## Van der Waals density functional: An appropriate exchange functional

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In this Rapid Communication, an exchange functional which is compatible with the nonlocal Rutgers-Chalmers correlation functional [van der Waals density functional (vdW-DF)] is presented. This functional, when employed with vdW-DF, demonstrates remarkable improvements on intermolecular separation distances while further improving the accuracy of vdW-DF interaction energies. The key to the success of this threeparameter functional is its reduction in short-range exchange repulsion through matching to the gradient expansion approximation in the slowly varying/high-density limit while recovering the large reduced gradient, *s*, limit set in the revised Perdew-Burke-Ernzerhof (revPBE) exchange functional. This augmented exchange functional could be a solution to long-standing issues of vdW-DF lending to further applicability of densityfunctional theory to the study of relatively large, dispersion bound (van der Waals) complexes.

DOI: 10.1103/PhysRevB.81.161104

PACS number(s): 71.15.Mb, 31.15.E-, 61.50.Lt

Van der Waals, or London dispersion, interactions have profound importance in bio-organic systems as well as many materials being investigated for energy applications. Unfortunately, the inability of traditional density-functional theory (DFT) exchange-correlation functionals to account for long ranged, van der Waals interactions has limited first-principles investigations of these systems to quantum chemical (QC) methods. Due to their computational expense, QC calculations have been limited to fragments of the true material; thus they may overlook some of the more salient features of these systems. As such, significant research efforts have been devoted to address the deficiencies of DFT. Many of the more popular and successful approaches, however, either require some empirical parametrization (e.g., dispersioncorrected DFT and pseudopotential-based correction schemes) or scale poorly with system size (e.g., N<sup>4</sup>-N<sup>5</sup> for symmetry adapted perturbation theory (SAPT)-DFT and N<sup>4</sup> for the random phase approximation).<sup>1,2</sup>

A promising solution to balancing speed and scalability with accuracy lies in the nonlocal correlation functional of the Rutgers-Chalmers collaboration, the aptly named van der Waals density functional (vdW-DF).<sup>3,4</sup> Here, long-range dispersion effects are included as a perturbation to the localdensity approximation correlation term. This method has been successful in describing a diverse group of materials properties: ranging from molecules to bulk polymers and the adsorption of molecules to surfaces and within bulk materials.<sup>5</sup> Recent developments show that self consistency gives no appreciable differences in computed interaction energies<sup>3,4</sup> and the vdW-DF can be incorporated in an extremely efficient manner.<sup>6</sup> However, the overwhelming success of vdW-DF is marred by its consistent overestimation of intermolecular distances.<sup>5</sup> Analysis of various generalized gradient approximation exchange functionals (GGA<sub>x</sub>) indicate that traditional functionals are either too repulsive at short distances or incorrectly exhibit some "correlation" binding at larger distances. The standard functional used within the vdW-DF, the revised Perdew-Burke-Erzenhoff functional (revPBE),7 unfortunately gives too much repulsion at short distances. Replacing revPBE with Hartree-Fock (HF) exchange improves vdW-DF interspecies separation distances but at the cost of overbinding, i.e., considerably larger interaction energies than obtained via coupled cluster, CCSD(T), calculations.<sup>8,9</sup> Recent work suggests that for many dispersion bound materials, the PW86 functional<sup>10</sup> most closely matches HF exchange.<sup>11,12</sup> Similarly when applied with vdW-DF, it also strongly overbinds (see Fig. 2).

In this Rapid Communication, a  $GGA_x$  that may be suitable for use with the vdW-DF correlation functional is proposed. This functional is derived through the introduction of an enhancement factor which obeys two specific constraints: (i) matching to the gradient expansion approximation (GEA) (Ref. 14) in the slowly varying/high-density limit and (ii) a smooth asymptote to the upper bound empirically set in revPBE. Initial results indicate dramatic improvements in vdW-DF separation distances and the chemical accuracy for a range of systems. Most notable are improvements in results obtained for S22 database structures<sup>15</sup> and graphite.

