A Genetic Algorithm for Training Recognizers of Latent Abnormal Behavior of Dynamic Systems

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- Keywords: Genetic Algorithm, Clustering, Machine Learning, Supervised Learning, Unsupervised Learning, Dynamic System, Abnormal Behavior, Training Set, Algebraic Approach, Axiomatic Approach.
- Abstract: We consider the problem of automatic construction of algorithms for recognition of abnormal behavior segments in phase trajectories of dynamic systems. The recognition algorithm is trained on a set of trajectories containing normal and abnormal behavior of the system. The exact position of segments corresponding to abnormal behavior in the trajectories of the training set is unknown. To construct recognition algorithm, we use axiomatic approach to abnormal behavior recognition. In this paper we propose a novel two-stage training algorithm which uses ideas of unsupervised learning and evolutonary computation. The results of experimental evaluation of the proposed algorithm and its variations on synthetic data show statistically significant increase in recognition quality for the recognizers constructed by the proposed algorithm compared to the existing training algorithm.

1 INTRODUCTION

Consider a dynamic system information about which can be accessed by reading data from sensors surrounding the system. The sensor readings are obtained from the sensors with a fixed frequency $1/\tau$.

A multidimensional phase trajectory in the space of sensor readings is an ordered set of vectors $X = (\overline{x_1}, \overline{x_2}, \dots, \overline{x_k})$, where $\overline{x_i} \in \mathbb{R}^s$ is a vector of sensor readings at $t = t_0 + i \cdot \tau$.

We assume that at any given moment of time the system can be in one of three states:

- *Normal state.* In this state, the system is fully functional.
- *Abnormal state.* In this state, the system is not fully functional or is going to lose some of its functions soon.
- *Emergency state*. In this state, the system is not functional.

The behavior that the system exhibits when it is in an abnormal state is called *abnormal behavior*. We assume that there are *L* classes of abnormal behavior, each of these classes is characterized by a phase trajectory X_{Anom}^{l} called *a reference trajectory*.

We assume that some period of time after exhibiting abnormal behavior of class l, the system enters an emergency state of class l. Our goal is to predict the emergency state of the system by recognizing the abnormal behavior that precedes it.

The observed phase trajectory X of the system can contain segments of abnormal behavior which are distorted compared to the reference trajectories. The distortions can be classified as amplitude distortions and time distortions. We say that a segment of abnormal behavior is distorted by amplitude compared to a reference trajectory if values in some points of the segment differ from those in the corresponding points of the reference trajectory. We say that a segment of abnormal behavior is distorted by time compared to a reference trajectory if there are missing or extra points in the segment compared to the reference trajectory. An example of an amplitude distortion is a stationary noise.

We need to recognize abnormal behavior of the system by finding abnormal behavior segments in the observed trajectory of the system and abnormal behavior class number for each segment found.

There are various problem settings that deal with recognition of abnormal behavior of dynamic systems. For example, (Yairi et al., 2001) considers recognition of anomalies in time series of housekeeping data of spacecraft systems. The problem setting in this paper differs from our paper: the authors build a model (a set of rules) for normal behavior of the system and consider any behavior deviating from

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this model abnormal. In our paper, we look for specific patterns of abnormal behavior in the observed trajectory of the dynamic system.

A variety of methods are used in the pattern recognition field, including the methods based on artificial neural networks (Haykin, 1998), k-nearest neighbor algorithm (Cover and Hart, 1967), algorithms based on Singular Spectrum Analysis (Hassani, 2007) and others. However, application of these methods and algorithms to this particular problem is complicated because of the presence of non-linear amplitude and time distortions of abnormal behavior segments in the observed phase trajectory X. To overcome these difficulties (emerging from the properties of dynamic systems in question) a parametric family of recognition algorithms based on axiomatic approach to abnormal behavior recognition was introduced in (Kovalenko et al., 2005). The idea of this parametric family is based on the idea of using algebraic approach to label planar configurations described in (Rudakov and Chekhovich, 2003). A genetic training algorithm for the parametric family was suggested in (Kovalenko et al., 2010) and improved in (Shcherbinin and Kostenko, 2013). Results from (Kostenko and Shcherbinin, 2013) show that this parametric family of recognition algorithms demonstrates high tolerance to non-linear amplitude and time distortions of abnormal behavior segments compared to other approaches.

