



An artificial immune algorithm for ergonomic product classification using anthropometric measurements



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ABSTRACT

Product classification using anthropometric measurements leads to ergonomic product design and user satisfaction. We propose an effective artificial immune algorithm (AIA) to classify ergonomic products with multi-criteria anthropometric measurements and tune the AIA parameters with a full factorial experimental design approach. We demonstrate the applicability and efficacy of the proposed algorithm by considering the anthropometric measurements of the hand, developing an ergonomic computer mouse, and classifying consumers into three categories. The resulting classifications are compared with expert opinions to facilitate the conformity of the computer mouse to user requirements.

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1. Introduction

Ergonomics is the science of fitting a job to a worker and a product to a user [59]. One of the basic challenges facing ergonomics is preventive health care. The goal of averting occupationally-generated musculoskeletal disorders and related illnesses can be attained by a suitable design of workplace and product [63]. Anthropometry is a field of science that deals with body measurements. It is associated with the physical characteristics of humans in different situations, especially with measurements of body size, shape, strength, and working capacity [59,55]. In the related literature, anthropometry is mentioned as a very consequential division of ergonomics because misalliances between human anthropometric dimensions and equipment dimensions may lead to inconvenience, plunging productivity, accidents, biomechanical stress, fatigue, injuries, and cumulative traumas [48,36,56]. Jung et al. [33] expressed that operator's anthropometry, task geometry, and design factors would be central components for an optimal ergonomic product design. There are a huge amount of hand tool

injuries each year worldwide which are partly related to worker-tool mismatches [1]. Pheasant [59] revealed that in order to design ergonomic products, as well as, to have compatibility between products and users, many factors should be considered. These factors include anthropometric characteristics of users, ways in which these characteristics impose constraints upon ergonomic product design process, and eventually, criteria that define a successful match between products and users. Hsu [29] developed accurate industrial standards and utilized an accurate girth ratio approach which is used to ascertain figure types enabling the production process in apparel manufacturing to be based on anthropometric data.

There are two types of anthropometric information including structural and functional information. Structural or static anthropometric information is linked to a body's measure or size in a fixed structural position and it is generally measured by marked anatomic points in a certain position. The second type of anthropometric information, which includes measures of access limits and is measured in practical conditions, is functional or dynamic anthropometric information [59]. Anthropometric measurements or morphologies of human body parts have been explored in a myriad of studies. Lacković et al. [43] presented a system for the purpose of force measurement in the legs and crutches based on a quantitative gait analysis of orthopaedic patients. Kieckhefer et al. [39] investigated ear print formation considering the morphology of an auricle and developed the appropriate optical equipment for this purpose.

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This article is primarily concerned with the anthropometric side of ergonomics, which considers matching the physical form and dimensions of a product to those of its users and, likewise, matching physical demands of a working task to the capacities of a workforce. In most real-world design problems, the concentration is on the users of products, while the structural or static anthropometric information is considered in this paper.

Humans in a population vary based on the proportion of their bodies. It has been shown in the literature that despite the goal of having 95% of the user population in the ergonomic product design interval, in fact 25% of the population will fall out of the design limit which means that only 75% of the population will be in the design limit [59,6]. This result indicates the significance of taking into account anthropometric data in the real ergonomic product design, and consequently the probable effects of the ergonomic product design on humans. For instance, Lacko et al. [42] utilized 3D anthropometric data as a feasible design method for brain-computer interfacing (BCI) headgear to create a one-size BCI headset prototype which fits all individuals. Chen et al. [10] have taken into account ergonomic considerations in order to design a wearable device for frozen shoulder rehabilitation. Garcia and Garcia-Mendoza [22] proposed an approach based on optimal ergonomic specifications in order to design mobility devices. Vincent et al. [69] investigated the constraints that influence the medical device design and development process. Wu et al. [71] applied a user-centered design approach to develop a hair washer for disabled people for use in normal postures. Martin et al. [49] reviewed the literature on methods for evaluating user requirements in ergonomics as well as examining factors that affect user requirements for medical device design.

Meanwhile, the classification of items can assist designers with conformity of tools/products and users/clients. Item classification has a variety of applications (e.g., clothing, desks, chairs, and other instruments humans utilize) which leads to a more comfortable user experience. Classification of technologies and products/services should facilitate the decision making process regarding realization of the system functionalities [57]. Classification in its most straightforward form attempts to assign some unknown objects into a known class of objects. Customarily, these classes are stereotypes which are hierarchically organized [12].