The general formula of a GGA<sub>x</sub> can be written as

$$E_{\rm x}^{\rm GGA} = \int d^3 r n \,\epsilon_{\rm x}^{\rm unif}(n) F_{\rm x}(s), \qquad (1)$$

where  $\epsilon_x^{\text{unif}}(n)$  is the exchange energy per particle in a uniform gas,  $\epsilon_x^{\text{unif}}(n) = -3ek_F/4\pi$  with  $k_F = [3\pi^2 n]^{1/3}$ , and  $F_x(s)$  is the enhancement factor which is a function of  $s = |\nabla n|/(2k_F n)$ . This form of exchange ensures proper, uniform density scaling<sup>16</sup> where  $F_x(s) = 1$  simply gives LDA exchange. In general, the enhancement factor is chosen such that  $F_x(0) = 1$ . Here we design an  $F_x(s)$  to fulfill two further criteria:

(i) to reduce the short-range exchange repulsion, in the limit of  $s \rightarrow 0$ , i.e., for slowly varying/high densities, the functional uses the GEA form

$$F_x^{\text{GEA}}(s) = 1 + \mu s^2 \tag{2}$$

with  $\mu = 0.0864$ .<sup>14</sup> (It should be noted that  $\mu_{\text{GEA}} = 10/81$ . The above value was used in PW86.<sup>10</sup>) Figure 1 depicts the enhancement factor of a number of GGA<sub>x</sub>. It can be clearly seen that many of these functionals deviate quite quickly from the GEA. Note that this constraint is similar to that used in the recent PBEsol exchange functional which was designed to restore the gradient expansion in order to remove



FIG. 1. (Color online) Enhancement factor for various  $GGA_x$  functionals. The blue (dotted), red (dashed), and green (solid) lines represent enhancement factors for PBE (Ref. 17), revPBE (Ref. 7), and PW86 (Ref. 10), respectively. The black dashed-dotted line is the GEA and the orange open circles are points from the proposed  $CO9_x$ . The solid black line is the revPBE upper bound.

artificial bias toward free atoms.<sup>18</sup> Here, decreasing the enhancement factor for small s (thus restoring the GEA) leads to a reduction in the short-range repulsion in the GGA<sub>x</sub>.

(ii) The second constraint used in the proposed GGA<sub>x</sub> is an asymptote to the revPBE upper bound of 2.245 in the large *s* limit.<sup>7</sup> This bound gave reasonable results in previous applications of vdW-DF employing the revPBE exchange functional. Here, we find that an empirical  $F_x(s)$  bound similar to revPBE gives the best interaction energies.

Using these constraints a simple, smooth, function can be constructed in the form

$$F_{\rm x}(s) = 1 + \mu s^2 e^{-\alpha s^2} + \kappa (1 - e^{-\alpha s^2/2})$$
(3)

with  $\mu$ =0.0617,  $\kappa$ =1.245, and  $\alpha$ =0.0483. Figure 1 displays the enhancement factor of Eq. (3) along with that for other GGAs. The parameters were determined by simultaneously fitting Eq. (3) to GEA for *s*<1.5 and to revPBE for 8.0<*s*<10.0. This fitting domain was arbitrarily chosen to allow for a decrease in  $F_x(s)$  for small *s* and a smooth recovery of revPBE for large values of *s*. (In accordance with previous naming conventions, this functional shall be referred to as C09<sub>x</sub>.) The complimentary exchange potential can be easily constructed from the functional derivative of Eq. (3) (see Ref. 10).

To test the compatibility of the proposed  $C09_x$  with the LDA correlation term plus the nonlocal vdW-DF long-range correlation term, self-consistent calculations within a modified version of the ABINIT plane-wave code<sup>19</sup> were performed. All calculations were carried out with a 30 Ha plane-wave cutoff and a single k point at  $\Gamma$ . To reduce the effects of periodic images, all simulation cells were padded with at least 10 Å of vacuum in all directions.

The interaction energy,  $\Delta E^{\text{int}}$ , as a function of separation distance,  $d_{\text{sep}}$ , for the benzene dimer stacked in the sandwich configuration is plotted in Fig. 2. A comparison of vdW-DF with the standard revPBE exchange functional



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FIG. 2. (Color online) Interaction energy as a function of separation distance for the benzene sandwich dimer. vdW-DF results with revPBE<sub>x</sub>, CO9<sub>x</sub>, and PW86<sub>x</sub> are represented by a black dashed line with open circles, a red solid line with closed circles, and a solid black line with closed triangles, respectively. CCSD(T) data [blue solid line (Ref. 13)] are plotted for reference.

