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Generalized Many-Body Dispersion Correction through Random-Phase Approximation for Chemically Accurate Density Functional Theory

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at a negligible cost, the DNN-MBDQ approach can be coupled with DFT functionals, such as PBE, PBE0, and B86bPBE (dispersionless). The DNN-MBQ-corrected functionals reach chemical accuracy while exhibiting lower errors compared to their dipole-only counterparts.

he importance of modeling matter through computer simulations has risen tremendously in the past decades in virtue of both the theoretical achievements and the advent of mass-produced computers whose performances have increased exponentially. Despite these tremendous achievements, the exact solution of the nonrelativistic electronic Schrödinger equation remains out of reach for multielectron systems and different approximations have been introduced in order to model and tackle systems of chemical relevance. In particular, Kohn-Sham Density Functional Theory (KS-DFT) has established itself as the most widely used electronic structure method being the cheapest way for introducing electronic correlation. KS-DFT is based on the idea of evaluating the kinetic energy from a single Slater determinant, thus assuming the electrons being noninteracting. The difference between the true and KS kinetic energies as well as the difference between the true electronic and exchange interaction ones is essentially embedded and modeled by the exchange correlation functional, key quantity in KS-DFT, however, in practice unknown.

The different approaches taken in modeling the exchangecorrelation functional define the plethora of KS-DFT variants which, however, are mainly capable of capturing local correlation effects. Dispersion interactions, on the other hand, are rooted in the long-range electron correlation that can clearly not be captured by the intrinsic locality of common exchange correlation functionals. This inadequacy of KS-DFT in modeling noncovalent dispersion interactions, ubiquitous in nature and materials, has risen the attention toward the development of dispersion correction models,¹ the most widely used being based on empirical pairwise terms, eq $1.^2$

$$\mathcal{E}_{\rm disp} = \sum_{i>j}^{N} - \frac{C_{6}^{ij}}{R_{ij}^{6}} \tag{1}$$

This pairwise approach coupled with KS-DFT has shown to provide very good accuracy despite its very simple nature that adds basically negligible computational time and, for this reason, it is also employed in the majority of force fields as attractive component of the Lennard-Jones potential.

Each of the above pairwise terms can be further expanded via a second-order perturbative approach³ (in the limit of large interatomic separation) to include the higher-order contributions shown in eq 2 where $\alpha_k^j(i\nu)$ is the *k*-pole polarizability at imaginary frequency of atom *j* (only the single pair *ij* is considered for the sake of simplicity).⁴

Received:December 7, 2022Accepted:January 26, 2023Published:February 7, 2023





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$$\mathcal{E}_{\text{disp}} = -\frac{C_6^{ij}}{R_{ij}^6} - \frac{C_8^{ij}}{R_{ij}^8} - \frac{C_{10}^{ij}}{R_{ij}^{10}} - \dots$$

$$C_{2n}^{ij} = \frac{(2n-2)!}{2\pi} \sum_{h=1}^{n-2} \frac{1}{(2h)!(2k)!} \int_0^\infty d\nu \alpha_h^i(i\nu) \alpha_k^j(i\nu),$$

$$k = n - h - 1$$
(2)

While providing the correct long-range asymptotic limit, C_6 terms alone are not enough to describe the short- and mediumrange dispersion effects and these higher-order terms have proven to improve significantly interaction energies near equilibrium regions as well as condensed phase properties obtained with molecular dynamics simulations based on force field potentials.⁵

Regardless of the higher-order terms inclusion discussed, this pairwise approach completely neglects the collective manybody dispersion (MBD) effects inherited from the intrinsic quantum mechanical nature of long-range electron correlation and their relevance has been proven in modeling extended systems, supramolecular complexes and proteins in solutions.^{6–9}

These nonadditive dispersion effects have been modeled via a set of coupled fluctuating dipoles^{10,11} as well as quantum Drude oscillators.¹² In recent years, Tkatchenko, DiStasio, Jr., and Ambrosetti have proposed the MBD@rsSCS introducing the range-separation of the self-consistent screening (rsSCS) of atomic polarizabilities based on Tkatchenko-Scheffler volume rescaling that relies on Hirshfeld's molecular electron density partitioning.¹³ This many-body dispersion approach is particularly elegant as it fully relies on ab initio-derived parameters (atomic polarizabilities) except for the unique range separation parameter governing the coupling to the chosen KS-DFT method, while providing chemical accuracy.¹³ A fractionally ionic variant of the MBD@rsSCS model (MBD@rsSCS/FI) has also been proposed by Gould et al. where iterative Hirshfeld (HI) density partitioning together with a charge-dependent atomic polarizability approach are embraced,¹⁴ making the partitioning more suitable than the original Hirshfeld scheme in the treatment of ionic compounds.¹⁵

Newer developments in the MBD modeling include the nonlocal many-body dispersion method (MBD-NL) of Hermann and Tkatchenko¹⁶ where the MBD approach is combined with the Vydrov and Van Voorhis (VV). A polarizability functional,¹⁷ as well as our recently proposed DNN-MBD model where atomic polarizabilities are obtained via deep neuronal network (DNN) model, thus bypassing the explicit electron density partitioning.¹⁸

