

Benchmarking the Plasmon-Pole and Multipole Approximations in the Yambo Code Using the GW100 Data Set

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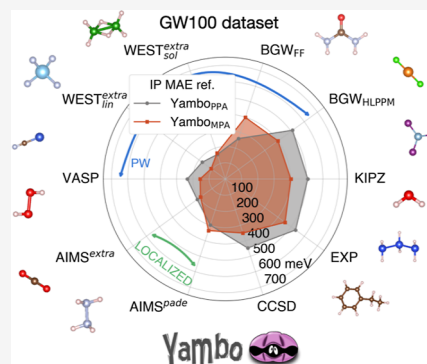


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ABSTRACT: Verification and validation of electronic structure codes are essential to ensure reliable and reproducible results in computational materials science. Besides the extensive work on density functional theory, in the field of many-body perturbation theory, the GW100 data set represents a key effort to benchmark and validate methods such as the GW approximation. In this work, we assess the numerical accuracy and convergence behavior of the GW implementation in the YAMBO code using both the Godby–Needs plasmon-pole model (GN-PPA) and the recently introduced multipole approximation (MPA). Quasiparticle energies are compared against GW100 reference data to evaluate the performance, numerical stability, and consistency of these approaches. Our results show that the GN-PPA approach is in very good agreement with full-frequency methods, further improved by the excellent agreement of the MPA scheme, especially when comparing with plane-wave codes.



1. INTRODUCTION

First-principles computational methods have become well-established in the realm of electronic structure and materials science, serving as invaluable tools that offer critical insight and steer experimental investigations, as well as playing a pivotal role in materials discovery and characterization.^{1–6} Once a theoretical framework is defined, such as density functional theory (DFT) or the GW approximation within many-body perturbation theory (MBPT), different numerical implementations should ideally yield equivalent results. In practice, however, algorithmic choices, numerical parameters, and approximations can introduce discrepancies. For this reason, verification (ensuring consistency across codes) and validation (assessing accuracy against experiments) have become key aspects of methodological development. Together, verification and validation (V&V) are essential to establish the reliability, reproducibility, and predictive power of computational techniques.

During the past decade, several systematic comparisons have demonstrated that state-of-the-art DFT codes yield highly consistent results for the ground-state properties of solid-state systems.^{7–9} On the other hand, excited-state properties, such as quasi-particle energy levels of materials calculated with GW in MBPT, have been primarily compared with experiments,^{10–16} while code-to-code V&V studies have emerged more recently.^{17–20} For example, the work by Rangel et al.¹⁷ compares G_0W_0 quasiparticles (QPs) of selected solids (Si, Au, TiO_2 , and ZnO), obtained with three different plane-wave (PW) codes: YAMBO,^{21,22} BERKELEYGW²³ (BGW), and ABINIT²⁴. The comparison shows that, when equivalent convergence parameters, numerical schemes, and pseudopotentials are

employed, QP energies agree within 100 meV. Their systematic analysis traced previously reported discrepancies to differences in the treatment of the Coulomb divergence and frequency-integration schemes. By making these aspects consistent across codes, they achieved consistent results even for challenging systems such as rutile TiO_2 and ZnO.^{25–27}

Concerning molecules, early studies have focused on benchmarking different flavors of the GW approximation, including different DFT starting points, with respect to small data sets of experimental results and accurate quantum chemistry calculations based, e.g., on coupled cluster methods.^{28–33} A significant community effort was made to V&V different GW implementations, using the larger GW100 data set, which includes 100 different molecules.^{34–40} This effort is primarily focused on the single-shot G_0W_0 calculation of the vertical ionization potential (IP) and electron affinity (EA),^{34,36,37} although results from self-consistent GW schemes have also been reported.³⁵ The chosen data set comprises only closed-shell molecules, thereby avoiding the well-known issues presented by open-shell systems.⁴¹

In terms of both single eigenvalues and statistical deviations (e.g., mean absolute errors), the reported differences between codes using PW and localized basis sets are of the order of 200 meV. The discrepancies are mainly due to (i) the size of the

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Table 1. Basis set and Approximations Used in Previous GW100 Benchmarks and in the Present Work^a

code	basis set	PP	empty states	χ -basis	Σ_c -integration	QP solutions	refs.
TM ⁵⁴	def2-QZVP	AE	all	LO	FA ⁵⁵	largest weight	34
FHI-aims ⁵⁶	def2-QZVP*	AE	all	LO	AC ^{56,57}	iterative	34
BGW ²³	PW	TM NC ⁵⁸	truncated ⁵⁹	PW	HL-PPA ⁴⁷ /FF ⁶⁰	lin./graphical	34
VASP ⁶¹	PAW*	PAW	truncated	PW*	AC ⁶²	linearized	36
WEST ⁶³	PW	ONCV	none ^{63,64}	PDEP* ⁶⁵	CD ⁶⁶	secant/lin.	37
YAMBO ^{21,22}	PW	ONCV	truncated ⁶⁷	PW	GN-PPA ⁴⁸ /MPA ⁴⁹	linearized	this work

^aThe codes TM and FHI-aims make use of the resolution-of-identity (RI) technique to compute the four-center Coulomb integrals.³⁴ In the column basis set, * indicates results obtained after extrapolation. In the column PP, we list the approach used for core and valence electrons: all electron (AE), Troullier–Martins norm conserving pseudopotentials (TM NC), projector augmented waves (PAW), and optimized norm-conserving Vanderbilt pseudopotentials (ONCV). In the column Σ_c -integration, the acronyms refer to the frequency integration methods used for the self-energy: fully-analytic (FA), analytic continuation (AC), contour deformation (CD), Godby–Needs plasmon-pole model (GN-PPA), and multipole approximation (MPA).

adopted basis sets and to (ii) the frequency-dependent treatment of the screened Coulomb potential W and of the self-energy Σ . Besides the basis sets, the treatment of core and valence electrons can also introduce some differences. Approaches based on pseudopotentials (PPs) reduce the number of explicitly treated electrons with respect to all-electron approaches. Nevertheless, different PP schemes (e.g., ultrasoft vs norm-conserving), the use of nonlinear core corrections (NLCC), or the inclusion of semicore states in valence can significantly affect GW QP energies.^{42–44} Concerning the frequency dependence, the performance of the different plasmon-pole approximation (PPA) models has been addressed in the literature for a few bulk materials.^{26,45,46} For the GW100 data set, however, only the Hybertsen and Louie (HL) model⁴⁷ has been benchmarked against full-frequency (FF) results, while a benchmark for the largely adopted Godby and Needs (GN) model⁴⁸ is still lacking.

In this work, we provide the results for IP and EA of all 100 molecules of the GW100 set as computed within the PW code YAMBO, using two different frequency dependence treatments: the Godby–Needs PPA (GN-PPA)⁴⁸ model and the recently developed multipole approximation (MPA).^{49,50} Our results show that GN-PPA, at variance with Hybertsen–Louie generalized PPA (HL-PPA), is in good agreement with GW FF data obtained from other PW codes, with an average deviation of only 190 meV. MPA further reduces the deviation to 143 meV, which is comparable to the relative deviation among other FF approaches.

This paper is organized as follows. In Section 2, we provide details about the GW100 set, the adopted GW methodology and implementation, the numerical parameters used in the simulations, and the techniques employed to automate the calculations. In Section 3, we present and discuss the IP and EA results for the GW100 data set, obtained with both GN-PPA and MPA. Our conclusions are drawn in Section 4.

