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# Path integral Monte Carlo determination of the fourth-order virial coefficient for unitary two-component Fermi gas with zero-range interactions

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The unitary equal-mass Fermi gas with zero-range interactions constitutes a paradigmatic model system that is relevant to atomic, condensed matter, nuclear, particle, and astro physics. This work determines the fourth-order virial coefficient  $b_4$  of such a strongly-interacting Fermi gas using a customized *ab initio* path integral Monte Carlo (PIMC) algorithm. In contrast to earlier theoretical results, which disagreed on the sign and magnitude of  $b_4$ , our  $b_4$  agrees within error bars with the experimentally determined value, thereby resolving an ongoing literature debate. Utilizing a trap regulator, our PIMC approach determines the fourth-order virial coefficient by directly sampling the partition function. An on-the-fly anti-symmetrization avoids the Thomas collapse and, combined with the use of the exact two-body zero-range propagator, establishes an efficient general means to treat small Fermi systems with zero-range interactions.

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*Introduction:* Strongly-interacting Fermi gases manifest themselves in nature in different forms, from neutrons in neutron stars [1] to electrons in solids [2]. These systems are generally deemed difficult to treat theoretically because of the lack of a small interaction parameter. Superconductivity [3] and exotic states such as fractional quantum hall [4] or Fulde-Ferrell-Larkin-Ovchinnikov [5–7] states have been observed or predicted to exist in these systems. Ultracold Fermi gases [8, 9], which can nowadays be produced routinely in table-top experiments, are ideal for studying strongly-interacting systems since (i) the van der Waals interaction is short-ranged, which means that it can be approximated by a contact potential that introduces a single length scale, i.e., the  $s$ -wave scattering length  $a_s$ ; and (ii)  $a_s$  can be tuned at will utilizing Feshbach resonance techniques [10]. When  $a_s$  diverges, i.e., becomes infinitely large, the two-body contact potential does not define a length scale [11]. Just like the non-interacting Fermi gas, the properties of the unitary Fermi gas (Fermi gas with infinite  $a_s$ ) are determined by two length scales, the de Broglie wavelength  $\lambda$  and interparticle spacing  $\bar{r}$  [12].

At high temperature,  $\lambda$  is much smaller than  $\bar{r}$  and the grand canonical thermodynamic potential  $\Omega$  can be expanded in terms of the fugacity [13, 14]. The  $n^{\text{th}}$ -order expansion or virial coefficient  $b_n$  is determined by the partition functions of clusters containing  $n$  or fewer fermions. Since all thermodynamic properties at high temperature can be derived from the virial coefficients  $b_n$  [15], the  $b_n$ 's are essential to understanding the normal state of strongly-interacting Fermi gases.

While the second- and third-order virial coefficients are well understood [13, 15–21], none of the theoretical calculations for  $b_4$  [22–25] agree with the experimental data [19, 26]. This letter rectifies this situation: our theoretically determined  $b_4$  agrees with the experimentally determined value. Our approach uses a trap regulator [27, 28] and employs the path integral Monte Carlo

