

New Algorithm for Electron Repulsion Integrals Oriented to the General Contraction Scheme

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ABSTRACT: An algorithm for computing electron repulsion integrals (ERIs) oriented to the general contraction scheme is presented. The accompanying coordinate expansion (ACE) method of Ishida is utilized to derive an efficient algorithm. The performance estimated with the floating-point operation (FLOP) count is about N^2 times and more as efficient as the conventional algorithm for the segmented contraction scheme, where N indicates the number of contracted Gaussian-type orbitals (GTOs) contained in a set of generally contracted GTOs. The efficiency is also confirmed by using a realistic molecular system, the benzene molecule, with C:14s9p/3s2p, H:8s4p/2s1p, and C:14s9p/6s5p, H:8s4p/4s3p basis sets. The measured central processing unit (CPU) time is in good agreement with the FLOP count estimation. © 2000 John Wiley & Sons, Inc. *Int J Quant Chem* 76: 396–406, 2000

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Introduction

Ab initio calculations on large molecular systems containing thousands of atoms or tens

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of thousands of basis functions are a challenging problem in current quantum chemistry. Such systems can be seen in many important chemical models, such as surface–molecular interface systems in reactions on a surface or in solution and in biological reactions. Although computational techniques, such as the direct method [1] and parallel computation, have made great progress, computation of the huge number of electron repulsion

integrals (ERIs) is still a major problem. Two aspects are crucial in achieving fast computation of ERIs. One is the selection of an efficient basis set, and the other is the adoption of a sophisticated computational scheme. In the present work, we propose an elaborate algorithm for ERIs involving the advantages of the general contraction scheme for Gaussian-type orbitals (GTOs) [2] and the accompanying coordinate expansion (ACE) method developed by Ishida [3].

The general contraction scheme was first proposed by Raffanetti [2] in which the contracted basis is treated as a linear combination of the primitive basis, with each primitive function contributing to several (or all) of the contracted functions. Explorations of general contraction schemes and applications to molecular problems were carried out by Raffanetti, Ruedenberg, and Bardo [4]. General contraction was also utilized in the well-tempered basis sets of Huzinaga [5], but wider use did not take place until two recent developments of atomic natural orbital (ANO) basis sets by Almlöf and Taylor [6] and correlation-consistent basis sets by Dunning [7]. Unlike most earlier basis sets, which were determined on the basis of self-consistent field (SCF) calculations, these new types of generally contracted basis sets are designed specifically to be effective in the treatment of correlation. Performance comparisons with other segmented contraction basis sets have been very favorable and confirm that these general contractions fulfill their design objectives. The efficient aspect of the general contraction scheme is that the basis sets can describe the accurate atomic properties with small one-particle space and are simple enough to treat the wave functions.

In spite of the advantages of the general contraction scheme, the computation of ERIs over generally contracted basis functions is very expensive if we use the conventional routines for the segmented contraction scheme. Since the degree of contraction of generally contracted basis sets is usually much higher than that of segmentally contracted basis sets, the calculation of ERIs over them is very costly even if we use the best algorithms. Another cause of the cost of the calculation is that the conventional routines cannot utilize the advantage of efficiency of the general contraction scheme, since they treat the basis set for the general contraction scheme in the same way as that for the segmented contraction scheme.

In the present work, we present a refined algorithm for calculating ERIs for the general contraction scheme so as to reduce significantly the cost over that of the conventional algorithms. Our algorithm employs the ACE-b3k3 formula derived by Ishida, which is well-suited for the calculation of ERIs over highly contracted GTOs [3, 8, 9]. The performance is supported by a theoretical measure of the FLOP count [10, 11].

In the following section, we show theoretically our refined algorithm to adapt the ACE-b3k3 formula to the general contraction scheme, comparing it with the conventional method. The computational efficiency is discussed in the third section. Our conclusion is given in the fourth section.

Method

ACE-b3k3 FORMULA

Several types of ACE formulas have been derived by Ishida, such as ACE-b1k1, ACE-b2k1, ACE-b2k3, and ACE-b3k3 [3]. The most efficient type, namely the formula that gives the least FLOP count, varies depending on the degree of the contraction. In the case of high contraction, the ACE-b3k3 formula is most effective, and thus it is suitable for applying to the general contraction scheme. In the present work, we focus on the ACE-b3k3 formula case.

First, we give a brief review of the ACE-b3k3 formulas. In the ACE-b3k3 method, a primitive ERI over primitive GTOs is expressed as the summation of the products of an accompanying coordinate part C_4^{ABCD} and a core part $H_{4\lambda\mu\nu\xi}^{ABCD}$,

$$\begin{aligned} & \left[\varphi_\lambda^A \varphi_\mu^B \mid \varphi_\nu^C \varphi_\xi^D \right] \\ & = S_{\lambda\mu}^{AB} S_{\nu\xi}^{CD} \sum_{\{\mathbf{N}_3\}} C_4^{ABCD} \{\mathbf{N}_3\} H_{4\lambda\mu\nu\xi}^{ABCD} \{\mathbf{N}_3\}. \quad (1) \end{aligned}$$

