Disease Track Record Analysis at Dr. Soetomo General Hospital Using PrefixSpan Algorithm

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Abstract— Dr. Soetomo's General Hospital has used computerized system to record their patient s diseases. With the large amount of data to be analyzed, Dr. Soetomo General Hospital needs to know the disease pattern to prevent and cure the disease. Based on the problem, the hospital needs to develop an application that generates sequences pattern of diseases so it could be used to predict sequence pattern of disease in later day.

This application is built with PrefixSpan algorithm to generate disease pattern in a particular region on particular time according Dr. Soetomo's General Hospital historical data. Output of this application is generates rules.

Keywords: data mining, frequent pattern-projected sequential pattern mining

I. INTRODUCTION

Dr. Soetomo's General Hospital in Surabaya is a national hospital which acts like reference from other hospital. Information system of the patients is stored using *Oracle Data and Application* [1].

The increasing of civilization in East Java province is increased the patients with various type of disease. The hospital needs tool to monitor this occurrence in order to anticipate the spread of the disease.

This research is offering PrefixSpan sequential pattern mining to discover disease pattern from inpatient history which are the disease's name and the occurrence region.

II. PREFIXSPAN ALGORITHM

Sequential pattern mining is method to discover the relation between items in a dataset [2].

Prefix-projected sequential pattern mining called PrefixSpan is a method to project sequence databases based on frequent prefixes because each subsequence frequent can be discover by growth frequent prefix PrefikSpan [3].

This research is using bilevel projection calculation, which is:

- Scan the sequence to get length-1 item.
- Create triangular matriks from length-1 item.

- For each length-2 sequential pattern, build aprojected database and count the occurrence item, then build s-matrix.
- Each item is put in the end of length-2 sequential pattern [4].

III. DATA PREPARATION AND PROCESS

This research is using patient table (regency, sex, date of birth, and other information), disease table (list of disease), diagnose table (time, doctor, patient, type of diagnose), province and regency table [5]. The algorithm to generate rules (Fig. 1) consists of:

- Sequence numbering which process based on id patient, time for in and out patient. Patient with the same id can have different sequence number because sequence number is based on period time of the last time patient out with the next time patient in.
- Sequence Pattern is created from item in sequence. If an item is occurrence in recurrent, it only write once. After the sequence pattern is created, the frequent of item (become frequent item list length-1) which fulfill the minimum support is counted. The result if descending from the frequent item.
- Triangular Matrix can be created by built matrix with item length-1 number x item length-1 number size, which each cell is contain of three data which is present length-2 sequence pattern. Each data from cell which fulfills minimum support will be frequent item list length-2.
- Each length-2 sequential pattern which fulfills minimum support will create projected database, length-1 in projected database and S-Matrix. If length-1 in projected database fulfills minimum support then S-Matrix for length-2 will be build. If S-Matrix fulfills minimum support, S-matrix will be stored and re-process. The process will be stop if the number of projected database less then minimum support. S-Matrix which fulfills minimum support will be place in the end of length-x and the result of length-1 from projected database will be place in the end of length-x. If all the recurrent process is finished, length-x Sequential Pattern is created.



Figure 1. PrefixSpan flowchart

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IV. DISCOVERING SEQUENTIAL DISEASE PATTERNS TESTING

Testing is using data from January 1, 2003 until December 31, 2003, with *minimum support* = 2, time period = 6 days, with all province (Fig. 2).

Sequence and sequence pattern is built from data in Fig. 1 based on sequence number. Fig. 3 is shown frequent item list length-1 which fulfills minimum support and Fig. 4 is shown the triangular matrix. Triangular matrix which fulfills minimum support becomes length-2 sequential pattern.

Keterangan	WKTMSK	WKTKLR	NO_SEQ
N81.3	2003-01-05	2003-01-14	1
E14.9	2003-01-07	2003-01-15	2
K74.6	2003-01-07	2003-01-15	2
R18.X	2003-01-07	2003-01-15	2
N18.9	2003-05-19	2003-05-22	3
Z51.9	2003-07-02	2003-07-11	4
J38.3	2003-05-04	2003-05-06	5
J38.3	2003-11-02	2003-11-04	6
D64.9	2003-06-02	2003-06-06	7
Z51.3	2003-06-02	2003-06-06	7
Z51.3	2003-09-19	2003-09-23	8
Z51.3	2003-11-10	2003-11-17	9
A30.4	2003-01-28	2003-01-31	10
A30.4	2003-06-09	2003-06-19	11
B35.4	2003-06-09	2003-06-19	11
110.X	2003-06-09	2003-06-19	11
A30.5	2003-09-11	2003-09-17	12
L52.X	2003-09-11	2003-09-17	12