The classification problem has been applied in varied realms such as computer science, engineering, image processing, and data mining [74]. Classification also has empirical applications in performing medical diagnosis [67,68,5,51], fault detection [44,30], assignment of personnel into proper occupational categories based on their qualifications [62,24], and customer satisfaction evaluation and characteristics analysis of different groups of customers [17,65]. Discerned human related emotions should be considered as pivotal components in the ergonomic product design process of a product. However, customers' affective needs are arduous to perceive. Therefore, product designers often incorrectly comprehend what customers really want most of the time. Bahn et al. [4] proposed a design framework by considering critical affective features of customers on products, and systematically, incorporated these types of features into ergonomic product design attributes. This broad span of practical applications of classification motivates researchers to develop various methodologies for the classification problem. Nagarajan and Balasubramanie [54] utilized a neural-based classifier approach for object classification which detects car objects in the midst of a cluttered background. Ghazali et al. [23] introduced the Scale Invariant Feature Transform (SIFT) technique to overcome the problem of weed classification in a hands-on application of image processing. Neural networks [3,47,26], evolutionary algorithms [14,7,18], and fuzzy logic [31,64] are some other soft computing approaches that have been extensively applied to adaptively evolve solutions for classification

issues. The Genetic Algorithm (GA) method has attracted much attention and has become one of the most popular techniques for classification [50]. However, GA suffers from a number of weaknesses, particularly in larger scale real world classification problems, such as encountering inefficiency in searching a large space, finding it difficult to break internal interference of training data, and getting trapped in local optima [74].

Producers need to draw the attention of more consumers in the present-day competitive markets, and on the other hand, consumers' views are vital motivating factors for product development and obtaining profit in the market place. Hence, producers are required to design products that satisfy the requirements of the majority of people and also provide comfort when they are used based on ergonomic criteria [46]. However, only very limited researches have been carried out analyzing hand sizing, which is essential for the design of hand tools. Kwon et al. [41] developed a glove sizing system that was based on three hand dimensions (length, circumference, and breadth). Pekelney and Chu [58] described the ergonomic design criteria and product attributes of a computer mouse device. Hsiao et al. [28] proposed an improved sizing scheme for fire-fighting gloves that appropriately fits the US firemen population. In another study, Liu and Fan [45] considered four anthropometric measurements of the hand (width and length of hand, palm, and index) in their study to design an ergonomic computer mouse for clients with wrist splints.

Thus, the significance of a technique which considers various anthropometric criteria in designing an ergonomic product-computer mouse with all real-world limitations is completely obvious. To tackle the above mentioned difficulties, a meta-heuristic method, called the artificial immune algorithm (AIA), is applied. Its name reflects the fact that it is driven by the natural immune system of the human body which is an origin of inventiveness for resolving optimization problems. There are other methods such as principal component analysis (PCA) and the discriminant analysis technique which may be utilized in this case, but the AIA has the capability of learning and mimicking the categorizing ability of a human expert in the field. An AIA imitates the learning ability carried out by the natural immune system [35].

In this paper, the classification of an ergonomic computer mouse is studied considering multi-criteria anthropometric measurements using an AIA. A problem of user classification into small (S), medium (M), and large (L) categories for validation is considered. Here, the artificial immune system algorithm is applied to learn the weights of the criteria along with SM (Small-Medium) and ML (Medium-Large) cut-off points from pre-classified items based on the research by Zandieh et al. [73]. Anthropometric measurements of five areas of the hand are considered in this study. The classifications resulting from the AIA is compared with an expert opinion, one who has the full authority and expertise to classify items on the basis of the mentioned anthropometric data. The main aim of this study is to provide a method to facilitate the conformity of products to users according to anthropometric measurements.

The rest of this article is organized as follows. The problem description is presented in Section 2. The general mechanism of the artificial immune system and the proposed AIA are elaborated in Section 3. The computational results and parameters tuning of the AIA are presented in Section 4. Finally, the conclusion highlights are presented in Section 5.

2. Problem description

Wrist pain is a frequent ailment and many kinds of wrist pains are caused by sudden injuries which lead to sprains or fractures. Wrist pain can be brought about by more chronic problems as well,

such as repetitive stress, arthritis and carpal tunnel syndrome as well. When hand tools are bigger or smaller than the hand size, long-term use can cause wrist pain. Liu and Fan [45] considered four anthropometric measurements of the hand (width and length, palm, and index) in their study to examine the effects of cutaneous feedback and hump position on effective use of the computer mouse with a splint. Furthermore, Kwon et al. [41] selected three pivotal hand dimensions (length, circumferences, and breadth) to design gloves for candidates.

