

On solving Schrödinger equation for the ground state of a two-electron atom using genetic algorithm

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A recipe is proposed for solving the radial Schrödinger equation (SE) for ground state of helium atom using genetic algorithm. A fitness landscape is generated and the problem of solving the radial SE is reduced to a search for the maximum on this landscape.

THE use of genetic algorithms (GAs)^{1,2} for solving the Schrödinger equation (SE) numerically, is of contemporary origin³⁻⁹. The primary motivation has been to look

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for a viable basis-set-free technique of solving the SE and generating optimal numerical wave functions. Indeed, much of quantum chemistry is dominated by the quality of basis sets, independent of whether one is using the Hartree–Fock, multiconfiguration Hartree–Fock, configuration interaction or coupled cluster methods. GA-based techniques, if proved viable, can change the scenario, for accurate numerical wave functions of atoms and few electron molecules could pave the way for designing optimal basis functions for different chemical environments. Secondly, GA is inherently parallelizable and can be therefore rather cost-effective. The basic philosophy of the method is to recast the energy eigenvalue problem in the form of a search for a global maximum on a defined fitness landscape. The targeted solution wave functions are defined as discrete functions representing the distribution of probability amplitudes in the coordinate space. A population of such wave function strings is created to start with and is made to evolve on the fitness landscape under the action of appropriately constructed genetic operators like selection, crossover, mutation, etc. In our formulation, the amplitudes are floating-point numbers and the wave functions are strings of floating-point numbers. The strings are normalized and obey appropriate boundary conditions of the problem.

The radial SE for He atom reads,

$$H\mathbf{y}_n(r_1, r_2) = E_n\mathbf{y}_n(r_1, r_2),$$

where

$$H = -\frac{\hbar^2}{2m_e} \sum_{i=1,2} \left(\frac{\partial^2}{\partial r_i^2} + \frac{2}{r_i} \frac{\partial}{\partial r_i} \right) - \sum_{i=1,2} \frac{ze^2}{r_i} + \frac{e^2}{r_{12}}. \quad (1)$$

From a definition of the Hamiltonian, it is evident that we are taking into account only radial correlation, leaving out the angular correlation altogether. The target is to reach the S -limit form of $\mathbf{y}_0(r_1, r_2)$.

We represent \mathbf{y} on a uniformly discretized two-dimensional coordinate space by n strings ($s_1, s_2, s_3, \dots, s_k, \dots, s_n$), each string representing a collection of probability amplitudes [$s(i, j)$] in a two-dimensional array of $n_1 \times n_2$ grid points. The square of $s(i, j)$ denotes the probability of finding one electron at r_1^i , while the other is at r_2^j , irrespective of spin. The permutation symmetry of the wave function \mathbf{y} now needs to be considered. Since helium ground state is spin-singlet (1s_0), the space part of the ground-state wave function of He [$\mathbf{y}(r_1, r_2)$] must be symmetric with respect to interchange of the coordinates of electrons 1 and 2. Therefore, the wave function strings are made to obey the following condition:

$$S_k(r_1^i, r_2^j) = S_k(r_2^i, r_1^j), k = 1, 2, \dots, n. \quad (2)$$

The fitness landscape is generated by defining a fitness function (f_k) for the k th string as follows:

$$f_k = e^{-S\Box_k}, \quad (3)$$

where

$$\Box_k = \left\{ \frac{\langle \mathbf{y}_k | H | \mathbf{y}_k \rangle}{\langle \mathbf{y}_k | \mathbf{y}_k \rangle} - E_l \right\}^2. \quad (4)$$

S is essentially a scaling parameter that takes care of dimensional requirements and prevents exponential overflow or underflow. E_l may be kept fixed, if a good estimate of lower bound is available or may be updated ($E_l^{i+1} = E_l^i \pm c \sqrt{\Box_i}$, $c = 0.25-0.75$). For actual calculation, \mathbf{y}_k in eq. (4) is replaced by the string S_k and the integrations are replaced by multidimensional quadrature. E_l is an estimated lower bound to the energy of the k th string. Fitness values for all the n strings are calculated and each string is subjected to a fitness-proportional roulette wheel selection procedure² that allows more copies of the better-solution string to pass into the mating pool. The average fitness of the population increases after selection – but no new information is created at this stage. For creating new information or new strings, two kinds of genetic operators are invoked, viz. crossover and mutation – the former occurring with a probability p_c and the latter with p_m . Let the strings s_k and s_l be randomly selected with probability p_c for undergoing crossover and let the i th row and the j th column of the arrays s_k, s_l be selected for crossover again with probability p_c . As a result of this operation, a pair of new strings – the children strings s'_k and s'_l are created, where

