

Insights into Drug Resistance Tuberculosis in Ghana / West Africa

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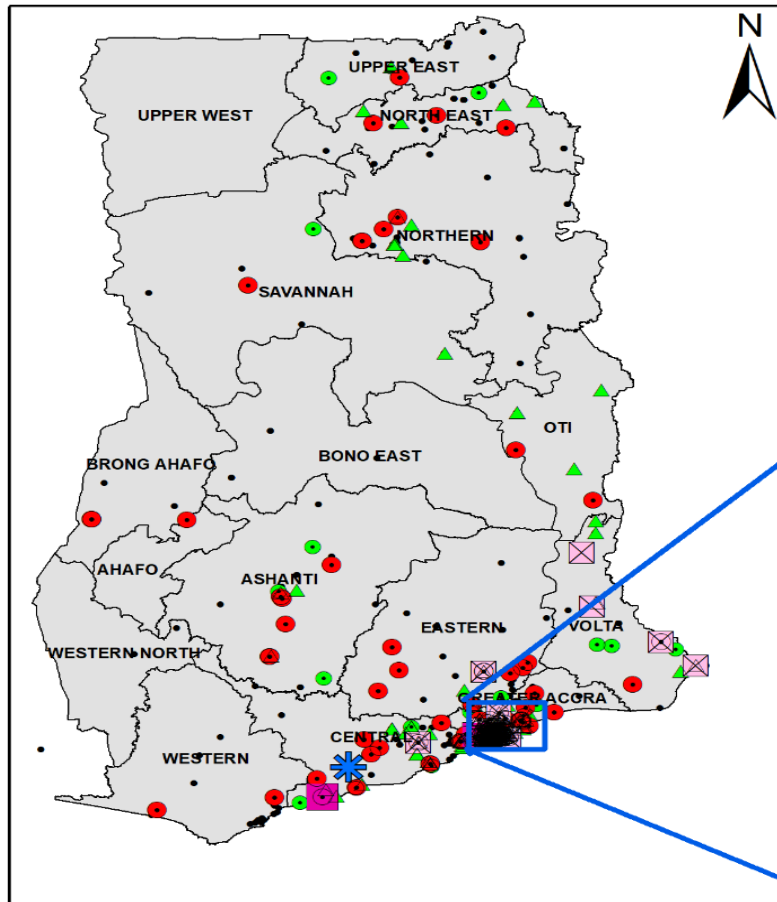
Background Information About Ghana

- WHO estimates for Ghana
- TB incidence of 133 per 100,000
- Part of the 30 high TB/HIV burden countries in 2021
- Estimated P% of new TB cases with MDR/RR-TB = 1.9%
- Estimated P% of previous TB cases with MDR/RR-TB = 3.6%
- More than 20% of TB is caused by *Mycobacterium africanum* (L5 and L6)



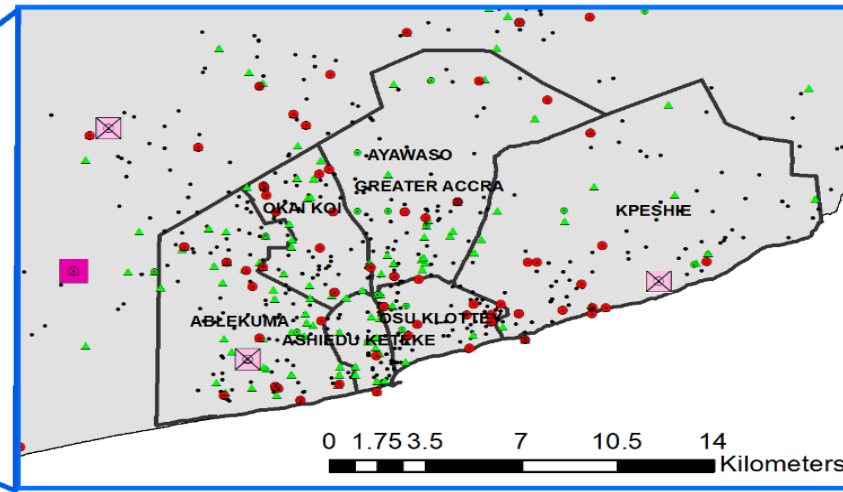
- BPaL and BPaLM rolled out in Ghana – since 2023
- All regions of Ghana currently on BPaL & BPaLM
- System developed for recording all adverse events related to MDR cases
- Approved by the FDA