The way the computer mouse and keyboard are used probably poses a risk of carpal tunnel syndrome and repetitive and sustained loading of the small muscles of the hand. It is highlighted in the literature that a vertical mouse and ergonomic mouse pads change wrist position; however, they do not alleviate carpal tunnel pressure in patients with carpal tunnel syndrome [60,61,21,38]. It is reported that mouse users adopt working postures of wrist extension, pronation, and ulnar deviation [37,19,13,8]. Liu and Fan [45] carried out a computer task by thirty participants utilizing two forms of computer mice (front hump and rear hump) and two types of wrist splints (dorsal and volar) and measured the movement times and satisfaction scores.

The computer mouse is one of the most functional devices that can influence the wrists of its users negatively. A study of the literature and industrial practices show that anthropometric measures of five important areas of the hand have been used to classify computer mouse users into small, medium and large categories and ultimately design an ergonomic product. The hand areas, which are presented in Fig. 1, include: A: wrist width; B: wrist length; C: palm width; D: middle finger length; and E: index finger length.

The seventy measured samples (both right and left hands) are selected from 35 individuals. Seventy hand dimensions were selected based upon the 1988 US Army hand anthropometric survey report [25], after which they are categorized into three dimensional groups (length, circumference, and breadth) [41].

3. Artificial immune algorithm

In this research, an artificial immune system algorithm is developed to categorize computer mouse users based on their hand size and finally produce computer mice which match effectively with anthropometric measurements of users' hands. The artificial immune system and its algorithm are presented as follows.

An AIA imitates a learning technique performed by the natural immune system [35] and is one of the methodologies inspired from biological systems. The natural immune system is an intricate adaptive pattern-recognition system that guards the body from bacteria or viruses (pathogens). It can classify all cells, i.e., or molecules, within the body as either those are members of its

own kind (self-cell) or those that have a foreign origin (non self-cell) [15].

The human body is under continuous attack from bacteria or viruses, or disease-carrying parasites. Immune system consists of two layers of protection, non-specific and specific defense. The first defensive line is to prevent pathogens from entering the body, called non-specific defense, which includes skin and mucous membranes. The specific defense is activated when these microorganisms get past the non-specific defense and invade the body. Two activities must occur to make the immune system work properly: First, the body must recognize that it has been invaded, either by pathogens or toxins. Second, before invaders destroy many body tissue cells, the immune response must be activated swiftly. In order to respond effectively, several conditions including the proper interaction of non-specific and specific defense must be in place. When non-specific defense cannot destroy microorganisms and is proven to be ineffective, specific defense goes into action.

According to the information processing perspective, an immune system is an exceptional adaptive system and can shed light on many significant aspects in the realm of computation [20,16]. The immune system provides a way to solve optimization problems more effectively. When the immune system combines with evolutionary algorithms, it can promote the search capability during the evolutionary process [32]. Artificial immune systems and their applications have been extensively investigated in the literature [40,66,9,52,70,72,34,27,35,11].

3.1. Implementation of immune system algorithm

The artificial immune system algorithm is applied to learn the weight of criteria along with SM (Small-Medium) and ML (Medium-Large) cut-off points from pre-classified items based on the research by Zandieh et al. [73]. A detailed description of the AIA is presented as follows.

3.1.1. Encoding mechanism

In a classification problem a set of antibodies representing one class or category, is used as a solution. These antibodies are randomly generated by computerized procedures. Antibodies will be replaced by others or random mutation will occur, at a time when a set of antibodies do not have high fitness. Replacement happens based on the replacement probability value. High fitness antibodies with high resemblance to each other result in a lack of uniformity of antibodies and also varieties in them. Thus an affinity function for antibodies fitness adjustment should be applied.

Points with a final degree greater than the cut-off value SM will be put in category small (S). Points with a final degree between the cut-off SM and cut-off ML will be in category medium (M), and other points with final degree less than cut-off value ML will be in category large (L). Here, coefficients of criteria and cut-offs in the segments with length of $K+2$ (with K sub-segment) are located. Sub-segments are coefficients (W_j) and cut-offs (X_{SM}, X_{ML}). Therefore, $C = (W_1, W_2, W_3, \dots, W_K, X_{SM}, X_{ML})$, where W_j is the weight of criterion j and $\sum_{j=1}^k W_j = 1, X_{SM} > X_{ML}$.

