

Supporting Information

A bench-top automated sputum-to-genotype system using a Lab-on-a-Film assembly for detection of multidrug-resistant *Mycobacterium tuberculosis*

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S1. Hybridization ratios for all markers for DNA-to-Genotyping tests.

Supporting Information S1

Sample		H37Rv	H37Rv	TDR-0015	TDR-0015	TDR-0018	TDR-0018			
Resistance Profile		Wild Type	Wild Type	INH Resistant	INH Resistant	RIF Resistant	RIF Resistant			
Amount		1 pg	1 pg	1 pg	1 pg	1 pg	1 pg			
RESULTS										
Final Results		Test Validity	VALID	VALID	VALID	VALID	VALID			
M. tuberculosis Complex		PRESENT	PRESENT	PRESENT	PRESENT	PRESENT	PRESENT			
Rifampin Resistance		Not Detected	Not Detected	Not Detected	Not Detected	DETECTED	DETECTED			
Isoniazid Resistance		Not Detected	Not Detected	DETECTED	DETECTED	Not Detected	Not Detected			
Suspected Mutation		rpoB	N/A	N/A	N/A	N/A	S531W			
		katG	N/A	N/A	S315T (ACC)	S315T (ACC)	N/A			
		inhA	N/A	N/A	-15T	-15T	N/A			
M13 (90, 103, 106)		Absent	Absent	Absent	Absent	Absent	Absent			
IS6110 (71)		PRESENT	PRESENT	PRESENT	PRESENT	PRESENT	PRESENT			
rpoB UN (137, 176)		PRESENT	PRESENT	PRESENT	PRESENT	PRESENT	PRESENT			
katG UN (178)		PRESENT	PRESENT	PRESENT	PRESENT	PRESENT	PRESENT			
inhA UN (139)		PRESENT	PRESENT	PRESENT	PRESENT	PRESENT	PRESENT			
rpoB		Not Detected	Not Detected	Not Detected	Not Detected	DETECTED	DETECTED			
katG		Not Detected	Not Detected	DETECTED	DETECTED	Not Detected	Not Detected			
inhA		Not Detected	Not Detected	DETECTED	DETECTED	Not Detected	Not Detected			
Discrimination Ratios (WT - MU) / (WT + MU)										
Target	Probe Description		WT	MU						
rpoB	507-wt	507-DEL	1	2	0.94	0.88	0.97	0.90	0.89	0.83
rpoB	510-wt	Q510H	3	4	0.73	0.69	0.78	0.75	0.76	0.70
rpoB	511-wt	L511P	5	6	0.64	0.58	0.66	0.49	0.56	0.62
rpoB	511-wt	L511R	5	7	0.76	0.65	0.65	0.57	0.61	0.69
rpoB	512-wt	S512T	8	9	0.67	0.62	0.59	0.58	0.57	0.49
rpoB	512-wt	S512R	8	10	0.85	0.72	0.78	0.74	0.78	0.71
rpoB	513-wt	Q513L	11	12	0.90	0.95	0.96	0.88	0.96	0.99
rpoB	513-wt	Q513K	11	13	0.91	0.94	0.86	0.86	0.87	0.88
rpoB	513-wt	Q513P	11	14	0.83	0.75	0.79	0.83	0.80	0.80
rpoB	515-wt	M515I	15	16	0.78	0.57	0.65	0.61	0.68	0.41
rpoB	516-wt	D516E	17	20	0.69	0.80	0.54	0.76	1.05	0.62
rpoB	516-wt	D516Y	17	23	0.50	0.24	0.65	0.41	0.53	0.36
rpoB	516-wt	D516G	17	24	0.69	0.54	0.25	0.81	0.83	0.50
rpoB	516-wt	D516V	17	25	0.56	0.53	0.65	0.60	0.14	0.54
rpoB	522-wt	S522L	26	28	0.79	0.86	0.91	0.83	0.84	0.77
rpoB	524-wt	L524S	29	30	0.87	1.05	0.92	0.92	0.84	0.83
rpoB	526-wt	H526D	31	32	0.87	0.83	0.83	0.85	0.87	0.90
rpoB	526-wt	H526R	31	34	0.92	0.93	0.87	0.84	0.97	0.93
rpoB	526-wt	H526L	31	35	0.90	0.92	0.85	0.86	0.91	0.89
rpoB	526-wt	H526Q (CAA)	31	36	0.89	1.00	0.99	0.86	0.97	0.98
rpoB	526-wt	H526Q (CAG)	31	37	0.95	0.98	0.90	0.94	0.96	0.98
rpoB	526-wt	H526C	31	38	0.97	0.98	0.89	0.93	0.96	0.98
rpoB	526-wt	H526N	31	39	0.91	0.92	0.92	0.94	0.97	0.95
rpoB	526-wt	H526P	31	41	0.75	0.82	0.83	0.87	0.90	0.84
rpoB	526-wt	H526Y	31	43	0.57	0.58	0.60	0.62	0.74	0.71
rpoB	531-wt	S531W	44	46	0.74	0.85	0.65	0.77	-0.96	-0.92
rpoB	531-wt	S531L	44	47	0.74	0.56	0.89	0.72	-0.42	-0.20
rpoB	531-wt	S531Q	44	48	0.86	1.08	0.93	0.99	0.76	0.47
rpoB	531-wt	S531C	44	50	0.61	0.75	0.56	0.56	-0.89	-0.79
rpoB	533-wt	L533P	51	52	0.83	0.79	0.75	0.76	1.18	0.78
katG	315-wt	S315T (ACC)	59	63	0.94	0.91	-0.79	-0.48	0.93	0.90
katG	315-wt	S315T (ACA)	59	78	0.93	0.92	-0.15	0.30	0.93	0.91
katG	315-wt	S315N	59	79	0.95	0.90	0.14	0.57	0.92	0.89
inhA	-8T-wt	-8A	82	83	0.82	0.82	0.50	0.63	0.90	0.85
inhA	-8T-wt	-8C	82	84	0.74	0.71	0.59	0.54	0.77	0.75
inhA	-15C-wt	-15T	85	86	0.94	0.92	-0.58	-0.49	0.97	0.95
inhA	-17G-wt	-17T	87	88	0.79	0.79	0.79	0.84	0.87	0.82

Ratios in black are called wild-type.

Ratios in red are called mutants.

Ratios in gray are not called in deference to more negative ratio in the same grouping.