Here we omit the normalization factors. The GTOs, $\varphi_\lambda^A, \varphi_\mu^B, \varphi_\nu^C, \varphi_\xi^D$, are centered at coordinates $\mathbf{A}, \mathbf{B}, \mathbf{C}, \mathbf{D}$ and have the exponents, $\alpha_\lambda^A, \alpha_\mu^B, \alpha_\nu^C, \alpha_\xi^D$, and the quantum numbers, $\mathbf{L}_A = (l_A m_A n_A)$, the sum of which, L_A , gives the angular momentum quantum number. The set $\{\mathbf{N}_3\}$ represents summation indices,

$$\{\mathbf{N}_3\} = \{M_A M_B M_C M_D i' j' k' h' i'' j'' k'' h''\}, \quad (2)$$

which obey the following restrictions:

$$\begin{aligned}
 0 \leq M_A \leq L_A, \quad 0 \leq i' \leq L_A - M_A, \\
 \hspace{15em} 0 \leq i'' \leq i', \\
 0 \leq M_B \leq L_B, \quad 0 \leq j' \leq L_B - M_B, \\
 0 \leq j'' \leq j', \hspace{15em} (3) \\
 0 \leq M_C \leq L_C, \quad 0 \leq k' \leq L_C - M_C, \\
 \hspace{15em} 0 \leq k'' \leq k', \\
 0 \leq M_D \leq L_D, \quad 0 \leq h' \leq L_D - M_D, \\
 \hspace{15em} 0 \leq h'' \leq h'.
 \end{aligned}$$

The symbols $S_{\lambda\mu}^{AB}$ and $S_{\nu\xi}^{CD}$ are the remaining parts that do not depend on $\{\mathbf{N}_3\}$,

$$S_{\lambda\mu}^{AB} = (2\pi)^{-1/4}(\pi/\gamma_1)^{3/2} \exp\left(-\alpha_\lambda^A \alpha_\mu^B \overline{AB}^2 / \gamma_1\right), \quad (4)$$

$$S_{\nu\xi}^{CD} = (2\pi)^{-1/4}(\pi/\gamma_2)^{3/2} \exp\left(-\alpha_\nu^C \alpha_\xi^D \overline{CD}^2 / \gamma_2\right), \quad (5)$$

with

$$\gamma_1 = \alpha_\lambda^A + \alpha_\mu^B, \quad \gamma_2 = \alpha_\nu^C + \alpha_\xi^D. \quad (6)$$

The accompanying coordinate part C_4^{ABCD} , which is independent of the exponents $(\alpha_\lambda^A, \alpha_\mu^B, \alpha_\nu^C, \alpha_\xi^D)$, is given by

$$C_4^{ABCD}\{\mathbf{N}_3\} = \sum_{\{\mathbf{M}\}} D_{i'i''}^{A3} D_{j'j''}^{B3} D_{k'k''}^{C3} D_{h'h''}^{D3}, \quad (7)$$

with

$$D_{i'i''}^{A3} = \sum_{\{i'\}} \sum_{\{i''\}} D_{i'_x i''_x}^{A3x} D_{i'_y i''_y}^{A3y} D_{i'_z i''_z}^{A3z} \quad (8)$$

$$\begin{aligned}
 D_{i'_x i''_x}^{A3x} = \begin{pmatrix} l_A \\ M_{Ax} \end{pmatrix} \begin{pmatrix} l_A - M_{Ax} \\ i'_x \end{pmatrix} \begin{pmatrix} i''_x \\ i''_x \end{pmatrix} \\
 \times AB_x^{l_A - M_{Ax} - i'_x} CD_x^{i'_x - i''_x} AC_x^{i''_x}. \quad (9)
 \end{aligned}$$

The sets $\{\mathbf{M}\}$, $\{i'\}$, and $\{i''\}$ are again summation indices,

$$\{\mathbf{M}\} = \{M_A M_B M_C M_D\}, \quad (10)$$

$$\mathbf{M}_A = (M_{Ax} M_{Ay} M_{Az}),$$

$$\{i'\} = \{i'_x i'_y i'_z\}, \quad \{i''\} = \{i''_x i''_y i''_z\}, \quad (11)$$

which obey the restrictions,

$$\begin{aligned}
 0 \leq M_{Ax} \leq l_A, \quad 0 \leq i'_x \leq l_A - M_{Ax}, \\
 \hspace{15em} 0 \leq i''_x \leq i'_x, \\
 0 \leq M_{Ay} \leq m_A, \quad 0 \leq i'_y \leq m_A - M_{Ay}, \\
 \hspace{15em} 0 \leq i''_y \leq i'_y, \hspace{15em} (12) \\
 0 \leq M_{Az} \leq n_A, \quad 0 \leq i'_z \leq n_A - M_{Az}, \\
 \hspace{15em} 0 \leq i''_z \leq i'_z, \\
 \hspace{15em} \dots \\
 M_{Ax} + M_{Ay} + M_{Az} = M_A, \quad i'_x + i'_y + i'_z = i', \\
 i''_x + i''_y + i''_z = i'', \\
 \hspace{15em} \dots
 \end{aligned}$$

$$M_{Ax} + M_{Bx} + M_{Cx} + M_{Dx} = \text{even}, \dots \quad (14)$$

Note that C_4^{ABCD} are determined only by the quantum numbers L_A, L_B, L_C, L_D , and the function centers $\mathbf{A}, \mathbf{B}, \mathbf{C}, \mathbf{D}$.