Being based on the interaction among coupled fluctuating dipoles, the MBD model (rsSCS is implicitly assumed and from now dropped) and its variations represent the many-body counterpart of eq 1, where the effect of higher-order fluctuating multipoles are neglected, although Massa et al. have recently proposed a beyond-dipole MBD model based on the Random Phase Approximation (RPA) formalism.¹⁹

In the same line, this work independently generalizes the MBD model to coupled fluctuating dipole and quadrupoles in order to improve the description of short and midrange manybody dispersion. Particularly, we will show how atomic quadrupole polarizabilities can be recursively derived from dipole ones, thus without the need on introducing further parameters. Furthermore, in doing so we will adopt our recently proposed DNN model approach that completely bypasses the explicit electron density partitioning. The outcoming density-free DNN-MBDQ model exhibits improved accuracy especially near equilibrium regions without the inclusion of any additional parameter, compared to other MBD-based models.

The correlation energy \mathcal{E}_c for a system of interacting electrons can be rigorously expressed with the adiabatic connection fluctuation-dissipation formula in eq 3, v being the Coulomb potential $|\mathbf{r} - \mathbf{r}'|^{-1}$ whose coupling strength is governed by λ while χ_{λ} and χ_0 are the interacting and noninteracting response functions where the latter can be evaluated from a set of single particle orbitals $\phi_i(\mathbf{r})$ with their corresponding energies ϵ_i and occupation numbers f_i via the Adler-Wiser formalism, eq 4.^{20,21}

$$\mathcal{E}_{c} = -\frac{1}{2\pi} \int_{0}^{\infty} d\nu \int_{0}^{1} d\lambda \operatorname{Tr}[\chi_{\lambda}(\mathbf{r}, \mathbf{r}', i\nu)\nu(\mathbf{r}, \mathbf{r}') - \chi_{0}(\mathbf{r}, \mathbf{r}', i\nu)\nu(\mathbf{r}, \mathbf{r}')]$$
(3)

$$\chi_0(\mathbf{r}, \mathbf{r}', i\nu) = \sum_{i,j} (f_i - f_j) \frac{\phi_i^*(\mathbf{r})\phi_i(\mathbf{r}')\phi_j^*(\mathbf{r}')\phi_j(\mathbf{r})}{\varepsilon_i - \varepsilon_j + i\nu}$$
(4)

By defining $\chi_{\lambda}(\mathbf{r}, \mathbf{r}', i\nu)$ via the self-consistent screening Dyson equation, eq 5, and by setting the exchange-correlation kernel $f_{\rm xc}^{\lambda} = 0$ as for the RPA,²² the correlation energy in eq 3 assumes the form in eq 6 where the last equality is obtained by anaytical integration over λ .²³

$$\chi_{\lambda} = \chi_0 + \chi_0 (\lambda \nu + f_{\rm xc}^{\lambda}) \chi_{\lambda}$$
⁽⁵⁾

$$\mathcal{E}_{c}^{\text{RPA}} = -\frac{1}{2\pi} \int_{0}^{\infty} d\nu \int_{0}^{1} \frac{d\lambda}{\lambda} \operatorname{Tr} \left[\frac{(\lambda \nu \chi_{0})^{2}}{1 - \lambda \nu \chi_{0}} \right]$$
$$= \frac{1}{2\pi} \int_{0}^{\infty} d\nu \operatorname{Tr} [\ln(1 - \chi_{0}\nu) + \chi_{0}\nu] \tag{6}$$

In virtue of its generality, the framework here discussed for the case of electronic correlation, can be used in different contexts and in particular in connection with the many-body dispersion model where the target energy arises from the correlation of coupled fluctuating dipoles.²³ In this case the response function assumes the form of the frequencydependent atom-in-molecule (AIM) isotropic dipole polarizability $\boldsymbol{\alpha}_{i}^{\mu}(i\nu)$ localized at the atomic position $R_{i\nu}$ while the Coulomb potential $\nu(\mathbf{r}, \mathbf{r}')$ is replaced by the (properly damped) dipole-dipole interaction tensor $\mathbf{T}_{ij}^{\text{LR}} = \mathbf{T}_{ij}^{\text{LR},\mu\mu}$.