(PIMC) technique [29, 30], with the contact interactions incorporated exactly via the two-body zero-range propagator [31]. The “post-anti-symmetrization” [29, 30], traditionally employed in PIMC calculations, does not work for the system with zero-range interactions, since the sampled paths shrink due to the Thomas collapse, a well known phenomenon for bosons [32, 33], to a single point. For bosons, the three-body Thomas collapse is cured by introducing an additional scale or three-body parameter [33]. For fermions, such a three-body parameter is not needed since the Pauli exclusion principle acts as an effective three-body repulsion [34, 35]. Thus, rather than the standard “post-anti-symmetrization”, we use an “on-the-fly scheme” [36, 37], which anti-symmetrizes at each imaginary time step. While the anti-symmetrization is, within Monte Carlo frameworks, usually associated with the infamous Fermi sign problem [38–40], in our case it stabilizes the simulation and affords the use of significantly smaller number of time slices than the use of finite-range interactions would. Our approach reproduces the trap regulated  $b_3$  over a wide temperature range. We determine the trap regulated  $b_4$  as a function of the temperature  $T$ . In the low-temperature regime, we find agreement with Ref. [22]. We separate the spin-balanced ( $b_{2,2}/2$ ) and spin-imbalanced ( $b_{3,1}$ ) sub-cluster contributions to  $b_4$ ,  $b_4 = b_{3,1} + b_{2,2}/2$ , and find  $b_{2,2} < 0$  and  $b_{3,1} > 0$  at all considered temperatures.  $b_{2,2}$  dominates at low  $T$  and  $b_{3,1}$  at high  $T$ . Converting the trap regulated virial coefficient  $b_4$  to that of the homogeneous system using the local density approximation (LDA) [16], we find agreement with the experimentally determined values [19, 26].

*Virial expansion framework:* The  $n^{\text{th}}$ -order virial coefficient  $b_n^{\text{hom}}$  of the homogeneous system at unitarity is related to the high-temperature limit  $b_n^0$  of the harmonically trapped unitary system via  $b_n^{\text{hom}} = n^{3/2}b_n^0$  [16, 28]. To determine  $b_n^0$ , we calculate the virial coefficient  $b_n$  of the harmonically trapped system for various tempera-

tures and then take the  $T \rightarrow \infty$  limit. The trap Hamiltonian  $H(n_1, n_2)$  for  $n_1$  particles of species 1 and  $n_2$  particles of species 2 with interspecies  $s$ -wave interactions reads

$$H(n_1, n_2) = \sum_{j=1}^{n_1+n_2} \left( \frac{-\hbar^2}{2m} \nabla_{\mathbf{r}_j}^2 + \frac{1}{2} m \omega^2 \mathbf{r}_j^2 \right) + \sum_{i=1}^{n_1} \sum_{j=n_1+1}^{n_1+n_2} V_{2b}(\mathbf{r}_i - \mathbf{r}_j), \quad (1)$$

where  $m$  denotes the mass of the particles,  $\mathbf{r}_j$  the position vector of the  $j^{\text{th}}$  particle,  $\omega$  the angular trapping frequency, and  $V_{2b}$  the regularized Fermi-Huang pseudopotential with infinite  $a_s$  [41]. The grand canonical thermodynamic potential  $\Omega$  can be written in terms of the fugacities  $z_i$  of species  $i$ ,

$$\Omega = -k_B T \ln \left( \sum_{n_1=0}^{\infty} \sum_{n_2=0}^{\infty} Q_{n_1, n_2} z_1^{n_1} z_2^{n_2} \right), \quad (2)$$

where  $z_i$  is equal to  $\exp[\mu_i/(k_B T)]$ ,  $\mu_i$  is the chemical potential of species  $i$ , and  $Q_{n_1, n_2}$  is the canonical partition function for  $H(n_1, n_2)$ ,

$$Q_{n_1, n_2} = \text{Tr} \exp[-H(n_1, n_2)/(k_B T)]. \quad (3)$$

Here,  $\text{Tr}$  is the trace operator. Defining  $\Delta\Omega = \Omega - \Omega^{\text{ni}}$ , where  $\Omega^{\text{ni}}$  is the grand canonical potential of the non-interacting system, and Taylor-expanding around  $z_1 = z_2 = 0$  [42, 43], one finds

$$\Delta\Omega = -k_B T Q_{1,0} \left( \sum_{n_1=1}^{\infty} \sum_{n_2=1}^{\infty} b_{n_1, n_2} z_1^{n_1} z_2^{n_2} \right). \quad (4)$$