2. COMPUTATIONAL APPROACH

2.1. The GW100 Data set

The GW100 set comprises a collection of 100 closed-shell molecules,³⁴ selected to include a wide range of IP energies (4–25 eV) as well as a variety of chemical bond arrangements. The set includes carbon-based covalently bonded compounds like C₂H₂, C₂H₄, C₂H₆, as well as ionically bonded molecules such as the alkaline metal halide LiF. There are also molecules with metal atoms, such as Ag₂, Li₂, K₂, or small metal clusters (Na₄, Na₆), and common molecules such as water and carbon mono- and dioxide.

All molecular structures considered in this study have been taken from the official GW100 repository,⁴⁰ derived either from experiments or optimized using DFT-PBE with the def2-QZVP basis set.³⁴ For the cases of CH₂CHBr and C₆H₅OH, for which there is more than one reported structure, we have considered the most recent ones, as described in ref 36. Following previous GW100 benchmarks, the present calculations employ a single-shot G_0W_0 scheme based on KS-DFT eigenvalues and eigenvectors obtained with the PBE functional,⁵¹ hereafter denoted as G_0W_0 @PBE. This choice allows us to bypass the well-known dependence of G_0W_0 results on the starting point,^{43,52,53} and to focus the discussion on the different GW implementations and underlying numerical approximations.

2.2. DFT Setup

Even with the same exchange–correlation functional and molecular geometries, the outcome of G_0W_0 calculations depends on several technical aspects of the underlying DFT setup. These include the adopted basis set, the treatment of core electrons (e.g., via PPs), the definition of the simulation cell and boundary conditions, and the treatment of the long-range Coulomb interaction between periodic replicas. In Table 1, we summarize the methodologies and approximations implemented in the codes benchmarked with the GW100 data set,^{34,37} as well as those implemented in YAMBO and used in the present work (see details below).

The GW100 data were originally obtained using a variety of basis sets, including projector augmented waves (PAW),^{68,69} linear augmented plane waves (LAPW),⁷⁰ linearized muffin-tin orbitals (LMTO),⁷¹ and local orbital (LO) basis sets. For a detailed comparison of IPs and EAs obtained using DFT-PBE with different basis sets, we refer the reader to ref 37. In this work, the Kohn–Sham (KS) wave functions and charge densities are expanded in plane waves (PWs) under periodic boundary conditions (PBCs), as implemented in the QUANTUM ESPRESSO simulation package.^{72,73}

The treatment of core and valence electrons varies across different codes. Approaches based on PPs reduce the number of explicitly treated electrons, simplifying the calculations while retaining accuracy for valence states. Nevertheless, different PP schemes (ultrasoft vs norm-conserving), the use of NLCC, or the inclusion of semicore states in valence can significantly affect GW QP energies.^{42–44} Here, we employ optimized norm-conserving Vanderbilt PPs from the SG15 library,^{74,75} without NLCC.⁷⁶ Details of the electronic configurations, including the semicore states treated as valence, are provided in the Supporting Information.⁷⁵

Each molecule is placed in a supercell with sufficient vacuum to suppress spurious interactions between replicas. To reduce the required vacuum spacing and the associated computational cost, the Martyna–Tuckerman method⁷⁷ is adopted to correct the error introduced by the periodic replica of the system. Choosing the lattice geometry in order to minimize the supercell volume is also essential for both numerical accuracy and computational efficiency. Previous GW100 studies³⁷ used simple cubic (SC) supercells with lattice parameters up to $a = 25$ Å. In the present work, we adopt a face-centered cubic (FCC) geometry with lattice parameter $a = 13$ Å, which reduces the supercell volume by a factor of $\sqrt{2}$ relative to the corresponding SC configuration. Tests performed on two molecular systems, with 10 and 56 atoms (the latter being the largest molecule in the GW100 set), show a residual IP/EA error of ~ 25 and ~ 180 meV, respectively (see Sec. II of the Supporting Information⁷⁸).

At the GW level, a real-space truncation of the Coulomb interaction is applied to both the screening and self-energy terms.⁷⁹ This spherical Coulomb cutoff, with a diameter of approximately 12 Å (i.e., 1 Å smaller than the lattice parameter), analytically removes the divergence of the long-range Coulomb interaction $v(q \rightarrow 0)$ and ensures rapid convergence with respect to the size of the supercell.

2.3. The QP Equation

Within the MBPT framework, QP energies are usually obtained perturbatively from DFT, as the solutions of the QP equation

$$\varepsilon_{nk}^{\text{QP}} = \varepsilon_{nk} + \langle nk | \Sigma(\varepsilon_{nk}^{\text{QP}}) - v_{xc} | nk \rangle \quad (1)$$

where Σ is the electron self-energy, $\{\varepsilon_{nk}, |nk\rangle\}$ are the reference Kohn–Sham (KS) eigenvalues and wave functions, and v_{xc} is the corresponding exchange–correlation potential. eq 1 originates from a nonlinear eigenvalue problem,^{80,81} where the self-energy has been assumed to be diagonal in the reference basis. This reduces the problem to a scalar but still nonlinear equation for each state, possibly leading to multiple solutions. To simplify the problem, one can linearize the self-energy as the first-order term of the Taylor expansion around ε_{nk} , yielding

$$\varepsilon_{nk}^{\text{QP}} = \varepsilon_{nk} + Z_{nk} \langle nk | \Sigma(\varepsilon_{nk}) - v_{xc} | nk \rangle \quad (2)$$

with the renormalization factor

$$Z_{nk} = \left[1 - \left\langle nk \left| \frac{\partial \Sigma(\omega)}{\partial \omega} \right| nk \right\rangle \right]^{-1}_{\omega=\varepsilon_{nk}} \quad (3)$$

eqs 1 or 2 can be solved numerically by using iterative root-finding methods. Multiple solutions have been reported, for instance, for some molecules in the GW100 benchmark.³⁴ When both the real and imaginary parts of Σ are considered, the different solutions correspond to the poles of the interacting Green's function G and can be assigned one to the main QP excitation and the others to satellites. The solutions can also be determined analytically using alternative approaches based on rational representations of Σ , through, for example, the algorithmic inversion method for sum-overpoles (AIM-SOP) reported in refs 82–85 or the MPA- Σ approach reported in ref 86.

In this work, we adopt the linearized form of eq 2, solved using Newton's method as implemented in the YAMBO code.

This approach assumes that Σ varies smoothly around the QP energy and therefore neglects secondary (satellite) solutions.

2.4. GW Self-Energy Evaluation

The state-of-the-art method for evaluating the self-energy Σ is the one-shot G_0W_0 approximation. In this approximation, Σ is given by a frequency convolution of the noninteracting KS Green's function G_0 and the screened Coulomb potential W_0

$$\Sigma(\omega) = \frac{i}{2\pi} \int d\omega' e^{i\omega'\theta^+} G_0(\omega + \omega') W_0(\omega') \quad (4)$$

In the expression above, W_0 is usually evaluated at the random-phase approximation (RPA) level and is typically split into two terms: the bare Coulomb interaction v , which is static, and the correlation part $W^c \equiv W_0 - v$, which contains all the frequency dependence. This, in turn, leads to a decomposition of the self-energy into exchange and correlation terms, $\Sigma(\omega) = \Sigma^x + \Sigma^c(\omega)$.