Eq 6 (last equality) is, thus, written in terms of the MBD model's quantities as shown in eq 7, $\mathbf{A}(i\nu)$ being the (3*N*, 3*N*) diagonal superpolarizability matrix having atomic polarizabilities as elements and \mathbf{T}^{LR} being the full dipole–dipole interaction tensor excluding self-interactions ($\mathbf{T}_{ii}^{LR} = \mathbf{0}$). We note that eq 7 does not include the $\chi_0 \nu$ term appearing in eq 6 due to the traceless property of the $\mathbf{A}(i\nu)\mathbf{T}^{LR}$ matrix product.

$$\mathbf{A}(i\nu) = \text{Diag}[\boldsymbol{\alpha}_{1}^{\mu}(i\nu), \, \boldsymbol{\alpha}_{2}^{\mu}(i\nu), \, ..., \, \boldsymbol{\alpha}_{N}^{\mu}(i\nu)]$$
$$\mathcal{E}_{\text{MBD}} = \frac{1}{2\pi} \int_{0}^{\infty} d\nu \text{Tr}[\ln(\mathbf{I} - \mathbf{A}(i\nu)\mathbf{T}^{\text{LR}})]$$
(7)

For MBD models based on coupled fluctuating dipoles, the RPA formula in eq 7 does not represent the most efficient

strategy to compute correlation energies solution as the problem can be equivalently solved via exact diagonalization of the MBD potential matrix for a system of coupled quantum harmonic oscillators.²³ However, the power and generality of the RPA formulation of the MBD model allows for the generalization of the model and the inclusion of higher-order polarizabilities as recently discussed by Massa et al.¹⁹

The matrix form of eq 7 can rather straightforwardly be generalized to higher-order moments gathered in $A(i\nu)$ by consistently augmenting the interaction tensor T with proper interaction blocks. In particular, for a model based on coupled fluctuating dipoles and quadrupoles (*Q*), the two matrices of dimension (12*N*, 12*N*), *N* being the number of atoms in the systems, take the form reported in eq 8 where a general out-of-

$$\mathbf{A}_{kk}(i\nu) = \mathbf{Diag}[\alpha_k^{\mu}(i\nu), \alpha_k^{\mu}(i\nu), \alpha_k^{\mu}(i\nu), \alpha_k^Q(i\nu), \dots, \alpha_k^Q(i\nu)]$$

$$\mathbf{T}_{ik}^{\mathrm{LR}} = \left(\begin{array}{c|c} \mathbf{T}_{ik}^{\mathrm{LR},\mu\mu} & \mathbf{T}_{ik}^{\mathrm{LR},\muQ} \\ \hline \mathbf{T}_{ik}^{\mathrm{LR},\muQ} & \mathbf{T}_{ik}^{\mathrm{LR},QQ} \\ \hline \mathbf{T}_{ik}^{\mathrm{LR},\muQ} & \mathbf{T}_{ik}^{\mathrm{LR},QQ} \end{array} \right) , \quad i \neq k$$
(8)

diagonal (thus nonvanishing) block of size (12,12) is shown. We will now focus our attention on **A** and, in particular, on its dipole and quadrupole dynamic polarizability entries being the model's key parameters.

As mentioned earlier, a pleasant feature of the MBD model is the fact that it only relies on *ab initio*-derived parameters. In particular, AIM dipole polarizabilities are typically obtained via the volume rescaling approach shown in eq 9, α_i^{μ} and V_i being the AIM static dipole polarizability and volume respectively of the *i*-th atom while the zero superscript denotes free-atom reference quantities.

$$\alpha_i^{\mu}(0) = \left(\frac{V_i}{V_i^0}\right) \alpha_i^{\mu,0}(0)$$
(9)

This volume/polarizability proportionality was discussed by Brinck et al.²⁴ and later used by Johson and Becke,²⁵ as well as Tkatchenko and Scheffler,²⁶ and further adopted in most of the MBD model's variations. Gould discussed a more complicated volume/polarizability relationship²⁷ while Szabo et al. recently suggested a four-dimensional scaling of dipole polarizability based on single-particle systems' analysis.²⁸ We will, in the present work, stick to the common volume/polarizability proportionality in eq 9 as this will allow us to make ready comparisons with our dipole-only DNN-MBD model.

The AIM volume V_i is commonly accessed by Hirshfeld partitioning the explicit electron density obtained via the solution of KS equations. Instead, we will adopt the recently proposed efficient and accurate 5-hidden layers DNN model trained with the minimal basis iterative Stockholder atom²⁹ (MBISA) volumes of approximately 4.6 millions molecules that provides AIM volumes bypassing the electron density partitioning.¹⁸

Compared to Hirshfeld partitioning as well as its Iterative HI variant more suitable for ionic systems,¹⁵ the MBISA scheme used to generate the training set does not suffer from asymmetric AIM densities arising from the use of free-atom reference densities that, especially for hydrogen atoms, lead to an overestimation of high radial moments, i.e., AIM volumes.²⁹

A further disadvantage related to the use of free-atom reference densities affecting the HI partitioning arises from the density interpolation for negatively charged atoms. In fact this procedure is, for some anionic species and for all doubly charged anions typically encountered in inorganic oxide clusters, ill-defined.