For spin-balanced systems,  $z_1$  and  $z_2$  are equal and Eq. (4) reduces to

$$\Delta\Omega = -2k_B T Q_{1,0} \left( \sum_{n=2}^{\infty} b_n z^n \right), \quad (5)$$

where  $b_2 = b_{1,1}/2$ ,  $b_3 = (b_{1,2} + b_{2,1})/2$ , and  $b_4 = (b_{1,3} + b_{3,1} + b_{2,2})/2$  (note, one has  $b_{2,1} = b_{1,2}$  and  $b_{3,1} = b_{1,3}$ ). It is convenient to write the virial coefficients  $b_{n_1, n_2}$  as

$$b_{n_1, n_2} = \Delta b_{n_1, n_2} + b_{n_1, n_2}^{\text{ref}}, \quad (6)$$

where  $b_{n_1, n_2}^{\text{ref}}$  is determined by the virial coefficients  $b_{j_1, j_2}$  and the canonical partition functions  $Q_{j_1, j_2}$  with  $j_1 + j_2 < n_1 + n_2$ . The term  $\Delta b_{n_1, n_2} = (Q_{n_1, n_2} - Q_{n_1, n_2}^{\text{ni}})/Q_{1,0}$ , where  $Q_{n_1, n_2}^{\text{ni}} = Q_{n_1, 0} Q_{0, n_2}$ , in contrast, accounts for the “new” physics introduced by the interacting  $(n_1, n_2)$  clusters [44].

*Contradicting literature results for  $b_4$ :* The literature results are summarized in Table I (see also the supplemental material [45]). Two independent experiments find consistent values for the fourth-order virial coefficient.

The theoretical literature results, however, disagree with these experimental results, reflecting the fact that the fourth-order problem is highly non-trivial analytically and numerically. Using a sum-over-states approach with an energy cutoff, Ref. [22] obtained the low-temperature behavior of  $b_4$ . Assuming a monotonic temperature dependence and extrapolating to the  $T \rightarrow \infty$  limit, Ref. [22] obtained  $b_4^0$ . It was concluded that more four-body energies would need to be calculated explicitly to obtain  $b_4$  reliably at high temperature. The fourth-order virial coefficient has also been obtained by a diagrammatic approach, which included only a subset of the four-body free-space diagrams [23], and by applying a conjecture inspired by three-body results [24, 25].

*Customized PIMC algorithm:*  $\Delta b_{n_1, n_2}$  is determined by the partition function  $Q_{n_1, n_2}$  of the interacting  $(n_1, n_2)$  system ( $Q_{n_1, n_2}$  is not known in general) and the partition function  $Q_{n_1, n_2}^{\text{ni}}$  of the non-interacting  $(n_1, n_2)$  system ( $Q_{n_1, n_2}^{\text{ni}}$  is known analytically). We calculate the ratio of the partition functions  $Q_{n_1, n_2}^{\text{ni}}/Q_{n_1, n_2}$  using the PIMC technique. Specifically, the simulation generates configurations according to  $Q_{n_1, n_2}$  and accumulates the ratio  $Q_{n_1, n_2}^{\text{ni}}/Q_{n_1, n_2}$  as a weight. The reason for using the partition function of the unitary Fermi gas and not that of the non-interacting gas as the “guiding function” is the following. The probability density to find two unlike particles with vanishing interparticle spacing is finite at unitarity and zero in the non-interacting limit. If we used  $Q_{n_1, n_2}^{\text{ni}}$  as the guiding function, configurations in which two unlike particles are at the same spatial position would be absent and the standard deviation of  $Q_{n_1, n_2}/Q_{n_1, n_2}^{\text{ni}}$  would be infinite, rendering the expectation value meaningless [46].

