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RESEARCH ARTICLE

NATURAL INSPIRED COMPUTING IN THE NETWORK SECURITY

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ABSTRACT

Traditional computing techniques and systems consider a main process device or main server, and technique details generally serially. They're non- robust and non-adaptive, and have limited quantity. Indifference, scientific technique details in a very similar and allocated manner, while not a main management. They're exceedingly strong, elastic, and ascendible. This paper offers a short conclusion of however the ideas from natural are will never to style new processing techniques and techniques that even have a number of the beneficial qualities of scientific techniques. Additionally, some illustrations are a device given of however these techniques will be used in details security programs.

INTRODUCTION

The power and recognition of current computing systems is basically as a effect of more agile and quicker CPUs and additional and additional memory convenience at low value. Yet, these "traditional" computing ways, architectures, systems, and networks largely think about a central process unit or a central server, they method data serially, and that they rely upon humans to be programmed and told what to try to (and how). It has some serious drawbacks. Foremost, the systems don't seem to be awfully secure. If one a part of a system breaks down, the complete system is useless. Second, they're not adaptations. Most computing systems don't read (or have solely restricted learning capability), and can't change or adjust to fresh or surprising things while not human intervention. Third, there's solely restricted measurability. The bigger the organization becomes, or the extra nodes are value-added to the network, the upper the employment of the C.P.U. Or server becomes, till it cannot method all directions or service requests during a cheap time any longer. In comparison, most scientific techniques process details in a similar and allocated way, without the lifestyle of a central control.

They usually involve a huge variety of relatively easy personal units, which act in similar and communicate only regionally. For example, the mind includes a huge variety of easy nerves (more or less comparative to on-off switches), each of which is connected only to a relatively small portion of all other nerves. Yet quantity of details handling is going on in the mind, where each neuron works only aspect of the handling, but they all do so in alike. In social pest hives, such as bugs and bees, a huge variety of relatively easy individuals manage to build complex nests or find the quickest path between the home and a food source, again in a similar and allocated way. The human immunity processes is another example, where (simple) personal defense cells perform only aspect of the complete task, but there are many of them working together in similar.

This parallel and distributed processing method makes these systems highly robust. If some individual units in the system break down, the system as a whole will still function. In fact, it is easy to repair or replace broken units without having to "shut down" the entire system. Furthermore, these systems are highly scalable. As many individual units can be added as desired, since there are only local interactions involved, and there will be no overload on one particular part of the system. Finally, most systems in nature are adaptive, either through learning (in individual organisms) or through evolution (at the level of populations or species). They can adjust to changing situations or even cope with entirely new situations. So, there

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are many advantages in biological systems that would be desirable to have in our computing systems. In this paper, a quick summary is given of however ideas from biology are wont to style new computing strategies. This can be typically mentioned as biologically galvanized computing (De Castro and Von Zuben, 2005). These strategies overcome a number of the disadvantages of ancient computing, creating them a lot of strong, adaptive, and ascendible. Especially, 3 examples are reviewed: (1) genetic algorithms, (2) neural networks, and (3) artificial immune systems. Moreover, for everyone of those 3 strategies, some actual applications within the space of knowledge security also are represented, especially in cryptography, life science for security, and laptop and network security. The biological ideas and concepts underlying the strategies represented here are often found in any commonplace textbook on biology, like (Alters, 1996; Campbell *et al.*, 1997).

Genetic Algorithms in Cryptography

1. Genetic algorithms were developed within the 60s and 70s by John Holland and his colleagues and students. They were used each as easy models of evolution and adaptation, and as new laptop algorithms to seek out sensible solutions to troublesome improvement issues. Later, they became very fashionable as a general improvement tool, and that they are applied with success to a good vary of issues. This section offers a quick summary of the algorithmic rule (more details are often found in (Holland, 1975; Holland, 1992; Goldberg, 1989; Mitchell, 1996)), and a few specific applications of genetic algorithms in cryptography and writing are described.

