



Effective Retrieval and Analysis of Uropathogens through NoSQL Database

Dr. V.T. Meenatchi, Dr. M. Thangaraj

Department of CA & IT, Thiagarajar, College, Department of Computer Science, M.K University, Madurai, India

S. Padmavathy, N. K. AshaDevi, K. Vignesh

Department of Zoology & Microbiology, Thiagarajar College, Madurai, India

ABSTRACT

In the today's web era, big data is emerging. The storage and retrieval of big data is becoming an issue. The database administrators are moving into new storage technology, the NoSQL database. This paper analyzes the predominant organisms causing the Urinary Tract Infection (UTI) based on gender wise and age-wise and also the antibiogram pattern of 3G and 4G antibiotics were analysed. The work is clinically proven using new methodologies and the data is then mapped into MongoDB, a NoSQL database. Through this type of mapping and analysis, the data retrieval becomes ease and simpler to manage huge data. The generated analytical report aids the medical practitioners to provide the needful therapy for UTI affected patients.

Keywords : NoSQL; MongoDB; Urinary Tract Infection; Antibiotics; Therapy

I. INTRODUCTION

One of the most prevalent problems faced by healthcare services is the increasing prevalence of antimicrobial resistance. Urinary tract infections (UTI) are the most common bacterial infections affecting humans throughout their lifetime. UTI is a serious ailment in human due to increasing frequency, recurrence and difficulty in eradication; it poses stiff challenge to the medical professionals. It is much more common in women than in men, due to anatomical and physiological reasons by virtue of its position urinogenital tract is more vulnerable to bacterial infections caused by both internal and external flora [3].

Every year, the rate of people affected by urinary tract infections is increasing. when the infections are not treated, it leads to serious health problems. This work aims to identify the uropathogens causing the infection. From the urine specimens, it was noted that the following uropathogens like Escherichia coli, Klebsiella pneumoniae, Citrobacter sp., Enterobacter sp., and Staphylococcus aureus are responsible main cause for infection. The NOSQL databases were created as a mean to offer high performance and high availability. NOSQL

is an unstructured Query Language [11] and data stores are widely used to store and retrieve large amounts of data.

Our paper is organized as follows, Section II discusses the related literatures, Section III deals about the proposed system and Section IV presents the Results and Discussion and Section V ends with Conclusion.

II. LITERATURE SURVEY

Urinary tract infection is one of a serious health problem affecting millions of people yearly. Infections of urinary tract are in the second most common type of infection in the body. Urinary tract infection has an important association in human female, the highest incidence of urinary tract infection occur in child bearing age and this has been linked to sexual activity and aging. Asymptomatic urinary tract infection occurs in 2 to 10% in women during pregnancy.

Urinary tract infection can be seen in three different forms in infected pregnant women, asymptomatic bacteriuria, acute cystitis or acute pyelonephritis. The incidence of asymptomatic urinary tract infection has

been reported between 2-13% however physiological changes in pregnancy lead to severe course of problem and also it has been reported that if untreated asymptomatic bacteriuria increase the frequency of premature delivery and neonates with low birth weights[7]. Urinary tract infection (UTI) is one of the most common and life threatening infection present in community practice.

Manifestations can vary from asymptomatic bacteriuria to symptomatic cystitis, pyelonephritis and blood stream infection[8]. The most common symptoms are pain, fever, or nausea and vomiting in addition to the classic symptoms of a lower urinary tract infection. Rarely the urine may appear bloody[9]. The general symptoms include a frequent urge to urinate a painful burning feeling during urination. These urinary infection are known to cause to a lot of problems in the physiological action of the urinary system. There are a number of reasons why bacterial resistance should be a concern for physicians. First resistant bacteria particularly Staphylococci, Enterococci, Klebsiella pneumoniae and Pseudomonas sp. are becoming common place in healthcare institutions. Bacterial resistance often results in treatment failure, which antibacterial therapy, defined as the initial use of an antibacterial agent to which the causative pathogen was not susceptible, has been associated with increased mortality rates in patients with bloodstream infections due to resistant Pseudomonas aeruginosa, Staphylococcus aureus, Klebsiella pneumoniae, Escherichia coli, Enterobacter sp., coagulase negative Staphylococci and Enterococci. Prolonged therapy with antimicrobial agents such as vancomycin or linezolid, may also lead to the development of low – level resistance that compromises therapy, but that may not be detected by routine susceptibility testing methods used in hospital laboratories. Many uropathogen are producing the Beta lactamase enzymes, so drugs resistant to particular pathogen for example Hydrolysis of beta lactam ring by beta lactamase enzymes produce by uropathogen.