The next step is to find a suitable expression for $\alpha_i^Q(0)$, possibly without the introduction of empirical parameters to preserve the original MBD spirit. At this point, it is possible to relate two consecutive dispersion coefficients by the scaled Starkschall–Gordon relation³⁰ shown in eq 10 for the specific case of homonuclear C_6^{ii} and C_8^{ii} coefficients, where Z_i is the atomic number, $\langle r^n \rangle$ are expectation values or multipole-type moments derived from atomic densities while γ scaling factor will be introduced shortly.³⁰ We note, in passing, that eq 10 represents a fundamental recursion relation for Grimme's successful DFT-D3 dispersion correction model.³¹

$$C_8^{ii} = 3C_6^{ii}Q_i$$
$$Q_i = \gamma \sqrt{Z_i} \frac{\langle r^4 \rangle_i}{\langle r^2 \rangle_i}$$
(10)

In order to retrieve $\alpha_i^Q(0)$ in terms of known quantities, we can now assume to model atom *i* via a quantum Drude particle characterized by the frequency ω_i as this allows us to express C_6^{ii} and C_8^{ii} in terms of dipole and quadrupole polarizabilities, eqs 11 and 12.³²

$$C_6^{ii} = \frac{3}{4} \omega_i [\alpha_i^{\mu}(0)]^2 \tag{11}$$

$$C_8^{ii} = 5\omega_i \alpha_i^{\mu}(0)\alpha_i^Q(0) \tag{12}$$

By inserting eqs 11 and 12 into eq 10, it is finally possible to isolate the static AIM quadrupole polarizability in eq 13.

$$\alpha_i^Q(0) = \frac{9}{20} Q_i \alpha_i^\mu(0) \tag{13}$$

The combination of the scaled Starkschall–Gordon relation with a quantum Drude particle expression for the C_8^{ii} dispersion coefficient was originally adopted by Carter-Fenk et al. in their extended symmetry-adapted perturbation theory with MBD (XSAPT+MBD) approach including pairwise dipole–quadrupole dispersion effects.³³ In that occasion, the γ scaling factor was successfully modeled according to eq 14, where γ^0 was chosen such that noble-gas quadrupole polarizabilities are reproduced and the same strategy is embraced in this work.

$$\gamma = \gamma_0 + \exp(-\sqrt{Z_i}/2) \tag{14}$$

A further discussion about the combination of the Starkschall– Gordon rule with a quantum Drude particle expression is found in the Appendix section at the end of the manuscript.

The above-discussed recursive relation represents an efficient strategy to derive quadrupole polarizabilities that can be readily applied to any AIM volume partitioning scheme as the atomic volume is the only density-related quantity which, however, in this work is retrieved from an accurate DNN model.

The solution of eq 7 requires frequency-dependent atomic polarizabilities while our discussion was, until now, restricted to the zero-frequency (static) case.

Dynamical dipole polarizabilities have been successfully modeled via a [0/2] Padé approximant form and Tang et al.

suggested the same form to be employed also in modeling higher-order multipole dynamical polarizabilities, as shown in eq 15, the superscript M denoting a general multipole level and $\omega_k^{M,0}$ the *k*-th free-atom multipolar-dependent characteristic excitation frequency.³⁴

$$\alpha_k^M(i\nu) = \frac{\alpha_k^{M,0}(0)}{1 - (i\nu/\omega_k^{M,0})^2}$$
(15)

In particular, we employ a common $\omega_k^{M,0}$ parameter for both dipole and quadrupole dynamical polarizabilities and we express it as a function of free atom C_6 and static dipole polarizability, eq 16.

$$\omega_k^{Q,0} = \omega_k^{\mu,0} = \frac{4}{3} \frac{C_6^{0,kk}}{[\alpha_k^{\mu,0}(0)]^2}$$
(16)

This assumption is, however, well-defined as it guarantees modeling dynamical quadrupole polarizabilities as lower bound quantities while the introduction of *ab initio*-derived scaling factors could be used to improve over this assumption.³⁴

We note that the dipole polarizabilities used in eq 13 can be either taken directly from eq 9 or can be further screened by solving Dyson-like self-consistent screening equations.^{13,35} In this work the latter is chosen and, as a consequence, these screening effects are thus transferred to quadrupole polarizabilities via eq 13.

The explicit expressions of the multipole interaction tensor T_{ik}^{LR} is given in eq 17 for the *ik* pair placed at distance R_{ik} , and greek letters are used to denote *x*, *y*, *z* Cartesian components.

$$\begin{aligned} \mathbf{T}_{ik,\tau\sigma}^{\mathrm{LR},\mu\mu} &= f_{\mu\mu} (R_{ik}, S_{ik}) \nabla_{ik,\tau\sigma}^{2} \left(\frac{1}{R_{ik}}\right) \\ \mathbf{T}_{\tau\sigma\theta}^{\mathrm{LR},\muQ} &= f_{\muQ} (R_{ik}, S_{ik}) \nabla_{ik,\tau\sigma\theta}^{3} \left(\frac{1}{R_{ik}}\right) \\ \mathbf{T}_{ik,\tau\sigma\theta\lambda}^{\mathrm{LR},QQ} &= f_{QQ} (R_{ik}, S_{ik}) \nabla_{ik,\tau\sigma\theta\lambda}^{4} \left(\frac{1}{R_{ik}}\right) \end{aligned}$$
(17)