The dynamical nature of the screened interaction W distinguishes the GW approximation from static theories like Hartree–Fock. A number of different strategies have been proposed to evaluate the frequency integral in eq 4, ranging from full-analytic⁵⁵ (FA) to full-frequency real-axis integration⁶⁰ (FF), analytic continuation⁸⁷ (AC), contour deformation (CD),⁶⁶ and multipole approaches (MPA).^{49,50} The earliest and simplest approach is the so-called PPA.^{47,48} Within PPA, the frequency dependence of W^c is simplified as a symmetric single pole function for each matrix element $\mathbf{GG}'$ and transferred momentum $\mathbf{q}$

$$W_{\mathbf{GG}'}^{c\text{PPA}}(\mathbf{q}, \omega) = \frac{2R_{\mathbf{qGG}'}\Omega_{\mathbf{qGG}'}}{\omega^2 - \Omega_{\mathbf{qGG}'}} \quad (5)$$

Among the different flavors of PPA,²⁶ in this work, we adopt the one proposed by Godby and Needs (GN-PPA),⁴⁸ which interpolates W^c from the values computed at two points, $\omega = 0$ and $\omega = iE_{\text{PPA}}$. Here, we set $E_{\text{PPA}} = 30$ eV, after having tested a subset of molecules by varying the sampling energy between 25 and 30 eV, without significant changes in the results. The HL-PPA⁴⁷ model follows a different approach and, in addition to evaluating the screening at zero frequency, imposes the fulfillment of Johnson's frequency sum rule (f -sum rule).⁸⁸ Although such a property can be mapped with the GN-PPA model, by increasing E_{PPA} to infinity,⁴⁹ the results of the standard GN-PPA and HL-PPA may differ significantly. For a detailed comparison of different PPA models, see, e.g., refs 26, 45, and 46.

Besides GN-PPA, we also adopt the MPA method,^{49,50} which generalizes eq 5 to multiple complex poles

$$W_{\mathbf{GG}'}^{c\text{MPA}}(\mathbf{q}, \omega) = \sum_p \frac{2R_{p\mathbf{qGG}'}\Omega_{p\mathbf{qGG}'}}{\omega^2 - \Omega_{p\mathbf{qGG}'}} \quad (6)$$

where n_p is typically around 10. Making use of the Lehmann representation for G_0 in a Bloch PW basis set, and the MPA model for W^c , the diagonal matrix elements of the correlation self-energy are integrated analytically and can be expressed as

$$\Sigma_{nk}^{\text{cMPA}}(\omega) = \sum_m \sum_{\mathbf{G}\mathbf{G}'} \sum_p^{n_p} \int \frac{d\mathbf{q}}{(2\pi)^3} S_{p\mathbf{G}\mathbf{G}'}^{nm}(\mathbf{k}, \mathbf{q}) \left[\frac{f_{m\mathbf{k}-\mathbf{q}}^{\text{KS}}}{\omega - \epsilon_{m\mathbf{k}-\mathbf{q}}^{\text{KS}} + \Omega_{p\mathbf{q}\mathbf{G}\mathbf{G}'}} + \frac{1 - f_{m\mathbf{k}-\mathbf{q}}^{\text{KS}}}{\omega - \epsilon_{m\mathbf{k}-\mathbf{q}}^{\text{KS}} - \Omega_{p\mathbf{q}\mathbf{G}\mathbf{G}'}} \right] \quad (7)$$

where we have defined

$$\begin{aligned} S_{p\mathbf{G}\mathbf{G}'}^{nm}(\mathbf{k}, \mathbf{q}) &= -2\rho_{nm}^{\text{KS}}(\mathbf{k}, \mathbf{q}, \mathbf{G}) R_{p\mathbf{q}\mathbf{G}\mathbf{G}'} \rho_{nm}^{\text{KS}*}(\mathbf{k}, \mathbf{q}, \mathbf{G}') \\ \rho_{nm}^{\text{KS}}(\mathbf{k}, \mathbf{q}, \mathbf{G}) &= \langle n | e^{i(\mathbf{q}+\mathbf{G})\cdot\mathbf{r}} | m \mathbf{k} - \mathbf{q} \rangle \end{aligned} \quad (8)$$

For each momentum transfer $\mathbf{q}$ and each $\mathbf{G}\mathbf{G}'$ matrix element, poles and residues are obtained through a nonlinear interpolation with frequency points z_i sampled in the complex plane

$$\sum_{p=1}^{n_p} \frac{2R_{p\mathbf{q}\mathbf{G}\mathbf{G}'}\Omega_{p\mathbf{q}\mathbf{G}\mathbf{G}'}}{z_i^2 - \Omega_{p\mathbf{q}\mathbf{G}\mathbf{G}'}} = W_{\mathbf{G}\mathbf{G}'}^{\text{c}}(\mathbf{q}, z_i) \quad (9)$$

where $i = 1, \dots, 2n_p$ and the set of complex frequencies z_i are conveniently selected according to the double parallel sampling defined in refs 49 and 50. Such efficient sampling allows the MPA method to achieve FF accuracy with a small number of poles or sampling frequencies, typically 1–2 orders of magnitude fewer than the number of frequency points required to converge FF-RA.

2.5. Convergences and Workflow

Accurate GW QP energies require convergence with respect to several computational parameters. In YAMBO, the most relevant parameters are the number of unoccupied states (N_b) that enter the sum over states and the kinetic-energy cutoff (G_{cut}) defining the number of reciprocal lattice vectors used in the evaluation of the self-energy in eq 7 and the screening matrix in eq 6. Within PPA, convergence with respect to N_b can be accelerated by using the Bruneval–Gonze (BG) terminator,⁶⁷ which replaces the contribution of high-energy states with an effective pole at an energy E_{GT} above the highest explicitly included band. For isolated systems, where unoccupied levels are densely spaced, effective E_{GT} values are typically smaller than for bulk systems. Tests on LiF for different E_{GT} values (Figure 1, left panel) show that 0.25 Ha yields the largest convergence acceleration, and this value is adopted for all molecules.

As discussed in the literature,^{17,26,89} the parameters N_b and G_{cut} are interdependent, as shown in the right panel of Figure 1, where the curves indicate that the QP energies have not yet reached convergence within the accessible range of parameters. In this work, the convergence with respect to these two parameters is assessed simultaneously via extrapolation. Specifically, QP corrections are computed on a two-dimensional grid, with N_b ranging from 1000 to 11000 (3000–13000 for MPA) and G_{cut} from 24 to 36 Ry. When possible, an additional point at $G_{\text{cut}} = 38$ Ry is included. The extrapolation to infinite cutoff values is obtained by fitting the QP corrections to

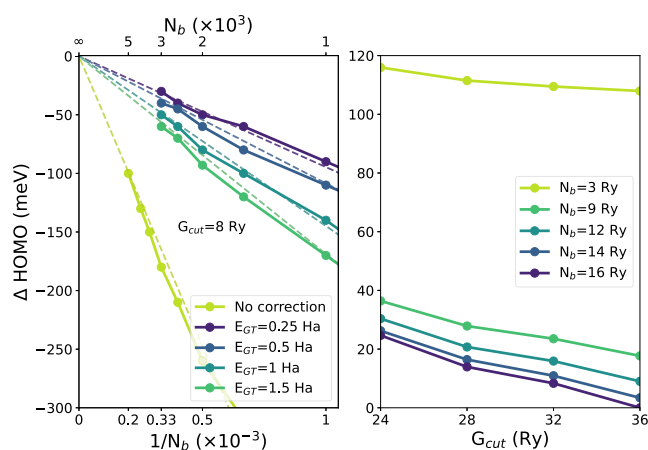


Figure 1. Convergence tests for the LiF molecule. Left panel: Effect of E_{GT} (see text) on the convergence acceleration of the HOMO with respect to empty states summation [the $N_b \rightarrow \infty$ (extrapolated) value for the noncorrected case is set to zero in the plot]. Right panel: HOMO energy versus the PW cutoff G_{cut} in the screening matrix, for different numbers of empty states, N_b . The result obtained with $N_b = 16$ Ry, $G_{\text{cut}} = 36$ Ry is set to zero. The values $N_b = 3, 9, 12, 14$, and 16 Ry correspond to 1000, 5000, 7000, 9000, and 11000 bands.