The methods described above are developed for the case when the reference abnormal behavior trajectory is known and we need to find it in the observed trajectory, taking into account possible amplitude and time distortions. In this paper we consider a more difficult problem, when the exact position of the abnormal behavior trajectories in the training set is not known. We only know the points of time when the system exhibited emergency state. We assume that the training set consists of trajectories of normal behavior and trajectories which contain segments of abnormal behavior, while the exact position of these segments is not known. We call the problem of recognizing such abnormal behavior *latent abnormal behavior recognition* problem.

For this problem a directed search algorithm for training recognizers based on axiomatic approach was proposed in (Kostenko and Shcherbinin, 2013). This paper introduces a new algorithm for training recognizers based on axiomatic approach. The proposed algorithm uses ideas of unsupervised learning and genetic algorithms.

2 LATENT ABNORMAL BEHAVIOR RECOGNITION

We assume that we always know if the system is in an emergency state, but it is not immediately obvious (without analyzing the trajectory of the system) if the system is in normal or abnormal state. That means that when the training dataset containing examples of the trajectories is formed, we can't label the positions of the segments of abnormal behavior. We can only label the points where the system is in an emergency state, i. e. the points of emergency.

We assume that our dataset TS has the following structure.

- For each class *l* of abnormal behavior, *TS* includes trajectories which contain exactly one segment of abnormal behavior of class *l* and no segments of abnormal behavior of other classes. Such trajectories are called *emergency trajectories*, since they can be acquired by taking a segment of a system's trajectory that lies directly before the emergency point.
- *TS* also includes trajectories which contain no segments of abnormal behavior, i.e. where the system exhibits only normal behavior. We call these trajectories *normal trajectories*.

The problem of constructing a dataset with such structure (i. e. ensuring that the emergency trajectories contain exactly one segment of abnormal behavior and the normal trajectories contain no segments of abnormal behavior) is a separate problem which we don't consider in this paper.

The dataset TS is divided into two nonoverlapping parts: the training set \widetilde{TS} and the validation set \widehat{TS} . The training set \widetilde{TS} and the validation set \widehat{TS} have the same size and contain emergency trajectories for each class of abnormal behavior as well as normal trajectories.

Suppose we are given an objective function $\varphi(e_1, e_2) : \mathbb{Z}_+ \times \mathbb{Z}_+ \to \mathbb{R}_+$ which is non-decreasing w.r.t. both its arguments. The problem of automatic construction of latent abnormal behavior recognition algorithm is formulated as follows (Kostenko and Shcherbinin, 2013). Given a training set \widetilde{TS} , a validation set \widehat{TS} and an objective function $\varphi(e_1, e_2)$, produce a recognition algorithm Al that satisfies the following conditions:

1. Al should show limited number of type I and type II errors on the training set \widetilde{TS} :

 $e_1(Al, \widetilde{TS}) \leq const_1, e_2(Al, \widetilde{TS}) \leq const_2$ (1)

Here $e_i(Al, TS)$ is the number of type *i* errors that Al makes on the trajectories from TS.

2. Al should minimize the objective function $\varphi(e_1, e_2)$ on the validation set \widehat{TS} :

$$Al = \arg\min_{Al}(\varphi(e_1(Al, \widehat{TS}), e_2(Al, \widehat{TS}))) \quad (2)$$

The problem definition described here corresponds to the classic definition of the problem of supervised learning described in (Vorontsov, 2004) and (Vapnik, 1998).

In this paper, our objective function is a linear combination of the numbers of type I and type II errors: $\varphi(e_1, e_2) = a \cdot e_1 + b \cdot e_2, a, b > 0.$

3 AXIOMATIC APPROACH TO ABNORMAL BEHAVIOR RECOGNITION

In this section we describe the parametric family of algorithms for recognition of abnormal behavior of dynamic systems introduced in (Kovalenko et al., 2005).

3.1 Basic Notions

Let $X = (x_1, x_2, ..., x_k)$, be a one-dimensional trajectory, $x_t \in \mathbb{R}$.

An elementary condition ec = ec(t, X, p) is a function defined on a point *t* and its neighborhood on a trajectory *X*. It depends on a set of parameters *p* and takes either true value or false value.

An example of an elementary condition is

$$ec(t,X,p) = \begin{cases} true, & \text{if } \forall i \in [t-l,t+r] \\ a \le x_i \le b, \\ false, & \text{otherwise.} \end{cases}$$
(3)

Here $p = \{a, b, l, r\}$ is the set of parameters of this elementary condition, $a, b \in \mathbb{R}$, $a < b, l, r \in \mathbb{N}^+$.