The general form of the Fermi damping function $f_{MM'}(R_{ik}, S_{ik})$ (MM' denoting $\mu\mu$, μQ , or QQ) is defined in eq 18 where R_i^{vdw} represents the AIM van der Waals radius here modeled as the ratio between the AIM dipole (screened) polarizability and the free atom one while β is the parameter modulating the range-separation of the multipole interaction tensor. Finally the $l_{MM'}$ term, represents a multipole-dependent universal factor that ensures a stronger damping as the multipole order increases ($l_{\mu\mu} < l_{\muQ} < l_{QQ}$), in line to the behavior of the popular Tang and Toennies damping functions based on the incomplete gamma function.³⁶

$$f_{MM'}(R_{ik}, S_{ik}) = \frac{1}{1 + \exp(-6(R_{ik}/S_{ik} - 1))}$$
$$S_{ik} = \beta l_{MM'}(R_i^{vdw} + R_k^{vdw})$$
$$R_i^{vdw} = \left(\frac{\alpha_i^{\mu}(0)}{\alpha_i^{\mu,0}(0)}\right)^{1/3} R_i^{0,vdw}$$
(18)

These $l_{MM'}$ terms could in principle be set as parameters and optimized together with β to reduce a target energy error. However, to avoid overfitting of these terms we will choose them once and for all regardless of the employed functional.

In particular, the $l_{\mu\mu}$ term was set to 1 as this follows the choice undertaken within dipole-only models. On the other hand, $l_{\mu Q}$ and l_{QQ} were chosen after a screening of a few test values, with the constraint being $l_{\mu Q} < l_{QQ}$ as this prevents short-range spurious effects. We note in passing that there is clearly room for improvements if a fully data-driven process was embraced in the optimization of these terms.

We solve the RPA formula in eq 7 by means of Gauss– Legendre quadrature, nevertheless, more sophisticated techniques such as the Clenshaw-Curtis quadrature^{37,38} could also be potentially employed.

The DNN-MBDQ model is implemented in the Tinker-HP package³⁹ where the extremely efficient linear-scaling stochatic Lanczos-based DNN-MBD model is also implemented.⁴⁰

In order to design an accurate and reliable dispersioncorrected KS-DFT model, both the functional and the dispersion correction must fulfill specific criteria. These requirements are essential for avoiding error cancellation effects in favor of more physically grounded achievement of small errors, as recently discussed by Price et al.⁴¹ While the dispersion correction should possibly include higher-order terms as well as many-body effects, the DFT functional should be dispersionless since by adding a dispersion correction to an exchange-correlation functional that somewhat partially includes dispersion, will lead to double counting at shorter ranges, with a consequent loss of accuracy. In fact, in order to reduce this artifact, dispersion corrections are often excessively damped with an inevitable deterioration of the midrange interactions' description. Among the possible dispersionless functionals, the B86bPBE has proven numerically stable and, coupled with the exchange-hole dipole moment dispersion model,^{42,43} has performed well for molecular and material applications.⁴⁴ On account of both the higher-order terms and the many-body nature of the proposed DNN-MBDQ dispersion correction, thus fulfilling the above-mentioned criteria, it is of interest to probe its performances in connection with the B86bPBE dispersionless functional. Furthermore, the DNN-MBDQ dispersion model is also coupled to the common PBE and PBE0 semilocal density functionals and this allows us to make comparisons with different MBD-based dispersion corrections including our recently proposed dipole-only DNN-MBD.

Pure energies have been calculated in all cases with Jensen's pcseg-3 basis set as this corresponds *de facto* to the complete basis set limit.⁴⁵

The S66x8 data set for noncovalent biologically relevant interactions⁴⁶ consisting of 66 dimers placed at 8 different intermolecular distances (CCSD(T) complete basis set interaction energies) was taken as reference to tune the β parameter in eq 18 for the three functionals here discussed and the optimal values relative to the DNN-MBDQ model are shown in Table 1 together with the ones for dipole-only based

Table 1. Range-Separation β Parameter Values Optimized by Minimizing the Mean Absolute Relative Error for the DNN-MBD and DNN-MBDQ Models Coupled with the PBE and PBE0 Functionals for the S66x8 Data Set

DFT functional	DNN-MBD	DNN-MBDQ
PBE	0.75	0.82
PBE0	0.77	0.83
B86bPBE	0.69	0.76



Figure 1. Interaction energy for the argon dimer computed with the PBE and B86bPBE pure functionals (pcseg-3 basis set) as a function of the interatomic distance.

DNN-MBD. $l_{\mu\mu\nu}$, $l_{\muQ2}$ and l_{QQ} factors are set to 1, 2.8, and 3.0, respectively, as these have proven to be good and transferrable values for different functionals. We observe that the PBE0/DNN-MBDQ method requires a larger range-separation parameter compared to the correspondent PBE-based model. This is consistent to what observed also in the PBE/DNN-MBD model due to the PBE0 functional's improved description of short-range exchange-correlation.