$$\left. \begin{aligned} S'_k(p, q) &= f S_k(p, q) + (1-f)(S_l(p, q)) \\ S'_l(p, q) &= f S_l(p, q) + (1-f)(S_k(p, q)) \end{aligned} \right\} \quad (5)$$

for $p = 1, 2, \dots, i; q = 1, j$,

while for $p > i, q > j$, however,

$$\left. \begin{aligned} S'_k(p, q) &= S_k(p, q) \\ S'_l(p, q) &= S_l(p, q) \end{aligned} \right\} \quad (6)$$

The mixing coefficient f is randomly chosen from a range ($0 < f < 1$). We may point out that the specific form of crossover operation used here is dictated by the physics of the problem. Since the region near the nucleus is energetically important, it is necessary to ensure that amplitudes for small values of r ($r = 0 \leftrightarrow \infty$) are frequently sampled by the crossover operator (CO). The redefinition of the CO used here has been found to be beneficial⁶. After crossover, each of the children strings is subjected to a process of mutation with probability p_m . The site for mutation is chosen by comparing a random number $r[0, 1]$ with p_m for each pair of column and row indices (i, j). If for the k th string $r < p_m$ for $i = p, j = q$, the corresponding amplitude is mutated as follows:

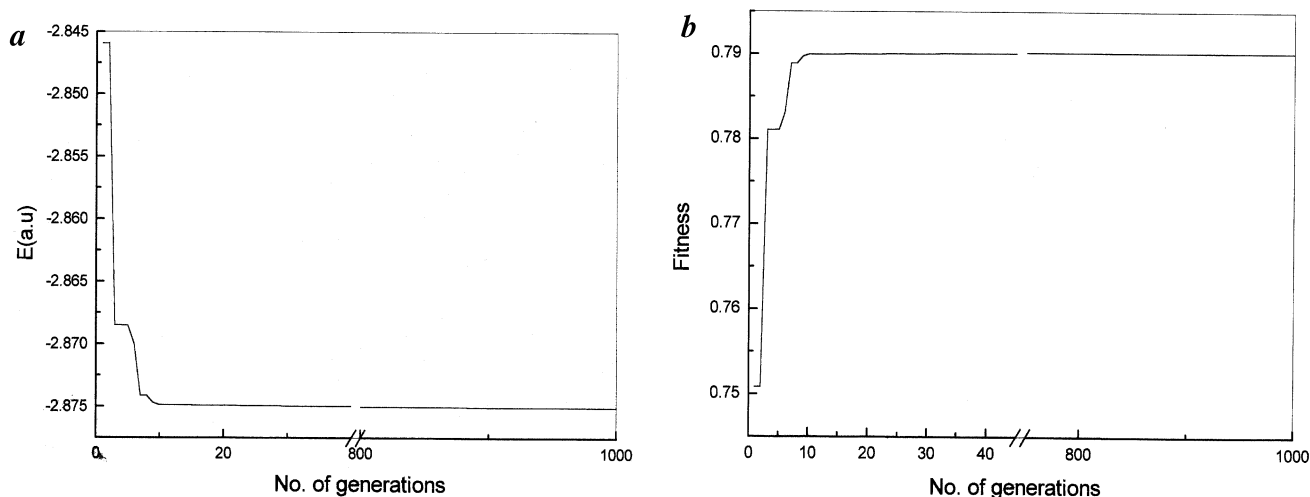


Figure 1. *a*, Evolution of energy of ground state of He atom during GA run. Energy refers to the string of the highest fitness in the population. *b*, Fitness profile during GA run for He atom in the ground state. Fitness corresponds to the string of the highest fitness in any generation.

$$S_k''(p, q) = S_k'(p, q) + (-1)^l r \Delta_m, \quad (7)$$

where r is a random number ($0 \leq r \leq 1$), l is a random integer and Δ_m is the intensity of mutation. It may vary with generation or remain static⁶.