In the PIMC formulation, the partition function  $Q_{n_1, n_2}^{\text{boltz}}(\beta)$  for Boltzmann particles (no exchange symmetries) at inverse temperature  $\beta$ ,  $\beta = (k_B T)^{-1}$ , is written in terms of a product over density matrices at imaginary time  $\tau$ ,

$$Q_{n_1, n_2}^{\text{boltz}}(\beta) = \int \dots \int \prod_{i=1}^N \rho(\mathbf{R}_i, \mathbf{R}_{i+1}; \tau) d\mathbf{R}_1 \dots d\mathbf{R}_N, \quad (7)$$

where  $\mathbf{R}_i$  collectively denotes the particle configurations at time slice  $i$ ,  $\mathbf{R}_N = \mathbf{R}_1$ , and  $N = \beta/\tau$ . For the two-component Fermi gas, the standard PIMC approach

TABLE I. Summary of literature and PIMC results.

$b_4^{\text{hom}}$	$b_4^0$	Ref.	comment
0.096(15)	0.01200(188)	[19]	ENS experiment
0.096(10)	0.01203(125)	[26]	MIT experiment
-0.016(4)	-0.0020(5)	[22]	sum-over-states approach
0.06	0.0075	[23]	diagrammatic approach
-0.063(1)	-0.007875(125)	[24]	3-body inspired conjecture
0.078(18)	0.0098(23)		PIMC, this work

writes the partition function as  $Q_{n_1, n_2} = \mathcal{A}Q_{n_1, n_2}^{\text{boltz}}$ , where  $\mathcal{A}$  is the anti-symmetrizer [29, 47]. For the two-component Bose gas, the anti-symmetrizer  $\mathcal{A}$  is replaced by the symmetrizer  $\mathcal{S}$ .  $\mathcal{A}$  and  $\mathcal{S}$  contain the same number and types of terms; however, while all terms in  $\mathcal{S}$  enter with a plus sign,  $\mathcal{A}$  contains alternating plus and minus signs. Since the symmetrizer and anti-symmetrizer are, in the standard PIMC approach, evaluated stochastically, the two-component Fermi and Bose gases are simulated using the same paths. Expectation values, however, are accumulated with plus and minus signs for fermions and with plus signs only for bosons. We refer to this standard approach as post-symmetrization. The bosonic system with interspecies two-body zero-range interactions but without a three-body regulator would collapse to a single point; this is the well-known Thomas collapse [32]. Correspondingly, the fermionic paths would also collapse, rendering the simulation meaningless. To get around this problem, we developed a customized on-the-fly anti-symmetrization scheme, which explicitly anti-symmetrizes the density matrix at each imaginary time step,

$$Q_{n_1, n_2}(\beta) = \int \dots \int \prod_{i=1}^N \mathcal{A}\rho(\mathbf{R}_i, \mathbf{R}_{i+1}; \tau) d\mathbf{R}_1 \dots d\mathbf{R}_N. \quad (8)$$

The observable is then calculated using

$$\frac{Q_{n_1, n_2}^{\text{ni}}}{Q_{n_1, n_2}} = \left\langle \frac{\prod_{i=1}^N \mathcal{A}\rho^{\text{ni}}(\mathbf{R}_i, \mathbf{R}_{i+1}; \tau)}{\prod_{i=1}^N \mathcal{A}\rho(\mathbf{R}_i, \mathbf{R}_{i+1}; \tau)} \right\rangle, \quad (9)$$

where  $\rho^{\text{ni}}$  denotes the density matrix for the non-interacting system and  $\langle \dots \rangle$  the thermal average using paths generated for the unitary Fermi gas using the on-the-fly anti-symmetrization scheme. Our simulation uses the pair-product approximation [29, 45] with the exact two-body density matrix for zero-range interactions. The on-the-fly scheme employed here is related to earlier works [36, 48], which anti-symmetrized, as we do, at each time slice. The key difference is that we employ a density matrix that accounts for the interactions while the earlier works employed the non-interacting density matrix together with the Trotter (or improved Trotter) formula.

