation curves of the \( \text{Ar}_2 \) dimer computed at the RPA level using PBE or fully scf orbitals are very close to each other, meaning that the PBE density is already a very good approximation for the scf-RPA density. On the other hand, passing from the PBE density to the full scf-RPA one has a larger effect in the case of \( \text{Kr}_2 \) dimer (right panel of Fig. 5) and a gain of \( \sim 1.3 \) meV can be obtained in the binding energy. We therefore argue a fully self-consistent RPAX calculation may lead to a similar gain in binding energy and ultimately to a

| TABLE I: Equilibrium properties of argon and krypton dimers computed within different functionals: LDA, PBE, rVV10, RPA and RPAX. Equilibrium bond length \( (R_0) \) in Å, binding energy \( (E_b) \) in meV, and vibrational frequency \( (\omega_0) \) in cm\(^{-1}\). Experimental data from Ref. 61 are also reported. |
| --- | --- | --- | --- | --- | --- |
| \( \text{Ar}_2 \) | LDA | PBE | RPA | RPAX | rVV10 | Exp. |
| \( R_0(\text{Å}) \) | 3.39 | 3.99 | 3.84 | 3.75 | 3.75 | 3.76 |
| \( E_b(\text{meV}) \) | 31.0 | 6.1 | 8.3 | 11.1 | 13.4 | 12.4 |
| \( \omega_0(\text{cm}^{-1}) \) | 58.3 | 23.4 | 26.8 | 30.8 | 32.3 | 31.2 |
| \( \text{Kr}_2 \) | \( R_0(\text{Å}) \) | 3.67 | 4.33 | 4.11 | 4.06 | 4.01 |
| \( E_b(\text{meV}) \) | 36.6 | 6.7 | 11.7 | 14.0 | 17.7 | 17.4 |
| \( \omega_0(\text{cm}^{-1}) \) | 39.8 | 15.4 | 20.1 | 22.8 | 23.8 | 23.6 |

FIG. 5: Relative importance of the scf for \( \text{Ar} \) and \( \text{Kr} \) dimers.
very good agreement with the experimental results also for the Kr$_2$ case.

Similar accuracy for the Ar and Kr dimers has been found in Ref. [64] using a range separated hybrid RPAx approach (RSH-RPAx). The RPAx functional of that work is defined in terms of the Hartree-Fock kernel and evaluated with Hartree-Fock orbitals. In general, at full-range, this procedure worsens the results compared to RPAx defined within DFT as done in this work. However, by introducing the range separation results of similar quality can be obtained for van der Waals systems and even provide a better description of the Be dimer which will be further discussed in the next section.

B. Alkaline earth dimers

Alkaline earth dimers represent a much more complex situation than rare-gas dimers. Both long-range van der Waals interaction and static correlation play an important role in these systems. They differ from the rare-gas atoms in having nearly degenerate $s-p$ sub-shells and hence multireference effects that are notoriously difficult to capture by a single Slater determinant approach.

Several theoretical investigations have been devoted to the simplest of these molecule, namely the beryllium dimer, using different $ab$ initio methods. Standard DFT calculations with local or semi-local functionals, predict bond lengths rather close to the experimental value but binding energies about 500% too large (see, e.g., Ref. [21]). Hartree-Fock or EXX does not even bind the molecule. Better results can be obtained resorting to Quantum Monte Carlo (QMC) techniques or high accuracy quantum chemistry methods such as the second-order Möller-Plesset perturbation theory, the couple-cluster approach, and the configuration interaction (CI) method. In addition, the study of dissociation energy curve and structure of Be dimer using RPA technique has also been carried out. Although nscf-RPA can predict $R_0$ and $\omega_0$ in rather good agreement with experiment, the binding energy is severely underestimated. Moreover in Ref. [24] the presence of an unphysical maximum in the nscf-RPA dissociation curve was found and the whole curve was shown to be very sensitive to the input orbitals used. The authors suggested a fully self-consistent treatment of the RPA density and potential as a possible solution for the hump puzzle. Recently Nguyen et al. have performed such a calculation for this system and pointed out that the scf treatment is indeed important and significantly lower the total energy of the system, yet it is not enough to fix the unphysical maximum problem and even leads to the metastability of the Be dimer. These findings, as already pointed out in the original work, indicate a real limitation of the RPA that calls for the inclusion of correlation contributions beyond the simple Hartree kernel.

