Advanced Human Machine Interaction

(Interaction) Data Analysis

Alexandre Pauchet

alexandre.pauchet@insa-rouen.fr - BO.B.RC18







IHME/IDA: objectives



Data analysis: goals



Problem modeling

Formal modeling of a problem

• Define formally the inputs

Ex: "input = $\{a_1, ..., a_n\}_{n < 20}$ a sequence of actions with each action a_i in $\{up, down, left, right, space\}$ "

Define formally the outputs

Ex: "output = a class among {beginner, advanced}"

Ex: "output = a sequence of words generated as an answer to the user's query $\{w_1, ..., w_n\}_{n>0}$ "

Define a class of problem / algorithm

- Ex (machine learning): classification problem, clustering problem, regression problem, ...
- Ex (logic): induction / deduction problem, ...
- Ex (algorithm): sorting, graph construction, ...

Analysis of interaction data



The data to analyze can be collected from:

- External/internal sensors (context of use)
- Interaction data (log of user actions and/or activity)
- External data requested from the interactive system

Input: discrete data (1/2)

• Discrete sequences

- Ex: sequence of visited web pages, of user actions
- Representation: ordered set $\{a_1, a_2, ..., a_n\}$

Discrete sequences of item-sets

- Ex: actions from left and right hands in a double-handed game or multiple paddles/keyboards
- Representations:
 - ordered set of item-sets

$$\{\{a_{1,1}, \ldots, a_{m,1}\}, \{a_{1,2}, \ldots, a_{m,2}\}, \ldots, \{a_{1,n}, \ldots, a_{m,n}\}\}$$

• matrix

$$\{a_{i,j}\}_{i=1..m; \ j=1..n}$$

Input: discrete data (2/2)

- Independent sequences
 - Ex: actions from two different players
 - Representation: set of ordered sets
 - $\{\{a_{1,1}, \ \dots, \ a_{m1,1}\}, \ \{a_{1,2}, \ \dots, \ a_{m2,2}\}, \ \dots, \ \{a_{1,n}, \ \dots, \ a_{mn,n}\}\}$

Remarks:

- Time is not considered and delays may be different (or not) between two actions
- Different sampling are possible in independent sequences

Input: continuous and mixed data

Continuous signal

- Ex: user's voice volume
- Representation: function f(t)

Continuous signals

- Ex: trajectories of two Wiimotes
- Representation: set of functions { $f_1(t)$, ... $f_n(t)$ }

Remarks:

- Any continuous signal can be discretized
- Mix of discrete sequences and continuous signals
- There always is sampling

Goal / Objective (class of problem)

Behavioral pattern extraction

- Intra- / Inter- user behavioral patterns
- (Multiple) Time scaling
- Frequent patterns / similar patterns

User classification

- Clustering of users' activity
- Segmentation and classification of parts of users' activity

Generation

- Simulation of user's behavior
- Generation of interactive behavior

Combination of problems



Exercise

- Let $A = \{a_1, ..., a_n\}$ be the set of *n* actions that users can perform. Let *T* be a set of *r* interaction traces from *s* different users $u_1, ..., u_s$:
 - $T = \{\{a_{1,x1}, \dots, a_{m1,x1}\}_{ux1} \dots \{a_{1,mr}, \dots, a_{mrxr}\}_{uxr}\}$
- eg. HMI with 3 possible actions: {{a1 a2 a1 a3}u1 {a2 a2 a2 a1 a3 }u2 {a2 a1 a2 a2 a1 a1 a3}u1 {a1 a1 a3}u2 {a1 a1 a2 a1 a3}u1 }
- Formalize the problem that, from a given sequence of actions, predicts the next action

Discrete sequences Pattern extraction Similarity-based approach

Sequence alignment



ATAGCTATAGTCATCTTACTCATCTTACGCT GCTACGAATCTCTGAATAACGCTACGAATCC...