This elementary condition is true whenever all values of the trajectory X in a specific neighborhood of point t lie between a and b.

Let $X = (\overline{x_1}, \overline{x_2}, \dots, \overline{x_k})$ be a multidimensional trajectory, $\overline{x_i} \in \mathbb{R}^s$.

An axiom a = a(t, X) is a function defined as a Boolean formula over a set of elementary conditions defined on a point t and its neighborhood on a multidimensional trajectory X:

$$a(t,X) = \bigvee_{i=1}^{p} \bigwedge_{j=1}^{q} ec_{ij}(t,X,p_{ij})$$
(4)

We call a finite collection of axioms $As = \{a_1, a_2, ..., a_m\}$ an axiom system if it meets the condition:

$$\forall X \,\forall \overline{x_t} \in X \quad \exists ! a_i \in As : a(t, X) = true \tag{5}$$

I. e. for any point t in any trajectory X there exists one and only one axiom a_i in axiom system As that is true on point t.

Any finite collection of axioms *as* can be transformed into an axiom system by:

1. Introducing an order in the collection by numbering axioms with consecutive integers:

$$as = \{a_1, a_2, \dots, a_M\}.$$
 (6)

We define that if an axiom with number *i* is true on a point *t* of a phase trajectory, then no other axiom of number j : j > i is true on *t*.

2. Adding to the set an axiom a_{∞} that has the lowest priority and is true at any point of any phase trajectory:

$$As = \{a_1, a_2, \dots, a_M, a_\infty\}.$$
 (7)

A marking of a trajectory $X = (\overline{x_1}, \overline{x_2}, ..., \overline{x_k})$ by an axiom system $As = \{a_1, a_2, ..., a_m\}$ is a finite sequence

$$J = (j_1, j_2, \dots, j_k)$$
 (8)

of numbers of axioms from as, such that a_{j_t} is true on the point t of trajectory X.

A marking of a reference abnormal behavior trajectory of class l is called *a model of abnormal behavior of class l*. We denote the model of abnormal behavior of class l as J_{Anom}^{l} .

3.2 The Recognition Algorithm

In accordance with (Kostenko and Shcherbinin, 2013), we define our parametric family of recognition algorithms S based on axiomatic approach as a family of algorithms, each of which is defined by a tuple

$$Al = (As, \{J_{Anom}^l\}_{l=1}^L, A_{search}),$$
(9)

where As is an axiom system, $\{J_{Anom}^l\}$ is a set of abnormal behavior models – one for each class of abnormal behavior, A_{search} is a fuzzy string search algorithm.

Al recognizes abnormal behavior segments in trajectory *X* by performing the following steps:

- 1. Perform marking of trajectory *X* by an axiom system *As*. We denote the marking of trajectory *X* as *J*.
- Perform fuzzy search for abnormal behavior models {J^l_{Anom}} in marking J using A_{search}.

The use of fuzzy search algorithms for searching for abnormal behavior models allows us to tackle time distortions. Algorithms based on DTW (Keogh and Pazzani, 2001) are used for marking search. To specify a recognition algorithm from parametric family *S* we need to construct an axiom system, construct an abnormal behavior model for each class of abnormal behavior and choose a fuzzy search algorithm and its parameters. Local optimization algorithms are used to adjust the parameters of marking search algorithm. The greatest difficulty is posed by construction of an axiom system and abnormal behavior models.

3.3 Existing Algorithm for Constructing Recognizers of Latent Abnormal Behavior

Here we give a brief description of the algorithm from (Kostenko and Shcherbinin, 2013) for construction of an axiom system and abnormal behavior models within the problem setting described in section 2.

The input of this algorithm includes a training set \widetilde{TS} , a validation set \widehat{TS} , an objective function $\varphi(e_1, e_2)$, the set of types of elementary conditions to use.

We define *intermediate objective function* $\psi(a)$ as follows:

$$\Psi(a) = \frac{freq_{Anom}^{TS}(a)}{(freq_{Norm}^{\widetilde{TS}}(a) + \delta)},$$
(10)

where $freq_{Anom}^{\widetilde{TS}}(a)$ is the frequency of fulfillment of an axiom *a* on the points of emergency trajectories of \widetilde{TS} , $freq_{Norm}^{\widetilde{TS}}(a)$ is the frequency of fulfillment of an axiom *a* on the points of normal trajectories of \widetilde{TS} , δ is a predefined small positive value.