The last operator to act on the post-crossover and post-mutation strings is the symmetrization operator, the action of which is defined as follows:

$$S_k'''(q, p) = S_k''(p, q), \text{ for } k = 1, 2, \dots, \text{npop}, \quad (8)$$

for $p = 1, 2, \dots, n_1$; $q = 1, 2, \dots, n_2$; where 'npop' is the number of strings in the population.

With symmetrization, one generation is said to have elapsed. From the post-symmetrization population, we choose 80% of the strings in order of fitness and the remaining 20% are randomly created. The sequence of operations outlined is repeated till the average fitness of the population does not improve any more.

We report here the results of application of the method to the ground state of He atom. For this case \mathbf{y} has been represented as amplitude distribution function on a two-dimensional grid of 20 a.u. of length in each dimension, each dimension having 500 uniformly distributed grid points. A string s_k thus carries 500×500 grid-point amplitudes which are allowed to evolve explicitly under the action of genetic operators, while the quadratures are performed over 1500×1500 grid-point amplitudes for more accurate evaluation of the integrals in eq. (4) for calculating the fitness of a string. The additional amplitudes are generated by two-dimensional bicubic interpolation. We have used a population size of ten throughout. The amplitude distributions on the 2D grid representing the strings were chosen from functions of the type

$$\mathbf{y} = N e^{-b(r_1+r_2)} \mathbf{c}(r_{12}), \quad (9)$$

with $\mathbf{c}(r_{12}) = 1 + \mathbf{g}r_{12} + \mathbf{d}r_{12}^2 + \dots$, with randomly chosen values of $\mathbf{b}, \mathbf{g}, \mathbf{d}, \dots$. After the selection phase of evaluation is executed, a pair of strings is chosen randomly with crossover probability $p_c = 0.75$, for crossover. The mutation operation is then carried out on the post-crossover strings, followed by spatial symmetrization. Figure 1 *a* and *b* shows the energy and fitness profiles during the evolution of the strings representing the ground state wave function of helium atom. The major lowering of energy takes place in the first 15 generations. The rapid improvement of energy and therefore fitness in the early stages of evolution is dominated by the crossover operator. Towards the end of the evolution, improvement in fitness is dominated by mutation which accounts for slow improvement of the fitness values. The two-dimensional contour plot of the converged ground-state radial wave function of He atom obtained by GA is shown in Figure 2. Since this is an *S*-state and only radial correlation is present, the probability density is high in the near nucleus region. The energy corresponding to the best string is -2.87505 a.u. (1 a.u. of energy = 27.209 eV), which matches with ground state *s*-limit energy of He atom¹⁰. The energy obtained by GA is a bit lower than the Hartree–Fock energy of the He atom in the ground state. Results obtained so far indicate that the quality of results does not deteriorate with the increase of nuclear charge (z). The GA-based recipe described here is computationally at least as viable as numerical Hartree–Fock method for two-electron atoms. When the number of electrons (n) rises, the problem of performing multidimensional quadratures may prove to be a bottleneck. For $n > 2$, therefore, we must explore Monte Carlo methods for evaluating multidimensional integrals along with parallelization.

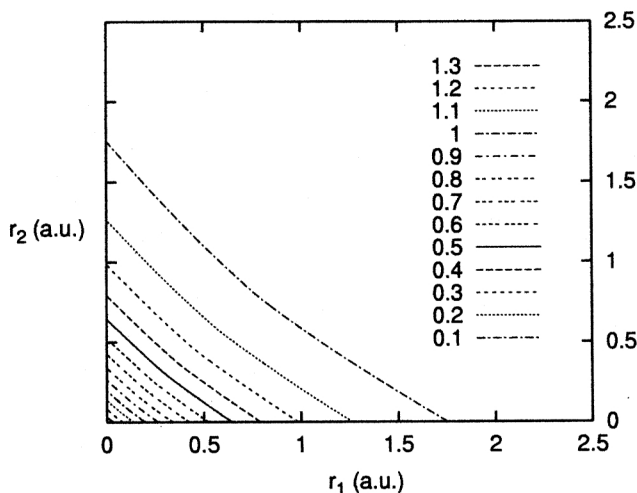


Figure 2. Two-dimensional contour plots of wave function amplitudes for He atom in the ground state.

We are in the process of completing the calculations on the entire helium sequence and extending the calculations to the fully correlated systems. We will shortly return to these results.

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