During cystitis, Uropathogenic Escherichia coli subvert innate defenses by invading superficial umbrella cells and rapidly increasing in numbers to form intracellular bacterial communities [2]. By working together, bacteria in biofilms build themselves into structures that are more firmly anchored in infected cells and are more resistant to immune-system assaults and antibiotic treatments.

This is often the cause of chronic urinary tract infections. Antibiotics are the main treatment for all UTIs. A variety of antibiotics are available, and choices depend on many factors, including whether the infection is complicated or uncomplicated or primary or recurrent.

Worldwide reports of antibiotic resistant E. coli isolates indicate the unwise and excessive consumption of antimicrobial drugs which in turn has brought about failure in treatment, and consequently concerns about the related issues in all nations including the developed and developing ones. As an example, TMP-SXT was conventionally used for uncomplicated cystitis in most regions, however, due to the resistance to it, fluoroquinolon and cephalosporin took its place and unfortunately after sometime resistance to these two drugs was also recognized and reported [4][5].

Reports of uropathogens resistant to previously effective antibiotics have emerged globally in recent years. The situation is especially dire in Africa where irrational antibiotic practices are common. Variations in antibiotic resistance patterns are known to occur across different geographical regions, even within the same country. Such variations must be well documented so as to inform local empirical treatment as well as foster rational antibiotic use. The treatment of bacterial infections is increasingly complicated by the ability of bacteria to develop resistance to antimicrobial agents [6].

The development of resistance to all available antibiotics in some organisms then precludes the effectiveness of any antibiotic region. Organisms that are resistant to all known effective antimicrobials pose a serious threat to hospitalized patients. Indiscriminate use of antibiotics leads to the development of resistance of initially sensitive strains of organisms and possibly destruction of the normal flora [1].

The traditional database system maintained are the relational database systems, in which all the database retrieval and transactions are done in (SQL). Since the database becomes huge in size, we have to retrieve databases through MongoDB [10]. The necessity of using MongoDB is to have document oriented storage. MongoDB is applied in Big Data, Content Management a Delivery, Mobile and Social Infrastructure, User Data Management, Data Hub. Database in MongoDB[12], is a physical container for collections. Each database gets its

own set of files on the file system. A single MongoDB server[13] typically has multiple databases.

III. PROPOSED SYSTEM

An architecture named as UTI_framework in Fig. 1 has been constructed with 6 main components namely,

- Sample collection
- clinical analysis with three subcomponents:
 - Morphological characterization
 - Culture characterization
 - Biochemical characterization
- Uropathogens Identification
- Mining through NoSQL
- Antibigram Pattern
- Analytical Report

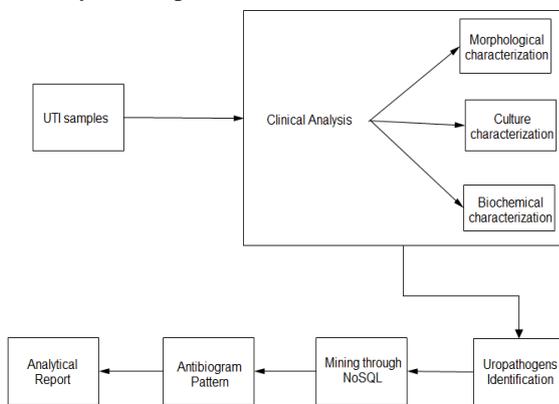


Figure 1. UTI Framework

All the components in the framework are interdependent. The collected urine samples were given for clinical analysis for uropathogen identification. The results were then mapped into MongoDB 3.2 document store environment for mining [14]. The Antibigram patterns were assessed for specific organism. Finally the reports were generated to depict organism predominance, proper antibiotic for treatment, its sensitive level to organism, gender and age wise analysis.

IV. RESULTS AND DISCUSSION

Following are the results derived through clinical study. For analysis, 500 samples were collected and major affected were female than male as shown in Table 1.

TABLE I. Sex distribution of UTI Patients

Sex	Total number
Male	150
Female	350
Total	500

Various organisms were identified and number of uropathogens with their corresponding percentage is shown in Table 2.