The algorithm consists of the following two stages:

- 1. Selection of axioms. We form a set of axioms *AX*, axioms from which are more frequently fulfilled on emergency trajectories and less frequently fulfilled on normal trajectories of the training set. To do this, we perform the following steps:
 - (a) Selection of elementary conditions. This step involves grid search for parameter values of each type of elementary condition. We select a predefined number of elementary conditions which have the highest value of intermediate objective function Ψ .
 - (b) Construction of axioms from elementary conditions. At this step the elementary conditions selected at the previous step are iteratively combined into axioms using OR and AND operations. Then a specified number of axioms with the highest value of ψ is selected.

- 2. Construction of an axiom system and models of abnormal behavior. Here we form a single-axiom axiom system from each axiom in AX and iteratively add to each of the axiom systems an axiom from AX while it decreases the objective function φ on the validation set \widehat{TS} . To calculate φ for an axiom system As, we do the following:
- (a) Construct the model of abnormal behavior for each abnormal behavior class l as the longest common subsequence (LCS) (Cormen et al., 2001) of the markings by As of the emergency trajectories of class l from the training set \widetilde{TS} .
- (b) Use As and constructed models of abnormal behavior to recognize abnormal behavior in the validation set \widehat{TS} .
- (c) Calculate the number of errors and the objective function $\varphi(e_1, e_2)$.

We stop when we can't decrease φ anymore, or when we exceed the predefined number of iterations. The axiom system with the lowest φ is the result of the algorithm.

The described algorithm has two distinctive features. Firstly, the algorithm relies on the frequency of fulfillment of axioms on training set trajectories to select axioms for recognition of abnormal behavior. This approach may not always produce axioms that constitute good models of abnormal behavior. Secondly, the algorithm uses LCS to construct models of abnormal behavior for a given axiom system. But there may be abnormal behavior models other than the LCS of the markings of emergency trajectories that produce better recognition quality.

4 GENETIC TRAINING ALGORITHM FOR LATENT ABNORMAL BEHAVIOR RECOGNIZERS

In this paper we propose a new algorithm for construction of an axiom system and a set of abnormal behavior models for recognition of latent abnormal behavior. The new algorithm, similarly to the existing one described in 3.3, has two stages: on the first stage we construct a set of axioms AX, on the second stage we select axioms from AX to form an axiom system and models of abnormal behavior. On the first stage we form AX in an unsupervised manner using clustering of the trajectories of the training set. On the second stage we employ a genetic algorithm to construct the models of abnormal behavior and the axiom system using axioms from AX.

4.1 Construction of the Set of Axioms

For this stage we use the idea of time series clustering from (Yairi et al., 2001). The purpose of this stage is to construct axioms that meaningfully represent the states of our dynamic system. This approach is different from the existing algorithm where we use frequency of fulfillment of axioms to construct the set AX.

We define a *feature* as a function that maps a one-dimensional segment of a trajectory to a real value. Examples of a feature are maximum, minimum, mean, standard deviation.

To form the set of axioms *AX*, for each dimension *s* of the trajectories we perform the following steps:

- 1. Randomly select a specified number of onedimensional segments with a specified length Nfrom the dimension *s* of trajectories of \widetilde{TS} .
- 2. Transform each selected segment into a vector of feature values calculated for this segment. The set of used features is a parameter of the algorithm.
- 3. Perform clustering of feature vectors using kmeans clustering (Hastie et al., 2001). After clustering we get *K* centroid vectors.
- 4. Add to *AX K* axioms, where *i*-th axiom is of the form:

$$a_{i}(t, X) = \begin{cases} true, & \text{if } F(X_{[t-\lceil \frac{N}{2} \rceil: t+\lfloor \frac{N}{2} \rfloor]}^{s}) \\ & \text{belongs to} \\ & \text{i-th cluster,} \\ false, & \text{otherwise.} \end{cases}$$
(11)

Here *F* is a function that maps a one-dimensional trajectory segment to a feature vector, $X_{[t_1:t_2]}^s$ is the one-dimensional segment of dimension *s* of trajectory *X* from point t_1 to point t_2 .

Thus each axiom corresponds to a cluster in the feature space.