The core part is given by

$$\begin{aligned}
 H_4^{ABCD}\{\mathbf{N}_3\} \\
 = \begin{bmatrix} M_A + M_B \\ 2 \end{bmatrix} \begin{bmatrix} M_C + M_D \\ 2 \end{bmatrix} \\
 = \sum_{i_1} \sum_{i_2} G_{i_1 i_2} \sigma_A^{b_A + d_A} \sigma_B^{a_B + c_B} \sigma_C^{d_C + b_C} \\
 \times \sigma_D^{c_D + a_D} \sigma_1^{M_A + M_B - i_1} \\
 \times \sigma_2^{M_C + M_D - i_2} / (\sigma_1 + \sigma_2)^{M - i_1 - i_2} \\
 \times \sum_{s_1}^{a_B + b_A} \sum_{s_2}^{c_D + d_C} (-1)^{s_1 + s_2} \begin{pmatrix} a_B + b_A \\ s_1 \end{pmatrix} \begin{pmatrix} c_D + d_C \\ s_2 \end{pmatrix} \\
 \times \left(\frac{\sigma_1}{\sigma_1 + \sigma_2}\right)^{i' + j' + s_1} \left(\frac{\sigma_2}{\sigma_1 + \sigma_2}\right)^{k' + h' + s_2} \\
 \times F_{\xi + s_1 + s_2}(z), \hspace{15em} (15)
 \end{aligned}$$

where

$$\sigma_1 = 1/(2\gamma_1), \quad \sigma_2 = 1/(2\gamma_2), \quad (16)$$

$$\begin{aligned}
 \sigma_A = \alpha_\lambda^A / \gamma_1, \quad a_B = L_A - M_A - i', \\
 \hspace{15em} a_D = i' - i'',
 \end{aligned}$$

$$\begin{aligned}
 \sigma_B = \alpha_\mu^B / \gamma_1, \quad b_A = L_B - M_B - j', \\
 \hspace{15em} b_C = j' - j'',
 \end{aligned}$$

$$\begin{aligned}
 \sigma_C = \alpha_\nu^C / \gamma_2, \quad c_D = L_C - M_C - k', \hspace{15em} (17) \\
 \hspace{15em} c_B = k' - k'',
 \end{aligned}$$

$$\begin{aligned}
 \sigma_D = \alpha_\xi^D / \gamma_2, \quad d_C = L_D - M_D - h', \\
 \hspace{15em} d_A = h' - h'',
 \end{aligned}$$

$$M = (M_A + M_B + M_C + M_D)/2, \quad (18)$$

$$\zeta = i' + j' + k' + h' + M - i_1 - i_2, \quad (19)$$

and

$$G_{i_1 i_2} = (-1)^{M_A + M_B + M - (i_1 + i_2)} \sum_{\{\mathbf{I}\}} g_x g_y g_z, \quad (20)$$

$$g_x = \binom{M_{A_x} + M_{B_x}}{2i_{1x}} \binom{M_{C_x} + M_{D_x}}{2i_{2x}} (2i_{1x} - 1)!! \\ \times (2i_{2x} - 1)!! (2M_x - 2i_{1x} - 2i_{2x} - 1)!! \quad (21)$$

The summation indices

$$\{\mathbf{I}\} = \{i_{1x} i_{1y} i_{1z} i_{2x} i_{2y} i_{2z}\} \quad (22)$$

obey the restrictions,

$$0 \leq i_{1x} \leq [(M_{A_x} + M_{B_x})/2], \\ \dots \quad (23)$$

$$0 \leq i_{2x} \leq [(M_{C_x} + M_{D_x})/2], \\ \dots$$

$$i_1 = i_{1x} + i_{1y} + i_{1z}, \quad i_2 = i_{2x} + i_{2y} + i_{2z}, \quad (24)$$

$$2M_x = M_{A_x} + M_{B_x} + M_{C_x} + M_{D_x}. \quad (25)$$

The function $F_m(z)$ in Eq. (15) is the molecular incomplete gamma function defined by

$$F_m(z) = \int_0^1 t^{2m} \exp(-zt^2) dt \quad (26)$$

with $z = \overline{PQ}^2/4\delta$, $\delta = 1/(4\gamma_1) + 1/(4\gamma_2)$, $\mathbf{P} = \sigma_A \mathbf{A} + \sigma_B \mathbf{B}$, $\mathbf{Q} = \sigma_C \mathbf{C} + \sigma_D \mathbf{D}$.

One important advantage of Eq. (1) is that the core part depending on the exponents (α_λ^A , α_μ^B , α_ν^C , α_ξ^D) is identical for all integrals for the shell-quadruplet, having a common set of angular momentum quantum numbers, (L_A , L_B , L_C , L_D). For example, 81 ($= 3^4$) individual integrals, ($p_x p_y | p_z p_x$), ($p_y p_z | p_y p_x$), and so on, have a common set of angular momentum quantum numbers. Once we calculate a core part for a set of angular momentum quantum numbers, (L_A , L_B , L_C , L_D), it can be used for all integrals for the shell-quadruplet having the same set of angular momentum quantum numbers. Utilizing this observation, we can reduce dramatically the cost of the calculation in the contraction step.

FLOP COUNT ESTIMATION FOR CALCULATION OF GENERALLY CONTRACTED ERIS

In this subsection, we give the FLOP count expression for generally contracted ERIs. First, let us consider the FLOP count expression for a usual contracted ERI (CERI).