... CTGTCCTGCATCACTGGATGTACCATCTTAC TATGGTACATCTATGTACTACTCAC-TTTTAAC TCTTACGCTATAGCTATAGTCTACGAATCAC... A)T

String mining and sequence alignment

- Alignment of two sequences of characters
 - Used to compare 2 sequences S_1 (I=m) and S_2 (I=n)
 - How S_1 can be transformed into S_2 ?
 - Based on a distance or a similarity measure
 - A sequence alignment can be computed using dynamic programming in O(mn)
- 2 types of alignments:
 - Global
 - Local (Smith & Waterman, 1981)

Sequence alignment

- (A T G C T A) is an alignment of the two sequences
 "ACGA" and "ATGCTA".
- Algorithmically, it corresponds to an edition script (i.e. a computer program)

Operation	Resulting sequence
Substitution of A by A	Α
Substitution of C by T	AT
Substitution of G by G	ATG
Insertion of C	ATGC
Insertion of T	ATGCT
Substitution of A by A	ATGCTA

Local alignments

- 3 editing operations
 - Substitution of a symbol from S_1 at a given position by a symbol from S_2
 - Deletion of a symbol from S_1 at a given position
 - Insertion of a symbol in S_2 at a given position
- Scores
 - Sub(a,b): score to substitute symbol a by symbol b
 - *Del(a)*: score to delete symbol *a*
 - Ins(a): score to insert symbol a

Similarity measure between 2 sub-sequences

- $E_{x,y}$: series of editing operations that transform x into y
- A score of *e* is computed as the sum of all its elementary editing operations

Dynamic programming



sub(x,x) = 1sub(x,z) = 0lns = del = -1

	A	С	G	Т	C	G	A	С	G
A	1	0	0	Q	0	0	1	0	0
С	0	2	1	0	1	0	0	2	1
Т	0	1	2	2	1	1	0	1	2
С	0	1	1	2	2	2	1	1	1
A	1	0	1	1	2	3	3	2	1
С	0	2	1	1	2	2	3	4	3
G	0	1	3	2	1	3	2	3	5

t[-1,-1]

t[i, -1]

t[-1, j]

t[i,j]

Smith & Waterman

$$= 0,$$

$$= 0,$$

$$= 0,$$

$$t[i-1, j-1] + Subs(x[i], y[j]),$$

$$t[i-1, j] + Del(x[i]),$$

$$t[i, j-1] + Ins(y[j]),$$

$$0$$

Remarks:

- Subs/Del/Ins have negative values
- The value at (i,j) position in table t only depends on the 3 adjacent positions

• An optimal alignment (i.e. of maximum score) can be produced by performing a trace back in the values of table t from the (maximal) values up to a position of 0.

Example

	A	С	G	T	С	G	A	С	G		A	С	G	Т	С	G	A	С	G
Α	1	0	0	0	0	0	1	0	0	A	1	0	0	0	0	0	1	0	0
С	0	2	0	0	1	0	0	2	0	С	0	2	1	0	1	0	0	2	1
Т	0	0	0	1	0	0	0	0	0	Т	0	1	2	2	1	1	0	1	2
С	0	1	0	0	2	0	0	1	0	С	0	1	1	2	3	2	1	1	1
Α	1	0	0	0	0	0	1	0	0	Α	1	0	1	1	2	3	3	2	1
C	0	2	0	0	1	0	0	2	0	С	0	2	1	1	2	2	3	4	3
G	0	0	3	1	0	2	0	0	3	G	0	1	3	2	1	3	2	3	5

Two accumulated similarity tables obtained using the Smith-Waterman algorithm. The left has been calculated using a similarity score of 1 for matches, and dissimilarity penalties of -2 for non-matching substitutions and indels. The right table has this penalty reduced to -1. In each case, the alignments with a similarity score of at least 3 have been highlighted. Note how the higher penalty leads to smaller, more local alignments.