$$f(N_b, G_{\text{cut}}) = \left(\frac{A}{N_b^\alpha} + b \right) \left(\frac{C}{G_{\text{cut}}^\beta} + d \right) \quad (10)$$

where α and β belong to $\{1, 2, 3\}$ and, for each molecule, the values that minimize the root-mean-square error are chosen. Similarly to the strategy adopted in refs 90–94, the fully converged QP energy reads

$$E_{\text{extra}}^{\text{QP}} = f(N_b \rightarrow \infty, G_{\text{cut}} \rightarrow \infty) = bd \quad (11)$$

All calculations were performed with a fully automated high-throughput (HT) scheme using the `aidda-yambo` plugin,^{89,95} which is part of the AiiDA workflow ecosystem^{96,97} and implements automated workflows and convergence protocols for MBPT YAMBO simulations. We note that related efforts have been undertaken by the community in recent years.^{98–100} For each of the 100 molecules, convergence workflows systematically explored the (N_b, G_{cut}) space, resulting in over 9000 QP evaluations across both PPA and MPA schemes. All input and output data are stored in the AiiDA database, ensuring full reproducibility. This work also serves as a large-scale validation of the `aidda-yambo` workflows. Computational performance statistics are reported in the Supporting Information.⁷⁸

3. RESULTS

To benchmark the accuracy of the GN–PPA and MPA implementations of the G_0W_0 method in YAMBO, as compared to other codes, we computed the IP and EA energies of the 100 molecules in the GW100 data set. These quantities are evaluated as

$$\begin{aligned} \text{IP} &= E^{\text{vac}} - E^{\text{HOMO}} \\ \text{EA} &= E^{\text{vac}} - E^{\text{LUMO}} \end{aligned} \quad (12)$$

where E^{vac} denotes the electrostatic potential in the vacuum region. E^{vac} is evaluated within the Martyna–Tuckerman method,⁷⁷ that automatically sets it to zero by shifting the energy scale (so that $\text{IP} = -E^{\text{HOMO}}$ and $\text{EA} = -E^{\text{LUMO}}$).

Table 2. Mean Error (ME), Mean Absolute Error (MAE), Mean Absolute Relative Error (MARE), and Standard deviation (σ) of the IPs Computed with YAMBO-GN-PPA and YAMBO-MPA with Respect to Reference GW100 Results^{34,36,37,101,103} Obtained with Different Codes and Approximations. We Also Included Experimental (EXP) Data^{34,102}

Code	ME (meV)		MAE (meV)		MARE (%)		σ (meV)	
	GN-PPA	MPA	GN-PPA	MPA	GN-PPA	MPA	GN-PPA	MPA
YAMBO _{GN-PPA}	-	77	-	168	-	1.8	-	114
WEST _{lin} ^{extra}	-117	-31	176	108	1.8	1.3	159	121
WEST _{sol} ^{extra}	9	96	159	167	1.8	1.9	165	123
VASP	-44	32	234	155	2.4	1.8	216	199
AIMS _{pade}	225	302	299	334	3.4	3.6	320	260
AIMS _{extra}	22	119	214	189	2.4	2.1	200	185
BGW _{HL-PPA}	-400	-323	511	400	4.7	3.8	336	321
BGW _{FF}	203	387	262	399	2.8	3.9	382	388
KIPZ	-375	-298	507	403	4.8	3.9	398	358
CCSD	-338	-259	447	350	4.1	3.3	374	310
EXP	-255	-175	534	454	5.0	4.3	401	387

The statistical accuracy of the computed IP energies is characterized by the deviations with respect to the results obtained with the codes we are comparing with. Using standard statistical measures, we present in Table 2 of Sec. 3.1 the values for the mean error (ME), the mean absolute error (MAE), the mean absolute relative error (MARE), and the standard deviation (σ). The IP values obtained for each molecule are reported in Table 3, together with reference GW results³⁴ from other codes, CCSD data,¹⁰¹ and experimental values from the NIST database.¹⁰² EA results are provided in Table 4 and discussed in Sec. 3.2. While the discussion below focuses primarily on IPs, which provide the most direct and comprehensive benchmark across the different implementations, the same conclusions apply to the EAs, analyzed in Sec. 3.2.

3.1. Ionization Potentials

We first examined the computed KS (PBE) HOMO energies, which serve as starting point for the subsequent GW QP corrections. These results are reported in Sec. II of the Supporting Information.⁷⁸ The agreement with previous QUANTUM ESPRESSO calculations³⁷ is very good; the remaining small discrepancies arise primarily from the different supercell shape adopted in this work (see Sec. 2.2). Compared with results obtained using localized basis set codes, such as FHI-AIMS or TM, the differences reflect both the different basis representations and the use of PPs, which influence the high-energy part of the spectrum. Interestingly, the present PBE HOMO values agree well (within 40 meV on average) with def2-QZVP GTO results (see ref 34 and the related Supporting Information), a few meV better than the corresponding extrapolation to the complete basis-set limit.

Moving to the discussion of GW QP-corrections for the HOMO and LUMO states, the situation becomes more complex. The GW QP corrections for HOMO and LUMO levels depend on several numerical parameters. As detailed in Sec. 2.5, we employed an extrapolation scheme with respect to the number of included empty states and the kinetic energy cutoff adopted to represent the response function. The accuracy of this procedure is analyzed in the Supporting Information.⁷⁸ Overall, calculations done with the tightest convergence parameters are close to full convergence. This is particularly true for the IP computed within PPA, for which the Bruneval–Gonze terminator⁶⁷ (available only for PPA) proved

essential for accelerating convergence due to the dense manifolds of empty states in finite systems.

Next, we analyze the G_0W_0 IPs obtained with the GN-PPA and MPA schemes of YAMBO. Table 2 summarizes the statistical indicators (ME, MAE, MARE, and σ) of YAMBO results with respect to other community codes. Beyond GW codes, we also include results from Koopmans spectral functionals (KIPZ)¹⁰³ and quantum chemistry (CCSD)¹⁰¹ methods, in such a way as to also provide a brief comparison with different theoretical level/approximations. The GN-PPA data exhibit slightly larger deviations, as expected from the simpler frequency treatment of the PPA, while MPA results tend to show smaller σ , overall indicating improved accuracy. When compared with other PW or mixed-basis implementations, such as WEST, VASP, and FHI-AIMS, both approaches attain comparable accuracy.

The individual IPs of all molecules are reported in Table 3, together with previously published GW100 results,³⁴ CCSD,¹⁰¹ and experimental data from the NIST database.¹⁰² The best agreement for the YAMBO GN-PPA results, as measured by the MAE, is obtained with WEST_{sol} and WEST_{lin}, followed by VASP and FHI-AIMS in the complete-basis-set limit. This level of consistency is remarkable, given the methodological differences among these codes, including the choice of basis sets (plane waves, augmented waves, or localized orbitals), the use of PPs, the different ways to solve the QP equation (full or linearized), and the different frequency-integration schemes. Specifically, the frequency integration is performed using AC (VASP), CD (WEST), and MPA (YAMBO). Overall, these differences have only a minor effect on the computed QP energies, confirming the robustness of modern G_0W_0 implementations.

