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Original Article

Application of Advanced Whole Genome Sequencing Technology for Tuberculosis Outbreak Investigation

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Abstract

Tuberculosis (TB) is an *aerosol-transmissible* disease caused by the *Mycobacterium tuberculosis* complex. Blocking the transmission chain is one of the key strategies to stop TB. The global tuberculosis control has entered the genomic era. For facilitating TB control, we used whole genome sequencing (WGS) to obtain single nucleotide polymorphism (SNP) information of *M. tuberculosis* to improve discriminative limitations of conventional genotyping methods and to strengthen the delineation of transmission networks. Based on a confirmed multidrug-resistant TB (MDR-TB) outbreak, we define SNPs differences for TB outbreak investigations. To confirm an outbreak, the transmission events between cases within clusters were calculated using thresholds of ≤ 5 (definite) and ≤ 15 (probable) SNPs difference between isolates. Depending on cost-effectiveness, an algorithm was established using mycobacterial interspersed repetitive units-variable number tandem repeat (MIRU-VNTR) and WGS as the primary and secondary genotyping, respectively. However, WGS still needs to be combined with classical epidemiological methods to improve outbreak investigations.

Keywords: Tuberculosis, *Mycobacterium tuberculosis* complex, whole genome sequencing, outbreak investigation

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week 42-week 43, 2023 (Oct. 15, 2023-Oct. 28, 2023)

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Weekly Data of Notifiable Inases (by week of diagnosis)

Case diagnosis year		Wee	ek 42★	Week 1-42				
Classification	Disease Diagnosed	2022	2022	2023 Total Imported Total			2022 Imported	
		2023	2022	cases★	cases	cases★	cases	
Category I	Plague	0	0	0	0	0	0	
	Rabies SARS	0	0 0	0	0	0	0	
	Smallpox	0	0	0	0	0	0	
	Cholera	0	0	1	0	0	0	
	Typhoid fever	0	0	8	6	3	1	
	Paratyphoid Fever	0	0	13	1	0	0	
	Epidemic Typhus Fever Shigellosis	0	0 0	0 55	0 12	0 59	0 5	
	Amoebiasis	3	4	224	89	168	58	
	Enterohemorrhagic E.coli Infection	0	0	0	0	2	0	
	Anthrax	0	0	0	0	0	0	
	Diphtheria Meningococcal Meningitis	0	0 0	0 3	0	0 1	0	
	Poliomyelitis	0	0	0	0	0	0	
	Acute Flaccid Paralysis	2	1	48	Ö	23	Ö	
	Measles	0	0	2	2	1	0	
	Rubella	0	0	0	0	0	0	
	Dengue Fever West Nile Fever	1,776 0	3 0	18,706 0	201 0	66 0	46 0	
	Acute Viral Hepatitis type A	3	0	73	4	114	1	
	Malaria	0	0	1	1	2	2	
	Chikungunya Fever	0	0	8	8	1	1	
	Hantavirus syndrome	0	0	6	0	3	0	
	Zika virus infection Mpox	0	0 0	3 345	3 13	0 4	0 4	
Category III	Acute Viral Hepatitis type B	4	2	110	6	81	0	
	Acute Viral Hepatitis type C	8	13	423	1	385	2	
	Acute Viral Hepatitis type D	0	0	0 12	0 4	0 10	0	
	Acute Viral Hepatitis type E Acute Viral Hepatitis, untyped	0	0 0	9	1	0	0	
	Congenital Syphilis	Ö	Ö	Ö	Ō	Ö	Ö	
	Congenital Rubella Syndrome	0	0	0	0	0	0	
	Enteroviruses Infection with Severe Complications Haemophilus Influenza type b Infection	0	0 0	13 1	0	0 2	0	
	Japanese Encephalitis	ő	Ö	23	0	19	Ö	
	Legionnaires' Disease	6	3	318	8	263	1	
	Mumps Neonatal Tetanus	6 0	7 0	230 0	7 0	195 0	0	
	Pertussis	0	1	0	0	1	0	
	Tetanus	0	0	5	0	7	0	
	Botulism	0	0	0	0	0	0	
	Brucellosis Complicated Varicella	0	0 0	0 36	0	0 26	0	
	Endemic Typhus Fever	1	Ö	22	ő	11	Ö	
	Herpesvirus B Infection	0	0	0	0	0	0	
	Influenza Case with Severe Complications Invasive Pneumococcal Disease	25 3	0 6	857 225	11 1	1 146	0	
Category IV	Leptospirosis	3	8	62	0	54	0	
	Listeriosis	0	5	157	2	119	0	
	Lyme Disease	0	0	0	0	1	1	
	Melioidosis Q Fever	0	1 0	19 3	0	19 3	0	
	Scrub Typhus	13	7	162	ő	233	Ö	
	Toxoplasmosis	0	0	23	2	20	0	
	Tularemia Severe Fever with Thrombocytopenia Syndrome	0	0 0	0	0	0 1	0	
	Severe Pneumonia with Novel Pathogens	288		1,393,788		7,433,034	-	
Category V	Ebola Virus Disease	0	Ó	0	Ő	0	Ô	
	Lassa Fever	0	0	0	0	0	0	
	Marburg Hemorrhagic Fever Middle East Respiratory Syndrome Coronavirus	0	0	0	0	0	0	
	Infections	0	0	0	0	0	0	
	Novel Influenza A Virus Infections	0	0	1	0	0	0	
	Rift Valley Fever	0	0	0	0	0	0	
	Yellow Fever	0	0	0	0	0	0	