Genetic Algorithms

A Genetic Algorithm (GA) could be a unique look for technique reinforced concepts from genetic and natural progress and choice. It's one among variety of process techniques generally mentioned as natural evolutionary computation (EC). rather than trying to straight fix a move, an answer is progressed over time by keeping a population of (initially random) candidate alternatives, creating subsequent generations by recombining completely different components of this best alternatives within the population. This way, new candidate evaluate tested reinforced based on current sample, wherever the look for is target-hunting by a range method that prefers the (currently) best alternatives within the population to use to make new ("offspring") alternatives.

Given some optimization disadvantage, first an appropriate cryptography for applicant alternatives has to be discovered. Usually, this cryptography requires the way of personality post like bit post (i.e., post of 0s and 1s). This is often analogous to the scientific difference between the genotype (the inherited encoding) and also the cosmetics (the real type and appearance) of a living thing. As an example, in chart problems wherever some the best possible set or partition of the nodes has to be discovered (such as a lowest protect or most cut set), a little sequence cryptography are often used wherever every bit place matches to 1 particular node within the chart. Development of the particular applicant quality (phenotype) from a given bit sequence (genotype) is finished as follows. For every bit with value one, the corresponding node within the chart is surrounded within the applicant set (or placed on one aspect of the applicant partition), and for every

bit with value zero, the corresponding node isn't surrounded within the set (or placed on the other aspect of the partition). This way, the GA will straight look for the (much simpler) place of bit post rather than the place of real applicant alternatives, even as organic progress happens at the level of genotypes. Next, a fitness function needs to be designed which can be used to evaluate candidate alternatives. The main idea is that this operate takes as its input an secured candidate solution (e.g., a bit string), converts this into an actual candidate solution (e.g., a partition of the nodes of a graph), and returns a variety according to how good this candidate solution is for the given issue (e.g., the count of sides between nodes from different sides of the partition for the maximum cut problem). This number, and fitness value, indicates the "goodness" of a candidate solution: higher fitness values mean better solution. This way, the GA can perform choice based on these fitness principles, just as natural choice happens at the level of the phenotypes. Given an appropriate development and fitness function (which have to be developed independently for each optimization problem that is considered), the real criteria is relatively simple. Supposing a bit sequence development is used, the primary GA performs as follows which is proven in Algorithm 1 (the choice and cross-over & mutation providers are described below):

Algorithm 1: Selection and Mutation process (Millan *et al.*, 1997)

- Initialize the population with N random bit strings ("individuals"), calculate their fitness values, and set $gen=1$.
- Create a "mating pool" by selecting (with replacement) N individuals from the current population based on fitness.
- While still individuals in the mating pool, do:
 - Remove the next pair of individuals ("parents") from the mating pool.
 - With probability pc perform crossover between the parents to create two "children".
 - With probability pm perform mutation on the children.
 - Place the children in the new population.
 - Replace the previous population with the new population, calculate the fitness of all individuals, and set $gen=gen+1$.
- If $gen < M$ go to step 2, otherwise stop.

There are many ways in which the selection operator can be implemented, but the main idea is that individuals with higher fitness values, compared to the rest of the population, have a higher chance of being selected than individuals with lower fitness values (i.e., fitness proportionate selection). In other words, the mating pool will (on average) contain multiple copies of the best individuals in the current population and no (or just a few) copies of the worst individuals. The crossover operator literally chops up the genotypes of the parent individuals and recombines them to create offspring genotypes. The most basic method is one-point crossover, in which a random crossover point is first chosen (somewhere between the first and last bit), and the first part of the first parent is recombined with the second part of the second parent to create the first child (and vice versa for the second child). Usually crossover is done with a certain probability pc (often set in the range [0.6;0.95]) for each pair of parents. If

crossover is not performed, the children will be identical to their parents. Finally, with a usually very low probability p^m , mutation is performed, where a bit is flipped at random. Examples of (one-point) crossover and mutation are shown below. In the crossover example, the crossover point is (randomly) chosen between the 3rd and 4th bit. In the mutation example, the 0 at the 9th position is mutated into a 1 (shown in bold). Finally, the creation of new generations of candidate solutions by selection and crossover & mutation is repeated for a set number M of generations. Other stopping criteria are possible, of course, such as reaching a certain level of fitness or a maximum amount of computing time. In short, the main idea of the algorithm is to evolve better and better solutions by repeatedly selecting the best candidate solutions from the current population and recombining parts of their genotypes to create subsequent generations of candidate solutions.