In contrast, larger discrepancies arise when comparing the YAMBO GN-PPA results with those obtained using the HL-PPA model of BGW. Despite both adopting the linearized QP equation, the two models differ in their parametrization of the dielectric response: the HL and GN versions of PPA can yield significantly different QP corrections depending on the system and the treatment of the so-called unfulfilled modes.^{17,26,45,46} A semianalytical comparison between the two has shown that the HL-PPA may systematically overestimate QP shifts with respect to GN-PPA.⁴⁹ Conversely, the FF BGW data show a noticeably better agreement with YAMBO, although the limited number of reported values (19 molecules) prevents a comprehensive comparison.

Table 3. Quasiparticle Ionization Potentials (IP) of the Molecules in the GW100 Data Set as Obtained Within This Work: Y_{PPA} and Y_{MPA} Refer to Y_{AMBO} Calculations Performed at the PPA and MPA Levels, Respectively; for Completeness, PBE IPs are Also Reported⁴⁴

index	formula	Y_{MPA}	Y_{PPA}	$\text{WEST}_{\text{lin}}^{\text{extra}}$	$\text{WEST}_{\text{sol}}^{\text{extra}}$	VASP	$\text{AIMS}_{\text{pade}}$	$\text{AIMS}_{\text{extra}}$	BGW_{HL}	BGW_{FF}	CCSD	EXP
1	He	23.72	23.33	23.65	23.42	23.62	23.48	23.49	24.1	-	24.51	24.59
2	Ne	20.14	20.06	20.52	20.33	20.36	20.38	20.33	21.35	-	21.32	21.56
3	Ar	15.56	15.34	15.5	15.37	15.42	15.13	15.28	15.94	-	15.54	15.76
4	Kr	13.9	13.79	13.87	13.76	14.03	13.57	13.89	14.0	-	13.94	14.0
5	Xe	13.34	13.52	13.38	13.22	12.22	12.02	-	12.08	-	-	12.13
6	H ₂	16.1	16.05	16.03	15.84	16.06	15.82	15.85	16.23	-	16.4	15.43
7	Li ₂	5.07	5.24	5.19	5.04	5.32	4.99	5.05	5.43	-	5.27	4.73
8	Na ₂	4.89	5.11	5.07	4.98	5.06	4.83	4.88	5.03	-	4.95	4.89
9	Na ₄	3.85	4.09	4.28	4.24	4.23	4.1	4.14	4.34	-	4.22	4.27
10	Na ₆	3.78	4.04	4.42	4.37	4.4	4.24	4.34	4.47	-	4.35	4.12
11	K ₂	3.96	4.3	4.21	4.14	4.24	3.98	4.08	4.02	-	4.06	4.06
12	Rb ₂	3.79	4.14	4.08	4.01	4.14	3.8	-	3.92	-	3.92	3.9
13	N ₂	15.16	14.84	15.08	14.94	15.06	14.89	15.05	15.43	14.72	15.57	15.58
14	P ₂	10.53	10.57	10.48	10.43	10.4	10.21	10.38	10.66	-	10.47	10.62
15	As ₂	9.62	9.7	9.58	9.55	9.62	9.47	9.67	9.67	-	9.78	10.0
16	F ₂	14.89	14.53	15.16	15.0	15.08	14.96	15.1	15.59	14.73	15.71	15.7
17	Cl ₂	11.61	11.45	11.5	11.41	11.4	11.1	11.31	11.85	-	11.41	11.49
18	Br ₂	10.62	10.5	10.52	10.44	10.65	10.22	10.56	10.64	-	10.54	10.51
19	I ₂	10.41	10.61	10.56	10.41	9.59	9.28	-	9.58	-	9.51	9.36
20	CH ₄	14.2	14.03	14.1	13.99	14.14	13.93	14.0	14.28	13.8	14.37	13.6
21	C ₂ H ₆	12.63	12.44	12.53	12.44	12.58	12.36	12.46	12.63	12.22	13.04	11.99
22	C ₃ H ₈	12.04	11.85	11.92	11.84	11.98	11.79	11.89	12.05	-	12.05	11.51
23	C ₄ H ₁₀	11.7	11.52	11.48	11.41	11.69	11.49	11.59	11.73	-	11.57	11.09
24	C ₂ H ₄	10.52	10.48	10.46	10.39	10.5	10.32	10.4	10.68	10.3	10.67	10.68
25	C ₂ H ₂	11.28	11.18	11.18	11.09	11.24	11.02	11.09	11.35	10.97	11.42	11.49
26	C ₄	11.11	10.98	10.97	10.9	10.97	10.78	10.91	11.49	-	11.26	12.54
27	C ₃ H ₆	10.78	10.63	10.73	10.67	10.78	10.56	10.65	10.93	-	10.86	10.54
28	C ₆ H ₆	9.17	9.1	9.13	9.08	9.16	8.99	9.1	9.21	-	9.29	9.23
29	C ₈ H ₈	8.21	8.09	8.2	8.16	8.24	8.06	8.18	8.47	-	8.35	8.43
30	C ₅ H ₆	8.56	8.48	8.49	8.44	8.51	8.35	8.45	8.77	-	8.68	8.53
31	C ₂ H ₃ F	10.4	10.28	10.36	10.29	10.36	10.2	10.32	10.8	10.14	10.55	10.63
32	C ₂ H ₃ Cl	10.09	9.97	10.0	9.94	10.0	9.76	9.9	10.32	-	10.09	10.2
33	C ₂ H ₃ Br	9.23	9.17	9.71	9.64	9.83	8.99	9.14	9.42	-	9.27	9.9
34	C ₂ H ₃ I	9.85	9.9	9.94	9.81	9.36	9.04	-	9.48	-	9.33	9.35
35	CF ₄	15.46	15.13	15.65	15.51	15.53	15.36	15.6	15.96	-	16.3	16.2
36	CCl ₄	11.49	11.35	11.41	11.29	11.31	10.98	11.21	11.77	-	11.56	11.69
37	CBr ₄	10.25	10.15	10.22	10.11	10.38	9.9	10.22	10.4	-	10.46	10.54
38	Cl ₄	9.82	10.02	-	-	9.23	8.82	-	9.23	-	9.27	9.1
39	SiH ₄	12.6	12.6	12.55	12.42	12.53	12.31	12.4	12.77	-	12.8	12.3
40	GeH ₄	12.47	12.53	12.44	12.32	12.24	12.02	12.12	12.28	-	12.5	11.34
41	Si ₂ H ₆	10.6	10.6	10.58	10.52	10.52	10.31	10.41	10.8	-	10.64	10.53
42	Si ₃ H ₁₂	8.69	8.7	9.25	9.19	9.19	8.94	9.05	9.45	-	9.27	9.36
43	LiH	7.31	7.55	7.2	6.62	7.2	6.54	6.58	7.85	6.67	7.96	7.9
44	KH	5.55	5.99	5.39	4.97	5.37	4.86	4.99	5.76	-	6.13	8.0
45	BH ₃	13.16	13.07	13.08	12.95	13.09	12.87	12.96	13.28	-	13.28	12.03
46	B ₂ H ₆	12.1	12.0	12.03	11.92	12.04	11.84	11.93	12.17	-	12.26	11.9
47	NH ₃	10.46	10.3	10.4	10.18	10.44	10.32	10.39	10.93	-	10.81	10.82
48	HN ₃	10.63	10.35	10.54	10.48	10.56	10.4	10.55	10.96	-	10.68	10.72
49	PH ₃	10.57	10.61	10.51	10.43	10.45	10.27	10.36	10.79	-	10.52	10.59
50	AsH ₃	10.43	10.45	10.4	10.33	10.36	10.12	10.21	10.45	-	10.4	10.58
51	SH ₂	10.43	10.37	10.36	10.23	10.3	10.03	10.13	10.64	-	10.31	10.5
52	FH	15.34	15.03	15.47	15.23	15.38	15.3	15.37	16.24	-	16.03	16.12
53	ClH	12.68	12.58	12.6	12.48	12.51	12.25	12.36	12.97	-	12.59	12.79
54	LiF	10.36	10.32	10.54	10.11	10.45	9.95	10.27	11.84	-	11.32	11.3
55	F ₂ Mg	12.65	12.59	12.84	12.46	12.77	12.32	12.5	13.73	12.44	13.71	13.3
56	TiF ₄	14.12	13.95	14.31	-	14.22	13.89	14.07	14.88	-	15.48	13.3
57	AlF ₃	14.45	14.22	14.63	14.4	14.53	14.25	14.48	15.11	-	15.46	15.45
58	BF	10.69	11.08	10.71	10.56	10.67	10.56	10.73	11.49	-	11.09	11.0
59	SF ₄	12.43	12.4	12.41	12.32	12.29	12.12	12.38	12.79	-	12.59	11.69