Moreover, the DNN-MBDQ model requires larger rangeseparation parameters compared to the DNN-MBD one, and this can be rationalized by virtue of the additional stabilizing dispersion contributions arising from the dipole-quadrupole and quadrupole-quadrupole interactions being the many-body analogues of the pairwise C_8 and C_{10} terms in eq 2. We note, in passing, that compared to the original MBD method based on Hirshfeld partitioning for the same functionals (PBE and PBE0), our DNN-MBD model requires smaller β values. The reason has to be sought in the relatively smaller AIM volumes predicted by the MBISA method used to build the DNN training set and the consequent effects in AIM volume-scaled polarizabilities for which less screening is necessary. Focusing now on the B86bPBE dispersionless case, we observe that both the DNN-MBD and DNN-MBDQ models require a smaller range-separation parameter (less damping) compared to their PBE/PBE0 counterparts. This is completely in line with the less stabilizing nature of the B86bPBE functional arising from its dispersion-free features as shown in Figure 1 where the pure potential energy surface of the argon dimer for the PBE and B86bPBE functionals are shown. In fact, a less damped DNN-MBDQ correction leads to model dispersion at shorter ranges where functionals such as PBE and PBE0 usually involve more damped dispersion corrections to avoid double counting, Figure 2.

It is now of interest to compare the DNN-MBDQ error for the present data set with the ones related to different dipoleonly MBD models, Table 2, where the DFT-D3 correction to the PBE functional is also included as a reference method considering its broad use and popularity. The proposed DNN-



Figure 2. Pictorial representation of the short-, mid-, and long-range interaction regimes. The dashed line represents the transition between the short-range part modeled by the KS-DFT functional (blue) and the long-range (yellow) regimes modeled by the DNN-MBDQ dispersion correction where the introduced quadrupoles improve the modeling of midranges near equilibrium distances. Smaller β parameters required by dispersionless functionals have the effect of pushing the yellow zone toward shorter ranges.

MBDQ model exhibits reduced mean absolute errors, although by a small margin, compared to its dipole-only based DNN-MBD version coupled with both PBE0 and PBE functionals while for the B86bPBE dispersionless functional, the DNN-MBDQ model introduces a more consistent improvement compared to the DNN-MBD and it reaches a notably small value of 0.19 kcal/mol. The MAE of the DNN-MBDQ model coupled to the PBE and PBE0 functionals is always lower than the one relative to other MBD-based models, for which reference data are found in literature, however, except for the B86bPBE functional the DNN-MBDQ model have an slightly higher MARE value. Even in the case of the DNN-MBDQ, dispersion corrected PBE0 provides the smallest error for the considered S66x8 data set, in line to what observed in the DNN-MBD as well as for other dipole-only MBD models.

As anticipated earlier, Massa et al. have recently proposed a beyond-dipole MBD model based on the RPA formalism termed MBD+Q@rsSCS, to be coupled to DFT,^{19,48} and it is thus of interest to compare it with our DNN-MBDQ model. In

Table 2. MAE (kcal/mol) and MARE (%) Relative to the S66x8 Data Set for Our Density Free DNN-MBD and DNN-MBDQ Models, As Well As Few Other Dipole-Only MBD Models Coupled with the PBE and PBE0 Functionals.^{*a*}

Model	MAE(kcal/mol)	MARE (%)
PBE	1.55	65
PBE/D3 ⁴⁷	0.44	n.a.
PBE/MBD@rsSCS ¹³	0.32	10.6
PBE/MBD@rsSCS/FI ¹⁴	0.28	9.0
PBE/DNN-MBD ¹⁸	0.25	9.0
PBE/DNN-MBDQ	0.24	10.5
PBE0	1.48	65
PBE0/MBD@rsSCS ¹³	0.30	9.2
PBE0/DNN-MBD ¹⁸	0.23	6.9
PBE0/DNN-MBDQ	0.22	8.3
B86bPBE	2.01	82
B86bPBE/DNN-MBD	0.31	9.9
B86bPBE/DNN-MBDQ	0.19	7.6

^{*a*}By virtue of its popularity and thus relevance, Grimme's D3 correction is also included. MAE and MARE are computed.

their approach the coupled quantum harmonic oscillators modeling many-body dispersion are parametrized by a mixed approach based on both the Tkatchenko–Scheffler polarizability rescaling and Johnson' exchange dipole model.⁴⁹ This differs substantially from our approach where quadrupole polarizabilities are obtained recursively from dipole ones, in turn derived from our DNN-based volume-rescaling scheme. Table 3 directly compares the MBD+Q@rsSCS and our DNN-

Table 3. MBD+Q@rsSCS and DNN-MBDQ Methods Compared in Terms of Mean Absolute Errors (kcal/mol), while MARE Are Reported in Parentheses for the S66, S66x7, and S66x8 Datasets^a

Model	S66	S66x7	S66x8
PBE/MBD+Q@rsSCS	0.45 (9.3%)	0.27 (7.6%)	0.34 (12.0%)
PBE/DNN-MBDQ	0.25 (5.9%)	0.22 (10.3%)	0.24 (10.5%)
PBE0/MBD+Q@rsSCS	0.28 (11.8%)	0.22 (7.1%)	0.37 (8.3%)
PBE0/DNN-MBDQ	0.23 (4.5%)	0.19 (7.8%)	0.22 (8.3%)

"In the MBD+Q@rsSCS model, two parameters are optimized the number in the damping function against the S66x8 set while for the DNN-MBDQ method a single parameter is optimized.