Applications of Genetic Algorithms in Cryptography and Coding

A little summary of the status of the art and of still start issuing in using transformative calculation methods (such as genetic algorithms) in secret writing is presented in (Isasi and Hernández, 2004). In cryptography, it is very important know how challenging it is to “break” a security technique. Obviously, methods that are very challenging to crack are recommended over methods that are more quickly damaged. Cryptanalysis is all about examining (or “attacking”) security methods to distinguish out how comfortable or challenging they are to break. Genetic methods have been applied efficiently in this field, for example in fighting replacement ciphers (Spillman *et al.*, 1993; Clark and Dawson, 1998), and transposition ciphers (Matthews, 1993). Although this does not straight cause better ciphers, it does display where their weak points are, which often can help in raising them. Furthermore, in (Millan *et al.*, 1997) an inherited criteria were utilized efficiently to discover Boolean features with good cryptographic qualities, thus displaying how these methods can also be used straight for building security methods.

An essential strategy that is frequently used in cryptography is that of producing pseudo random numbers. Hither, the aim is to get a random act (by some deterministic method) that is “every bit singular as possible”, and which bears a higher interval (i.e., it will not answer it again itself quickly). An exciting scheme, using a transformative strategy just like GAs, was presented in (Sipper and Tomassini, 1996), where mobile automata (simple identical and allocated processing devices) were advanced to generate pseudo unique figures with a higher degree of entropy.

As an example, consider programming methods for transmitting information. Next to offering information protection through protection, it is likewise important that information reduction is reduced during transmitting of secured data. Inherited methods have been applied efficiently to improve so-called “turbo codes” (Durand *et al.*, 1999). In this situation, the GA was able to find a small bit better rule than what was usable at the time. These programs are just a selection of the many opportunities of implementing inherited methods and other transformative calculations techniques in the area of information security. Next, an introduction to sensory systems, another naturally motivated processing method, is provided.

Neural Networks in Biometrics for Security

The research on neural networks was pioneered by McCulloch and Pitts in 1943 (McCulloch and Pitts, 1943). They gave a logical (mathematical) model of a simple neuron, and demonstrated that a suitably constructed network of such “artificial neurons” can, in principle, compute any computable function. Thusly, a neural network is equivalent (in terms of computational power) to a general Turing machine, but with a very different architecture. In this section, first the concept of neural networks is briefly surveyed. A full introduction to computing with neural networks is provided in (Lippmann, 1987), and more detailed info can be found in any standard textbook on neural networks, such as (Anderson, 1995). Next, an example of an application of neural networks in biometrics for security is traced.

Neural Networks

A neural networks (NN) is a parallel distributed processing (PDP) structure that is prepared after the working of the brain. It can perform calculations, in particular category of information, and provides an example of an alternative design of calculations as opposed to serially and centrally based calculations of conventional handling systems. Our minds involve many (around 10 billion) simple tissues known as nerves. Each neuron includes a mobile whole body, an axon (a pointed “transmission line” through which substance alerts can travel), and many dendrites (a treelike framework of many branching “tentacles”), which end in synapses which type relationships with the axons of other nerves. Basically put, each neuron gets information (the existence or lack of signals) from other nerves through the synaptic relationships, Our minds involve many (around 10 billion) simple tissues known as nerves. Each neuron includes a mobile whole body, an axon (a pointed “transmission line” through which substance alerts can travel), and many dendrites (a treelike framework of many branching “tentacles”), which end in synapses which type relationships with the axons of other nerves. Basically put, each neuron g ets information (the existence or lack of signals) from other nerves through the synaptic relationships, which journey down the dendrites of the mobile whole body. Here, the information are “added up”, and if a certain limit is obtained the neuron delivers out an indication itself through its axon, which is then developed an feedback to yet other nerves which are linked with its axon. Nevertheless, not all synaptic relationships are equivalent. Some are more powerful than others, and then some information has a greater “weight” than others. Studying is obtained by modifying the strong point s (weights) of current synaptic relationships, or by pre paring new or removing old relationships.