These segments are determined as antibodies of the artificial immune system. In this case, the weight of each criterion can be coded as an exact value in an antibody. The value of each sub-segment is called the allele and it represents the weight of each criterion which is independent of the other alleles. For a known antibody C , categorizing consumer j by the weight sum $ws(c, j)$ is determined as Eq. (1) [2]:

$$ws(c, j) = \sum_{j=1}^k W_j \frac{i_j - \min j}{\max j - \min j} \tag{1}$$

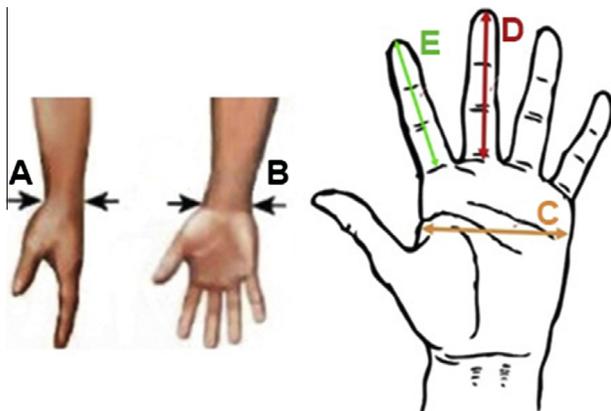


Fig. 1. Five anthropometric measurements of hand.

where k is the number of criteria, i_j is the anthropometric measurement of person i for criterion j , and $\max j$ and $\min j$ are the maximum and minimum criterion values for all people, respectively. Classification of person j according to the antibody C is presented as Eq. (2) [2]:

$$\text{Classification}(c, j) = \begin{cases} S, & \text{if } X_{SM} \leq ws(c, j) \\ M, & \text{if } X_{ML} \leq ws(c, j) < X_{SM} \\ L, & \text{otherwise.} \end{cases} \quad (2)$$

With this clear encoding method, the classification manner uses standard genetic algorithm functions (reproduction, crossover, and mutation) for the antibodies population. Also, the immune system algorithm transfers the best antibodies of each generation to the next generation. The fitness evaluation of an antibody, crossover, and mutation operators, which are used in this method, is described as follows.

3.1.2. Fitness function

Antibody fitness in the classification problem shows the ability of a test set's correct classification by that antibody. As a result, each user who is not classified correctly will receive a penalty. Due to the linearity of the class order, it is necessary to separate classification error of class S person to class M from classification error of class S person to class L . In this approach, fitness of antibody C is presented as Eqs. (3) and (4) [2]:

$$\text{Fitness}(c) = \frac{\sum_{i=1}^t P_i}{t} \quad (3)$$

$$p_i = \begin{cases} 1, & \text{classification}(i, c) = \text{class}(i) \\ 0.4, & |\text{classification}(i, c) - \text{class}(i)| = 1 \\ 0, & \text{otherwise,} \end{cases} \quad (4)$$

In Eq. (3), t is the size of the test set. Eq. (4) shows the penalty function of a misclassified person and class (i) is the class of person i which is determined by the expert. If the classification specified for the i -th training instance by the expert (i.e., class (i)) is equal to classification of person i according to antibody c (i.e., classification (i, c)), the penalty value is equal to 1. It should be noted that this fitness function prefers the antibody that makes one error from two class differences to the antibody that makes two errors from one class difference. Therefore, when the difference of class (i) and classification (i, c) is 1, the penalty value is considered to be 0.4.

3.1.3. Crossover operator

The crossover operator is the most important operator in the artificial immune system algorithm. The crossover operation is conducted for the selected pairs of antibodies (which are randomly selected by the roulette wheel method). The feasibility condition of each antibody is described as follows: the value of each sub-segment must be between 0 and 1; the summation of weights should be 1; and the value of cut-off SM is greater than the value of cut-off ML . Although the initial population is made in a way that all antibodies are feasible and correct, using the standard crossover operator leads to defective sub-segments. An example is provided in Appendix A.

In this research, a recently developed format of the crossover operator, called the "continuous uniform crossover", is utilized. This operator guarantees the feasibility of the offspring [2]. For two parent antibodies $x = (x_1, x_2, \dots, x_n)$ and $y = (y_1, y_2, \dots, y_n)$, the offspring antibodies are: $x' = (x'_1, x'_2, \dots, x'_n)$ and $y' = (y'_1, y'_2, \dots, y'_n)$ in a way that $x'_i = Sx_i + (1 - S)y_i$, $y'_i = (1 - S)x_i + Sy_i$; where S is a constant for each crossover operator. This operator keeps the sum of any subset of sub-segments. On the basis of classification,

if the sum of m sub-segments in an antibody segment is equal to 1 ($\sum_{i=1}^m x_i = 1$), after the crossover operator it would result in Eq. (5):

$$\sum_{i=1}^m x'_i = s \sum_{i=1}^m x_i + (1 - s) \sum_{i=1}^m y_i = s + (1 - s) = 1 \quad (5)$$

It should be mentioned that Eq. (5) is valid for each of the two offspring. In addition, this operator will keep the greater-than relation between sub-segments. That is if $X_{SM} > X_{ML}$ then $X'_{SM} > X'_{ML}$. Hence, the continuous uniform crossover operator keeps antibodies' feasibility for multi criteria SML classification [2]. Appendix B provides a more detailed mathematical treatment of the selection of S .