We define that a feature vector v belongs to the *i*th cluster if the centroid of *i*-th cluster is closer to v in Euclidean metric than centroids of other clusters.

In this paper, the following features were used:

- 1. Minimum value.
- 2. Maximum value.
- 3. Standard deviation.
- 4. Linear regression coefficient (Hastie et al., 2001) calculated using the following formula:

$$f(x_1, x_2, \dots, x_N) = \frac{\sum_{i=1}^{N} (x_i - \overline{x})(i - \frac{1+N}{2})}{\sum_{i=1}^{N} (i - \frac{1+N}{2})}, \quad (12)$$

where $x_1, x_2, ..., x_N$ are the values in the points of the segment, $\overline{x} = \frac{1}{N} \sum_{i=1}^{N} x_i$

4.2 Construction of an Axiom System and the Models of Abnormal Behavior

We construct the models of abnormal behavior and the axiom system by means of a genetic algorithm.

The individual of the algorithm is a pair $(As, \{J_{Anom}^{l}\}_{l=1}^{L})$, where As is an axiom system and $\{J_{Anom}^{l}\}_{l=1}^{L}$ is the set of abnormal behavior models which use axioms from As. These can be viewed as strings in the alphabet As.

The fitness function of the algorithm is the objective function $\varphi(e_1, e_2)$ calculated on the validation set \widehat{TS} . The goal of the algorithm is minimization of the fitness function.

This stage consists of the following steps:

- 1. Generate initial population of size PSize.
- 2. Mutate each individual in the population, while retaining the original individual, an add the mutated individual to the population. Now the population has size $2 \cdot PSize$.
- 3. Select $2 \cdot PSize$ pairs of individuals randomly from the current population, perform crossover on them (which yields 2 more individuals) and add the resulting individuals to the population. Now the population has size $4 \cdot PSize$.
- 4. Calculate the fitness function by running recognition on validation set \widehat{TS} for each member of the population.
- 5. Perform roulette wheel selection with elitism. Form a new population of size *PSize* by selecting a specified fraction of the individuals of the current population with the lowest fitness value and selecting the rest of the new population using roulette wheel selection.
- 6. Check the stopping criteria. If the specified number of iterations *IMax* is exceeded or the best fitness function did not decrease for *IMaxNonDecrease* iterations, stop. Otherwise, return to step 2.

The result of the algorithm is the individual with the lowest fitness value among the population after stopping.

In this algorithm we evolve an axiom system and the models of abnormal behavior. This is different from the existing algorithm where axiom systems are constructed separately and for each axiom system only models of abnormal behavior based on the LCS of the markings of emergency trajectories are considered.

In the rest of this section we describe the operations of the proposed genetic algorithm.

4.3 Generation of the Initial Population

Each individual $(As, \{J_{Anom}^l\}_{l=1}^L)$ in the initial population is generated in the following way:

- 1. As consists of a specified number of axioms randomly chosen from AX.
- 2. Each model of abnormal behavior J_{Anom}^l is formed as the marking of a randomly chosen emergency trajectory of class *l* from the training set \widetilde{TS} .

4.4 Mutation

To mutate an individual $(As, \{J_{Anom}^l\}_{l=1}^L)$, we first randomly select a class of abnormal behavior $l, 1 \le l \le L$. Mutation is performed on the model of abnormal behavior of class $l J_{Anom}^l$. We randomly select and perform one of the following actions:

- 1. Insert a randomly chosen axiom *a* from *As* at a random position into J_{Anom}^{l} .
- 2. Insert a randomly chosen axiom *a* from $AX \setminus As$ at a random position into J_{Anom}^{l} and add *a* to *As* with a random priority.
- 3. Replace an element of J_{Anom}^l at a randomly selected position with a randomly chosen axiom *a* from *As*.
- 4. Replace an element of J_{Anom}^{l} at a randomly selected position with a randomly chosen axiom *a* from $AX \setminus As$. The axiom *a* is also added to As with a random priority.
- 5. Remove an axiom at a random position of J^l_{Anom} . If the axiom does not occur in any of the models of abnormal behavior anymore, remove it from *As*.

