

Introduction

Mycobacterium tuberculosis (*M.tb.*) is an aerobic, gram positive, and acid-fast bacillus that leads to the complex disease manifestation of TB. Multidrug resistance (MDR) is a growing concern in endemic countries as a result of poor compliance, length of treatment, and adverse medication effects. Current diagnostic methods for detecting active *M.tb* infection are slow, unreliable and are not cost effective for high-risk populations. The Khan laboratory is currently developing novel strategies for studying blood based immune-biomarkers as well as intracellular signaling proteins and pathways in cell lysates using high-throughput multiplex microbead immunoassays. The goal of my summer project was aimed at optimizing the detection of active Mycobacterium tuberculosis infection and multidrug resistance by multiplex suspension arrays.



Materials

- Multiplex kits for measuring cytokines, chemokines and growth factors, for use on the Luminex platform (Luminex Corp, Austin, Tx), were obtained from BioRad, Hercules, CA. Assays were performed per manufacturer's instructions.
- Plasma samples from healthy individuals in the same TB endemic areas were used to compare confirmed TB patient samples.
- Primers targeting variable genomic targets observed in multi-drug resistant *M.tb.* strains were designed from genomic biomarkers for TB-MDR.













Perform fluorescent Analyze data

Detection of Active Mycobacterium Tuberculosis Infection and Multidrug Resistance by Multiplex Suspension Arrays

Emmanuel Mendoza, Peter Nham, Resmi Ravindran and Imran Khan University of California, Davis Department of Pathology and Laboratory Medicine *Correspondence to Imran Khan (ihkhan@ucdavis.edu)



Methods										
DNA extracted from sputum samples of TB patients was used to identify TB-MDR through multiplex PCR.										
Multiplex-PCR was used to amplify seven <i>M.tb</i> . genes in DNA isolated from purified TB patient sputum samples to identify multidrug resistance.										
Multiplex MDR TB Panel										
	Step DNA Extr (3 hr		Step Multiplex 7 genes for line dru (2hr)	PCR r 4 first- ugs		Step 3 Multiplex icrobead assay for detection (30 min)				
Results										
Sample	P38-PN F 100ug	P38-PN F 25ug	Rv0934-P38 4ug	P38-PN 25ug	P38-PN 12.5ug	Antihuman IgG 20ug	188 BSA 100ug pool			
HBP 1/12	55	17	21	23	14	6965	25			
	63 114	17 21	21 30	36	14 20	6692 6838	25 33			
HBP 3/12	104	22	34	33	18	6745	30			
TB-159/11	3305	97	2384	468	35	6006	18			
	3342	91	2333	413	36 24	6306 6778	19			
TB-138/11	313	33	609	95 102	24	6778	11			
HBP 1/12 HBP 3/12 TB-159/11 TB-138/11 • Figure 1 microbe	63 114 104 3305 3342 313 327 : Optimiza ad susper	17 21 22 97 91 34 33 33	21 30 34 2384 2333 619 609 9 blasma a s to study	24 36 33 468 413 95 102 htibody	14 20 18 35 36 24 28 multip	6692 6838 6745 6006 6306 6778 6778 6778	25 33 30 18 19 11 11 11			
associated with active TB. P38 and Rv0934 are <i>M.tb</i> specific antigens used in the serodection of TB in healthy blood plasma (HBP) against plasma from culture confirmed <i>M.tb</i> infected patients.										
		Antibiotic	Genes	Genes Multi Pi						
Rifampin		rpoB		٥ ٥						
		Pyrazinamie	katG, inh de none	۱A	9					
		Ethambutc	ambutol ombR		5					
		Streptomyc	rpsL, rr	S	0					

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tion of the plasma antibody multiplex method using sion arrays to study blood-based biomarkers tive TB. P38 and Rv0934 are <i>M.tb</i> specific antigens ection of TB in healthy blood plasma (HBP) against re confirmed <i>M.tb</i> infected patients.									
Antibiotic	Genes	Mult	iplex-PCF Probes	2					
Rifamnin	rpoB		18						
leoniazid		• ^	Q						
Pyrazinamio	de none	IA	0						
Ethambutc	embB		5						
Streptomyc	in rpsL, rr	S	8						

Figure 2: First line anti-*M.tb*. drugs with their corresponding genes identified in MDR-*M.tb*. strains. Multiplex-PCR probes corresponds to the available targets used in the identification of MDR-TB.

Sample	katG Ser WT	katG 315 Thr	rpoB 531 Ser WT	rpoB 531 Leu	rpsL 43 Lys WT	rpsL 43 Arg	EMB 306 Met WT	EMB 306 Val	IS6110
Blank	104	86	101	99	177	96	148	133	56
	96	71	54	96	205	147	156	91	73
H37RV	2911	104	959	205	2596	95	1495	664	2776
	3283	96	920	211	2826	140	1535	688	2806
MDR isolate	325	4002	190	2322	181	2883	225	900	2339
	349	4103	109	2362	156	2458	187	871	2190

Figure 3: Multiplex-PCR assay detection of genomic targets observed in multidrug resistant *M.tb*. in comparison to corresponding wild-type genomic targets. H37RV is a wild-type *M.tb*. strain used to differentiate between the culture confirmed, isolated MDR-*M.tb.*

- Continue to validate and identify more biomarkers for multidrug resistance *M.tb*. Verify results via a field validation study for sensitivy and specificy.
- Ongoing developments for commercialization of blood-based diagnostic assay.

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Results

Future Directions

Optimize TB-MDR assay for fresh patient sputum samples.

References

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