3.1.4. Probability function

The probability value of the selection antibodies can be computed according to the following procedure [2]:

Step 1: Compute the fitness value of each antibody in the population.

Step 2: Compute the value of fitness function for each antibody based on Eqs. (3) and (4).

Step 3: Compute the selection probability for each antibody according to Eq. (6):

$$\text{Probability}(c) = \frac{\text{Fitness value}(c)}{\text{Sum}(\text{FitnessValue})} \quad (6)$$

3.1.5. Mutation operator

The mutation operator randomly alters the value of a sub-segment. After applying the mutation operator, the normalization of the changed antibody is required. The normalization will be done in a similar way as the crossover operator with $S < 0$.

3.1.6. Affinity value evaluation using entropy theory

In this section, entropy theory is applied for the evaluation of the affinity value. In immune algorithms, each candidate solution will be compared with the best-known solution obtained up to that point. This will allow an affinity value to be computed using the entropy theory. The value of the affinity function expresses the similarity between a candidate solution and the best-known solution obtained up to that point. Mori et al. [53] defined information entropy $H(x)$ for a discrete random variable X as shown in Eq. (7):

$$H(x) = - \sum_{i=1}^n p_i \cdot \text{Log} p_i \quad (7)$$

where $X = \{x_1, x_2, x_3, \dots, x_n\}$ is a discrete random variable with probability mass function of $P(X = x_i) = p_i$, $i = 1, 2, \dots, n$.

Using information entropy, the similarity of a sequence in relation to a reference sequence can be expressed as Eq. (8):

$$\text{aff}(i) = \frac{1}{1 + \frac{1}{k} \sum_{j=1}^k h_{ij}} \quad (8)$$

where $\text{aff}(i)$ is the measure of similarity, k presents the measure of the antibody ($k = \text{number of criteria} + 2$), and $h_{ij} = -p_{ij} \text{Log} p_{ij}$. If $x_j = x_j \text{ref}$, then $p_{ij} = 1$, and therefore, $h_{ij} = -\text{Log} 1 = 0$. In other words, if $x_j \neq x_j \text{ref}$, then $p_{ij} = 0.5$; this indicates that $h_{ij} = -0.5 \text{Log} 0.5 = 0.151$. It should be noted that two identical sets of numbers will be given an affinity value of 1. Conversely, two entirely unlike sets of numbers would result in an affinity value equal to: $1/(1 + 0.151)$, which is approximately equal to 0.87.

3.2. Proposed method

The steps of the proposed AIA are presented in Fig. 2.

The proposed artificial immune system algorithm used in this research is summarized in the following steps:

Step 1. Initialization: Select values for the number of initial population (np), the number of generations (ng), the crossover probability (pc), the mutation probability (pm), the affinity threshold (at) and the affinity adjustment (aa). These values are only for initialization of the algorithm and in later stages

the values of the parameters will be set. Generate the initial population randomly. Generate feasible antibodies in the initial population randomly.

Step 2. Objective function value evaluation: Compute the fitness of each antibody with the fitness function.

Step 3. Antibody probability assignment: Assign a probability value to each antibody. In the probability assignment, an antibody with the higher degree of fitness will receive greater