Table 3. continued

index	formula	Y_{MPA}	Y_{PPA}	$\text{WEST}_{\text{lin}}^{\text{extra}}$	$\text{WEST}_{\text{sol}}^{\text{extra}}$	VASP	$\text{AIMS}^{\text{spade}}$	$\text{AIMS}^{\text{extra}}$	BGW_{HL}	BGW_{FF}	CCSD	EXP
60	BrK	8.04	8.1	7.96	-	8.04	7.3	7.57	7.99	-	8.13	8.82
61	GaCl	10.35	10.29	10.27	10.19	9.99	9.55	9.74	10.24	-	9.77	10.07
62	NaCl	8.98	9.02	-	-	8.76	8.1	8.43	9.6	-	9.03	9.8
63	MgCl ₂	11.46	11.45	11.47	11.25	11.41	10.99	11.2	11.98	-	11.66	11.8
64	AlI ₃	10.12	10.3	10.59	10.31	9.69	9.32	-	9.67	-	9.82	9.66
65	BN	11.6	11.34	-	-	10.61	11.03	11.15	12.19	9.68	11.89	11.5
66	NCH	13.5	13.3	13.41	13.22	13.43	13.21	13.32	13.87	-	13.87	13.61
67	PN	11.51	11.28	11.43	11.26	11.41	11.14	11.29	12.13	-	11.74	11.88
68	H ₂ NNH ₂	9.48	9.21	9.42	9.27	9.45	9.28	9.37	9.78	9.1	9.72	8.98
69	H ₂ CO	10.57	10.24	10.56	10.41	10.57	10.33	10.46	11.02	-	10.84	10.88
70	CH ₄ O	10.64	10.39	10.74	10.6	10.72	10.56	10.67	11.14	-	11.04	10.96
71	C ₂ H ₆ O	10.36	10.0	10.36	10.21	10.33	10.16	10.27	10.57	-	10.68	10.64
72	C ₂ H ₄ O	9.84	9.46	9.81	9.61	9.8	9.55	9.66	10.16	9.43	10.21	10.24
73	C ₂ H ₁₀ O	9.46	9.13	9.52	9.4	9.52	9.32	9.42	9.7	-	9.82	9.61
74	CH ₂ O ₂	10.87	10.72	11.01	10.82	10.98	10.73	10.87	11.39	-	11.42	11.5
75	HOOH	11.18	10.79	11.16	11.0	11.12	10.99	11.1	11.58	10.82	11.59	11.7
76	H ₂ O	12.08	11.81	12.09	11.87	12.05	11.97	12.05	12.75	11.68	12.56	12.62
77	CO ₂	13.45	13.1	13.46	13.37	13.44	13.25	13.46	13.81	13.17	13.71	13.77
78	CS ₂	10.15	10.09	10.1	10.05	10.01	9.75	9.95	10.37	-	9.98	10.09
79	OCS	11.26	11.15	-	-	11.13	10.91	11.11	11.49	11.02	11.17	11.19
80	OCS _e	10.49	10.45	10.43	10.37	10.5	10.2	10.43	10.55	-	10.78	10.37
81	CO	13.81	13.77	13.79	13.66	13.76	13.57	13.71	14.33	-	14.21	14.01
82	O ₃	12.18	11.84	-	-	12.07	11.39	11.49	13.05	12.0	12.55	12.73
83	SO ₂	12.08	11.81	12.08	11.96	12.04	11.82	12.06	12.55	-	13.49	12.5
84	BeO	9.57	9.57	-	-	9.5	8.58	8.6	10.66	9.68	9.94	10.1
85	MgO	7.34	7.32	-	-	7.1	6.68	6.75	8.51	7.08	7.49	8.76
86	C ₇ H ₈	8.69	8.65	8.75	8.71	8.79	8.61	8.72	8.97	-	8.9	8.82
87	C ₈ H ₁₀	8.61	8.52	8.7	8.66	8.73	8.55	8.66	8.92	-	8.85	8.77
88	C ₆ F ₆	9.61	9.46	9.7	9.65	9.69	9.49	9.74	10.04	-	9.93	10.2
89	C ₆ H ₅ OH	8.4	8.36	8.42	8.37	8.43	8.37	8.51	8.72	-	8.7	8.75
90	C ₆ H ₅ NH ₂	7.79	7.64	7.8	7.73	7.84	7.64	7.78	7.98	-	7.99	8.05
91	C ₅ H ₅ N	9.32	9.1	9.28	9.13	9.31	9.04	9.17	9.5	-	9.66	9.66
92	C ₅ H ₅ N ₅ O	7.71	7.54	7.86	7.82	8.18	7.69	7.87	7.92	-	8.03	8.24
93	C ₅ H ₅ N ₅	8.08	7.9	8.14	8.09	8.18	7.98	8.15	8.35	-	8.33	8.48
94	C ₄ H ₅ N ₃ O	8.46	8.26	8.49	8.4	8.5	8.29	8.44	8.77	-	9.51	8.94
95	C ₃ H ₆ N ₂ O ₂	8.81	8.62	8.88	8.82	8.89	8.71	8.87	9.19	-	9.08	9.2
96	C ₄ H ₄ N ₂ O ₂	9.38	9.22	9.26	9.19	9.55	9.22	9.38	9.94	-	10.12	9.68
97	CH ₄ N ₂ O	9.58	9.33	9.6	9.4	9.59	9.32	9.46	9.94	-	10.05	9.8
98	Ag ₂	8.11	8.27	8.12	8.04	7.95	7.07	-	8.57	-	7.49	7.66
99	Cu ₂	7.24	7.66	-	-	7.4	7.54	7.78	8.6	-	7.57	7.46
100	NCCu	9.93	10.12	-	-	9.99	9.42	9.56	10.91	-	10.85	-

^aReference GW results from other codes are also provided (AIMS, TM, and BGW from ref 34 and WEST from ref 34), together with experimental data.¹⁰² For a full comparison with respect to the KIPZ results, we redirect the reader to ref 103.