MBDQ model based on the complete S66x8 data set, the S66x7 subset where the shortest intermolecular distances are disregarded as well as the S66 data set for both the PBE and PBE0 functionals. In the data sets considered, and for both PBE and PBE0 cases, the DNN-MBDQ exhibits lower MAE and MARE compared to MBD+Q@rsSCS, except for the S66x7 where the MBD+Q@rsSCS model exhibits a lower MARE. In both models, the removal of the shortest intermolecular distance (S66x7) involves a lowering of the errors compared to the full S66x8 set and this denotes the challenge related to modeling mid- to short-range intermolecular distances where dispersion models smoothly turn into local electron correlation governed by the semilocal density functionals. The overall lower errors exhibited by the DNN-MBDQ model can be explained in terms of the differences between two models. In fact, we note that while in the DNN-MBDQ multipole dependent damping functions are employed, the MBD+Q@rsSCS model relies on a single damping function for the different multipole-interaction tensors. At shorter intermolecular distances this may not be optimal as the interaction among higher multipole moments, such as quadrupole-quadrupole grows 2 orders of magnitude faster than the dipole-dipole ones and a consistent differentiation of their damping functions is required. While the choice of a multipole-dependent damping functions is essential for multipole interaction tensors obtained from the differentiation of the 1/R Coulomb term (Eq. 18), it becomes less crucial if the intrinsically damped $\operatorname{erf}(R/\sigma)/R$ term characterizing the interaction of Gaussian-distributed charge densities of width σ is differentiated instead as assumed in the MBD+Q@rsSCS model.¹⁹ This explains the possibility of using a single Fermi damping function in connection to the MBD+Q@rsSCS model that, although not reaching the accuracy of the DNN-MBDQ model, provides chemically accurate interaction energies.

The dipole contribution to dispersion interactions dominates the long-range asymptotic limit while inclusion of quadrupole contributions improves the description at medium ranges near equilibrium distances. The S66x8 is an optimal data set to optimize the range-separation parameters as it is not biased toward equilibrium distances; however, in virtue or its wide range of intermolecular distances, it does not represent a fully optimal set to investigate the effects of quadrupole terms. In that regard, results on the S22 data set, which is composed of model complexes at equilibrium distances, are a better judge of the accuracy of our model since S22 includes more interactions in the full quadrupole (and mixed dipolequadrupole) operational regime. The S22 validation set thus enables us to better understand the effects of the added quadrupole terms as well as to investigate the transferability of the DNN-MBDQ model by employing the same rangeseparation parameters reported in Table 1. Table 4 shows the

Table 4. MAE (kcal/mol) and MARE (%) Relative to the S22 Data Set^{50,51} for Some Dipole-Only Based MBD Models, As Well As for Our DNN-MBDQ Model^a

Model	MAE (kcal/mol)	MARE (%)
PBE	2.66	58
PBE/D3 ¹	0.48	9.9
PBE/MBD@rsSCS	0.49	8.9
PBE/DNN-MBD	0.41	6.6
PBE/DNN-MBDQ	0.27	6.0
PBE0	2.44	55
PBE0/MBD@rsSCS	0.55	8.5
PBE0/DNN-MBD	0.43	5.6
PBE0/DNN-MBDQ	0.32	5.9
B86bPBE	3.29	72
B86bPBE/DNN-MBD	0.50	10.4
B86bPBE/DNN-MBDQ	0.39	9.7

^{*a*}For the MBD-based models, the range-separation parameter was optimized against the S66x8 dataset.

performances for some of the dispersion correction models reported in Table 2 for which results relative to the S22 set are available in literature. Compared to the S66x8, the MAE values reported in Table 4 for the DNN-MBDQ model coupled to the chosen DFT functionals are higher (although abundantly below the chemical accuracy threshold), however, this is in line with the fact that no β optimization was carried out this time. All the PBE, PBE0, and B86bPBE functionals coupled to the

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DNN-MBDQ model exhibit markedly reduced MAE values compared to the already accurate DNN-MBD dipole-only model while MARE values are not significantly affected. If the DNN-MBD model is compared directly to the MBD@rsSCS model coupled with the PBE and PBE0, the MAE decreases by 45% and 42% respectively. These noteworthy performances stem certainly from the additional quadrupolar terms as this is readily observed by a direct comparison between the DNN-MBD and DNN-MBDQ models. The pleasant recursive relation for quadrupole polarizabilities, eq 13, benefits from the DNN model's AIM volumes accuracy arising from the MBISA volumes' data used in the training process whose advantages, compared to the common Hirshfeld partitioning, have been discussed profusely in references and previously recalled in this work.^{18,29}

Among MBD-based models, the DNN-MBDQ coupled with the functionals here considered provides one of the lowest errors for the S22 set without an *ad hoc* parameter optimization. This is rather significant considering that the model fully relies on *ab initio* quantities. The accuracy performances of the DNN-MBDQ model could be potentially increased further by adding an extra parameter in the Fermi damping function (eq 18), as well as by performing a complete optimization for the choice of the $l_{MM'}$ terms and future efforts will be spent in this direction.