 ^{1. ★}The weekly and cumulative total numbers include indigenous and imported cases of notifiable infectious diseases.
 2. MDR-TB, Tuberculosis, Syphilis, Gonorrhea, HIV Infection, AIDS, Hansen's Disease and Creutzfeldt-Jakob Disease are excluded from the table.
 3. Numbers of Mumps and Tetanus are based on reported cases and summed up by week of report.
 4. "Mpox" has been listed as a Notifiable Infectious Disease since June 23, 2022.
 5. "Severe Pneumonia with Novel Pathogens": The case definition has been revised to include patients who have both a positive test for SARS-CoV-2 and associated complications since March 20, 2023. Additionally, it has been modified from Category IV since May 1, 2023.

Suspected Clusters

● Fifty-one clusters related to Upper respiratory tract infection (32), Diarrhea (12), Enterovirus (3), TB (2) and Varicella (2) were reported during week 42.

Imported Infectious Diseases

● There were 15 imported cases from at least 6 countries / areas during week 42.

Dengue Fever: 12 cases from Vietnam (8), India (2), Indonesia (1), and Cambodia (1).

Severe Pneumonia with Novel Pathogens: 2 cases from China (1), and Japan (1). **Amoebiasis**: 1 case from Vietnam.

- ●During week 1-42, there were 18,522 imported cases of notifiable diseases. The top three were Severe Pneumonia with Novel Pathogens (18,137), Dengue Fever (201) and Amoebiasis (89).
- During week 1-42, imported cases of notifiable diseases were from at least 48 countries/areas. The top three were China (3,179), Japan (728) and Thailand (206).

Summary of Epidemic

- ●Influenza: In the midst of the epidemic period, the epidemic has decreased recently.
- **Enterovirus**: The epidemic has decreased.
- **Dengue Fever:** The epidemic is similar to the previous week, the risk of transmission remains high.

Weekly Data of Notifiable Inases (by week of diagnosis)