The YAMBO-MPA results generally improve the agreement with all other codes relative to GN-PPA, as reflected by the smaller MAE and MARE values in Table 2. The improvement is particularly significant when comparing with WEST_{lin} and VASP, confirming that MPA provides an efficient description of frequency dependence, with an accuracy comparable to that of FF-RA, CD, and AC approaches.^{49,50,86} The only notable exception is the comparison with FHI-AIMS results, based on localized def2-QZVP basis sets, prior to complete-basis-set extrapolation, for which slightly larger deviations persist. It is worth noting that, despite the absence of the Bruneval–Gonze terminator⁶⁷ in the MPA implementation, which, at the PPA level, has been shown to accelerate convergence, the overall quality of the extrapolated MPA data remains unaffected. Finally, the relatively large MAE observed between the YAMBO GW data and experimental IPs can be attributed to several factors beyond the computational framework, including the

neglect of finite-temperature effects and coupling with ionic degrees of freedom, as well as higher-order corrections such as self-consistency and vertex corrections.³⁴ This is also consistent with the discrepancy observed with respect to the KIPZ method, which indeed can be considered as including approximate vertex corrections. For more details on the Koopmans functionals approach, we refer to refs 103–106.

A more detailed assessment of the deviations across different implementations can be obtained from the violin plots in Figure 2, which combine box- and density-plot representations. The vertical asymmetry of each distribution indicates whether the IPs computed with YAMBO are systematically over- or underestimated relative to the reference data, while the width of the main peak reflects the standard deviation σ reported in Table 2. The values plotted in Figure 2 are summarized in Table 3, with PPA (yellow) and MPA (red) results shown side by side. For VASP, 16% (36%) of the molecules deviate by

Table 4. Quasiparticle Electron Affinity for the Selected Molecule of the GW100 Data Set as Obtained Within This Work, Compared with the Available Results of West_{lin}^{extra37,a}

index	formula	PBE (this work)	PBE (ref 37)	YAMBO _{MPA}	YAMBO _{PPA}	WEST _{lin}
7	Li ₂	1.75	1.78	0.73	0.76	0.58
8	Na ₂	1.7	1.77	0.73	0.75	0.58
9	Na ₄	1.8	2.09	1.0	1.03	1.04
10	Na ₆	1.43	1.89	0.87	0.93	1.0
11	K ₂	1.33	1.6	0.72	0.79	0.72
12	Rb ₂	1.17	1.55	0.59	0.65	0.71
14	P ₂	3.42	3.42	1.16	1.08	1.08
15	As ₂	3.39	3.4	1.17	1.08	1.08
16	F ₂	5.94	5.94	0.7	0.32	1.05
17	Cl ₂	4.22	4.22	1.5	1.25	1.37
18	Br ₂	4.49	4.49	2.0	1.79	1.87
19	I ₂	4.44	4.45	3.15	3.18	3.21
26	C ₄	6.05	6.05	3.18	2.93	3.11
29	C ₈ H ₈	2.29	2.31	0.22	−0.04	0.05
35	CCl ₄	2.69	2.7	0.55	0.31	0.39
36	CBr ₄	3.48	3.5	1.62	1.42	1.44
37	Cl ₄	4.11	4.21	3.1	3.06	3.03
41	Si ₃ H ₁₂	1.22	1.68	−0.12	−0.17	0.05
42	LiH	1.59	1.58	0.1	0.1	0.05
43	KH	1.6	1.61	0.32	0.3	0.22
54	F ₂ Mg	2.64	2.64	0.31	0.25	0.32
55	TiF ₄	4.07	4.08	0.7	0.29	0.82
56	AlF ₃	2.61	2.61	0.12	0.03	0.14
58	SF ₄	3.57	2.96	3.13	3.35	0.05
59	BrK	1.82	1.87	0.45	0.42	0.39
60	GaCl	2.39	2.39	0.46	0.45	0.42
61	NaCl	2.22	2.22	0.49	0.47	0.45
62	MgCl ₂	2.57	2.59	0.76	0.72	0.68
63	AlI ₃	2.58	2.82	1.57	1.54	1.65
66	PN	3.41	3.41	0.58	0.35	0.49
77	CS ₂	2.79	2.83	0.56	0.33	0.48
81	O ₃	6.17	6.17	2.56	1.87	2.6
82	SO ₂	4.4	4.42	1.37	1.02	1.36
83	BeO	4.84	4.84	2.25	2.27	2.23
84	MgO	4.29	4.29	1.95	1.71	2.03
93	C ₅ H ₆ N ₂ O ₂	2.24	2.29	0.19	−0.11	0.08
94	C ₄ H ₄ N ₂ O ₂	2.44	2.45	0.2	−0.05	0.13
96	Ag ₂	3.08	3.1	1.54	1.52	1.47
97	Cu ₂	3.09	3.1	1.3	1.3	1.29
98	NCCu	4.12	4.12	1.91	1.85	1.92

^aBoth DFT and QP values are indicated.

approximately −95 (30) meV for PPA (MPA), forming the largest peak in each respective distribution. When comparing YAMBO–PPA with WEST_{lin}, 19% of the molecules cluster around −80 meV, while 32% form a shoulder at about −300 meV. In contrast, the MPA results show a narrower and more symmetric distribution, with 35% of the molecules deviating only by 19 meV from West_{lin}, without any shoulder. Overall, YAMBO–MPA consistently improves the agreement with all other reference codes, further demonstrating the reliability of the MPA formulation.

Most distributions in Figure 2 are asymmetric toward a slight overestimation of the IPs computed with YAMBO relative to the other methods. The exceptions are the comparisons with BGW_{HL-PPA}, CCSD, and experimental data, where the deviation signs reverse. Outliers are present in almost all

distributions. The largest deviation from VASP corresponds to the Xe molecule with differences of 1.30 eV (PPA) and 1.12 eV (MPA). For WEST_{lin}, the largest outlier is the F₂ molecule, with a PPA deviation of 633 meV, while other fluorine-containing molecules such as CF₄, FH, and AlF₃ show absolute deviations of 400–500 meV. These discrepancies are probably due to the presence of strongly localized 2p and 3d electrons, which are challenging to treat accurately in standard one-shot G_0W_0 calculations,¹⁰⁷ particularly within PPA.²⁶ Importantly, the MPA approximation substantially mitigates these discrepancies, reducing, for example, the F₂ difference to 273 meV. The improved performance of the MPA approach is further supported by the cumulative statistics: the number of molecules for which the difference between YAMBO and VASP (WEST_{lin}) is smaller than 150 meV increases from 42 (50) in PPA to 72 (73) in MPA.

Finally, we briefly discuss the main factors influencing the residual discrepancies observed in the present data set: (i) the reduced finite size of the simulation cell, and (ii) the linearization of the QP equation. Regarding the supercell size, as discussed in Sec. II of the Supporting Information,⁷⁸ the use of a 13 Å FCC cell—smaller than the 25 Å cubic cells employed in WEST³⁷—introduces a finite-size error in the G_0W_0 IPs. For molecules of average size, this amounts to roughly 25 meV, while for the largest molecules, it increases to about 180 meV. Consistently, when comparing with codes that use similar computational setups (e.g., WEST_{lin} or VASP), the MAEs lie within the expected uncertainty range.

Regarding the linearization of the QP equation, it shows a moderate impact when compared with WEST data obtained with and without linearization, changing the MAEs relative to YAMBO–PPA and MPA by about 20 and 60 meV, respectively. Finally, we did not find any significant correlation between the residual extrapolation error (with respect to the most converged YAMBO results) and the deviations from other codes, as shown in Figure S5 of the Supporting Information.⁷⁸ This indicates that differences in the final IPs are only marginally influenced by the extrapolation procedure.