A contracted GTO (CGTO) is a linear combination of primitive GTOs with all primitives having the same angular momentum and function center but different exponents,

$$\phi_p^A(\mathbf{r}) = \sum_{\lambda}^{K_A} d_{p\lambda}^A \phi_{\lambda}^A(\mathbf{r}), \quad (27)$$

where K_A is the degree of contraction, and $d_{p\lambda}^A$ are the contraction coefficients. Combining Eqs. (1) and (27), we obtain an expression for a contracted ERI:

$$(\phi_p^A \phi_q^B | \phi_r^C \phi_s^D) = \sum_{\mathbf{N}_3} C_4^{ABCD} \{\mathbf{N}_3\} \ddot{H}_{4pqrs}^{ABCD} \{\mathbf{N}_3\}, \quad (28)$$

where \ddot{H}_{4pqrs}^{ABCD} is the contracted core part including the factors $S_{\lambda\mu}^{AB}$ and $S_{\nu\xi}^{CD}$,

$$\ddot{H}_{4pqrs}^{ABCD} \{\mathbf{N}_3\} = \sum_{\lambda}^{K_A} \sum_{\mu}^{K_B} d_{p\lambda}^A d_{q\mu}^B S_{\lambda\mu}^{AB} \\ \times \sum_{\nu}^{K_C} \sum_{\xi}^{K_D} d_{r\nu}^C d_{s\xi}^D S_{\nu\xi}^{CD} H_{4\lambda\mu\nu\xi}^{ABCD} \{\mathbf{N}_3\}. \quad (29)$$

The quartet summation in Eq. (29) can be regarded as the double summation for (ν, ξ) in the double summation for (λ, μ) . These double summations are named the K_{BRA} and K_{KET} summations, where $K_{\text{BRA}} = K_A K_B$ and $K_{\text{KET}} = K_C K_D$. Then the K_{BRA} and K_{KET} summations can be split into two steps:

$$\ddot{H}_{4pqrs}^{ABCD} \{\mathbf{N}_3\} = \sum_{\lambda}^{K_A} \sum_{\mu}^{K_B} d_{p\lambda}^A d_{q\mu}^B S_{\lambda\mu}^{AB} \dot{H}_{4\lambda\mu,rs}^{ABCD} \{\mathbf{N}_3\}, \quad (30)$$

$$\dot{H}_{4\lambda\mu,rs}^{ABCD} \{\mathbf{N}_3\} = \sum_{\nu}^{K_C} \sum_{\xi}^{K_D} d_{r\nu}^C d_{s\xi}^D S_{\nu\xi}^{CD} H_{4\lambda\mu\nu\xi}^{ABCD} \{\mathbf{N}_3\}. \quad (31)$$

Considering the FLOP count in Eqs. (28), (30), and (31), the total FLOP count for calculating a CERI can be expressed as a function of K_{BRA} and K_{KET} ,

$$xK_{\text{BRA}} K_{\text{KET}} + yK_{\text{BRA}} + z. \quad (32)$$

The coefficients of x , y , and z are called FLOP count parameters [11]. The term, $xK_{\text{BRA}} K_{\text{KET}}$, is the FLOP count for K_{KET} summations for all in-

dices in $\{N_3\}$, λ ($1 \leq \lambda \leq K_A$) and μ ($1 \leq \mu \leq K_B$). The term yK_{BRA} is the FLOP count for K_{BRA} summations for all indices in $\{N_3\}$ after K_{KET} summations are finished. The constant value z , depending on neither K_{BRA} nor K_{KET} , is the FLOP count for calculating a CERI after K_{BRA} and K_{KET} summations for all indices in $\{N_3\}$ are finished. Suppose that all contraction lengths are equal, i.e., $K_A = K_B = K_C = K_D = K$, the FLOP count expression can then be given with K by

$$xK^4 + yK^2 + z, \quad (33)$$

which gives a plain representation of the order for the FLOP count expression.

Again, note that the ACE-b3k3 formula gives the best FLOP count parameters concerned with the contraction length, that is, the values of the x and y parameters in Eq. (33) are small, and thus the total FLOP count for large K is reduced.

Now, let us consider the FLOP count expression for generally contracted ERIs (GCERIs). We here introduce FLOP count expressions for the conventional algorithm and for our refined algorithm.

A set of generally contracted GTOs (GCGTOs),

$$\{\phi_p^A(\mathbf{r}) \mid 1 \leq p \leq N_A\} \quad (34)$$

is composed of N_A 's GCGTOs. Each is a linear combination of the same set of primitive GTOs with a different set of contraction coefficients. A set of GCERIs is given by

$$\left\{ (\phi_p^A \phi_q^B \mid \phi_r^C \phi_s^D) \mid 1 \leq p \leq N_A, 1 \leq q \leq N_B, 1 \leq r \leq N_C, 1 \leq s \leq N_D \right\}, \quad (35)$$

which contains $(N_{\text{BRA}} N_{\text{KET}})$'s GCERIs. ($N_{\text{BRA}} = N_A N_B$ and $N_{\text{KET}} = N_C N_D$.)

In the conventional algorithm, to obtain a set of GCERIs, all the GCERIs are calculated one by one, with a routine for computing a single CERI. Thus the FLOP count for calculating all the GCERIs is given by

$$N_{\text{BRA}} N_{\text{KET}} (xK_{\text{BRA}} K_{\text{KET}} + yK_{\text{BRA}} + z) \quad (36)$$

$$= N^4 (xK^4 + yK^2 + z). \quad (37)$$

However, if we use the ACE-b3k3 formulas, we can utilize the fact that $S_{\nu\xi}^{CD} H_{4\lambda\mu\nu\xi}^{ABCD}(\mathbf{N}_3)$ in Eq. (31) is identical for each GCERI, since it includes no parameters depending on the contraction coefficients. Therefore the above FLOP count expression

in Eqs. (36) and (37) can be reduced to

$$(x_2 N_{\text{KET}} + x_0) K_{\text{BRA}} K_{\text{KET}} + (y_4 N_{\text{BRA}} N_{\text{KET}} + y_2 N_{\text{BRA}} + y_0) K_{\text{BRA}} + z_4 N_{\text{BRA}} N_{\text{KET}} \quad (38)$$

$$= (x_2 N^2 + x_0) K^4 + (y_4 N^4 + y_2 N^2 + y_0) \times K^2 + z_4 N^4. \quad (39)$$