Spread of DR-TB across Ghana



DR_status

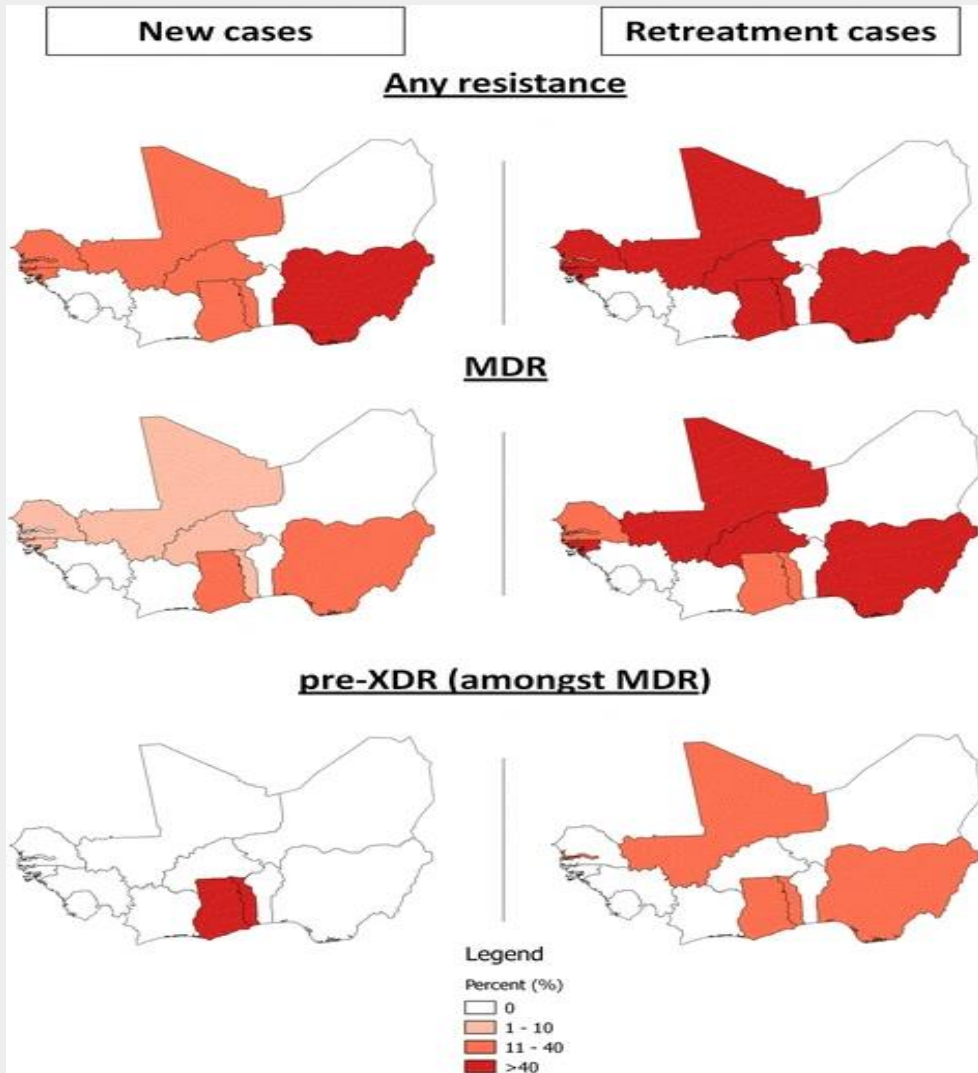
- ✠ XDR
- ✠ Pre XDR
- ★ Poly Resistance
- MDR
- RIFr
- ▲ INHr
- sensitive