A simplified example of a real neuron is illustrated in the figure below. A neuron receives inputs (x^i) from other neurons, which are weighted (w^i) and then added (\sum). The production of a neuron is a function of this weighted sum of inputs, and can in turn constitute the input to other nerve cells input to other neurons. In the simplest case, each input can be either 0 (absence of signal) or 1 (presence of signal), and the output function is a step function such that the output is 0 if the weighted sum of inputs is below a certain threshold value, and 1 if it is above the threshold value. In more realistic cases, the inputs and outputs are real valued number s within some range,

and the output function is for example as shape which is shown in figure 1.

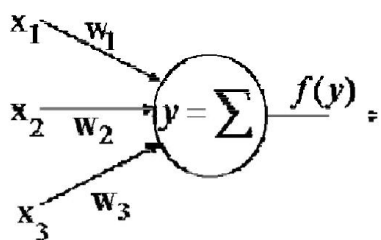


Figure 1. Neural Network Architecture

go “forward” and provide as information to the first invisible part. These nerves then generate their results which provide as information to the next invisible part (if present), until the ultimate, or outcome, part is achieved. The condition of the nerves in the outcome part can then be considered as the “answer”. For example, in category issues, if the condition of the first outcome neuron is 1 and that of the second one is 0, the feedback linked with one category. If their last declares are changed (i.e., 0 and 1, respectively), then the feedback linked with the other category (assuming there are two sessions into which to partition the inputs). Other system architectures are of course also possible, such as repeated systems, where relationships can nourish returning to past levels as well, or lines systems, where the nerves are organized in a lines with relationships between nearby nerves. This will be explained in figure 2.

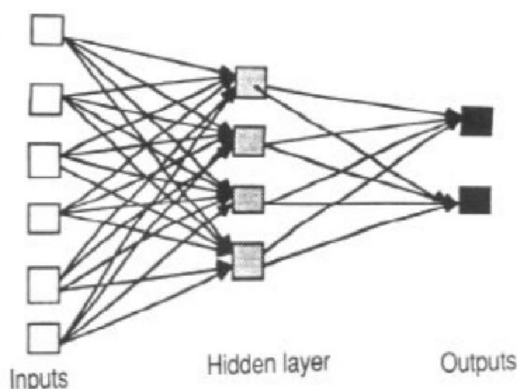


Figure 2. Internal Architecture of Neural Network

Given some neural network structure, it is not straight apparent how to set the loads on the relationships to get a certain system actions. However, several coaching methods have been developed to improve these loads. The primary concept of these methods is to continuously existing the system with example information for which the appropriate response is known. The loads in the system are then modified based on the quantity of mistake between the appropriate response and that given by the system. This is recurring until no more mistakes are created, or the quantity of mistake drops below a certain limit. The system can now be said to have discovered the given process. At the next level, the system can be used to execute the process on new information which it might not have seen before. Any variety of such nerves can be linked with each other to type an synthetic sensory system. A conventional system structure that is often used is a nourish

ahead system. In such a NN, there is one part of feedback nerves, one or more levels of “hidden” nerves, and one part of outcome nerves, as shown in the determine on the next web page. The nerves in the feedback part are initialized with some feedback design, and the results from this part

Applications of Neural Networks in Fingerprint

Recognition

One place where neural networks have become very popular is picture handling, such as design identification and category, disturbance filtration, advantage recognition, etc. As an program in biometrics for protection, they can be used efficiently for finger marks identification. Finger print identification is often divided up in two stages: (1) function removal, and (2) category. In the first level, certain functions from a finger marks picture are produced, such as variety guidelines, archways and whorls, delta points, etc. In the second level, these functions are used to identify (or classify) the given finger marks picture.