In Fig. 6 and Table I we report our results for the binding energy curve of Be$_2$ from standard PBE, rVV10, RPA and RPAx calculations and compare them with an accurate model potential-energy-surface fitted on experimental data. The results from the simple modifications (tRPAx and t′RPAx) of the original RPAx proposed to fix its pathological behavior in the limit of small gaps are also shown. The dimer and the corresponding isolated atom have been simulated using a simple-cubic super cell with a linear size $a = 25$ bohr and a kinetic energy cut-off of 40 Ry. Up to 200 lowest-lying eigenpairs of the generalized-eigenvalue problem in Eq. (10) have been used to compute the RPA and RPAx correlation energies. All the RPA and RPAx calculations have been performed starting from well converged PBE orbitals.

The PBE calculation predicts a bond length rather close to the experimental value but too large a binding energy (400% overestimate). The rVV10 functional that
The RPA severely underestimates the binding energy but still gives equilibrium bond length and vibrational frequency in good agreement with experimental data. Passing from RPA to RPAX leads to a worse description of the dissociation curve near the equilibrium position and even to the metastability of the Be dimer. However we notice that in the dissociation region the RPAX curve approaches the experimental potential energy surface faster than the RPA one in agreement with the largely improved description of the van der Waals forces within RPAX. The results obtained from the alternative re-summations tRPAX and t’RPAX defined in Ref. 18 recover the RPA performance near the minimum, reduce the height of the hump for intermediate values of the bond length and approach the correct asymptotic behavior of the Be-Be interaction potential much faster than all the other approximations. The hump problem has been previously analyzed in the literature, but is not yet fully understood. A spurious maximum in the exchange energy as a function of atomic separation appears when the EXX energy (Eq. (2)) is evaluated with orbitals coming from a local potential which partly explains the hump in the total energy. The use of self-consistent EXX orbitals reduces the total energy hump but does not completely remove it while the use of HF or hybrid orbitals generated by a non-local potential is sufficient to remove the hump for both RPA and RPAX dissociation curves but produces a very shallow minimum at large interatomic distance. No energy hump and an underestimated binding energy is obtained for Be2 in the recently developed r2PT approach that otherwise poorly describes the dissociation curve of H2, that is instead satisfactorily described by RPAX.

The same basic perturbative ingredients, consistently treated in RPAX via Görling-Levy perturbation theory, are treated in r2PT starting from a Rayleigh-Schrödinger perturbation theory perspective, leading to the need to add single-excitation correction terms to account for the inconsistency between the DFT reference and the HF-like perturbative approach used. Higher order contributions from selected classes of energy diagrams amenable to resummation are also included. RPAX and r2PT are therefore expected to contain very similar physical effects and the difference in performance of the two methods springs from the different approximate treatment of the high order contributions to the correlation energy.

As a second example of alkaline-earth dimer we studied the Mg2 molecule. The dimer and the corresponding isolated atom have been simulated using a simple-cubic super cell with a linear size a = 25 bohr and a kinetic energy cut-off of 35 Ry. Up to 180 lowest-lying eigenpairs of the generalized-eigenvalue problem in Eq. (10) have been used to compute the RPA and RPAX correlation energies. All the RPA and RPAX calculations have been performed starting from well converged PBE orbitals.

Our results for the PBE, rVV10, RPA and RPAX dissociation curves are shown in Fig. 7 and Table II together with a model potential fitted on spectroscopic data. As already seen in the Be2 molecule, also in this case the PBE functional greatly overbinds the system still giving a reasonable value for the equilibrium distance. The rVV10 functional slightly modifies the PBE curve for large inter-atomic distances with no significant improvement in the equilibrium region. The RPA curve still exhibits a hump at intermediate distances, although much less pronounced than in the case of Be2. As a result an almost flat curve is obtained for intermediate and large inter-atomic separations with a very shallow global minimum appearing at distances much greater than the experimental equilibrium one. Including the exchange contribution to the kernel improves the van der Waals forces which leads to a qualitative improvement in the description of the bond. The bump is no longer visible and a proper behavior for the dissociation curve is obtained with one minimum located in between the PBE and the RPA one. The larger improvement within RPAX for the Mg2 bond compared to the Be2 bond suggest that this bond has a larger van der Waals character. Similarly to the Be molecule the alternative re-summations produce a better agreement with experiment.