Accra metropolitan assembly

The Emerging Threat of DR-TB in West Africa

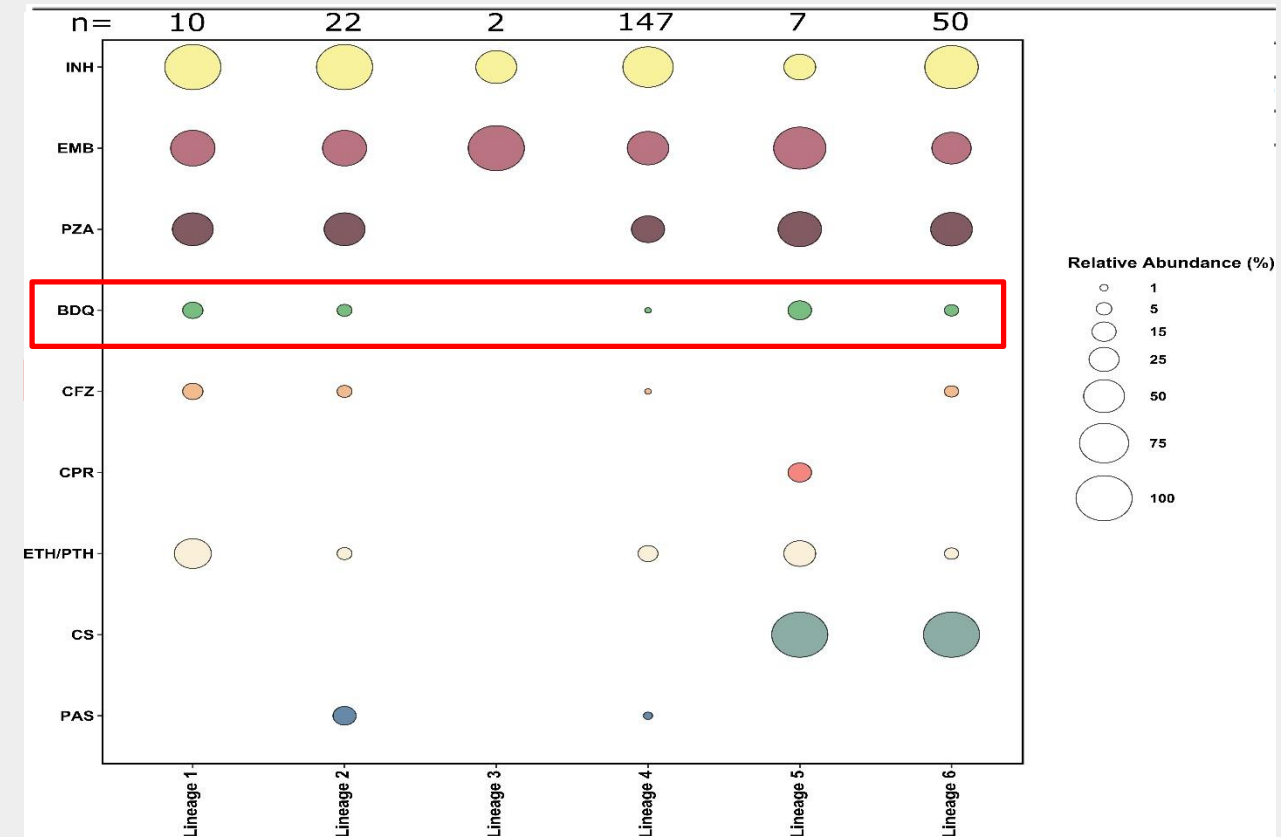
DR-TB in WANETAM study sites