The on-the-fly anti-symmetrization scheme treats the  $n_1!n_2!$  permutations explicitly at each time slice, eliminating the need of the standard stochastic “permute move”. As a consequence, the scheme is computationally prohibitively demanding for large systems. For small systems, however, it is quite efficient for three reasons: (i) The number of permutations is manageable for small  $n_1 + n_2$ . (ii) The use of the zero-range interactions eliminates the need to perform calculations for several different ranges of the underlying two-body potential. (iii) Compared to finite-range interactions [49], the number of time slices needed to reach convergence for the zero-range interacting systems considered here is rather small; e.g.,

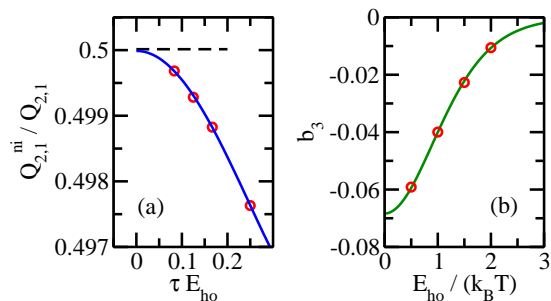


FIG. 1. (Color online) Benchmarking our PIMC results (circles) for the (2,1) system at unitarity through comparison with sum-over-states results. (a) The observable  $Q_{2,1}^{\text{ni}}/Q_{2,1}$  as a function of the imaginary time step  $\tau$  at  $k_B T = E_{\text{ho}}$ . The circles show our PIMC results. The error bars (not shown) are smaller than the symbol size. The solid line shows the fourth-order polynomial fit of the form  $a + b\tau^2 + c\tau^4$ . The dashed line shows the sum-over-states results. (b)  $b_3$  as a function of  $1/(k_B T)$ . The circles show our PIMC results while the solid line shows the sum-over-states results.

our scheme yields  $Q_{3,1}^{\text{ni}}/Q_{3,1}$  at  $E_{\text{ho}}/(k_B T) = 0.8$  with 0.1% error using only  $N = 9$  imaginary time slices (here,  $E_{\text{ho}} = \hbar\omega$ ). Within our approach, the key challenge in determining  $b_4$  reliably at high temperature comes from the fact that  $\Delta b_{2,2}$ ,  $\Delta b_{3,1}$ ,  $b_{2,2}^{\text{ref}}$ , and  $b_{3,1}^{\text{ref}}$  diverge, to leading order, as  $(k_B T/E_{\text{ho}})^6$ . This implies that  $b_{2,2}$  and  $b_{3,1}$  are, at high temperature, obtained by adding two numbers of opposite sign and nearly equal magnitude. Thus, to obtain reliable values at high temperature, we need to determine our observables with high accuracy. In practice, our available computer time limits us to  $k_B T \leq 2E_{\text{ho}}$  for the (2,2) and (3,1) systems.

*PIMC results:* To benchmark our customized PIMC algorithm, we apply it to the (2,1) system at unitarity, for which  $Q_{2,1}^{\text{ni}}/Q_{2,1}$  and  $b_3$  can be calculated with high accuracy for all temperatures using the sum-over-states approach [16]. As an example, circles in Fig. 1(a) show the quantity  $Q_{2,1}^{\text{ni}}/Q_{2,1}$  for  $k_B T = E_{\text{ho}}$ , obtained using our PIMC algorithm, as a function of the imaginary time step  $\tau$ . The  $\tau$  considered correspond to between  $N = 4$  and 10 time slices. The simulation is exact in the  $\tau \rightarrow 0$  (or equivalently,  $N \rightarrow \infty$ ) limit. To extrapolate to the  $\tau \rightarrow 0$  limit, we fit a fourth-order polynomial of the form  $a + b\tau^2 + c\tau^4$  to the PIMC data [solid line in Fig. 1(a)]. Our extrapolated result of 0.499989(26) agrees within error bars with the value of 0.500014 [dashed line in Fig. 1(a)] obtained by the sum-over-states approach. Using the extrapolated  $\tau \rightarrow 0$  values for  $Q_{2,1}^{\text{ni}}/Q_{2,1}$  at various temperatures  $T$ , we obtain  $b_3$  as a function of  $T$  [circles in Fig. 1(b)]. The agreement with the sum-over-states results [solid line in Fig. 1(b)] is excellent for all  $T$  considered, demonstrating the reliability and accuracy of our PIMC approach.