(vdW-DF<sup>revPBE</sup>) and the exchange functional (vdW-DF<sup> $C09_x$ </sup>) using the enhancement factor of Eq. (3) shows a significant shortening of the separation distance from 4.07 to 3.87 Å. The vdW-DF<sup> $C09_x$ </sup> is now in much better agreement, with regards to both interaction energy and separation distance, with both the benchmark CCSD(T) extrapolated to the complete basis set limit (CBS) ( $\Delta E^{int}$ =1.70 kcal/mol and  $d_{sep}$ =3.90 Å) 13) as well as SAPT (Ref. (DFT)  $(\Delta E^{\text{int}}=1.67 \text{ kcal/mol and } d_{\text{sep}}=3.80 \text{ Å})$  (Ref. 20) calculations. As previously mentioned, the PW86 exchange functional, which was recently reported to mimic Hartree-Fock exchange for dispersion bound complexes, gives excellent separation distances but significantly over estimates the interaction energy.



FIG. 3. (Color online) Interaction energy as a function of interplanar separation distance for graphite. The vdW-DF results with revPBE<sub>x</sub> and  $CO9_x$  are represented by a black dashed line with open circles and a solid black line with closed triangles, respectively. A recent experimental value (Ref. 21) (blue circles) is plotted for reference.

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TABLE I. Computed  $\Delta E$  for the S22 benchmark data (Ref. 15). C09<sub>x</sub> values in parentheses are for full geometry optimizations.  $\Delta E$  for vdW-DF with revPBE<sub>x</sub> and PBE<sub>x</sub> are also listed (structures are fully optimized and taken from Ref. 23, unless otherwise noted). Deformation energies are not included. Energies are in kilocalorie per mole.

		vdW-DF			
No.	Complex	revPBE <sub>x</sub>	PBE <sub>x</sub>	C09 <sub>x</sub>	Benchmark CCSD(T)/CBS
Hydrog	en-bonded complexes				
1	$(NH_3)_2(C_{2h})$	2.44	3.71	2.88 (2.92)	3.17
2	$({\rm H_2O})_2(C_{\rm s})$	4.08	5.58	4.99 (5.15)	5.02
3	Formic acid dimer $(C_{2h})$	14.07	18.28	20.12 (24.59)	18.61
4	Formamide dimer $(C_{2h})$	12.50	16.10	16.15 (17.15)	15.96
5	Uracil dimer $(C_{2h})$	16.17	20.59	20.93 (22.36)	20.65
6	2-pyridoxine $\cdot$ 2-aminopyridine ( $C_1$ )	14.02	17.94	17.67 (19.66)	16.71
7	Adenine $\cdot$ thymine WC ( $C_1$ )	15.19 (Ref. 24)	17.57	17.25 (19.16)	16.37
Comple	xes with predominant dispersion contribution	on			
8	$(CH_4)_2(D_{3d})$	0.88	1.55	0.51 (0.51)	0.53
9	$(C_2H_4)_2(D_{3d})$	1.41	2.68	1.16 (1.16)	1.51
10	Benzene $\cdot$ CH <sub>4</sub> (C <sub>3</sub> )	1.57 (Ref. 25)	2.51	1.50 (1.71)	1.50
11	Benzene dimer $(C_{2h})$	2.74 (Ref. 8)	4.96	3.32 (3.40)	2.73
12	Pyrazine dimer $(C_s)$	3.87	6.25	4.74 (4.75)	4.42
13	Uracil dimer $(C_2)$	9.41 (Ref. 26)	12.91	10.31 (10.41)	10.12
14	Indole · benzene $(C_1)$	4.34	6.25	5.44 (5.48)	5.22
15	Adenine thymine stack $(C_1)$	10.60	14.74	12.79 (12.73)	12.23
Mixed o	complexes				
16	Ethene $\cdot$ ethine $(C_{2v})$	1.55	2.38	1.61 (1.60)	1.53
17	Benzene $H_2O(C_s)$	2.72 (Ref. 27)	4.15	3.25 (3.24)	3.28
18	Benzene · NH <sub>3</sub> ( $C_{\rm s}$ )	1.87	3.18	2.28 (2.27)	2.35
19	Benzene · HCN $(C_s)$	3.87	5.488	4.51 (4.48)	4.46
20	Benzene dimer $(C_{2v})$	2.05 (Ref. 26)	3.98	2.85 (2.84)	2.74
21	Indole · benzene T-shape $(C_1)$	4.72	6.90	5.75 (5.71)	5.73
22	Phenol dimer $(C_1)$	5.81	8.51	7.00 (7.20)	7.05
Average percent deviation		17	36	5 (9)	
Mean unsigned error (kcal/mol)		1.22	1.14	0.31 (0.82)	