1	Case diagnosis year		eek 43	201	Week 1-43 2023 2022			
Classification	Disease Diagnosed	2023	2022	Total cases★	Imported cases	Total cases★	Importe d cases	
Category I	Plague	0	0	0	0	0	0	
	Rabies SARS	0	0 0	0	0	0	0	
	Smallpox	ő	0	Ö	Ö	0	Ö	
	Cholera	0	0	1	0	0	0	
	Typhoid fever	0	0	8	6	3	1	
	Paratyphoid Fever	0	3	13	1	3	0	
	Epidemic Typhus Fever Shigellosis	0	0 1	0 55	0 12	0 60	0 5	
	Amoebiasis	2	5	226	91	173	59	
	Enterohemorrhagic E.coli Infection	ō	Ő	0	0	2	0	
	Anthrax	0	0	0	0	0	0	
	Diphtheria	0	0	0	0	0	0	
	Meningococcal Meningitis	0	0	3	0	1	0	
Category II	Poliomyelitis Acute Flaccid Paralysis	0 2	0 0	0 50	0	0 23	0	
	Measles	0	0	2	2	1	ő	
	Rubella	Ō	Ö	0	0	0	Ō	
	Dengue Fever	1,525	4	20,230	210	70	50	
	West Nile Fever	0	0	0	0	0	0	
	Acute Viral Hepatitis type A Malaria	2	0 0	75 1	5 1	114 2	1 2	
	Chikungunya Fever	0	0	8	8	1	1	
	Hantavirus syndrome	ő	Ö	6	Ö	3	Ō	
	Zika virus infection	0	0	3	3	0	0	
	Мрох	5	0	350	13	4	4	
	Acute Viral Hepatitis type B	3	4	113	6	85	0	
Category III	Acute Viral Hepatitis type C Acute Viral Hepatitis type D	8	8 0	431 0	1 0	393 0	2 0	
	Acute Viral Hepatitis type E	ő	0	12	4	10	Ö	
	Acute Viral Hepatitis, untyped	0	Ö	9	1	0	Ö	
	Congenital Syphilis	0	0	0	0	0	0	
	Congenital Rubella Syndrome	0	0	0	0	0	0	
	Enteroviruses Infection with Severe Complications Haemophilus Influenza type b Infection	0	0 0	13 1	0	0 2	0	
	Japanese Encephalitis	1	0	24	0	19	Ö	
	Legionnaires' Disease	6	9	324	8	272	1	
	Mumps	9	17	239	7	212	0	
	Neonatal Tetanus	0	0	0	0	0	0	
	Pertussis Tetanus	0	0 0	0 5	0	1 7	0	
	Botulism	0	0	0	0	0	0	
	Brucellosis	0	0	Ö	0	0	Ō	
	Complicated Varicella	2	2	38	0	28	0	
	Endemic Typhus Fever	1	0	23	0	11	0	
Category IV	Herpesvirus B Infection Influenza Case with Severe Complications	0 18	0 0	0 875	0 11	0 1	0	
	Invasive Pneumococcal Disease	3	2	228	1	148	ő	
	Leptospirosis	1	0	63	0	54	0	
	Listeriosis	5	1	162	2	120	0	
	Lyme Disease	0	0	0	0	1	1	
	Melioidosis Ω Fever	1 0	1 0	20 3	2 0	20 3	2 0	
	Scrub Typhus	6	7	168	Ö	240	ő	
	Toxoplasmosis	0	1	23	2	21	0	
	Tularemia	0	0	0	0	0	0	
	Severe Fever with Thrombocytopenia Syndrome	0 225	0 239.986	1 204 012	10 120	7 672 020	0 33,935	
Category V	Severe Pneumonia with Novel Pathogens Ebola Virus Disease	0	239,986	1,394,013	18,138 0	7,673,020	33,935	
	Lassa Fever	0	0	0	0	0	0	
	Marburg Hemorrhagic Fever	0	Ö	ő	Ő	ő	Ö	
	Middle East Respiratory Syndrome Coronavirus	0	0	0	0	0	0	
	Infections	_		1	0	_	0	
	Novel Influenza A Virus Infections							
	Novel Influenza A Virus Infections Rift Valley Fever	0	0 0	0	0	0	0	

- 1. ★The weekly and cumulative total numbers include indigenous and imported cases of notifiable infectious diseases.
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Suspected Clusters

• Fifty-seven clusters related to Upper respiratory tract infection (36), Diarrhea (14), TB (6), and Enterovirus (1) were reported during week 43.

Imported Infectious Diseases

There were 15 imported cases from at least 5 countries / areas during week 43.

Dengue Fever: 12 cases from Vietnam (7), Thailand (4), and Indonesia (1).

Severe Pneumonia with Novel Pathogens: 1 case from Turkey.

Acute Viral Hepatitis type A : 1 case from USA.

Amoebiasis: 1 case from Indonesia.

- During week 1-43, there were 18,535 imported cases of notifiable diseases. The top three were Severe Pneumonia with Novel Pathogens (18,138), Dengue Fever (210), Amoebiasis (91).
- During week 1-43, imported cases of notifiable diseases were from at least 48 countries/areas. The top three China (3,178), Japan (728), Thailand (209).

Summary of Epidemic

- ●Influenza: The epidemic has decreased recently.
- **Dengue Fever:** The epidemic has decreased, but the risk of transmission remains high.

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