Gehre et al

Blankson et al

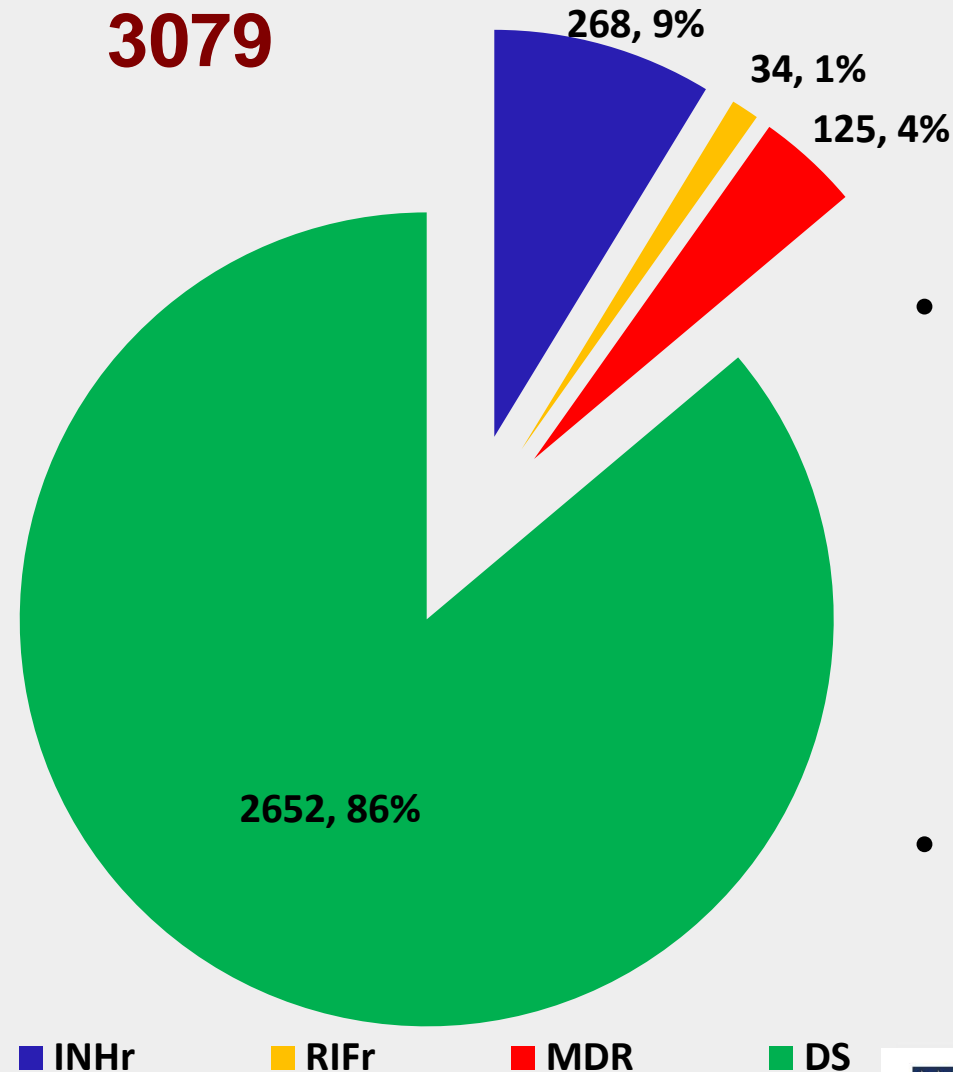
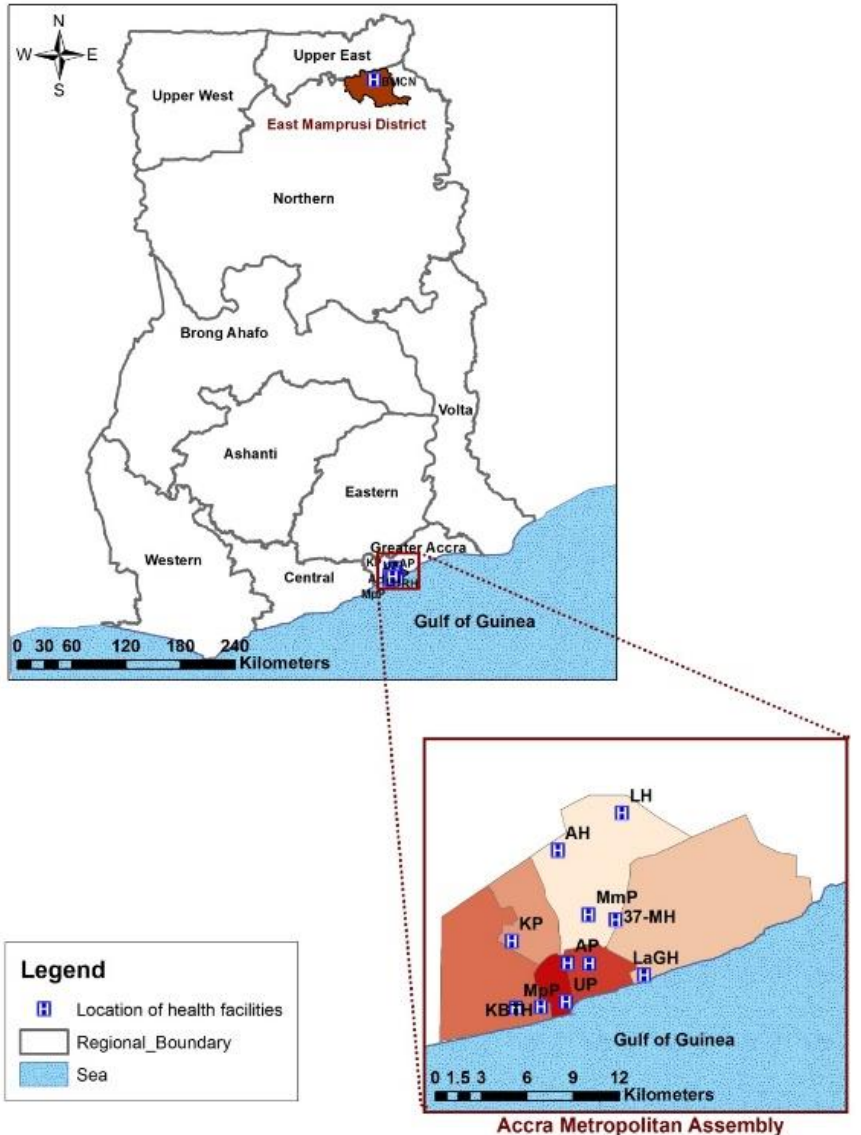
Second-line DR in West Africa