To further illustrate the value of the  $\text{C09}_x$ , the interaction energy as a function of *c* lattice parameter for graphite is plotted in Fig. 3. Here,  $\text{C09}_x$  offers significant improvements in both the value of graphite lattice constants as well as the interplanar interaction energy. Our computed lattice constant of 6.56 Å is within 2% of experiment (6.70 Å).<sup>21</sup> The interplanar binding energy (-59 meV/atom) is also greatly improved (-52±5 meV/atom for experiment<sup>21</sup>). Similar results were obtained using the DFT-D approach.<sup>22</sup> Note, vdW-DF<sup>revPBEx</sup> gives an interaction energy and lattice constant of -39 meV/atom and 7.35 Å, respectively.

A more stringent evaluation of the accuracy of the functional can be gained through comparison with the benchmark S22 database of Jurečka *et al.*<sup>15</sup> This database contains the interaction energies and structures of 22 structures with varying degrees of hydrogen bonding and vdW interactions computed with CCSD(T) extrapolated to the complete basis-set limit and is currently accepted as the gold standard for theoretical methods used to study systems with significant dispersion interactions. Recently, Gulans *et al.*<sup>23</sup> examined the S22 database using vdW-DF with both the revPBE and PBE exchange functionals. In general, they found reasonable agreement with the interaction energies of the S22 database; with PBE producing better results for hydrogen-bonded complexes and revPBE showing less deviation for dispersion dominated interactions. However, their results were all for vdW-DF optimized structures; which always give too large separation distances.

Table I lists the computed interaction energies for the S22 database. Here, it is evident that across the board vdW-DF<sup>C09</sup><sub>x</sub> is in much better agreement with the CCSD(T) benchmark values than when vdW-DF is used with either the PBE or revPBE functionals. In fact, vdW-DF<sup>C09</sup><sub>x</sub> has an average percent deviation of 5% (9% if full geometry optimizations were performed); far less than revPBE and PBE (17% and 36%, respectively). Even more important is the

fact that these interaction energies were obtained using the published S22 geometries—without any adjustment of dimer separation distances, demonstrating once again the improvement that this functional offers with regards to both interaction energies and determining optimum separation distances. The mean absolute error (0.31 kcal/mol) is also in remarkable agreement with other approaches for incorporating dispersion interactions (e.g., M06-2X: 0.47 kcal/mol,  $\omega$ B97-D: 0.22 kcal/mol, and B97-D: 0.50 kcal/mol).<sup>28,29</sup>

All 22 structures were subsequently relaxed such that the forces on all the atoms were less than 0.02 eV/Å. These values are listed in parentheses in Table I. Analysis of the relaxed geometries indicate that the majority of the deviations are related to changes in internal bond lengths. This is evident in the larger changes in the vdW-DF<sup>C09</sup><sub>x</sub> interaction energies for hydrogen-bonded structures.

In summary, an exchange functional that is compatible with the Rutgers-Chalmers van der Waals correlation functional is proposed. This functional was derived to closely match the enhancement factor F(s) of the gradient exchange approximation for values of 0 < s < 1.5 while having an asymptote to the revPBE bound of 2.245. In general, this functional shows significant improvements over the previous revPBE exchange. In particular, vdW-DF<sup>C09</sup><sub>x</sub> offers better agreement with the benchmark S22 database with an average deviation of only 5% at the intermolecular separation distances of the published geometries. This is a feat which far surpasses that of previous vdW-DF calculations which produced larger separation distances. These results highlight the promise of this functional for use with the vdW-DF method and may offer a pathway to even more accurate firstprinciples calculations of dispersion bound systems.

V.R.C. would like to thank David C. Langreth for many valuable discussions. V.R.C. also acknowledges comments from Kieron Burke and discussions with Alaska Subedi. This work was supported by U.S. Department of Energy, Division of Materials Sciences and Engineering. This research used resources of the National Energy Research Scientific Computing Center, which is supported by the Office of Science of the U.S. Department of Energy under Contract No. DE-AC02-05CH11231.

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