4.5 Crossover

To cross two individuals $(As_1, \{J_l^1\}_{l=1}^L)$ and $(As_2, \{J_2^l\}_{l=1}^L)$, we first randomly select a class of abnormal behavior $l, 1 \leq l \leq L$. The result of the crossover operation is two new individuals: $(As_1', \{J_1^1, J_1^2, \ldots, J_1^{l-1}, J_1'^l, J_1^{l+1}, \ldots, J_1^L\})$ and $(As_2', \{J_2^1, J_2^2, \ldots, J_2^{l-1}, J_2'^l, J_2^{l+1}, \ldots, J_2^L\})$. The first individual inherits all models of abnormal behavior except the model of abnormal behavior of class l from the first parent, similarly with the second

individual. For parents' models of abnormal behavior of class l we perform one-point crossover. Denoting

$$J_{1}^{l} = (j_{1,1}^{l}, j_{1,2}^{l}, \dots, j_{1,p}^{l}),$$

$$J_{2}^{l} = (j_{2,1}^{l}, j_{2,2}^{l}, \dots, j_{2,q}^{l}),$$
(13)

we select two random integers: $r, 1 \le r \le p$ and $s, 1 \le s \le q$. New abnormal behavior models $J_1^{\prime l}$ and $J_2^{\prime l}$ are formed as follows:

$$J_1^{\prime l} = (j_{1,1}^l, j_{1,2}^l, \dots, j_{1,r}^l, j_{2,s+1}^l, \dots, j_{2,q}^l),$$

$$J_2^{\prime l} = (j_{2,1}^l, j_{2,2}^l, \dots, j_{2,s}^l, j_{1,r+1}^l, \dots, j_{1,p}^l).$$
(14)

The axiom system As'_i , $i \in \{1, 2\}$ of the offspring individual is inherited from the *i*th parent and then adjusted so that if an axiom is not present in any models of abnormal behavior anymore, it is removed from the axiom system, an if a new axiom that is not present in As_i is added to J'_i , it is added to the axiom system with a random priority.

5 EXPERIMENTAL EVALUATION

During experimental evaluation we compared the existing algorithm described in section 3.3 (denoted as A_{orig}) with the proposed algorithm described in section 4 (denoted as $A_{clust_genetic}$). We also experimented with the following two algorithms:

- An algorithm where the first stage is from the existing algorithm (i. e. we construct the set of axioms *AX* using the frequency of fulfillment of axioms on the trajectories of the training set), and the second stage is from the proposed algorithm (i. e. the genetic algorithm described in section 4.2). We denote this algorithm as *A*_{genetic}.
- An algorithm where the first stage is from the proposed algorithm (i. e. we construct the set of axioms AX using clustering of the trajectories of the training set, as described in section 4.1) and the second stage is from the existing algorithm (i. e. the axiom system and the models of abnormal behavior are constructed using simple directed search). We denote this algorithm as A_{clust}.

The objective function used during the experiments was $\varphi(e_1, e_2) = e_1 + 20 \cdot e_2$. The coefficient for type II errors is greater because type II errors are usually more costly than type I errors.

The experiments were conducted using synthetic data.

5.1 Synthetic Data

Consider a finite alphabet Σ where each symbol $x \in \Sigma$ corresponds to a segment of a one-dimensional trajectory. Then a string *s* in this alphabet corresponds to a

Table 2: Summary of the results. For each of the considered algorithms the table shows maximum, minimum and average numbers of type I and type II errors and the value of the objective function ϕ for the recognizer trained with the corresponding algorithm.

	A _{orig}			A _{clust}			Agenetic			$A_{clust_genetic}$		
	e_1	e_2	φ	e_1	e_2	φ	e_1	e_2	φ	e_1	e_2	φ
Maximum	1872	8	1872	70	6	174	82	3	104	77	1	97
Minimum	2	0	10	1	0	1	2	0	2	0	0	0
Average	111.7	1.1	133.8	10.3	0.1	11.8	15.4	0.1	17.1	5.4	0	5.7

Table 1: The functions used to generate segments of experimental data alphabet.

A	$y = x^3$
B	$y = -x^3$
C	$y = -(x-5)^2 + 25$
D	y = -x
E	y = x
F	$y = (x-5)^2 - 25$
G	$y = x^2$

trajectory X which is a result of concatenation of the segments corresponding to the symbols of the string. We call such a string s a *signature* of the trajectory X.

For generation of the data for the experiments, we consider an alphabet of segments which correspond to values of functions shown in table 1 in the points x = 0, 1, ..., 9. Each segment therefore has length 10.