- Resistance prevalence:
 - INH= 82% ($n = 196$)
 - EMB= 53% ($n = 126$)
 - PZA= 39% ($n = 92$)
- 5 strains were resistant to BDQ/clofazimine (CFZ) based on mutations in **Rv0678**

Distribution of Drug resistance among MTBC Isolates

Urban vs Rural Setting



- 14% of the isolates were resistant to at least one of the drugs tested
- Approximately 4% MDR-TB cases

Drug resistant conferring and compensatory mutations

RIF Resistance Isolates

Target	Mutation	Number	Other mutations
<i>rpoB</i>	Q432P	1;1.5%	
	Q432P & I491S	1;1.5%	
	Q432K	1;1.5%	
	D435V	11;16.7%	
	D435Y	1;1.5%	
	S441L	1;1.5%	
	H445R	3;4.5%	
	H445C	2;3.0%	
	H445D	4;6.1%	
	H445Y	6;9.1%	
	<u>S450L</u>	32;48.5%	1: <i>rpoB</i> S388L, 1: <i>rpoB</i> Q409R* , <i>rpoC</i> ;2:G332R, 3:V483G
<i>rpoC</i>	G332R	2;3.0%	<i>rpoB</i> ; S450L
	V483G	3;4.5%	<i>rpoB</i> ; S450L

INH Resistant Isolates

Target	Mutation	Number
<i>inhApro</i>	-8T/C	7;3.5%
	-15C/T	28;14.6%
	-17G/C	1;0.5%
<i>katG</i>	S315T & I317V	1;0.5%
	<u>S315T</u>	142;70.3%
<i>inhA</i>	G204D*	16;7.2%
	V78A	1;0.5%
<i>ahpCpro</i>	-54C/T	1;0.5%
	-88G/A*	2;1.0%
	-142G/A*	1;0.5%
<i>ndh</i>	V117I*	2;1.0%

* Represents Novel mutations

DR and resistance conferring mutations are driven by specific MTBC genotypes

Association of drug resistance with Specific MTBC genotypes

Resistance	Cameroon (806)	Ghana (195)	OR	95% CI	p-value
INH	54 (6.7%)	43 (22.1%)*	0.25	0.16–0.40	0.0000
RIF	19 (2.4%)	12 (6.2%)*	0.37	0.17–0.85	0.0103
MDR	13 (1.6%)	9 (4.6%)*	0.34	0.13–0.91	0.0240
ANY	60 (7.4%)	46 (23.6%)*	0.26	0.17–0.41	0.0000
	MAF WA 1 (165)	Ghana (195)			
INH	25 (15.2%)	43 (22.1%)	0.63	0.35–1.12	0.1059
RIF	10 (6.1%)	12 (6.2%)	0.98	0.37–2.56	0.9998
MDR	10 (6.1%)	9 (4.6%)	1.33	0.47–3.81	0.6384
ANY	25 (15.2%)	46 (23.6%)*	0.58	0.32–1.02	0.0472
	MAF WA 2 (107)	Ghana (195)			
INH	6 (5.6%)	43 (22.1%)*	0.21	0.07–0.52	0.0001
RIF	1 (0.9%)	12 (6.2%)*	0.144	0.00–1.00	0.0372
MDR	1 (0.9%)	9 (4.6%)	0.19	0.00–1.45	0.1040
ANY	6 (5.6%)	46 (23.6%)*	0.19	0.06–0.48	0.0000