Exercise

 Formalize the problem corresponding to the pattern extraction from discrete sequences using a similarity-based approach

From patterns to frequent patterns



Discrete sequences Pattern extraction Frequency-based approach

- Frequent sequential patterns
 - Ex: users that perform action 'A', often perform action 'B' shortly after
 - Association rules ('A' => 'B' shortly after)
 - Gap: number of elements (actions) between 'A' and 'B'
 - Confidence: how often a rule has been found to be true
- A simple structure: suffix tree

Suffix trees

- The **non-compact suffix tree** of a word *y* is the deterministic finite automaton, having a single initial condition called *root* and where the terminal states correspond to the suffix of the word. The language recognized by this automaton is all suffixes of y.
- In practice a terminator is added at the end of the word (usually denoted \$).
- Leaves are numbered according to the starting position of the suffix they recognize.
- To **compact the tree**, the internal nodes having only a single outgoing branch are removed and the branches are concatenated.

Example of suffix tree (single word)



Generalized suffix tree (multiple words)



Application: CISMeF

 Extraction of recurrent behaviors in the navigations within an online health catalog (CISMeF)



Application: CISMeF

- Data preparation
 - Episode extraction: IP + semantic distance between documents + time between requests
 - Resource identification: unique ID + delimiter
 example of session: /59451/ /303901/ /170702/
- Recurrent pattern extraction
 - Generalized suffix tree
 - Longest repeated substrings

Application: CISMeF

Nombre de liens visités	Nombre d'épisodes	Proportion	22 days of log analysis:10mn max for an episod	е			
1	34 005	70,6 %	 48 168 episodes 17mn of data processing 	n (2 30G	H7/512	MO)	
2	8 254	17,1 %		J (2,000	112/012	.1010)	
3	2 940	6,1 %					
4	1 284	2,7 %					
5	658	1,4 %					
6	346	0,7 %					
7	216	0,4 %					
8	139	0,3 %					
9	91	0,2 %	I an guarra dag matifa	2	2	1	5
10	60	0,1 %	Longueurs des mours	2	3	4	5
>10	175	0,4 %	Nombres de motifs	1557	146	20	4

	Episodes contenant un motif							
	longueur 2 longueur 3 longueur 4 lo							
Effectifs	4127	326	42	8				
% épisodes	8,568	0,677	0,087	0,017				
% épisodes (l>1)	29,139	2,302	0,297	0,056				

Discrete sequences

- Pattern extraction:
 - Sequence alignments

 \rightarrow similarity OK, but frequency is difficult to evaluate (should be paired with pattern clustering)

- Prediction:
 - Generalized suffix trees
 - Seq2seq, CRF, HMM, ...

 \rightarrow frequency OK, but only slight variations (similarity) can be taken into account

Exercise

- Formalize the problem that consists in predicting the most probable action of user given the set of previous actions performed?
- What would be an algorithm to construct a (compact) suffix tree?

Discrete sequences of item-sets Pattern extraction Similarity-based approach

Approach (similar to sequences)



Decomposition in two separated steps

- Extraction of (pair of) interaction patterns
- Clustering of interaction patterns



Rick Moritz (PhD thesis), Orange labs - Meylan







sim(a,b) =





Evaluation: usual measures



Model evaluation:
 – Precision

 $precision = \frac{|\{relevant \ documents\} \cap \{retrieved \ documents\}|}{|\{retrieved \ documents\}|}$

- Recall

 $\operatorname{recall} = \frac{|\{\operatorname{relevant} \operatorname{documents}\} \cap \{\operatorname{retrieved} \operatorname{documents}\}|}{|\{\operatorname{relevant} \operatorname{documents}\}|}$

- F-measure
$$F = 2 \cdot \frac{\text{precision} \cdot \text{recall}}{\text{precision} + \text{recall}}$$

