RELATIONAL SEQUENCE BASED CLASSIFICATION IN MULTI-AGENT SYSTEMS

Grazia Bombini, Nicola Di Mauro, Stefano Ferilli and Floriana Esposito

Dipartimento di Informatica, Università degli Studi di Bari, via Orabana 4, Bari, Italy {gbombini, ndm, ferilli, esposito }@di.uniba.it

Keywords: Group Behavior, Adversary Classification, Sequence Learning, Relational Sequence Similarity

Abstract: In multiagent adversarial environment, the adversary consists of a team of opponents that may interfere with

the achievement of goals. In this domain agents must be able to quickly adapt to the environment and infer knowledge from other agents' deportment to identify the future behaviors of opponents. We present a relational model to characterize adversary teams based on its behavior. A team's deportment is represent by a set of relational sequences of basic actions extracted from their observed behaviors. Based on this, we present a similarity measure to classify the teams' behavior. The sequence extraction and classification are implemented

in the domain of simulated robotic soccer, and experimental results are presented.

1 INTRODUCTION

In the Agent Modeling field an essential task consists of observing other agents and modeling their behavior. The main idea is to infer knowledge from other agents' deportment to identify the future behaviors of opponents. This model could be used to predict future behavior of the other agents, in order to coordinate and cooperate with them or counteract their actions. In multiagent adversarial environment, the adversary consists of a team of opponents that may interfere with the achievement of goals. In this domain agents must be able to adapt to the environment, especially to the current opponents' sequences of actions. The observed data from this kind of environments are inherently sequential, and hence it is necessary to have a mechanism able to handle sequential data.

In this paper we consider the problem of identifying agent behavior in complex domains where it is necessary to consider enormous and possibly continuous state and action spaces. In dynamic environments, the agents have a limited time to reason before to choose an action. For an effective adaptation, it is necessary to capture the similarity between observed behaviors, in order to adopt the most effective strategy. Hence, a mechanism for distinguishing the

different manners and classify them to recognize different adversary classes is essential. From this point of view, a key challenge is to determine what are the sequential behaviors characterizing a team.

Our proposal is to extract from row multi-agent observations of a dynamic and complex environment, a set of relational sequences describing an agent behavior. Then, the main aim is to propose a general method to classify different agents using those relational sequences. The goal of this paper has several aspect: abstracting useful atomic actions (*events*) from the multi-agent system log files; recognizing relational sequences of events that characterize the behavior of a team; extracting useful features from relational sequences; defining a similarity value between two feature-based sequence descriptions and comparing different teams' behavior by means of classification.

2 RECOGNIZING SEQUENCES OF ATOMIC BEHAVIORS

In an adversarial environment, the predicted behavior of other agents is referred to as an opponent model. One setting in which opponent modeling research has been conducted is the RoboCup Simulation league in the Robot World Cup Initiative¹ (Kitano et al., 1997). In this league technologies like multi-agent collaboration and adversary classification must be exploited.

RoboCup use Soccer Server System, a clientserver system for simulating soccer. The server keeps track of the current state of the world, and stores all the data for a given match in a log file. This represents a stream of consecutive raw observations about each soccer player's position, the position of the ball, the ball possessor, etc. at each moment of the match. From this log streams it is possible to recognize basic actions (high-level events). Each team have sequences of basic actions used to form coordinated activities which attempt to achieve the team's goals. In our work, we identify the following high-level events: - $catch(Player_n, T)$: $Player_n$ is a goalkeeper and catches

- the ball close to the penalty box at time T;
- $pass(Player_n, Player_m, T)$: $Player_n$ kicks the ball and the $Player_m$ gains possession, and both the players are of the same team at time T;
- **dribble**($Player_n, T$): $Player_n$ moves a significant distance since an opponent gains possession of the ball at time T;
- $intercept(Player_n, T)$: $Player_n$ gains possession at time T, and the previous ball owner belongs of the opponent team:
- **shoot**($Player_n, T$): $Player_n$ kick the ball close to the penalty box of the opposite team at time T;
- **outside**($Player_n, T$): $Player_n$ kick the ball at time T, and the ball go out of the bounds;
- **goal**($Player_n, T$): $Player_n$ kick the ball at time T, and the ball go in the goal.