Each dataset generated for the experiments had 1 dimension, 1 class of abnormal behavior, 20 emergency trajectories and 20 normal trajectories. Emergency trajectories were generated in the following way:

- 1. A signature s for the abnormal behavior segment was generated. The signature contained from 3 to 6 symbols.
- 2. Signatures containing 10 symbols were generated randomly for each emergency trajectory, then the signature of abnormal behavior segment was inserted at a random position in each signature of emergency trajectory. We also ensured that each signature of emergency trajectory had only one substring corresponding to the signature of abnormal behavior segment.
- 3. Signatures containing 10 symbols were generated randomly for each normal trajectory. We took care to generate normal trajectory signatures that don't contain abnormal behavior segment signature as a substring.
- 4. Each signature of emergency and normal trajectory was converted to the trajectory itself, during this process distortions were added:
 - Non-linear time distortion was added by ran-

domly and independently shrinking each segment up to 50% or growing it up to 200% of the original size.

• Amplitude distortion was added by applying Gaussian noise with $\sigma = 3$ to the resulting trajectory.

Note that because of the time distortion the length of each resulting trajectory could vary from 50 to 200 points.

5.2 Results

A total number of 132 experiments were conducted. The results of the experiments are summarized in table 2.

Using the binomial test (Conover, 1971), we were able to prove the following hypotheses with significance level $\alpha = 0.05$:

- Each of the algorithms A_{clust} , $A_{genetic}$, $A_{clust_genetic}$ with probability at least 0.9 trains a recognizer that delivers a lower value of the objective function φ on validation set than the best recognizer constructed by A_{orig} . We considered the alternative hypothesis that the probability of the event that each of the algorithms A_{clust} , $A_{genetic}$, $A_{clust_genetic}$ trains a recognizer whose objective function value is less than the objective function value of the recognizer trained by A_{orig} is less than 0.9. The *p*-value for the binomial test was 0.007.
- The algorithm $A_{clust_genetic}$ with probability 0.8 trains a recognizer that has 50% less type I errors and no more type II errors than the recognizer trained by A_{orig} . The *p*-value for the binomial test was 0.022.
- The algorithm A_{clust} with probability 0.8 trains a recognizer that has 10% less type I errors and no more type II errors than the recognizer trained by A_{orig} . The *p*-value for the binomial test was also 0.022.

Running times for each algorithm on a machine with processor Intel(R) Core(TM) i5-2520M CPU 2.50 GHz, 64 L1-cache and 4GB RAM is shown in Table 3.

	A _{orig}	A _{clust}	Agenetic	A _{clust_genetic}
Maximum	195 min	61 sec	80 min	26 min
Minimum	1.6 min	2 sec	6 min	2 min
Average	7.8 min	9.8 sec	19.7 min	8.7 min

Table 3: Average, maximum and minimum running times for each algorithm.

6 CONCLUSION

This paper considers the problem of automatic construction of algorithms that recognize segments of abnormal behavior in multidimensional phase trajectories of dynamic systems. The recognizers are constructed using a training set of example trajectories of normal and abnormal behavior of the system. The notable feature of the problem setting considered by this paper is that the exact position of the segments corresponding to abnormal behavior in the trajectories of the training set is unknown.

This paper proposes a two-step algorithm for training recognizers of abnormal behavior of dynamic systems. On the first step, axioms corresponding to typical patterns of the trajectories are constructed by clustering the trajectories of the training set. On the second step, genetic algorithm is used to construct the models of abnormal behavior of the dynamic system from the axioms obtained on the first step.

The proposed algorithm and its variations were empirically evaluated on synthetic data. The results of conducted experiments show that the proposed algorithm is able to improve recognition quality of trained recognizers compared to the existing training algorithm. On synthetic data, we were able to prove with significance level 0.05 a statistical hypothesis that the recognizer trained by the proposed algorithm with probability 0.8 makes 50% less type I errors and no more type II errors than the one trained by the existing algorithm.

For the variation of the proposed algorithm based on clustering and directed search, we were able to prove a statistical hypothesis that the recognizer trained by this algorithm with probability 0.8 makes 10% less type I errors and no more type II errors than the one trained by the existing algorithm. The advantage of this variation of the algorithm is that it runs considerably faster than the existing algorithm or the algorithm based on clustering and genetics (for synthetic data, average time was 9.8 seconds vs 7.8 minutes for the existing algorithm and 8.7 minutes for the algorithm based on clustering and genetics).

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