- Remarks:
 - Unbalanced classes!
 - Evaluation by class

Evaluation

- Data and ground truth
- Computation time
- Precision and recall for each pattern alignment
- Alignment size
- Number of alignments



Total pattern cells: 24 Total aligned cells: 20 Aligned pattern cells = 16 Precision = 16/20 = 4/5Recall = 16/24 = 2/3Size ratio = 20/24 = 5/6

(False Positives) Aligned cells outside pattern Unaligned pattern cells *(False Negatives)* *(True Negatives)* Unaligned non-pattern cells Aligned pattern cells *(True Positives)*

• Synthetic data generation



• Scenarios:

- 1) Regular activity, no noise
- 2) Regular activity, noise between patterns
- 3) Noisy patterns
- 4) Irregular pattern apparition
- 5) Faulty sensors (3/4 random data)

Measure	Dataset 1	Dataset 2	Dataset 3	Dataset 4	Dataset 5
$\frac{\#\text{alignments}}{\#\text{pairs of pat.}}$	0.31 ± 0.26	0.69 ± 0.29	0.041 ± 0.054	0.31 ± 0.26	0.22 ± 0.32
precision	0.54 ± 0.22	0.77 ± 0.20	0.13 ± 0.06	0.52 ± 0.24	0.54 ± 0.18
recall	1.00 ± 0.00	1.00 ± 0.00	0.56 ± 0.19	1.00 ± 0.01	0.20 ± 0.06
$\frac{\text{alignment size}}{\text{pattern size}}$	5.66 ± 4.48	2.45 ± 2.51	6.72 ± 3.96	6.65 ± 6.08	1.60 ± 0.24

Exercise

 Formalize the problem of behavior pattern extraction from discrete sequences of set of actions, based on a similarity approach (looking for longest and more similar behaviors) Discrete sequences of item-sets Pattern extraction Frequency-based approach

Data-mining sequences of item-sets

- Frequent item-sets
 - Ex: users usually select A & B keys
 - Association rules ('A' => 'B' in the same item-set)
 - Support: how frequently an item-set appears in the dataset
 - Confidence: how often a rule has been found to be true
- Frequent sequential patterns
 - Ex: users that perform action 'A', often perform action 'B' shortly after
 - Association rules ('A' => 'B' in two following item-sets)
 - Gap: item-sets appearing inside the association rules

Definitions

- Item: minimal element
- **Item-set**: ordered (preference, ...) set of items
- Sequence of item-sets: ordered set of item-sets
- **Transaction**: tuple in the database; a set of items or a sequence of item-sets.
 - training set = set of transactions
 - $\ \{\{a_{1,1}, \ \dots, \ a_{1,m1}\}, \ \{a_{2,1}, \ \dots, \ a_{2,m2}\}, \ \dots, \ \{a_{n,1}, \ \dots, \ a_{n,mn}\}\}$
 - Or a sparse matrix...
- Association rule: application $X \to Y$ where X and Y are disjoint set of items or set of item-sets

Evaluation of association rules (X \rightarrow **Y)**

- **Support**: absolute probability $P(X \cup Y)$ $||X \cup Y||/||BD|| = \%$ of transactions satisfying the rule
- **Confidence**: conditional probability P(Y/X)

 $||X \cup Y|| / ||X|| = \%$ of transactions

verifying the implication

= support(XY)/support(X)

 An interesting association rule is a rule whose Confidence > Minconf and Support > Minsup

Apriori algorithm (Agrawal & Srikant, 1994)

- Idea: if an item-set is not frequent, then all its supersets are not frequent:
 - If {A} is not frequent then {AB} cannot be frequent
 - if {AB} is frequent then {A} and {B} are frequent
- Process:

1)Generate iteratively candidate item-sets:

- First pass: search for frequent 1-sets
- Generate a candidate of size k from two candidates of size k-1 differentiated by the last element
- Filter the sets of items with minimum support (keeping frequent item-sets)