We now discuss the determination of  $b_4$ . The extrapolation of the raw data to the  $\tau \rightarrow 0$  limit is discussed

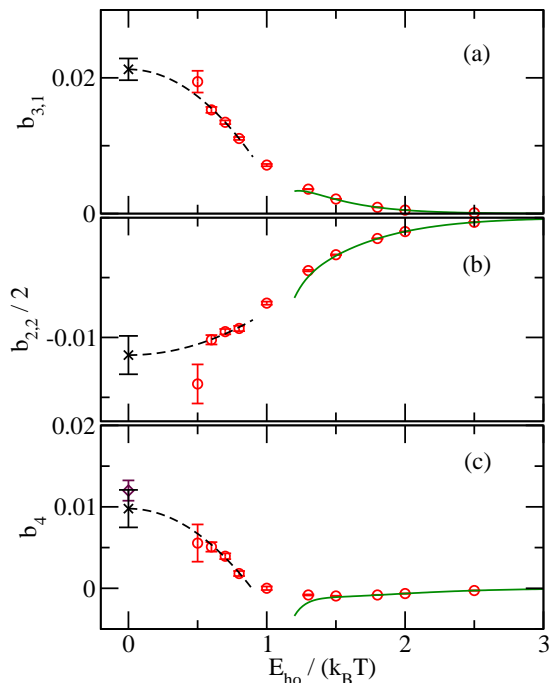


FIG. 2. (Color online) PIMC determination of the fourth-order virial coefficient. Circles in panels (a), (b), and (c) show  $b_{3,1}$ ,  $b_{2,2}/2$ , and  $b_4$ , respectively, determined by our PIMC approach. The crosses in (a) and (b) show the  $T \rightarrow \infty$  limit of the two-parameter fit (dashed line) to the PIMC data at the four highest temperatures. The dashed line and the cross in (c) show the sum of the fits from (a) and (b). The error bar in (c) is obtained by error propagation. The diamond with error bar shows the experimental result from Ref. [26].

in the supplemental material [45]. Circles in Figs. 2(a) and 2(b) show our PIMC results for  $b_{3,1}$  and  $b_{2,2}$ , respectively, as a function of the inverse temperature. At low temperature, the PIMC results agree with the sum-over-states results (solid lines), obtained using the data provided in Ref. [22]. At all temperatures,  $b_{3,1}$  is positive and  $b_{2,2}$  is negative. It has been shown that  $b_{1,1}$  and  $b_{2,1}$  are even functions of  $E_{ho}/(k_B T)$  [16, 20, 43], and the conjecture of Ref. [25] implies that  $b_{3,1}$  and  $b_{2,2}$  are also even functions of  $E_{ho}/(k_B T)$ . Thus, to obtain  $b_{3,1}$  and  $b_{2,2}$ , we fit the data points for the four highest temperatures to the form  $a + b[E_{ho}/(k_B T)]^2$ . The dashed lines in Figs. 2(a) and 2(b) show the fits. Since the data points at  $k_B T = 2E_{ho}$  have much larger error bars than those at lower temperatures, the data points contribute comparatively little to the fit, which weighs each data point by the inverse of the square of its error bar. We find  $b_{3,1}^0 = 0.0212(8)$  and  $b_{2,2}^0/2 = -0.0115(8)$ , where the error bars reflect the uncertainty of the fit. We unfortunately do not have sufficiently many data to include a  $(k_B T)^{-4}$  term in the fit. Since the inclusion of a  $(k_B T)^{-4}$  term in the fit could alter the  $T \rightarrow \infty$  result, we add an ad-hoc systematic error of 0.0008 to  $b_{3,1}^0$  and  $b_{2,2}^0/2$ , yielding  $b_{3,1}^0 = 0.0212(16)$  and  $b_{2,2}^0/2 = -0.0115(16)$  [crosses