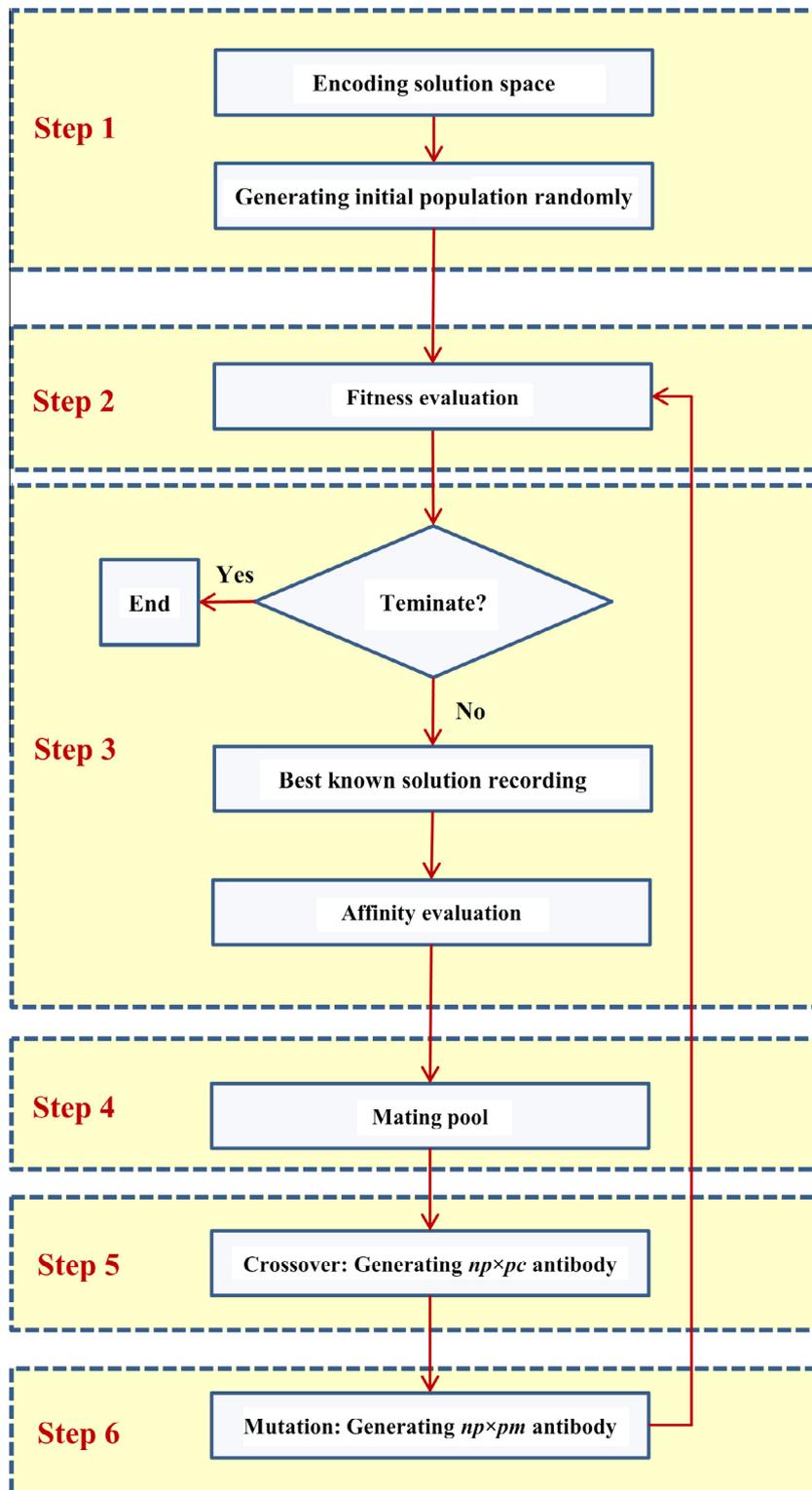


Fig. 2. The proposed artificial immune algorithm procedure.

probability, and this probability value will determine the chance of each antibody to exist in the next generation. In other words, antibodies with greater probability have more opportunity to propagate.

Step 4. Mating pool generation: The best known antibody according to the fitness function will be selected from the mating pool (according to the accelerating mechanism). Then determine the affinity value between each antibody and the best known antibody according to the fitness function. If some antibodies have a similarity probability higher than the predetermined threshold, then decrease the probability value (i.e., the fitness function) that is assigned to them. This work will be done by multiplying an assigned probability to each antibody with expressed conditions by a number less than 1 (i.e., affinity adjustment) and the normalization of the probability values of other antibodies (i.e., restraining mechanism). To expand the mating pool, select $(np - 1)$ antibodies from the population (including the best antibody) with replacement. The chance of selection of each antibody depends on its fitness value. In other words, antibodies with greater fitness value have more chance to be selected.

Step 5. Crossover operation: Select $(np \times pc)$ pairs of mating pool antibodies as parents. Produce offspring with the crossover operator randomly. These antibodies belong to the next generation.

Step 6. Mutation operation: Select $(np \times pm)$ pairs of mating pool antibodies as parents. Change one of the bits with the mutation operator randomly. These antibodies belong to the next generation.

Step 7. Replacement: While keeping the best antibody from the previous generation, replace other $(np - 1)$ antibodies that are obtained from the operators.

Step 8. Fitness evaluation: Compute the fitness value of each antibody with the fitness function.

Step 9. Termination test: If stopping criteria are achieved, then terminate the algorithm; otherwise, return to Step 3. The termination condition is considered as follows: if the fitness value has not changed in consistency for more than 10 iterations, the algorithm will stop.