2)Use frequent item-sets to generate association rules

Apriori

Apriori

Input: L₁={frequent 1-itemsets}; Output: L_k={frequent k-itemsets};

```
for (k=2; L_{k-1} \neq \emptyset; k++) do {

C_{k}=apriori-gen(L_{k-1});

// Generate new candidates

}

L_{k}={ c\inC_{k} | numberOf(c,DB) >=

minsup };

// Filter candidates

}

return L_{k};
```

Apriori-gen

Input: L_{k-1}={frequent (k-1)-itemsets};
Items of L_{k-1} are ordered lexicographically
Output: C_k={candidates frequent (k)itemsets};

Step 1: Self-join on L_{k-1}
 For each (p_{k-1},q_{k-1}) so that p_{k-1}<q_{k-1} do {
 C_k={ c_k | lexicographically ordered combinaison of p_{k-1} and q_{k-1}}
 }

```
    Step 2: Pruning
        foreach c<sub>k</sub> in C<sub>k</sub> do {
        foreach (k-1)-subsets t<sub>k-1</sub> of C<sub>k</sub> do {
        if (t<sub>k-1</sub> is not in L<sub>k-1</sub>) then
        delete t<sub>k-1</sub> from C<sub>k</sub>
        }
```

Apriori (exemple)

min_support=2



Apriori (generating association rules)

```
//Input: MinConf, L<sub>k</sub> (frequent item-sets)
//Output: R, set of association rules
```

Example :	
$\begin{array}{l} \{2 \ 3\} \rightarrow \{5\} \\ \{2 \ 5\} \rightarrow \{3\} \end{array}$	confidence=2/2 confidence=2/3
… {2} → {3 5} …	confidence=2/3

Exercise

 Formalize the problem of behavior pattern extraction from discrete sequences of set of actions, based on a frequency approach (looking for most frequent behaviors) Continuous signal(s)

(Single) continuous signal

Continuous numeric acquisition is impossible

- Discretize an analogical signal to numerical values along 2 dimensions:
 - Continuous / discrete (alphabet)
 - Time (sampling)

NB: a re-sampling can be necessary according to the goal

Discretization process

- Heterogeneous sensors (unnormalized)
- Semantic information (analyze & user's feedback)

Example: GPS signal



Daniel Ashbrook & Thad Starner : « Using GPS to learn significant locations and predict movement across multiple users », Personal and Ubiquitous Computing, volume 7, number 5, pp. 275-286, Springer, 2003.

- Finding **significant places** *I* **positions** (time dependence: time threshold)
- Clustering places into **locations / keypoints** (spacial dependence: cluster radius)



57/59

Example: GPS signal



Example: GPS signal



Transition	Relative Frequency	Probability
$A \to B$	14/20	0.7
$A \to B \to A$	3/14	0.2142
$A \to B \to C$	2/14	0.1428
$A \to B \to D$	3/14	0.2142
$A \to B \to E$	1/14	0.0714
$A \to B \to F$	1/14	0.0714
$A \to B \to G$	1/14	0.0714
$A \to B \to H$	1/14	0.0714
$A \to B \to I$	1/14	0.0714
$B \to A$	16/77	0.2077
$B \to A \to B$	13/16	0.8125
$B \to A \to J$	3/16	0.1875
$B \to C$	10/77	0.1298
$B \to C \to A$	6/10	0.6
$B \to C \to K$	4/10	0.4
$D \to B$	5/7	0.7142
$D \to B \to A$	2/5	0.4
$D \to B \to L$	2/5	0.4
$D \to B \to M$	1/5	0.2

Probabilities for transitions in Markov models Key: A = "Home"

Exercise

- Formalize the problem of finding keypoints from GPS data?
- Formalize the problem of behavior pattern extraction from a sequence of visited places? From GPS data?

(Multiple) Continuous signals



Multiple sensors: (discrete) approaches