C. The LiH molecule

The stretched H2 molecule with its single covalent bond represents the simplest example of a strongly correlated system. Standard DFAs such as the LDA or GGA fail but also more sophisticated approaches such as the MP2 and the GW approximation. A good total energy description is usually recovered resorting to an unrestricted calculation, however at the price of having the wrong spin symmetry. On the contrary, both RPA and

|                            | Be2               | Mg2               |
|-----------------------------|-------------------|-------------------|
|                            | PBE RPA RPAX tRPAX rVV10 Exp. | PBE RPA RPAX tRPAX rVV10 Exp. |
| \( R_0 (\text{\AA}) \)     | 2.46 2.40 2.49 2.46 2.44 2.45 | 3.51 – 4.31 4.30 3.56 3.89 |
| \( E_b (\text{meV}) \)     | 399.4 25.1 -13.1 20.0 387.7 115.3 | 137.1 – 22.3 30.9 131.1 52.57 |
| \( \omega_0 (\text{cm}^{-1}) \) | 336 293 221 231 326 276 | 101.5 – 27.0 31.2 90.2 51.1 |

Experimental data from Ref. 70,71 are also reported.
RPAX are able to capture almost the full dissociation curve, as first shown in Ref. [29] More recently, it was shown that also the strong triple bond of the N₂ dimer can be accurately captured by the tRPAX [25] for which only CCSD(T) has been applied successfully so far [27]. The common difficulty of H₂ and N₂ is the fact that they are composed of atoms with open shells, which results in a nearly degenerate situation as the bond is stretched. A similar situation occurs in the case of the LiH molecule which is described by a covalent/ionic bond. In addition, LiH is hetero-nuclear and as such the problem of charge delocalisation, as predicted by most functionals, is largely enhanced. As the LiH bond breaks the electrons should, in an exact treatment, localize and form two neutral atoms. However, standard approximations allow the electrons to delocalise and form fractionally charged dissociation fragments. This failure of common DFAs was first discussed in Ref. [78] where it was linked to a missing derivative discontinuity of the xc functional and an associated step in the corresponding xc potential.

In order to estimate this type of delocalisation error one can determine the amount of fractional charge a given functional predicts by means of a fractional charge analysis [59]. The functional is extended to allow for densities that integrate to a non-integer particle number and the energy of the independent atoms are calculated as a function of particle number. The total energy of the combined system (i.e. the molecule at infinite separation) can then be determined as a function of the amount of charge on e.g. the H atom and the minimum can be located. We carried out such a fractional charge analysis for LiH using the PBE, RPA and RPAX functionals. The results can be found in Fig. 8. Notice that the pseudo potential approximation has been applied to the Li-atom (middle panel of Fig. 8) so we cannot compare the total energy to the exact result. This can, however, be done in the case of the H-atom, see left panel of Fig. 8. As expected, at integer particle number RPA and RPAX outperform the PBE. However, we also see that as soon as we deviate from an integer both RPA and RPAX are smooth with a rather large curvature. As already found within the RPA [22] also within the RPAX the derivative discontinuity is missing meaning that the inclusion of the exchange kernel does not improve the qualitative behavior of the RPA for fractional particle numbers. In the third panel of Fig. 8 we see that the RPAX minimizes at 1.2 electrons on the H-atom which is a result comparable to those of the RPA and PBE. This result is perhaps not surprising since for two electrons the EXX kernel is Hartree-like and thus does not introduce any new correlation effects. If the RPAX calculation for the LiH molecule was performed self-consistently we would obtain 1.2 electrons on the H-atom in the dissociation limit. To be consistent when calculating the binding energy we should therefore subtract the value of the energy at this minimum and not the energy of the two atoms with integer particle numbers. In Fig. 9 we have plotted the binding energy curve of LiH with the energy in RPA and RPAX evaluated using PBE orbitals. Since the PBE functional minimizes at a value close to RPA(RPAX) we expect the result to be close to the self-consistent result. We see that, although RPAX gives a better total energy (not shown), the binding energy is slightly worse than RPA, still giving an improvement compared to PBE. The use of tRPAX does not improve the results much. The fact that RPA and RPAX perform worse for LiH as compared to the H₂ is a result of the additional delocalisation error. We also notice a very slow convergence to the dissociation limit. The maximum of the hump has not been reached at 5 Å. This slow convergence can also be attributed to the delocalisation error as we will now further discuss.