The following courses of action are considered in the AIA to overcome the local optimum solutions. First, by using the roulette wheel method, candidate solutions with a higher fitness value have a higher probability to be selected for the production of the next generation of candidate solutions. Second, in order to diversify the search space in the AIA, an accelerating mechanism enhanced with a restraining mechanism is utilized. In the accelerating mechanism, a candidate solution with the best fitness is recorded, which will be used to seed the mating pool. This mechanism ensures that the mating pool contains a large proportion of candidate solutions with good fitness. In addition, in the restraining mechanism, if a candidate solution has a higher affinity value than a prescribed threshold value, then the probability value, obtained from the fitness, is multiplied by a number which is less than one; and the smaller probability value will be assigned to it. This will decrease the selection probability of that solution and prevent overall dominance of a candidate solution. Without the restraining mechanism, a good candidate solution may be pervaded very fast and may be dominated in the mating pool rapidly.

4. Computational results

The AIA, explained in the previous section, is coded by the MATLAB programming software to solve the problem. The problem is surveying the classification of consumers into three categories: S,

M, and L. Anthropometric measurements of five areas of the hand for designing an ergonomic product are also measured. Thirty-five individuals are classified into three categories: S, M, and L. Each person's left and right hands are measured according to criteria and effective measures in design. Each area is considered as a criterion in the proposed algorithm. These criteria include wrist width, wrist length, palm width, middle finger length, and index finger length. They are measured by calipers with accuracy up to 0.01. The 70 measured samples (both right and left hands) are categorized into the S group (12 samples), the M group (38 samples), and the L group (20 samples) based on the expert opinion. The expert opinion is expressed based on their knowledge and expertise using related tools and checklists. After each run of the program the correct proportion is assessed and shown as a percentage. The fitness value is calculated based on Eqs. (3) and (4). This value shows the ability of a test set's correct classification by that antibody. The probability value is calculated according to Eq. (6). This value determines the chance of each antibody to exist in the next generation. The affinity value expresses the similarity between a candidate solution and the best-known solution obtained up to that point. This value can be computed on the basis of Eq. (8).

4.1. The Artificial immune algorithm (AIA) parameters tuning

This section is aimed at studying the behavior of the various operators and parameters of the proposed AIA to achieve better robustness of the algorithm. There are many approaches in the literature to calibrate the algorithm, i.e., parameter tuning of the algorithm; yet, the most commonly practiced method is a full factorial design of experiment.

Many levels for the parameters of the algorithm are considered which are near optimal. Different combinations of the parameter values will constitute a plan (also known as a treatment) for the experiment. The results of applying the algorithm to each of these plans are used to select the appropriate levels that maximize the final solution for the parameters. According to the research by Zandieh et al. [73], the number of runs should be chosen to be eight. Different levels of these factors are shown in Table 1. In conducting our experiments to test the performance of the algorithm, the number of the initial population (np) is set to be 100.

The full factorial design to find the optimal value of the aforementioned five factors, i.e., parameters of the AIA algorithm, requires $3^4 + 4^1 = 324$ experiments. Therefore, 324 combinations of control factors were taken into account. For each trial, eight

Table 1
Various levels of AIA factors.

Factors	Symbol	Index of level	Levels
Crossover probability	pc	1	0.40
		2	0.50
		3	0.80
Mutation probability	pm	1	0.05
		2	0.10
		3	0.20
		4	0.30
Affinity threshold	at	1	0.85
		2	0.90
		3	0.95
Affinity adjustment	aa	1	0.50
		2	0.40
		3	0.30
Number of generations	ng	1	30
		2	50
		3	70

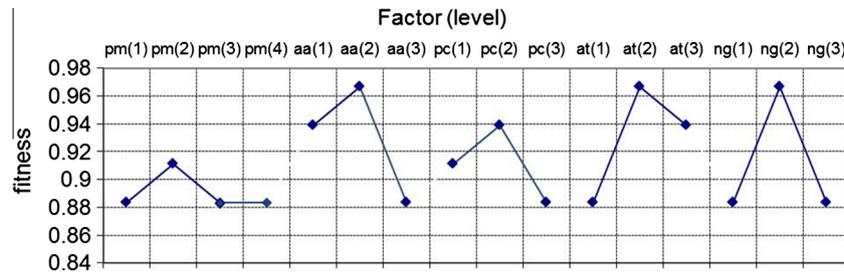


Fig. 3. Fitness values of various factors' levels.

Table 2

Comparison of AIA and the expert classification on the hand measured items.

