RANKING EPISODES USING A PARTITION MODEL

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RANKING EPISODES

Episodes are

(i) patterns occurring in sequences, (*ii*) order of events is described by DAGs, (*iii*) gap events are allowed.

Example:

Episode G occurs in a sequence if and only if (i) a occurs, (ii) then b and c, in any order, (iii) and then *d*. Gap events are allowed.



INDEPENDENCE MODEL

We need

p(episode occurs in a sequence of length n)

No closed form but can be computed.

Example:

Episode *G* occurs in a sequence if and only if we reach H_6 from from H_1 in M.

PARTITION MODEL

General idea:

(*i*) split the episode into two subepisodes, *(ii)* study how often these episodes occur, (iii) incorporate this into a model, (*iv*) try all splits, and use the best.

Example: Split *G* into G_1 and G_2



For a set of sequences S, the support is

 $supp(G) = |\{S \in S \mid G \text{ occurs in } S\}|$.

Mining using support counter-productive: (*i*) output is humongous, *(ii)* pattern are redundant

Rank patterns based on expectation. Computing expectation is more intricate than with itemsets: *(i)* models are difficult to compute, *(ii)* depends on the sequence length

Assume we can get

p(n) = p(G occurs in a sequence of length n)

Then the expected support is



To compute the probability use

$$p(H,n) = q \times p(H,n-1) + \sum_{e=(F,H)\in E(M)} p(e)p(F,n-1)$$

where q is the probability of being stuck in Hfor a single event

$$q = 1 - \sum_{e \in E(M)} p(e) \quad .$$

FREE-RIDER EPISODES

Independence model does not get rid of freeriders.

 G_1 and G_2

Model how likely G_i is discovered once we see at least one event in G_i .

In practice, boost the probabilities p(e) in



Boosted probabilities leads to higher expected support, and lowered rank.

Edge *e* with label *l* is modeled as

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p(e) \propto \begin{cases} \exp(u_l + t_1), & \text{if } e \in C_1, \\ \exp(u_l + t_2), & \text{if } e \in C_2, \\ \exp(u_l), & \text{otherwise} \end{cases}
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Parameters u_l and t_1 can be learned by maximizing likelihood; gradient descent will converge to the global optimum.

 $\mu = \sum p(|S|) \quad .$ $S \in S$

Compare supp(G) and μ : The larger the difference, the more important is the episode.

Example: If *G* is significant, then G' is also significant even if x is independent of G.



EXPERIMENTS

Top episodes in *Plant* dataset. The symbols *x* and *y* represent noise events.

Indepe	endence model		Partition model				
Rank	Episode type	$r_{ind}(G)$	Rank	Episode type	$r_{prt}(G)$		
1.	$a \rightarrow b \rightarrow c \rightarrow d$	∞	1.	$a \rightarrow b \rightarrow c \rightarrow d$	10^{308}		
2.	$k \xrightarrow{h} n \longrightarrow l$	249	2.	$e { ightarrow} f$	128		
3.–7.	$a \rightarrow b \rightarrow c \rightarrow d \rightarrow x$	184 - 185	3.	$k \stackrel{n}{\checkmark} l$	78		
8.	$e {\rightarrow} f$	128	4.–	$a \rightarrow b \rightarrow c \rightarrow d \rightarrow x$	0–14		
9.–	$x \rightarrow y$ or x, y	2-14		or $x \rightarrow y$ or x, y			

SUPEREPISODES

Similar to partition model except now test for superepisodes.

Example: F is a superepisode of G



Model how likely *F* is discovered once we see at least one event in F.

In practice, boost the probabilities p(e) in



Episodes discovered from *JMLR* abstracts:

ranked by $r_{ind}(G)$		r_{ind}	r_{prt}	ranked by $r_{prt}(G)$	r_{ind}	r_{prt}	Episodes with high r_{ind} and low r_{prt}	
1.	support→vector→machin	∞	357	support→vector	440	440	G_1 : support \rightarrow vector \rightarrow machin	regress
2.	support→vector	440	440	support→vector→machin	∞	357	95.1 regress	
3.	support→vector→machin→svm	404	90	support→machin	324	324	G_2 :support-vector-machin	
4.	support→vector→machin svm	356	10^{-3}	vectormachin	306	306	90.4	
5.	reproduc-kernel-hilbert-space	341	73	data→set	284	284	G_3 : support \rightarrow vector \rightarrow machin	number
6.	support→machin	325	325	real→world	260	260	52.0 regress	
7.	vector→machin	306	306	real→data	213	213	G_4 :support-vector-machin	
8.	data→set	284	284	state→art	191	191	86.4	
9.	real→world	260	260	machin→learn	190	190	G_5 : support \rightarrow vector \rightarrow machin	space
10.	support→vector→svm	250	85	bayesian→network	166	166	51.6	_