Moreover, for each event the system takes into account some description related to the actions and its players:

- $dash(Player_n, T)$: if the $Player_n$ at time T dashes.
- $\mathbf{neck}(Player_n, T)$: if the $Player_n$ at time T looks around.
- $turn(Player_n, T)$: if the $Player_n$ at time T changes direc-
- **kick**($Player_n, T$): if the $Player_n$ at time T is a ball owner and kick the ball.
- **chngview**($Player_n, T$): if the $Player_n$ at time T changes

The log stream is processed to infer the high-level events occurred during a match. An event takes place when a ball possession changes or the ball is out of bounds. The event sequence of one team is separated from the events of its opponent team. Each next recognized event performed by the same team, forms the sequence until the opposing team gains the ball possession or the ball is out of bounds. An interesting sequence is composed at least by two successive actions performed by players of the same team. Sequences represent a symbolic abstraction of the row observation. A set of sequences is created for each team. This set characterizes the observed team behavior and separated models are learned for each team. The result of this phase is a set of the most meaningful relational sequences of recognized events that describes each team.

CLASSIFYING BEHAVIOUR BY RELATIONAL SEQUENTIAL **PATTERNS**

In this section we present a method based on relational pattern mining, to extract meaningful features able to represent relational sequences and a distance function to measure the dissimilarity between two corresponding feature vectors. Finally, those distances will be used in the k-nearest neighbor (k-NN) algorithm to classify the adversary behavior.

Given an alphabet of symbols \mathcal{A} , and let be $k \geq 1$ a positive integer, then a k-gram (k-mers), is a sequence σ of symbols over \mathcal{A} of length k ($\sigma \in \mathcal{A}^k$, $|\sigma| =$ k). For a given sequence $\sigma = (s_1 s_2 \dots s_t)$, the k-grams of interest are all subsequences $\sigma' = (s_i s_{i+1} \dots s_{i+k-1})$ of length k occurring in σ .

Now we can translate the concept of k-grams to the relational case. Given an alphabet of atoms \mathcal{A} , a **relational** k**-gram** is a relational sequence σ of length k defined over \mathcal{A} . Given a set of relational sequences $S = {\sigma_i}_{i=1}^n$, K is the set of all relational k-grams on all the sequences belonging to S:

$$\mathcal{K} = \bigcup_{i=1}^{n} K_{\sigma_i}.$$
 (1)

where K_{σ_i} is the set of all relational k-grams over the sequence σ_i . In particular, \mathcal{K} represents the set of all relational features over S. We define $K(\alpha) \subseteq K$, the set of relational k-grams having a support greater than $\alpha - 1$: $\mathcal{K}(\alpha) = {\sigma | \sigma \in \mathcal{K} \land support(\sigma) \ge \alpha}.$

In order to select the best set of features, we use an Inductive Logic Programming (ILP) (Muggleton and De Raedt, 1994) algorithm, based on (Esposito et al., 2008), for discovering relational patterns from sequences. It is based on a level-wise search method, known in data mining from the APRIORI algorithm (Agrawal et al., 1996). It takes into account the sequences, tagged with the belonging class, and the α parameter denoting the minimum support of the patterns. It is essentially composed by two steps, one for generating pattern candidates and the other for evaluating their support. The level-wise algorithm makes a breadth-first search in the lattice of patterns ordered by a specialization relation. Starting form

¹ http://www.robocup.org/

the most general pattern, at each level of the lattice the algorithm generates candidates by using the lattice structure and then evaluates the frequencies of the candidates.