Note that the FLOP count for the K_{KET} summation has been effectively reduced by $1/N^2$ order. The loop structures being used in conventional algorithms and our algorithm are illustrated in Figures 1 and 2. It can be easily seen that the conventional algorithm shown in Figure 1 wastes time by performing the same K_{KET} summations N_{BRA} times. On the other hand, in our refined algorithm, shown in Figure 2, the necessary K_{KET} summations are performed just once outside the N_{BRA} loop and stored in the computer memory.

To show that the ACE-b3k3 method is sufficiently adaptable to obtain GCERIs more efficiently, we present some numerical estimations of the FLOP count. Tables I and II list the FLOP count parameters of all classes of four-centered GCERIs over s and p GCGTOs for our algorithm and the conventional algorithm with the ACE-b3k3 formula. The FLOP count for the molecular incomplete gamma function [Eq. (26)] is not included.

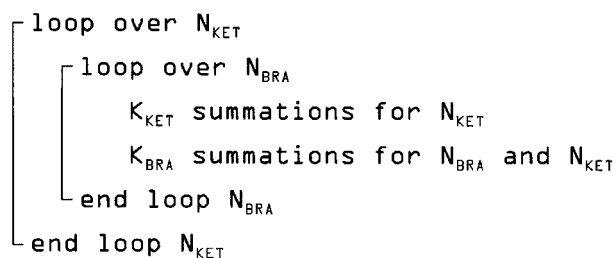


FIGURE 1. Loop structure of the conventional algorithm to obtain GCERIs.

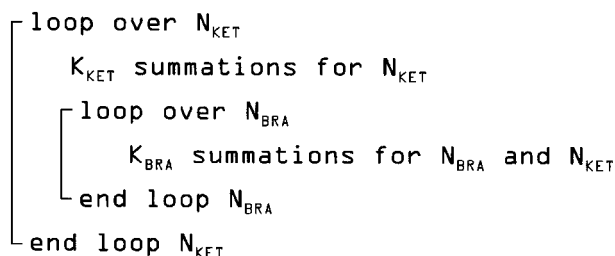


FIGURE 2. Loop structure of our algorithm to obtain GCERIs.

TABLE I
FLOP count parameters of algorithm with ACE-b3k3 formula.^{a,b}

ERI class	x_2	x_0	y_4	y_2	y_0	z_4
(ss ss)	2	0	2	0	0	0
(ps ss)	4	4	6	1	0	15
(ss ps)	6	3	6	2	0	15
(pp ss)	10	12	18	4	0	93
(ss pp)	12	7	18	10	1	93
(ps ps)	12	12	20	8	0	93
(pp ps)	24	26	54	27	0	439
(ps pp)	24	22	54	29	3	439
(pp pp)	44	41	134	80	3	1908

^aIgnoring the calculation of $F_m(z)$.^bTotal FLOP count is given by Eq. (38).**TABLE II**
FLOP count parameters of conventional algorithm with ACE-b3k3 formula.^{a,b,c}

ERI class	x	y	z
(ss ss)	3	0	0
(ps ss)	8	3	15
(ss ps)	7	7	15
(pp ss)	19	11	93
(ss pp)	14	26	93
(ps ps)	19	22	93
(pp ps)	38	61	439
(ps pp)	34	69	439
(pp pp)	63	167	1908

^aIgnoring the calculation of $F_m(z)$.^bTotal FLOP count is given by Eq. (36).^cTwo FLOPs for $(d_{\rho\lambda}^A d_{q\mu}^B)(d_{r\nu}^C d_{s\xi}^D)F_m(z)$ are included into x , while these are counted for making $F_m(z)$ in a previous work (Ref. [3]).**TABLE III**
Total FLOP count of algorithm in case $K = 10$ and $3 \leq N \leq 6$.^{a,b}

ERI class	$N = 3$	$N = 4$	$N = 5$	$N = 6$
(ss ss)	196000 (12.4)	371000 (20.7)	625000 (30.0)	979000 (39.7)
(ps ss)	451000 (12.7)	839000 (21.5)	1427000 (31.0)	2281000 (40.2)
(pp ss)	1177000 (9.8)	2211000 (16.5)	3813000 (23.4)	6188000 (29.9)
(ps ps)	1377000 (11.3)	2589000 (19.0)	4448000 (27.0)	7181000 (34.7)
(pp ps)	2880000 (9.8)	5601000 (15.9)	9942000 (21.8)	16532000 (27.2)
(pp pp)	5682000 (9.2)	11498000 (14.4)	21181000 (19.1)	36384000 (23.1)

^aIgnoring the calculation of $F_m(z)$.^bThe rate of speed-up over the conventional algorithm is given in parenthesis.

Table III shows the FLOP counts estimated by Eq. (39) for our algorithm in the case $K = 10$ and $3 \leq N \leq 6$. The rates of speed-up over the conventional algorithm are given in parentheses in the table.