TABLE II. Percentage of Isolated Uropathogens using Selective and Differential Medium

Name of the organisms	Number of Uropathogens	Percentage of Uropathogens
<i>E.coli</i>	88	31.10
<i>K.pneumoniae</i>	55	19.43
<i>Staphylococcus aureus</i>	42	14.84
<i>Enterobacter sp.,</i>	32	11.31
<i>Pseudomonas sp.,</i>	27	9.54
<i>Streptococcus sp.,</i>	20	7.07
<i>Citrobacter sp.,</i>	13	4.59
<i>Proteus sp.,</i>	6	2.12
Infected	283	-
Non infected	217	-
Total	500	100

The age wise and gender wise analysis were made and shown in Table 3.

From the table, one could identify the maximum percentage affected in male falls in 50 years and above, whereas in case of female, the maximum percentage falls under 30-39 years age group.

TABLE III. Distribution of the UTI Cases due to *E.coli* with reference to Age and Gender Status

Age groups (in Years)	Male		Female	
	Total	Percentage	Total	Percentage
Less than 1	2	4.34	7	2.37
1-19	8	17.39	29	10.14
20-29	3	6.53	58	20.39
30-39	5	10.87	82	28.94
40-49	10	21.74	57	20.14
50 and above	18	39.13	51	18.02

The antibiogram pattern for the infected organisms with their sensitivity levels are depicted in Table 4.

TABLE IV. Antibiogram Pattern of Uropathogenic *E.coli* before Plasmid Curing

Antimicrobial Agents	Sensitivity Isolates	Intermediate Isolates	Resistant Isolates
Amikacin (30 µgms)	222	51	10
Nitrofurantoin (300 µgms)	185	55	43
Ampicillin/Sulbactam (10/10 µgms)	86	145	52
Kanamycin (30µgms)	40	165	78
Norfloxacin(10 µgms)	92	80	113
Meropenem (10 µgms)	260	15	8
Cefixime (5µgms)	37	79	167
Ciprofloxacin (5 µgms)	85	96	102
Ofloxacin (5µgms)	96	99	88
Co-Trimoxazole (23.75/1.25 µgms)	77	40	166
Pristinamycin (15µgms)	46	104	133
Vancomycin (30 µgms)	34	32	217
Gentamicin (10µgms)	53	195	35
Cefoperazone (75 µgms)	36	115	132
Chloramphenicol (30µgms)	177	51	55
Linezolid (30 µgms)	42	34	207
Azithromycin (15 µgms)	108	114	61
Erythromycin (15 µgms)	39	84	160
Piperacillin/Tazobactam(100/10µgms)	184	63	36
Sulphafurazole (300µgms)	65	57	161
Tobramycin (10 µgms)	40	215	28
Oxytetracycline (30µgms)	66	41	176
Clarithromycin (15 µgms)	24	86	173
Doxycycline Hydrochloride (30 µgms)	65	141	77
Moxifloxacin (5 µgms)	103	131	49

After clinical analysis, the results were mapped into MongoDB as shown in Fig. 2 , for effective retrieval.

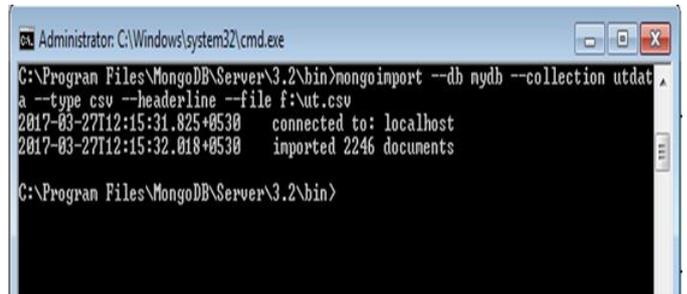


Figure 2. Mapping of UTI data into NoSQL

After mapping the data is depicted as shown in Fig. 3.From the UTI dataset, out of 200, 138 number of female were affected from Escherichia Coli as shown in Fig. 4.