3.2. Electron Affinities

Electron affinities are significantly more difficult to compare than IPs. As noted in the original GW100 paper,³⁴ the local orbital description of the unbound states, particularly for small molecules, is not well-converged, leading to large deviations with respect to results obtained with plane waves. For this reason, we compare the Yambo results only with the WEST code and the linearized G_0W_0 QP solution, which showed the best agreement for the HOMO levels. The analysis of the EA energies follows the same protocol as for the IPs. A summary of the computed values is reported in Table 4, while Figure 3 displays the statistical differences between YAMBO and WEST. We first notice that the PBE results are in very good agreement with the corresponding results obtained in ref 37, showing a small MAE of 80 meV.

As observed for the IPs, MPA systematically improves the agreement with the reference data compared to that of PPA. The MAE amounts to 163 meV for MPA and 217 meV for PPA, confirming that the MPA representation yields a more accurate frequency dependence of the self-energy, including for unoccupied states. The largest deviation in the data set corresponds to the SF₄ molecule, for which the LUMO exhibits an anomalously high discrepancy (approximately 600 meV) already at the DFT level.³⁷ This deviation propagates to

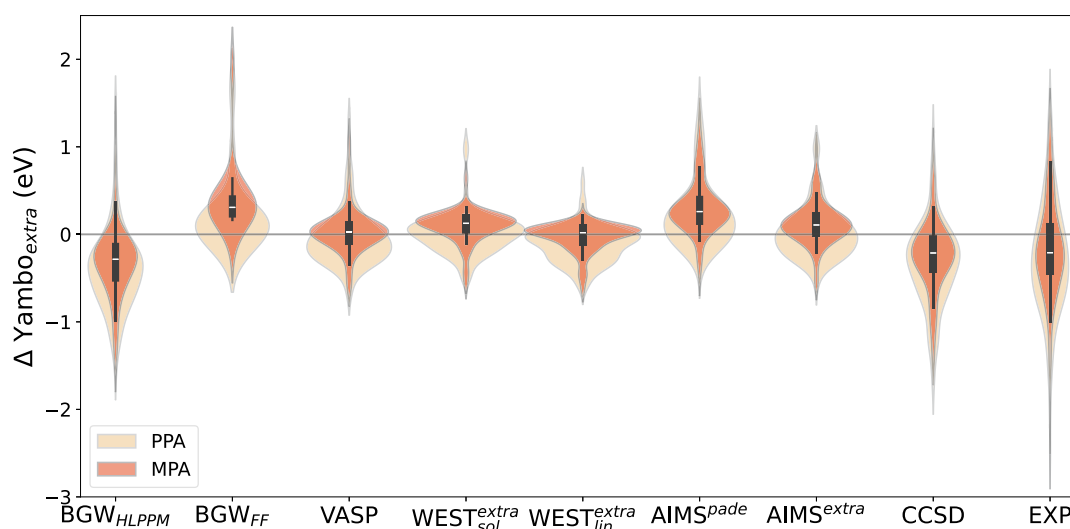


Figure 2. Violin plots representing the distribution of the IP deviation between YAMBO PPA and MPA results and other GW codes. The width of each curve reflects the density of data points. In the MPA case, the vertical black box represents the interquartile range, and the white horizontal line indicates the median.

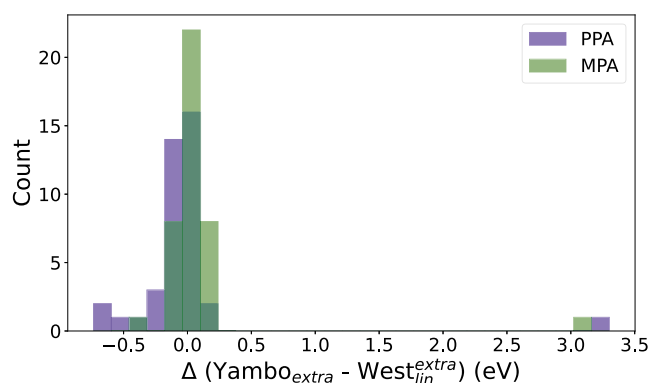


Figure 3. Deviation of the quasiparticle electron affinity (EA) between the YAMBO and the WEST linearized extrapolated results.

the QP correction and thus accounts for the outlier observed in Figure 3. Overall, the EA data set reinforces the conclusions drawn from the IP analysis: the MPA approach achieves accuracy comparable to that of FF methods at a substantially reduced computational cost. A comprehensive table of QP EA as obtained within this work is provided in the [Supporting Information](#).⁷⁸

4. CONCLUSIONS

In this work, we present a comprehensive benchmark of IPs and EAs for the 100 molecules of the GW100 data set, computed at the G_0W_0 level with the YAMBO code. The study provides the first systematic assessment of the GN-PPA and the recently introduced MPA, within a PW PP framework. All calculations were performed using GPU-accelerated resources and automated workflows via the `aiida-yambo` plugin, ensuring reproducibility and high-throughput efficiency.

Overall, our results show very good agreement with reference data from other GW implementations, including VASP, WEST, BGW, and FHI-aims, with mean absolute errors of typically below 0.25 eV. The best agreement is found with WEST_{lin} and VASP, codes that also use the linearized solution of the QP equation and plane waves and PAW basis sets, respectively. On the other hand, the largest deviations, which

are found for codes with localized basis sets (AIMS, Turbomole), remain moderate. Comparison with KIPZ, CCSD, and experimental IPs confirms trends previously reported for the GW100 set, with residual differences largely due to the lack of finite-temperature and electron–phonon effects, as well as the absence of self-consistency and vertex corrections in the present one-shot G_0W_0 framework.

Importantly, the comparison between different frequency-integration schemes highlights the reliability of GN-PPA and its partial improvement over the Hybertsen–Louie PPA version used in BGW, which shows larger deviations with respect to full-frequency results. MPA further enhances the accuracy of the frequency treatment, providing good agreement with full-frequency approaches (analytic continuation or contour deformation) at a fraction of their computational cost. MPA extrapolated results also seem not to suffer from the lack of the BG terminator,⁶⁷ used here with PPA.

The remaining discrepancies can be mainly attributed to the use of norm-conserving PPs and the finite size of the supercells. The linearization of the QP equation contributes only marginally (20–60 meV) to the differences with respect to other GW implementations. The error due to the extrapolation scheme with respect to the number of included empty states and the kinetic energy cutoff does not show a significant correlation with the deviations from other code results, suggesting that the extrapolation scheme is a valid method to improve accuracy with respect to the results obtained with the largest feasible convergence parameters.

In summary, the present benchmark demonstrates that GN-PPA and, in particular, MPA provide accurate and efficient treatments of G_0W_0 QP calculations, enabling verification against more demanding full-frequency methods. These results consolidate the reliability of the YAMBO code for large-scale molecular GW simulations. Future developments will target the implementation of self-consistent GW and beyond-GW schemes on GPU architectures, further enhancing both the accuracy and the scalability of excited-state calculations.

■ ASSOCIATED CONTENT

Data Availability Statement

The data presented in this work are available on the Materials Cloud Archive¹⁰⁸ at the URL: [10.24435/materialscloud:4a-d7](https://pubs.acs.org/doi/10.24435/materialscloud:4a-d7).

SI Supporting Information

The Supporting Information is available free of charge at <https://pubs.acs.org/doi/10.1021/acs.jctc.6c00239>.

Additional details on computational resource requirements, supercell choice, and the extrapolation procedure used to obtain the results of this work (PDF)

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Notes

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