Our algorithm gives small values of the x and y parameters throughout all classes of GCERIs. This result is advantageous for GCGTOs, since they are contracted to a high degree in many cases. In the case of low-angular-momentum GCGTOs, the calculation of the GCERIs speeds up by almost N^2 order. For the (ss | ss) class, the calculation of which is fastest, the FLOP count expression by Eq. (39) for the refined algorithm is $(2N^2K^4 + 2N^4K^2)$, while the FLOP count expression by Eq. (37) for the conventional algorithm is $(3N^4K^4)$. As seen from these FLOP count expressions, most of the calculation of (ss | ss) is spent on K_{KET} summations. Therefore the performance by N^2 order works more effectively when the values of the y and z parameters are small. On the other hand, when the angular momentum becomes higher, the y and z parameters increase so sharply that the term $(yN^4K^2 + zN^4)$, which is included in both of the FLOP count expressions by Eqs. (39) and (37), becomes too large to be ignored.

Tables IV and V show the FLOP count parameters of (ps | ps) and (pp | pp) for the case that some of the four function centers, **A**, **B**, **C**, **D**, have the same coordinates, namely overlap. There are 14 cases: the one-center case is (AA | AA); two-center cases are (AA | CC), (AB | AB), (AB | BA), (AA | AD), (AA | CA), (AB | AA), and (AB | BB); three-center cases are (AA | CD), (AB | AD), (AB | CA), (AB | BD), (AB | CB), and (AB | CC). For (AA | CD), the relation **AB** = 0 can be substituted for **AB** in the accompanying coordinate part in Eq. (7).

TABLE IV
FLOP count parameters of our algorithm for $(ps | ps)$ considering the overlaps of function centers.^{a,b}

Overlap type		x_2	x_0	y_4	y_2	y_0	z_4
4 centers	(AB CD)	12	12	20	8	0	93
3 centers	(AA CD)	4	4	10	6	0	48
	(AB AD)	10	13	10	5	1	49
	(AB CA)	12	12	10	9	0	48
	(AB BD)	10	13	10	7	1	48
	(AB CB)	12	12	10	12	0	48
	(AB CC)	10	10	10	1	0	48
2 centers	(AA CC)	4	4	4	2	0	12
	(AB AB)	8	13	4	5	0	12
	(AB BA)	10	13	4	7	0	13
	(AA AD)	4	4	4	4	1	12
	(AA CA)	4	4	4	6	0	12
	(AB AA)	4	8	4	2	0	12
	(AB BB)	4	9	4	2	0	12

^aIgnoring the calculation of $F_m(z)$.

^bTotal FLOP count is given by Eq. (38).

This simplifies the formula and, particularly, makes the values of the y and z parameters much smaller for high angular momentum. The case of (AA | CC) is most effective as shown in Tables VI and VII. For $(pp | pp)$, the FLOP count of (AA | CC) is about six times smaller than that of the four-center case. For $(ps | ps)$ the FLOP count is about three times smaller than that of the four-center case. To calculate GCERIs faster, it should be effective, especially for high-order GCGTOs, to provide individual routines for each case of overlapping.

TABLE VI
Total FLOP count of algorithm for $(ps | ps)$ considering the overlaps of function centers in case $K = 10$ and $3 \leq N \leq 6$.^a

Overlap type		$N = 3$	$N = 4$	$N = 5$	$N = 6$
4 centers	(AB CD)	1377000	2589000	4448000	7181000
3 centers	(AA CD)	490000	958000	1710000	2860000
	(AB AD)	1119000	2007000	3298000	5108000
	(AB CA)	1121000	2009000	3303000	5113000
2 centers	(AA CC)	435000	788000	1302000	2021000
	(AB AB)	887000	1523000	2400000	3561000
	(AB BA)	1070000	1847000	2906000	4290000
	(AA CA)	439000	795000	1312000	2035000
	(AA AD)	436000	790000	1305000	2025000

^aIgnoring the calculation of $F_m(z)$.

TABLE V
FLOP count parameters of algorithm for $(pp | pp)$ considering the overlaps of function centers.^{a,b}

Overlap type		x_2	x_0	y_4	y_2	y_0	z_4
4 centers	(AB CD)	44	41	134	80	3	1908
3 centers	(AA CD)	12	11	42	33	3	570
	(AB AD)	44	42	66	91	3	885
	(AB CC)	40	40	22	13	0	576
2 centers	(AA CC)	12	11	12	8	1	87
	(AB AB)	38	46	18	79	0	142
	(AA AD)	12	11	12	25	5	101
	(AB AA)	14	35	12	7	1	103

^aIgnoring the calculation of $F_m(z)$.

^bTotal FLOP count is given by Eq. (38).

Computational Results and Discussion

A program code based on our algorithm was implemented and tested using some practical molecular calculations. The loop structure of our program is shown in Figure 3. The routines listed in Figures 3 and 2 give the same total FLOP count. However, considering memory caching, the amount of memory required in what we call the K^4 step has been reduced in the routine of Figure 3. In our program we use Ishida's algorithm shown in Ref. [3] to compute the molecular incomplete gamma function.

The routines for calculating GCERIs for $(ss | ss)$, $(ps | ss)$, $(pp | ss)$, $(ps | ps)$, $(pp | ps)$, and $(pp | pp)$ have been implemented in the present case.