MAF WA 1: *M. africanum* West Africa 1 (Lineage 5).

MAF WA 2: *M. africanum* West Africa 2 (Lineage 6).

* Significantly higher.

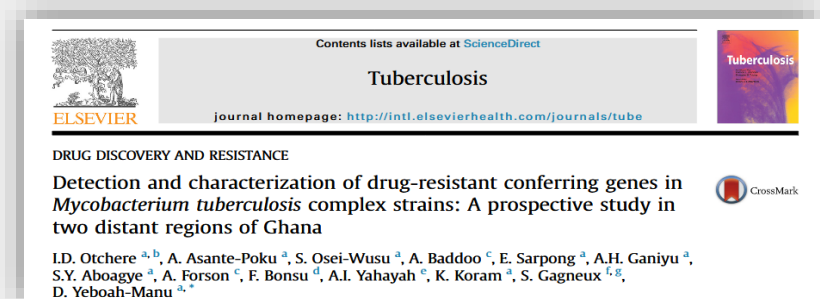
Ghana genotype of L4 causing about 13% of TB in Ghana is associated with drug resistance

• Bedaquilin-resistance?

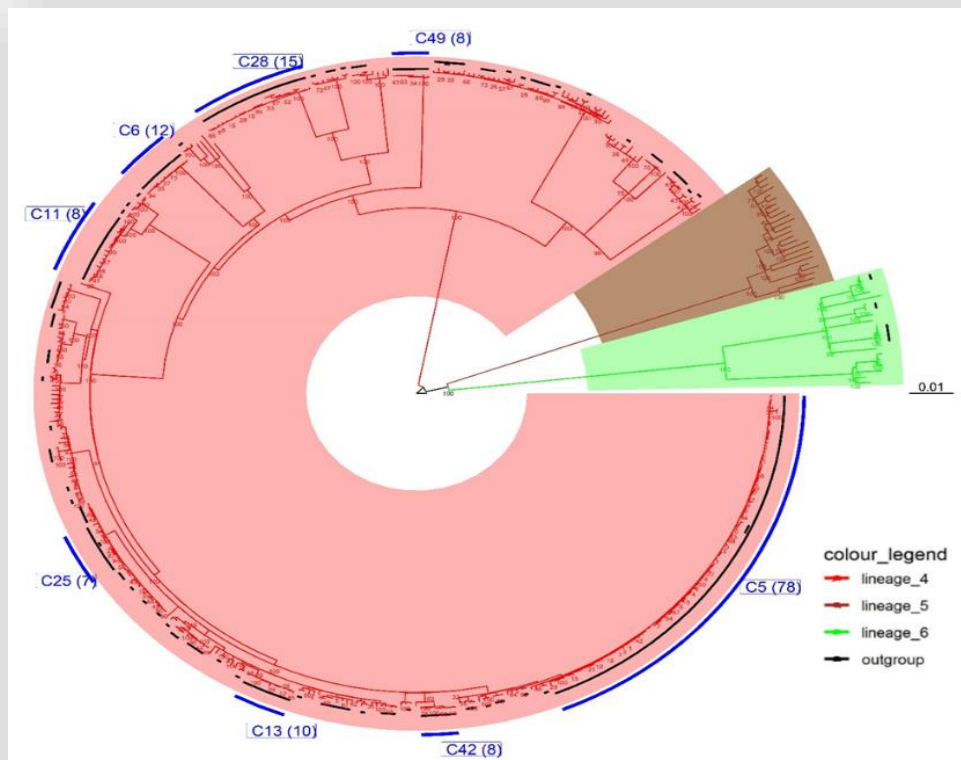
- *Rv0678*: No mutation seen
- *atpG*: Y220S (L5-specific)
- *atpH*: Q40K (2 isolates) & E351D (2 isolates).
- *pepQ*: G96R (1 isolate)
- *Rv1979c*: L14R (3 isolates), A50T (2 isolates) & D286G (5 isolates)

• Pretomanid resistance?