We propose the DNN-MBDQ model where the densityfree/deep learning-aided many-body dispersion model is extended to quadrupolar polarizability terms thanks to a Random Phase Approximation formalism. Quadrupole polarizabilities are recursively retrieved from free-atom dipole polarizabilities and atom-in-molecule volumes modeled via the accurate and transferable neuronal network recently proposed.¹⁸ The described strategy can be readily applied to MBD model based on the volume rescaling polarizability partitioning as it does not require any additional electron density-derived quantity. The density-free DNN-MBDQ model implemented in the Tinker-HP package exhibits improved accuracy compared its dipole-only DNN-MBD counterpart, as well as reference MBD-based models. In particular, for the widely used S22 data set of dimers placed at equilibrium intermolecular distances, the DNN-MBDQ shows a remarkable improvement compared to dipole-only reference MBD models.

In addition to the common and successful PBE and PBE semilocal functionals, the DNN-MBDQ model is also coupled with the B86bPBE one, to explore the possibility of employing dispersionless functionals, preventing double counting dispersion, especially at shorter ranges. For this case, the errors are abundantly below the chemical accuracy threshold and similar (in some cases even lower) to the ones obtained for the PBE/PBE0 case.

We believe that the high accuracy exhibited by the DNN-MBDQ model coupled with semilocal and dispersionless functionals will be beneficial in pushing Kohn–Sham Density Functional Theory a step closer to post-Hartree–Fock reference standards which, due to their high computational cost cannot be employed to explore larger systems of relevance in bio- and material modeling. Further work will be, therefore, dedicated to the evaluation of possible couplings of the DNN-MBDQ approach with existing modern DFT functionals toward generalized chemical accuracy. Such a strategy opens the door to designing accurate, large-scale, energies databases toward various machine learning applications. Finally, the

DNN-MBDQ energy correction can be directly added as an *a posteriori* term to pure KS-DFT energies in virtue of the model's density-free features arising from the transferable deep neuronal network employed.

APPENDIX

An analytical connection between the Starkschall–Gordon rule and the quantum Drude oscillator (QDO) expression is nontrivial to obtain; however, to verify the consistency of the two expressions, it is possible to evaluate the C_8/C_6 proportionality factor for the hydrogen atom, for which α^Q , α^{μ} , $\langle r^2 \rangle$, and $\langle r^4 \rangle$ are analytically available.^{30,52}

We start by taking the QDO expression for the C_8 coefficient where we introduce $\omega = \frac{4}{3} \frac{C_6}{(\alpha'')^2}$. The resulting expression is the QDO-analogue of the Starkschall–Gordon rule, see equation below where the second equality represents the unscaled Starkschall–Gordon rule for hydrogen.

$$C_8 = 203 \left(\frac{\alpha^Q}{\alpha^{\mu}}\right) C_6 = f_{\text{QDO}} C_6$$
$$C_8 = 3 \left(\frac{\langle r^4 \rangle}{\langle r^2 \rangle}\right) C_6 = f_{\text{SG}} C_6$$

It is now possible to evaluate $f_{\rm QDO}$ and $f_{\rm SG}$ from the analytical α^Q , α^μ and $\langle r^2 \rangle$, $\langle r^4 \rangle$ expressions respectively to get $f_{\rm QDO} = 22.22$ and $f_{\rm SG} = 22.5$ (atomic units). The $\approx 1\%$ difference among the two factors denotes a marked consistency of the two formulations while the scaling version of the Starkschall–Gordon rule simply improves the results for light elements as discussed by Carter-Fenk and Herbert et al. in connection to their XSAPT+MBD scheme.³³

ASSOCIATED CONTENT

Data Availability Statement

Data can be download directly via the Zenodo repository located at DOI: 10.5281/zenodo.6397639. It contains raw PBE, PB0, and B86bPBE interaction energy data for the S66x8 and/or S22 data sets, as well as their relative DNN-MBDQ dispersion corrections for the different β values discussed.

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Notes

The authors declare no competing financial interest.

ACKNOWLEDGMENTS

This work has been funded by the European Research Council (ERC) under the European Union's Horizon 2020 research and innovation program (grant No 810367), project EMC2 (JPP). Computations have been performed at GENCI (IDRIS, Orsay, France and TGCC, Bruyères le Chatel) on grant no A0070707671.

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