TABLE VII
Total FLOP count of algorithm for $(pp | pp)$ considering the overlaps of function centers in case $K = 10$ and $3 \leq N \leq 6^a$

Overlap type	$N = 3$	$N = 4$	$N = 5$	$N = 6$
4 centers $(AB CD)$	5682000	11497000	21177000	36377000
3 centers $(AA CD)$	1606000	3304000	6174000	10731000
$(AB AD)$	5068000	9522000	16325000	26286000
2 centers $(AA CC)$	1302000	2372000	3934000	6127000
$(AB AB)$	4108000	7163000	11371000	16941000
$(AA AD)$	1318000	2404000	3986000	6207000

^aIgnoring the calculation of $F_m(z)$.

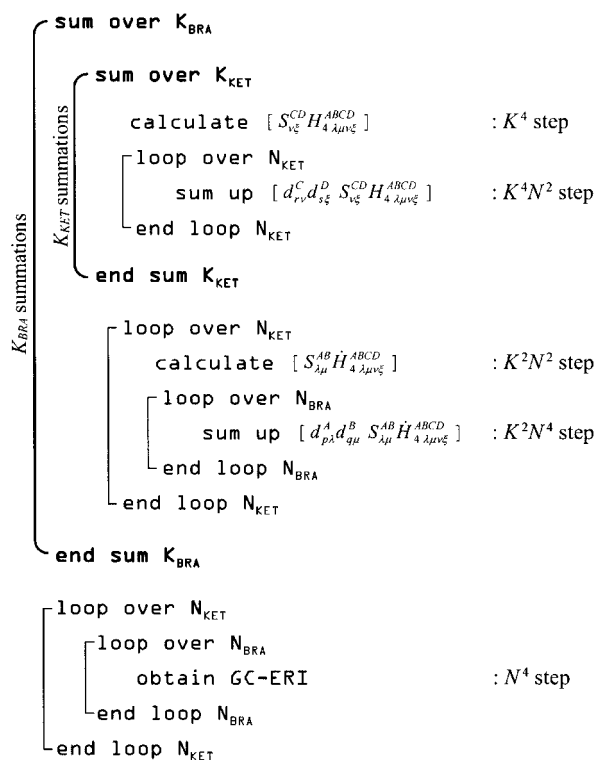


FIGURE 3. Loop structure of our program to obtain GCERIs.

We measured the central processing unit (CPU) time for calculating a set of GCERIs over GCGTOs to exhibit the computational performance of our algorithm. The size of the GCGTOs used in the calculations is the same as that of the GCGTOs used in the FLOP count estimation. By doing so, the theoretical and realistic computational efficiency could be compared directly. All the calculations were carried out on a single node of IBM SP2(133 MHz) clusters. The CPU time does not include input/output (I/O) operation time.

The four-center GCERIs are evaluated with the GCGTOs $K = 10$ and $3 \leq N \leq 6$. Table VIII lists the CPU times of our algorithm and the rates of speed-up (in parentheses) over the conventional algorithm. It can be seen that the efficiency measured by CPU time far exceeds N^2 order, which is the suggestion from theoretical assessments in the previous section. Especially for $(ss | ss)$, the rates of speed-up obey the order of almost N^4 . A similar behavior is seen in the other results of CPU time. This bonus improvement of the computational efficiency originates from the cost of computing the molecular incomplete gamma function, which is ignored in the FLOP count estimation in the previous section. In our refined algorithm, the

TABLE VIII
CPU times of algorithm in case $K = 10$ and $3 \leq N \leq 6$ (in millisecond).^a

ERI class	$N = 3$	$N = 4$	$N = 5$	$N = 6$
$(ss ss)$	22 (70.3)	26 (188)	31 (385)	40 (618)
$(ps ps)$	49 (54.6)	66 (128)	116 (177)	189 (226)
$(pp pp)$	139 (27.3)	300 (40.0)	595 (49.3)	1090 (55.8)

^aThe rate of speed-up over the conventional algorithm is given in parenthesis.

necessary molecular incomplete gamma function is computed just once in the K^4 step in Figure 3. On the other hand, the conventional algorithm carries out the same computation of the molecular incomplete gamma function N^4 times. If we use the FLOP count for the molecular incomplete gamma function, it can be added into the parameters x_0 and y_0 for Eq. (39) and the parameters x and y for Eq. (37). In Ishida's algorithm the FLOP count for the molecular incomplete gamma function is given to be $(16 + S)K^4 + 5K^2$ for $(ss | ss)$, $(29 + S)K^4 + 5K^2$ for $(ps | ps)$, and $(35 + S)K^4 + 5K^2$ for $(pp | pp)$, where the parameter S is the FLOP count for one square-root calculation and dependent on computer environments. We measured a CPU time for one square-root calculation and for one floating-point multiplication, and then estimated a condition $S = 65$ on our computer environment. To show the numerical effect on the FLOP count, including the calculation of the molecular incomplete gamma function, Table IX lists the rate of speed-up for the total FLOP count, assuming the condition $S = 65$ compared with that in the measured CPU time. It can be seen that the rate of speed-up in FLOP count estimation show good agreement with that in the computational measurement. As this result shows, the computation of the incomplete gamma function is a vital part of evaluating the ERIs for all segmented contraction methods. However, it is a trivial part in the present method. This feature confers a very important advantage, which can give an efficiency of almost N^4 order for $(ss | ss)$, when the GCGTOs have low angular momentum, i.e., the values of the y and z parameters are small.

Tables X and XI show the computation times of GCERIs $(ps | ps)$ and $(pp | pp)$ considering the overlaps of the function centers. Similarly to the

TABLE X
CPU times of our algorithm for $(ps | ps)$ considering the overlaps of the function centers in case $K = 10$ and $3 \leq N \leq 6$ (in milliseconds).