Figure 2. Data January 1, 2003 until December 31, 2003

sequent	e dan Sequence Pattern		
NO SEQ	SEQUENCE	SEQUENTIAL PATTERN	
1	<n80.4></n80.4>	[7841]	▲
2	<(E14.1, K73.8,R11.X)>	[4326, 6597, 9380]	
3	<n16.8></n16.8>	[7663]	
4	<z51.1></z51.1>	[11696]	
5	<j35.8></j35.8>	[6174]	
6	<j35.8></j35.8>	[6174]	
7	<(D63.0, Z50.6)>	[4142, 11691]	
8	<z50.6></z50.6>	[11691]	
9	<z50.6></z50.6>	[11691]	
10	<a28.1></a28.1>	[2705]	
11	<(A28.1, B34.3,I08.3)>	[2705, 3071, 5717]	
12	<(A28.2, L50.6)>	[2706, 6825]	
13	<(A28.1, H65.3,L50.6)>	[2705, 5603, 6825]	
14	<d13.2></d13.2>	[3874]	
15	<d13.2></d13.2>	[3874]	

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Item Length-1	Jumlah Frequent
11689	694
11690	180
11696	171
11691	170
4142	169
3745	156
5717	127
4326	118
11688	110
4072	94
3576	88
3735	88
11692	82
2753	71
2617	69

Figure 3. Sequence and sequence pattern (left), frequent item list length-
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Triangula	Friangular Matriks Length-2 Sequential Patter							iential Pattern				
A	Z50.4	Z50.5	Z51.1	Z50.6	D63.0	C88.1	108.3	E14.1	Z5		Item Length-2	Jumlah Fregu
Z50.4	5										<11689, 1168	5
Z50.5	1,2,0	6									<11690, 1168	2
Z51.1	0,0,0	0,0,0	6							=	<11690, 1169	6
Z50.6	2,0,0	1,0,0	0,0,0	1						H	<11696, 1169	6
D63.0	0,0,11	0,0,1	0,2,27	0,1,49	0					1	<11689, 1169	2
C88.1	6,8,0	2,3,0	1,1,0	0,1,0	0,0,2	45				1	<(4142, 11689	11
108.3	1,0,2	0,0,0	1,0,13	0,0,1	1,0,8	2,0,2	3				<4142, 11696>	2
E14.1	0,1,4	0,1,1	2,1,11	0,1,1	1,1,6	0,0,0	1,0,22	5		1	<(4142, 11696	27
Z50.3	6,0,0	0,0,0	0,0,0	1,1,0	1,0,4	0,0,0	0,0,0	0,0,0	1	1	<(4142, 11691	49
D48.1	2,0,17	0,0,1	0,1,2	0,1,30	2,1,0	0,0,0	0,0,1	0,0,0	0,0,4		<3745, 11689>	8
C50.9	0,0,0	0,0,1	0,0,0	0,2,0	0,1,5	0,0,0	0,0,0	0,0,3	0,1,0	1	<11689, 3745>	6
C84.1	0,0,0	0,0,0	0,0,0	0,0,0	0,0,0	2,1,0	0,0,0	0,0,3	0,0,0	1	<3745, 11690>	3
Z50.7	0,0,0	0,0,0	0,0,0	0,0,0	1,0,4	0,1,0	1,0,3	1,0,2	0,0,0	1	<11690, 3745>	2
A40.9	0,1,3	2,1,2	1,1,7	0,0,1	0,0,3	1,0,9	0,0,6	0,0,9	0,0,0	1	<(3745, 4142)>	2
A07.9	0,0,1	0,0,0	0,0,1	0,0,0	0,0,2	0,0,1	0,1,5	0,0,6	0,0,0		<3745, 3745>	45

Figure 4. Triangular matrix (left), length-2 sequential pattern (right)

Hasil Sequential	Jumlah Sequence : 126			
No	Prefiks	Hasil dalam Kode	Hasil dalam ICDX	Jumlah
17	C88.1	<3745, 11689>	<c88.1, z50.4=""></c88.1,>	8
18		<3745, 11689, 3745>	<c88.1, c88.1="" z50.4,=""></c88.1,>	2
19		<3745, 11689, 11689>	<c88.1, z50.4="" z50.4,=""></c88.1,>	2
20		<3745, 11690>	<c88.1, z50.5=""></c88.1,>	3
21		<3745, 3745>	<c88.1, c88.1=""></c88.1,>	45
22		<3745, 3745, 3745>	<c88.1, c88.1="" c88.1,=""></c88.1,>	24
23		<3745, 3745, 11689>	<c88.1, c88.1,="" z50.4=""></c88.1,>	5
24		<3745, 3745, 5717>	<c88.1, c88.1,="" i08.3=""></c88.1,>	2
25		<3745, 3745, 3745, 3745>	<c88.1, c88.1="" c88.1,=""></c88.1,>	8
26		<3745, 3745, 3745, 3745, 3745, 3745>	<c88.1, c88.1="" c88.1,=""></c88.1,>	3
27		<3745, 3745, 11689, 3745>	<c88.1, c88.1="" c88.1,="" z50.4,=""></c88.1,>	2
28		<3745, 3745, 11689, 11689>	<c88.1, c88.1,="" z50.4="" z50.4,=""></c88.1,>	2
29		<3745, 5717>	<c88.1, i08.3=""></c88.1,>	2
30		<3745, 3735>	<c88.1, c84.1=""></c88.1,>	2
31	108.3	<5717, 5717>	<108.3, 108.3>	3
32	E14.1	<(4326, 4142) 11696>	<(E14.1, D63.0) Z51.1>	2
33		<4326, 4326>	<e14.1, e14.1=""></e14.1,>	5