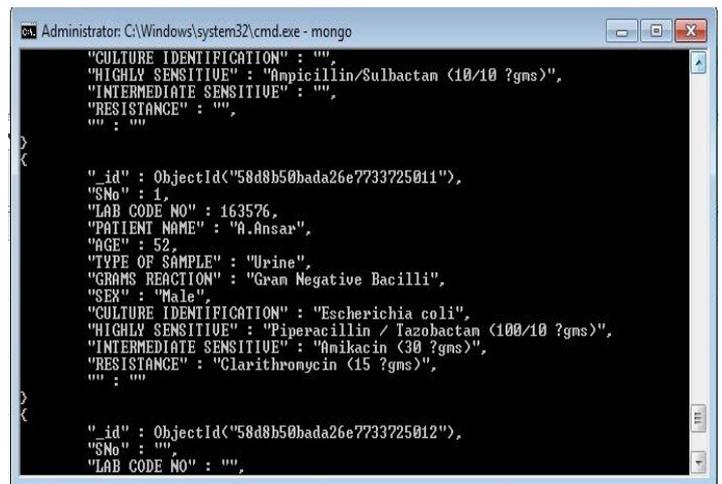


Figure 3. Display of documents

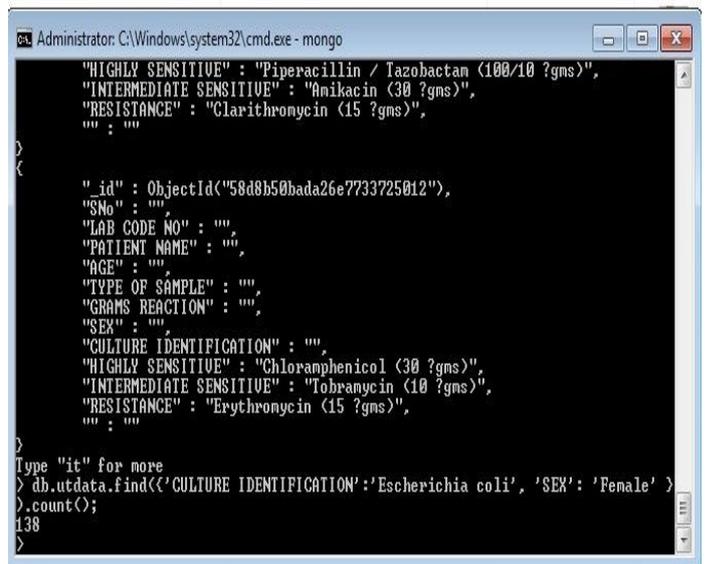


Figure 4. Culture Identification

Fig.5 shows the number of isolates affected through E. Coli from 100 samples.

```
Administrator: C:\Windows\system32\cmd.exe - mongo
{"CULTURE IDENTIFICATION": "Escherichia coli",
"HIGHLY SENSITIVE": "Piperacillin / Tazobactam (100/10 ?gms)",
"INTERMEDIATE SENSITIVE": "Amikacin (30 ?gms)",
"RESISTANCE": "Clarithromycin (15 ?gms)",
"" : ""
}
{
  "_id": ObjectId("58d8b50bada26e7733725012"),
  "SN": "",
  "LAB CODE NO": "",
  "PATIENT NAME": "",
  "AGE": "",
  "TYPE OF SAMPLE": "",
  "GRAMS REACTION": "",
  "SEX": "",
  "CULTURE IDENTIFICATION": "",
  "HIGHLY SENSITIVE": "Chloramphenicol (30 ?gms)",
  "INTERMEDIATE SENSITIVE": "Tobramycin (10 ?gms)",
  "RESISTANCE": "Erythromycin (15 ?gms)",
  "" : ""
}
Type "it" for more
> db.utdata.find({'CULTURE IDENTIFICATION':'Escherichia coli'}).count()
89
>
```

Figure 5. E. Coli affected samples

Thus the desired reports can be generated through NoSQL database and any useful information can be tracked anytime when needed.

V. Conclusion

It is concluded from the present study that routine microbiological analysis for antibiotics sensitivity tests for urinary tract infected patients and other patients to be carried out before administration of drugs for treatment and management of urinary tract infections, since resistance to these drugs are developing in the community. This study appears to suggest a need for a continuous monitoring of bacterial antibiotics susceptibility before antibiotics prescription in order to ensure adequate treatment for urinary tract infection and reduction in the spread of bacteria resistant strain. Identification and proper treatment of UTI infections will lead to 10 fold decrease in the occurrence of acute pyelonephritis. After clinical study, data is computerized through NoSQL database. This makes the overall retrieval and mining easier. Moreover, self-medication should be avoided in order to prevent spread of drug resistant strains of bacteria.

VI. References

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