Overlap type		$N = 3$	$N = 4$	$N = 5$	$N = 6$
4 centers	$(AB CD)$	49	66	116	189
3 centers	$(AA CD)$	38	45	64	103
	$(AB AD)$	46	58	80	130
2 centers	$(AB CA)$	45	58	80	123
	$(AA CC)$	38	43	52	70
	$(AB AB)$	42	50	64	86
	$(AB BA)$	44	55	72	94
	$(AA CA)$	37	41	53	66
	$(AA AD)$	36	43	52	68

TABLE XI
CPU times of algorithm for $(pp | pp)$ considering the overlaps of function centers in case $K = 10$ and $3 \leq N \leq 6$ (in milliseconds).

Overlap type		$N = 3$	$N = 4$	$N = 5$	$N = 6$
4 centers	$(AB CD)$	139	300	595	1090
3 centers	$(AA CD)$	62	104	204	359
	$(AB AD)$	122	226	418	697
2 centers	$(AA CC)$	56	76	113	188
	$(AB AB)$	101	154	255	400
	$(AA AD)$	57	74	113	174

FLOP count estimation, we examine the case of $(AA | CC)$. The CPU time shows a similar efficiency as the FLOP count estimation. The case of $(AA | CC)$ is the fastest. For $(pp | pp)$, the CPU time of $(AA | CC)$ is about six times as fast as that of the four-center case. For $(ps | ps)$ the CPU time is about three times as fast as that of the four-center case. These results also show that the programs

TABLE IX
Comparison with theoretical efficiency, including the molecular incomplete gamma function calculation and computational efficiency in case $K = 10$ and $3 \leq N \leq 6$.

ERI class		$N = 3$	$N = 4$	$N = 5$	$N = 6$
$(ss ss)$	FLOP rate	67.6	182	365	608
	CPU time rate	70.3	188	385	618
$(ps ps)$	FLOP rate	39.6	82	131	180
	CPU time rate	54.6	128	177	226
$(pp pp)$	FLOP rate	20.0	33.8	46.4	57.1
	CPU time rate	27.3	40.0	49.3	55.8

specific to each case may work with the good performance predicted from the FLOP count.

Finally, we measured the CPU time for computing all the ERIs in a benzene molecule, as an example of a realistic molecular orbital (MO) calculation. Table XII shows the results given for two kinds of basis sets for carbon and hydrogen atoms: (A) C:14s9p/3s2p, H:8s4p/2s1p, and (B) C:14s9p/6s5p, H:8s4p/4s3p. Here the basis sets specified with the format, $(K_s)s(K_p)p/(N_s)s(N_p)p$ have (N_s) 's s -type GCGTOs containing (K_s) 's primitive GTOs and (N_p) 's p -type GCGTOs containing (K_p) 's primitive GTOs. The number of functions in the basis sets is (A) 84 and (B) 204. The calculations were also performed with the ERI routine of Gaussian94 [12] (using the PRISM [11] method) and HONDO [13] (using the PH method [10]). Since these routines can treat only a segmented contraction scheme, we give these basis sets as segmentally CGTO to the program. In Table XII, the ratio of CPU time of the present routine to that of the routine in Gaussian94 is (A) 2.5 and (B) 28. The performance of our program is very good compared with conventional programs, especially when the numbers of GCGTOs, namely N_s and N_p , are large. The ratio of the CPU time for the two basis sets, (A) and (B), is about 2 in the present program, about 23 in Gaussian94 and HONDO. With our routine, fast calculation of ERIs over high-quality basis sets can be performed.

Summary

We proposed a new algorithm and developed programs for the efficient computation of a set of GCERIs over s and p GCGTOs. We utilized the

ACE-b3k3 method developed by Ishida, which can rapidly calculate a highly contracted ERI. In our routine, the necessary K_{KET} summation, or K^4 summations, are performed just once outside the N_{BRA} loop, and stored in computer memory. The conventional calculation of GCERIs used in many ab initio packages unnecessarily calculates the identical K^4 terms N^2 times. By the described refinement, we obtain efficiency by N^2 order for the FLOP count estimation with s and p GCGTOs. The FLOP count estimation shows that the calculation of GCERIs is effective when the GCGTOs have low angular momentum.

The performance measured by CPU time exceeds that obtained by the theoretical estimation and shows an efficiency of almost N^4 order in $(ss|ss)$. Our program can work more effectively than the theoretical estimation does, since it is a trivial cost to compute the molecular incomplete gamma functions in our algorithm.

When the function centers of GCGTOs overlap each other, the relations of coordinates can simplify the formulas and make the values of the y and z parameters small. When the GCGTOs have high angular momentum, this simplification of the y and z parameters is very effective.

Very recently, Ishida has shown that the ACE-b3k3 method becomes more effective with a high order of GTO [8, 9], while the present work demonstrates that the effectiveness with the general contraction scheme works well even for a low order of GTO.

We expect that our new algorithm and routine for GCERIs will open a new world in the ab initio molecular orbital calculation for large-scale systems.

TABLE XII
CPU times of algorithm for computing all ERIs in benzene molecule (in seconds).

	Basis sets	Present	Gaussian ^{a, b}	HONDO ^{a, c}
(A)	C:14s9p/3s2p H:8s4p/2s1p	600	2450	5740
(B)	C:14s9p/6s5p H:8s4p/4s3p	1180	57100	137800

^aIncluding prescreening, indexing, packing, and sorting.

^bUsing the PRISM method [11].

^cUsing the PH method [10].

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