Class	Expert	AIA		
		S	M	L
S	12	9	3	0
M	38	0	38	0
L	20	0	2	18
Total	70	9	43	18

replications of AIA were performed to yield more reliable information, and then the results of the total fitness are recorded. The fitness value of the total number of generated solutions is averaged in each level. Fig. 3 represents these values plotted against each control factor. Based on Fig. 3, which shows the fitness of the algorithm, better fitness is acquired when the parameters are set as follows: $pc = 0.5$, $pm = 0.1$, $at = 0.90$, $aa = 0.4$, and $ng = 50$.

Anthropometric measurements of five areas of the hand for designing an ergonomic product are also measured. The person's left and right hands are measured according to the criteria and effective measures in ergonomic product design. Each area is considered as a criterion in our proposed algorithm. Then, the remaining 70 items (35 items for the left hand and 35 items for the right hand) are classified utilizing the weights and cut-off values learned by the AIA into 3 categories S, M, and L. In this experiment, we take advantage of a human expert, who has the full authority and expertise, to classify items based on the mentioned anthropometric data.

The results of our experimental investigation are reported in Table 2. The expert classification did not agree with the result obtained by AIA on 5 items, i.e., based on Table 2, three items in class S; and two items in class L. As can be seen, the proposed AIA confirms the expert classification at least in 93% of the cases.

5. Conclusion

Designing ergonomic products has been proven to have an important effect on users' behavior and reaction. In this paper, an AIA was proposed as a meta-heuristic method in order to classify ergonomic products. A case study in this paper, which considers anthropometric measurements for the computer mouse, makes a significant contribution to the ergonomic computer mouse literature. Classification with this method overcomes inefficiency in searching a large space, and results in a higher accuracy in classification ergonomic products regarding anthropometric data compared to the classification of experts. The AIA has a learning mechanism from anthropometric measurements of samples by classifying them into three categories: small (S), medium (M), and large (L), similar to what the PCA and the discriminant analysis techniques do. An expert advisory opinion was also utilized in the case study of this research based on their knowledge and expertise

using related tools and checklists. After the learning step, weights for each criterion and cut-off points were determined. Thus, any other samples can be allocated to one of the categories based on the weights and cut-off points obtained from prior steps. To obtain the precise calibration of various operators and parameters of the AIA, a comprehensive comparison by using full factorial design of experiment was performed. To evaluate the effectiveness and robustness of the proposed AIA, it was compared against the expert classification outcomes. The comparative results showed that the proposed method confirms the classification of the expert. For future research explorations, other clustering and classification methods, such as K-Means, PCA, or discriminant analysis, can be applied and compared in the case problem for comparison purposes and to provide more reliable outcomes. In addition, the presented approach can be applied to other classification problems such as text categorization (e.g., spam filtering), optical character recognition, and machine vision (e.g., face detection).

Appendix A

A.1. Crossover operator in AIA-example

Consider the following two antibodies as parents: $X = (0.1, 0.2, 0.3|0.4)$ and $Y = (0.2, 0.2, 0.1|0.5)$. With the single-point classic crossover operator, the following offspring will be produced: $X' = (0.1, 0.2, 0.3, 0.5)$ and $Y' = (0.2, 0.2, 0.1, 0.4)$; where $\sum_{j=1}^4 X'_j$ is equal to 1.1, which is greater than one, and $\sum_{j=1}^4 Y'_j$ is equal to 0.9, which is less than one. Therefore, X' , Y' have defective or infeasible weights.

Appendix B

B.1. Continuous uniform crossover - mathematical treatment of the selection of S

The selection of S is critical. If $S = 0$, then the offspring will be exactly similar to the parents. If $S > 0$, then the sub-segments will be between 0 and 1. However, sub-segments over production of consecutive generations will be similar to each other. If $S = 0.5$, then both offspring will be the same and the values of the sub-segments will be the average of the corresponding sub-segments values in the parents. On the other hand, if $S > 0$, then the sub-segments will be distant from the corresponding sub-segments in the parents and may go out of the determined limits. For instance, the sub-segment value may be negative or more than 1, although the sum of sub-segments is still equal to 1. In this case, antibodies should be normalized and for $S < 0$, the value of each sub-segment after the crossover operation will be surveyed until it is not below 0. If a sub-segment with the value greater than 1 exists, certainly a sub-segment with negative value will exist. In the case of a negative sub-segment, at first the minimum sub-segment value is subtracted from all the sub-segments, and then,

the value of each sub-segment will be equal to $x_i = \frac{x_i}{\sum_{j=1}^k x_j}$; where k is the number of criteria. If $S < 0$, it may result that $X_{SM} < X_{ML}$ and for resolving this, X_{SM} and X_{ML} are interchanged.

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