Given a set of sequences \mathcal{S} , we apply the algorithm previously described (Esposito et al., 2008), to find all the relational k-grams $\mathcal{K}(\alpha)$ over the set \mathcal{S} with a support at least equal to α . $\mathcal{K}(\alpha)$ is the ordered set of *features* \mathcal{F} that will be used to compute the boolean vector representation of each sequence in the following way. Given a sequence $\sigma \in \mathcal{S}$, and $\mathcal{F} = \mathcal{K}(\alpha) = \{\omega_i\}_{i=1}^n$ the set of relational k-grams over \mathcal{S} , the *feature vector* of σ is

$$V_{\mathbf{\sigma}} = (f_1(\mathbf{\sigma}), f_2(\mathbf{\sigma}), ..., f_n(\mathbf{\sigma}))$$

where

$$f_i(\sigma) = \begin{cases} 1 & if \ \omega_i \sqsubseteq \sigma \\ 0 & otherwise \end{cases}$$

Now, the function distance $d_r(\cdot, \cdot)$ between two relational sequences σ_1 and σ_2 is computed using the classical Tanimoto measure (Duda et al., 2000):

$$d_{r_1}(\sigma_1, \sigma_2) = \frac{n_{1\sigma_1} + n_{1\sigma_2} - 2n_{1\sigma_{12}}}{n_{1\sigma_1} + n_{1\sigma_2} - n_{1\sigma_{12}}} = \frac{2(n - n_{1\sigma_{12}})}{2n - n_{1\sigma_{12}}}$$
(2)

where $n_{1\sigma_i} = n = |\mathcal{F}|$ is the number of the features, and $n_{1\sigma_{12}} = |\{f_i|f_i(\sigma_1) = f_i(\sigma_2)\}|$ is the number of features with the same value in both σ_1 and σ_2 . However, this basic formulation takes into account features not appearing (with value 0) in the sequences, and in case of a lot of feature this can lead to underfitting. Equation (2) may be extended in the following way:

$$d_{r_2}(\sigma_1, \sigma_2) = \sum_{i=1}^{n} \frac{f_i(\sigma_1) + f_i(\sigma_2) - 2f_i(\sigma_1)f_i(\sigma_2)}{f_i(\sigma_1) + f_i(\sigma_2) - f_i(\sigma_1)f_i(\sigma_2)}$$
(3)

where $n_{2\sigma_i} = \sum_{j=1}^n f_j(\sigma_i)$ is the number of the features holding in the sequence σ_i , and $n_{\sigma_{12}} = |\{f_i|f_i(\sigma_1) = f_i(\sigma_2) = 1\}|$ is the number of features that hold both in σ_1 and σ_2 .

4 EXPERIMENTS

In order to evaluate our approach we analyze log files of soccer games of the RoboCup 2008 Exercise Competitions². This is a preceding event for RoboCup initiative, and includes a 2D simulation league. We have implemented a system that is able to identify and extract the interesting sequences of coordinated team behaviors using the recorded observations (logs) of this simulation games. There is an underlying assumption, that the strategy of a team does not change during the competition. We have analyzed the log files for 4 teams, concerning to 4 matches of the competition, 2 matches for each team. One adversary

class was created for each team by analyzing the log files of two matches of the same team, producing a set of relational sequences. Each sequence is made up of interesting uninterrupted consecutive actions performed by player of the same team and represents its characteristic behavior. From the row observations of the log files we have obtained the dataset. It is made up of 443 sequences, defined on 7 atomic behaviors (catch, pass, dribble, etc.) and 5 action descriptions (neck, turn, kick, etc.). In particular, we have 112 sequences for the first team C0, 106 sequence for the second team C1, 93 sequence for the third team C2 and 132 sequence for the fourth team C3.

After creating these adversary classes, the goal was to identify which teams were playing based on one sequence of its actions. A weighted 10-NN classifier was constructed and tested using the 10-fold cross-validation to find the classification accuracy. In the first step, the set $\mathcal{K}(\alpha)$ of frequent k-grams has been mined. Here, α denotes the support of each k-gram $\sigma \in \mathcal{K}(\alpha)$ corresponding to the ratio $sup-port(\sigma)/|\mathcal{S}|$, where \mathcal{S} is the set of sequences in the training set for each fold.