- *fbiA*: I208V (1 isolate)
- *fbiB/C*: No mutation
- *fgb1*: K270M (5 isolates)
- *ddn*: D113N (L5)



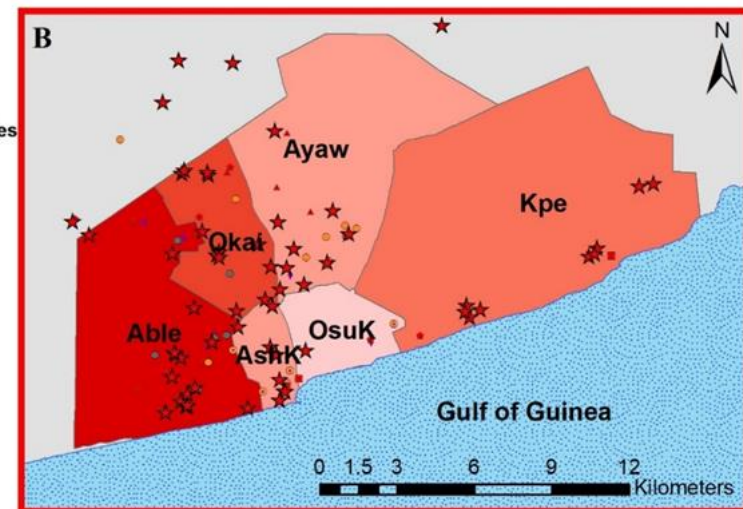
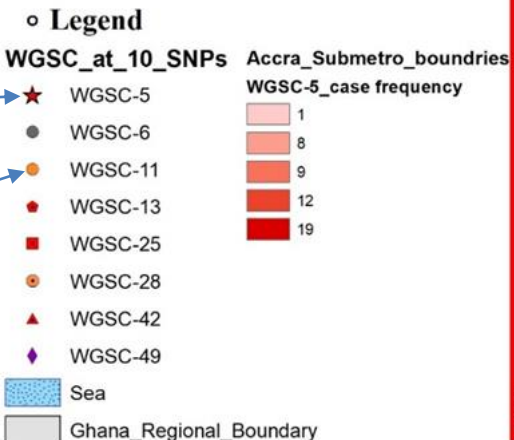
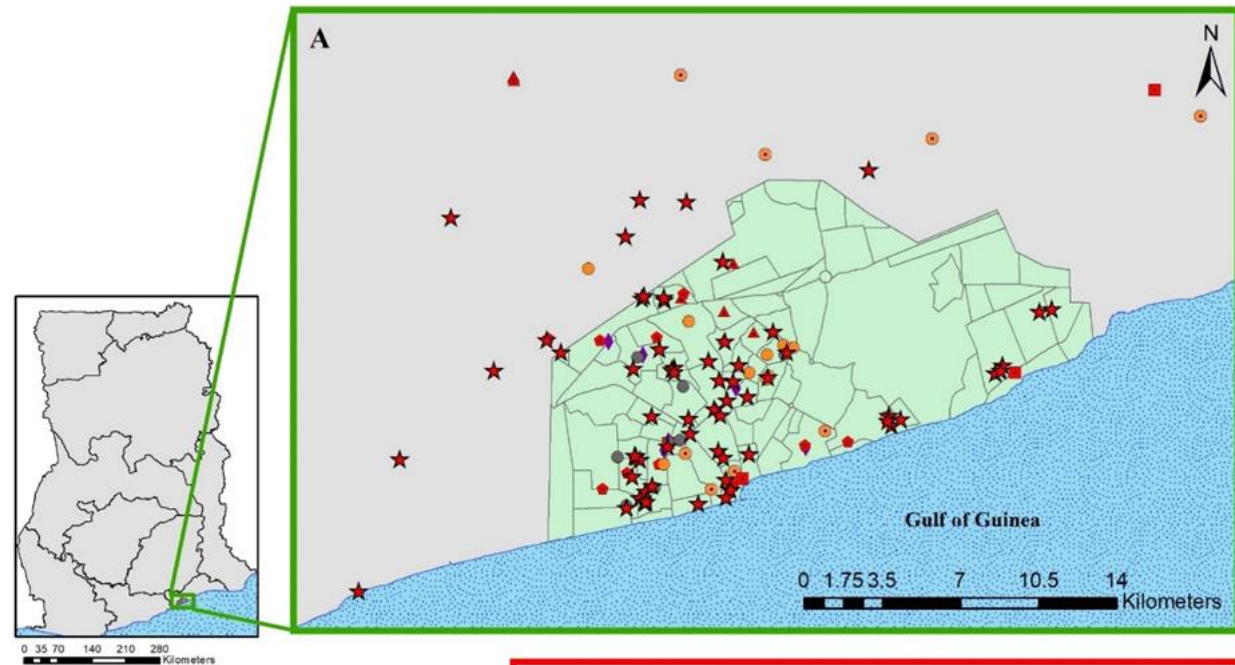
Hotspots of recent TB transmission in Accra, Ghana