in Figs. 2(a) and 2(b)]. To obtain  $b_4$  [see Fig. 2(c)], we combine  $b_{3,1}$  and  $b_{2,2}$ . Specifically, the circles and the fit are obtained by adding the data of Figs. 2(a) and 2(b) while the error bar of the cross at  $T \rightarrow \infty$  is obtained using standard error propagation.  $b_4$  displays an interesting temperature dependence: It is negative at low temperature due to the dominance of  $b_{2,2}$ , vanishes at  $k_B T \approx E_{ho}$  due to a cancellation of  $b_{3,1}$  and  $b_{2,2}/2$ , and is positive at high temperature due to the dominance of  $b_{3,1}$ . Our results resolve the discrepancy of the sign of  $b_4$  between Ref. [22] and the experiments [19, 26]. Our extrapolated  $b_4$  at infinite temperature is  $b_4^0 = 0.0098(23)$ , which agrees with the experimental results of  $b_4^0 = 0.01203(125)$  [26] [diamond in Fig. 2(c)] and  $b_4^0 = 0.01200(188)$  [19] (see also Table I). Using the LDA, we find  $b_4^{hom} = 0.078(18)$ .

We now compare our results for  $b_{3,1}^0$  and  $b_{2,2}^0$  with the literature. The diagrammatic approach [23] yields  $b_{3,1}^0 = 0.025$ , which is within 2.5 standard deviations of our value, and  $b_{2,2}^0/2 = -0.018$ , which differs by a factor of about 1.5 (or many standard deviations) from our value. This comparison suggests that the convergence of the diagrammatic approach is slower for the (2,2) system than for the (3,1) system. The conjecture-based approach [24, 25] yields  $b_{3,1}^0 = 0.02297(4)$ , which agrees within error bars with our value, and  $b_{2,2}^0/2 = -0.0309(1)$ , which differs by about a factor of 3 from our value.

*Conclusion:* This letter presented the PIMC determination of the fourth-order virial coefficient of the trapped unitary two-component Fermi gas. Our extrapolated infinite temperature result was found to agree with experiments within error bars, which, to the best of our knowledge, is the first numerical confirmation of the experimental determination of  $b_4$ . The customized PIMC scheme, which allows for the treatment of Fermi gases with zero-range interactions, can be applied to a variety of other situations. Since the zero-range density matrix can be constructed for arbitrary  $s$ -wave scattering length  $a_s$ , the scheme can be used to study the finite-temperature characteristics of the BEC-BCS crossover of few-body Fermi gases. Moreover, the algorithmic developments can be integrated into PIMC ground state calculations, providing a viable alternative to basis set expansion approaches.

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*Note added:* After submission of this paper, Endo and Castin revised their conjecture presented in Ref. [25].

The new calculation yields  $b_4^0 = 0.00775(10)$ ,  $b_{3,1}^0 = 0.02297(4)$ , and  $b_{2,2}^0/2 = -0.0152(1)$  [50].

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- [45] The supplemental material at TO.BE.INSERTED.BY.THE.EDITOR contains (i) a summary of the literature values of the fourth-order virial coefficient; (ii) information on the pair product approximation and the two-body density matrix for zero-range interactions; and (iii) PIMC simulation details and tables containing selected raw data.
- [46] For a simulation of finite length  $l$ , the standard deviation would not be following a Gaussian distribution, implying that the standard deviation would not decrease as  $1/\sqrt{l}$  with increasing simulation length  $l$ .
- [47] For the (2,2) system, e.g.,  $\mathcal{A}$  reads  $(1 - P_{12} - P_{34} + P_{12}P_{34})/4$ , where  $P_{ij}$  permutes the coordinates of particles  $i$  and  $j$ .
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