In this experiment, α has been set to 0.10, 0.15, 0.20 and 0.25, and the algorithm extracted, respectively, 1095.2, 720.3, 501.4 and 315.2 k-grams on average on the 10 fold. The average accuracy results, respectively, 68.39, 66.17, 68.63 and 63.9. The results for different value of α are shown in Table1. Since in this experiment there are 4 classes to be distinguished, the accuracy on guessing should be equal to 25%, proving that our accuracy is better then guessing.

5 CONCLUSIONS AND RELATED WORKS

In competitive domains, the knowledge about the opponent can be very advantageous. In the area of Agent Modeling, Kaminka et al. (Kaminka et al., 2002) focus on the unsupervised autonomous learning of the sequential behaviors of agents, from observations of their behavior using a hybrid approach to produce time-series of recognized atomic behaviors. These time-series are then analyzed to find sub-sequences characterizing each agent behavior.

Similarly to the previous approach, Lattner et al. (Lattner et al., 2005), present a symbolic approach based on association rule mining for pattern matching on qualitative representations. The process creates patterns in dynamic scenes, based on the qualitative information of the environment, producing a set of prediction rule. However, these previous works

²http://robocup-cn.org/en/exercise/08/

	Class	F1	F2	F3	F4	F5	F6	F7	F8	F9	F10	Mean
$\alpha = 0.25$	C0	54.55	40	40	10	60	60	36.36	30	75	30	39.42
	C1	70	66.67	72.73	60	22.22	10	30	60	10	30	43.16
	C2	100	100	75	88.89	100	88.89	100	88.89	88.89	90	92.06
	C3	66.67	76.92	53.85	84.62	84.62	100	92.31	38.46	100	85.71	78.32
$\alpha = 0.20$	C0	36.36	50	40	20	50	70	72.73	40	33.33	40	42.24
	C1	90	75	72.73	40	22.22	20	20	50	10	40	43.99
	C2	100	100	83.33	88.89	100	88.89	100	88.89	100	90	94
	C3	80	92.31	100	76.92	92.31	100	92.31	46.15	100	85.71	88.75
$\alpha = 0.15$	C0	54.55	40	50	30	70	80	72.73	30	33.33	40	50.06
	C1	70	66.67	63.64	50	22.22	20	20	20	10	30	36.25
	C2	100	100	83.33	88.89	100	88.89	88.89	88.89	88.89	90	91.78
	C3	86.67	84.62	61.54	76.92	92.31	83.33	76.92	84.62	84.62	92.86	82.44
$\alpha = 0.10$	C0	63.64	20	50	30	60	50	72.73	40	33.33	30	44.97
	C1	80	66.67	36.36	50	44.44	30	30	50	20	30	43.75
	C2	88.89	100	58.33	88.89	100	88.89	100	100	88.89	90	91.39
	C3	80	92.31	84.62	69.23	100	83.33	92.31	100	100	85.71	88.75

Table 1: Classification accuracy using 10-fold cross-validation.

focused on unsupervised learning, with no ability to classify behaviors into classes.

In this area, Riley and Veloso (Riley and Veloso, 2000) present an approach that model high-level adversarial behavior by classifying the current opponent team into predefined adversary classes. It is assumed that opponent teams do not change strategies during the league. Their system could classify fixed duration windows of behavior using a set of sequence-invariant action features. The authors use a windowing approach to extracting useful feature removing time sequencing from data, but this length affect the accuracy of the classifier and its performance. To classify the instance of observations decision tree on flat symbols are used.

In our work, we proposed a relational model to characterize adversary teams based on its behavior. A team's deportment is represent by a set of relational sequences of basic actions extract to their observed behaviors. Based on this, a similarity measure for classify the teams' behavior has been presented.

The log files used to extract the dataset, are the results of the matches of a RoboCup competition. In order to create winner teams, many people working together using a great variety of technique and strategies. Moreover, we create an adversary class for each team. If two teams adopt a similar strategy, we request to the classification method to distinguish them. A more refined method to define adversary classes, likely could improve the classification accuracy. Experimental results proved the validity of the proposed approach. As a future work, we will investigate methods for extracting patterns with a high discriminative power, and we will compare different similarity func-

tions.

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