Neural systems have been used efficiently in both of these levels, often providing increase to high correct category prices and low incorrect being rejected prices, and frequently outperforming more conventional methods. Furthermore, sensory systems can be used in the same way for other picture identification projects in biometrics protection, such as retina or eye check out categories, or for speech identification.

Finally, as a last example of naturally motivated handling in the place of information protection, a brief summary of synthetic immunity processes for computer protection is provided in the next section.

4. Artificial Immune Systems for Computer Security

A very recent concept that is still being developed is that of building a pc defense mechanisms. The task of such a program is to provide pc and network security based on the technicalities of the individual defense mechanisms. This area first provides a high-level and somewhat simple summary of the individual defense mechanisms. Next, an example of an execution of a simple pc defense mechanisms is given to demonstrate the usefulness of the concept.

The human Immune System

The individual defense mechanism is a complicated and multi-layered program. The aspect that is of most attention here is the flexible defense reaction. A brief summary of this is given below, with many information remaining out. However, the common qualities of this aspect of the defense mechanisms provide as a place to start for the style of a synthetic immunity processes for computer and system protection. The body system includes many different kinds of elements (mostly proteins), which are generally known as “self”. Everything else, such as things that make us ill, is generally known as “non-self”. So, the main process of the defense mechanisms is to differentiate “non-self” from “self”, and induce a reaction whenever “non-self” necessary protein are recognized. However, this is not always easy as there are an approximated “non-self” necessary protein that the immunity processes

needs to identify, in comparison to about “self” necessary protein. The way the defense mechanisms resolve this problem is by using a powerful and allocated program.

Whenever you want, many “detector” tissues, such as so-called T-cells, flow through our organizations. These tissues older in a body known as the thymus, where they are revealed to most of the “self” necessary proteins that create up our systems. If any of the growing T-cells holds to any of these “self” necessary proteins, that T-cell is transferred. Hence, the only T-cells that prevent the thymus are those that do not unite to “self” necessary proteins. Therefore, if a grew up T-cell does combine to a protein, it suggests this must be a “non-self” proteins, and an appropriate defense reaction will be triggered. Nevertheless, not all T-cellular telephones are capable to merge two (or “recognize”) entirely possible “non-self” necessary proteins, but some T-cells combine to some “non-self” necessary proteins, other T-cells to others, etc. In this way, the defense mechanisms is an allocated program. It is also powerful, as T-cells are consistently changed through an inherited process such as the difference (or unique “mutations”). This way, the set of “non-self” necessary protein that the defense mechanisms is able to identify, changes eventually. Since it is impossible to identify all possible “non-self” necessary protein at any once, this powerful program is the next best solution. Furthermore, because of this, no two individuals will have exactly the same set of T-cells at some point, so what might make me fed up, my next door neighbor might be safe from, and the other way around. Lastly, the defense mechanisms also has a “memory”. It is able of keeping in mind illness-causing “non- self” necessary protein (antigens), so that when a person gets contaminated with the same antigen, it is identified instantly and an appropriate defense reaction can be activated, avoiding the real sickness from happening again.

Computer Immunology

Forrest and learners were some of the officers of using concepts from the human immunity processes to design an attack recognition program for computer systems and systems . In particular, they show the results of a basic execution based on checking short series of program phone calls. Temporarily, the idea is as follows. In the first level, a data source of program contact series during “normal” activities is built. This data source thus contains the series that represent “self”. In the next level, program contact series are examined during program function that might contain attack efforts. These series are then as opposed to available data source, and any series that is not present in the data source (“non- self”) activates an “alarm”. This way, irregular activities can be easily recognized, and appropriate activities can be conducted if necessary. Obviously, the data source containing regular actions have to be modified regularly. For example, including new customers or application and components to the program will modify the regular actions, or a user’s actions might modify over time (different projects, different main concerns, etc.). However, with this style, the attack recognition program becomes more flexible, as it is able of acknowledging irregular actions that has not been noticed before. In other terms, the program can recognize, for example, new malware or new fighting systems, without the need for installing new malware “signatures” from some main server first. Furthermore, different computer systems will have different data source of