Recently, fully self-consistent RPA was implemented within QUANTUM ESPRESSO [29]. However, in order to compare with exact results we decided to simplify the problem and model the LiH molecule as a one-dimensional system with a softened Coulomb interaction. For this problem we could use an implementation
FIG. 10: (Left) The KS potentials of a model LiH at $R = 3.5\ \text{Å}$. $x = 0$ corresponds to the bond midpoint. Inset shows the corresponding potential energy curves. (Right) The dipole moment as a function of nuclear separation of the exact density and the self-consistent densities within EXX, RPA and RPAx.

based on cubic splines as basis functions which is known to solve the optimized effective potential equation with high precision.\cite{20030880,20030880} To obtain the exact potential we used OCTOPUS.\cite{20030880} In Fig. 10 (left) we plot the full KS potential at a bond distance of $R = 3.5\ \text{Å}$ in RPA, RPAx and EXX and compare the results to the exact potential. The bond midpoint is located at $x = 0$ and the corresponding potential energy curves are shown in the inset. The RPA is in rather good agreement with the exact potential. It exhibits the correlation peak at the bond midpoint but it misses the step feature. The RPAx is similar to RPA but with an overestimated bond midpoint peak. Again the step feature is missing which is consistent with the missing derivative discontinuity in the xc energy. The large humps of the EXX potential is a result of the large delocalisation error in EXX which has been estimated to 1.4 electrons on the H-atom in previous works.\cite{20030880,20030880}

In order to understand the huge impact of the missing step or derivative discontinuity on the density we have also calculated the dipole moment of the LiH molecule, see Fig. 10 (right). In the exact treatment the dipole moment should tend to zero in the dissociation limit since the molecule dissociates into neutral atoms. Due to the fact that approximate functionals produce fractional charges on the dissociation fragments the dipole moment will instead tend to infinity.\cite{20030880} Indeed, this is exactly what we see for the approximations studied and the bond breaking transition is missing in the self-consistent density of the EXX, RPA and RPAx.

IV. CONCLUSIONS

We presented an assessment of RPA and RPAx functionals on a set of diatomic molecules having chemical bonds of different nature. For weakly interacting systems we found that the inclusion of the exact-exchange kernel leads to a quantitative improvement with respect to the RPA results. The binding energy and the structural properties of Ar$_2$ and Kr$_2$ molecules show a significant improvement when moving from the RPA to the RPAx approximation due to an improved description of the van der Waals forces. For the binding energy curve of Be$_2$ even if a better description of the dissociation region and a reduction of the hump at intermediate inter-atomic distances can be achieved within the tRPAx approximation, the binding energy is still far from the experimental results and a potential energy surface with two minima is obtained. On the other hand, the dissociation curve of the Mg$_2$ molecule shows a much larger improvement passing from the RPA to the RPAx approximation due to the increased van der Waals character of the bond. The double minimum structure found at the RPA level is correctly removed by including the exact-exchange contribution to the kernel and a well behaved potential energy surface is recovered. A comparison between results for purely van der Waals and mixed covalent-van der Waals molecules reveals the difficulty of RPA and RPAx functionals in describing situations where the energetic and structural properties of the system result from a delicate balance between bonding forces of different nature. Nevertheless also in these difficult cases the long range part of the dissociation curves display a sensible improvement passing from the RPA to the RPAx. Finally, we addressed the problem of strong correlation by analysing the dissociation of the LiH molecule. The RPA gives a qualitative improvement of the total energy as compared to standard density functionals. The inclusion of the exchange kernel within RPAx further corrects the correlation energy and gives results in good agreement.
with experiment. The hump at intermediate bond distances remains, however, and is shown to be a symptom of a large error in the corresponding RPA/RPAX densities. This error originates from a poor description of fractional charges and a missing derivative discontinuity within RPA and RPAX.

Although in the present work we focused on molecular systems, for which exact reference data are available for comparison, the current approach can easily be extended to solids and surfaces. The additional computational workload required by the Brillouin zone integration (see eq. 9) should be largely compensated by the reduced number of plane-waves associated to a much smaller unit cell. We therefore believe the application of the method to extended systems to be as computational demanding as the calculations presented here. Work toward this goal is underway.

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