Figure 5. Sequential

Hasil Ru	le :	Jumlah Rule :	327	
Rule ke -	Rule (dalam ICD_X)	Rule (dalam KAT3)	Jumlah Kemunculan	
28	<c88.1=>C88.1=>Z50.4=>C88.1></c88.1=>	<alpha chain="" disease="" heavy="">Alpha heavy chain disease=>Psychotherapy, nec=>Alpha he</alpha>	2	
29	<c88.1=>C88.1=>Z50.4=>Z50.4></c88.1=>	<alpha chain="" disease="" heavy="">Alpha heavy chain disease=>Psychotherapy, nec=>Psychoth</alpha>	2	
30	<c88.1=>I08.3></c88.1=>	<alpha chain="" disease="" heavy="">Combined disorders of mitral, aortic and tricuspid valves></alpha>	2	
31	<c88.1=>C84.1></c88.1=>	<alpha chain="" disease="" heavy="">Sezary's disease></alpha>	2	
32	<108.3=>108.3>	<combined and="" aortic="" disorders="" mitral,="" of="" tricuspid="" valves="">Combined disorders of mitral,</combined>	3	
33	<(E14.1->D63.0)=>Z51.1>	<(With Ketoacidosis->Anaemia in neoplastic disease (C00-D48+))=>Chemotherapy sessio	2	
34	<(D63.0->E14.1)=>Z51.1>	<(Anaemia in neoplastic disease (C00-D48+)->With Ketoacidosis)=>Chemotherapy sessio	2	
35	<e14.1=>E14.1></e14.1=>	<with ketoacidosis="">With Ketoacidosis></with>	5	
36	<c50.9=>Z50.6></c50.9=>	<breast, unspecified="">Orthoptic training></breast,>	2	
37	<c84.1=>C84.1></c84.1=>	<sezary's disease="">Sezary's disease></sezary's>	5	
38	<c84.1=>C79.7></c84.1=>	<sezary's disease="">Secondary malignant neoplasm of adrenal gland></sezary's>	2	
39	<(A40.9->108.3)=>N36.0>	<(Streptococcal Septicaemia, Unspecified->Combined disorders of mitral, aortic and tricus	3	
40	<(I08.3->A40.9)=>N36.0>	<(Combined disorders of mitral, aortic and tricuspid valves->Streptococcal Septicaemia, Un	3	
41	<(A40.9->E14.1)=>N36.0>	<(Streptococcal Septicaemia, Unspecified->With Ketoacidosis)=>Urethral fistula>	2	
42	<(E14.1->A40.9)=>N36.0>	<(With Ketoacidosis->Streptococcal Septicaemia, Unspecified)=>Urethral fistula>	2	

Figure 6. Rules

Fig. 5 is the result of *length-x* process in code form and Fig. 6 is the result with the diagnosis's name. For example, alpha heavy chain disease is followed by psychotherapy nec.

V. CONCLUSION

PrefixSpan method can be used for mining sequential diseases pattern from database sequential. The generated patterns can be used a knowledge to predict sequential diseases. As a result the medical representative can take preventive and curative action more precisely. The rule will increase as the smaller *minimum support*, but it will cost time processing.

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REFERENCES

- [1] Indonesia, Departemen Kesehatan, "Sistem informasi rumah sakit di Indonesia, Jakarta, 2000.
- [2] Han, Jiawei, Kamber, Micheline, "Data mining : Concept and techniques", Canada: Simon Fraser University, 2000.
- [3] Han, Jiawei, et al, "Mining sequential patterns by pattern-growth : The PrefixSpan Approach", IEEE Transactions on Knowledge and Data Engineering, ol. 16, November 2004, pp 1424-1140.
- [4] Han, Jiawei, et al, "PefixSpan : Mining sequential patterns efficiently by prefix-projected pattern growth", Proceedings of the 17th International Conference on Data Engineering, 2001, pp 215-224.
- [5] Kusuma, Yinsi R., Renov Yapola, "Implementasi star schema dan data warehouse pada RSU Dr. Soetomo, unpublised literature, Petra Christian University, Surabaya, 2009.