“self” actions, so a malware that infects one pc, might not be able to contaminate every other pc. This way, the system as a whole also has a better (distributed) security

The (small-scale) illustrations and simulator that have been applied so far indicate the stability of these concepts, and display a appealing upcoming. Currently, the concepts and styles are still being designed further, and are also being grabbed by others. Pc immunology and synthetic immunity processes are now an effective area of analysis.

Conclusion

Traditional handling techniques have several drawbacks, such as a lack of sturdiness and flexibility, and limited scalability. In contrast, scientific techniques, being mostly similar allocated handling techniques, are highly effective, convenient, and scalable. Naturally motivated handling includes the design, execution, and application of new pc techniques and techniques that integrate these beneficial qualities of scientific techniques. In this paper, a brief summary of biologically motivated handling has been presented, with some specific examples of how these techniques can be used in details protection in particular. Many of these techniques have already been applied efficiently, such as inherited methods and sensory networks, and some are still being further developed, such as pc immunology. It is clear that the area of details protection can benefit greatly from these new and interesting handling techniques.

References

- Alters, S. 1996. *Biology: Understanding Life*. Mosby.
- Anderson, J. A. 1995. *Introduction to Neural Networks*. MIT Press.
- Campbell, N. A., L. G. Mitchell, and J. B. Reece, 1997. *Biology: Concepts & Connections*, 2nd edition. Benjamin Cummings.
- Clark, A. and E. Dawson, 1998. “Optimisation heuristic for the automated cryptanalysis of classical ciphers,” *Journal of Combinatorial Mathematics and Combinatorial Computing*, vol. 28, pp. 63-86.
- De Castro, L. N. and F. J. Von Zuben, 2005. *Recent Developments in Biologically Inspired Computing*. Idea Group Publishing.
- Durand, N., J.M. Alliot, and B. Bartolomé, 1999. “Turbo codes optimization using genetic algorithms,” in *Proceedings of the Congress on Evolutionary Computation*.
- Goldberg, D. E. 1989. *Genetic Algorithms in Search, Optimization, and Machine Learning*, Addison-Wesley.
- Holland, J. H. 1975. *Adaptation in Natural and Artificial Systems*. University of Michigan Press.
- Holland, J. H. 1992. “Genetic Algorithms,” *Scientific American*, vol. 267 (1), pp. 66-72.
- Isasi, P. and Hernández, J. C. 2004. “Introduction to the applications of evolutionary computation in computer security and cryptography,” *Computational Intelligence*, vol. 20 (3), pp. 445-449.
- Lippmann, R. P. 1987. “An introduction to computing with neural nets,” *IEEE ASSP Magazine*, vol. 4, pp. 4-22.

- Matthews, R. A. J. 1993. The use of genetic algorithms in cryptanalysis," *Cryptologica*, vol. 17 (2), pp. 187-201, 1993.
- McCulloch, W. S. and W. Pitts, 1943. "A logical calculus of the ideas immanent in nervous activity," *Bulletin of Mathematical Biophysics*, vol. 5, pp. 115-133.
- Millan, W., A. Clark, and E. Dawson, 1997. "An effective genetic algorithm for finding Boolean functions," in *Proceedings of the International Conference on Information and Communications Security*.
- Mitchell, M. 1996. *An Introduction to Genetic Algorithms*. MIT Press.
- Sipper, M. and M. Tomassini, 1996. "Co-evolving parallel random number generators," in *Proceedings of the Parallel Problem Solving from Nature Conference*, pp. 950-959.
- Spillman, R., M. Janssen, B. Nelson, and M. Kepner, 1993. "Use of a genetic algorithm in the cryptanalysis of simple substitution ciphers," *Cryptologica*, vol. 17 (1), pp. 31-44.