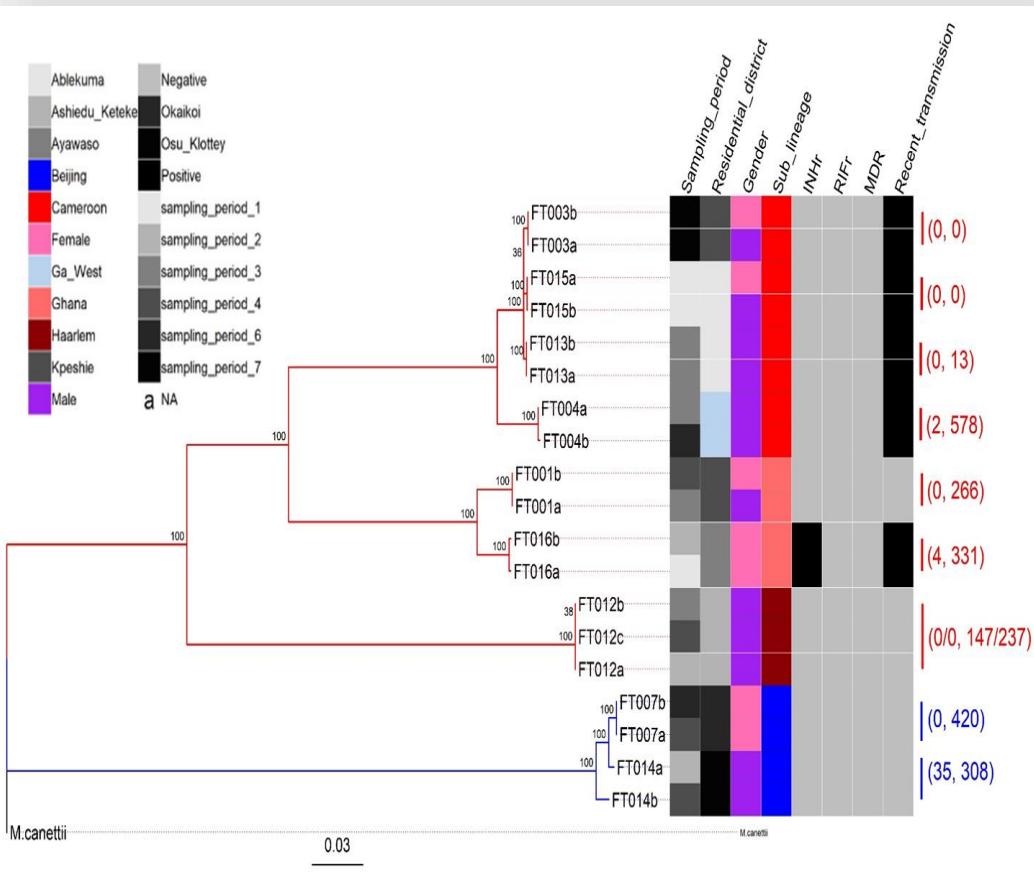
Largest recent transmission cluster predominantly in Ablekuma sub-district

Isoniazid transmitting cluster predominantly in Ayawaso sub-district

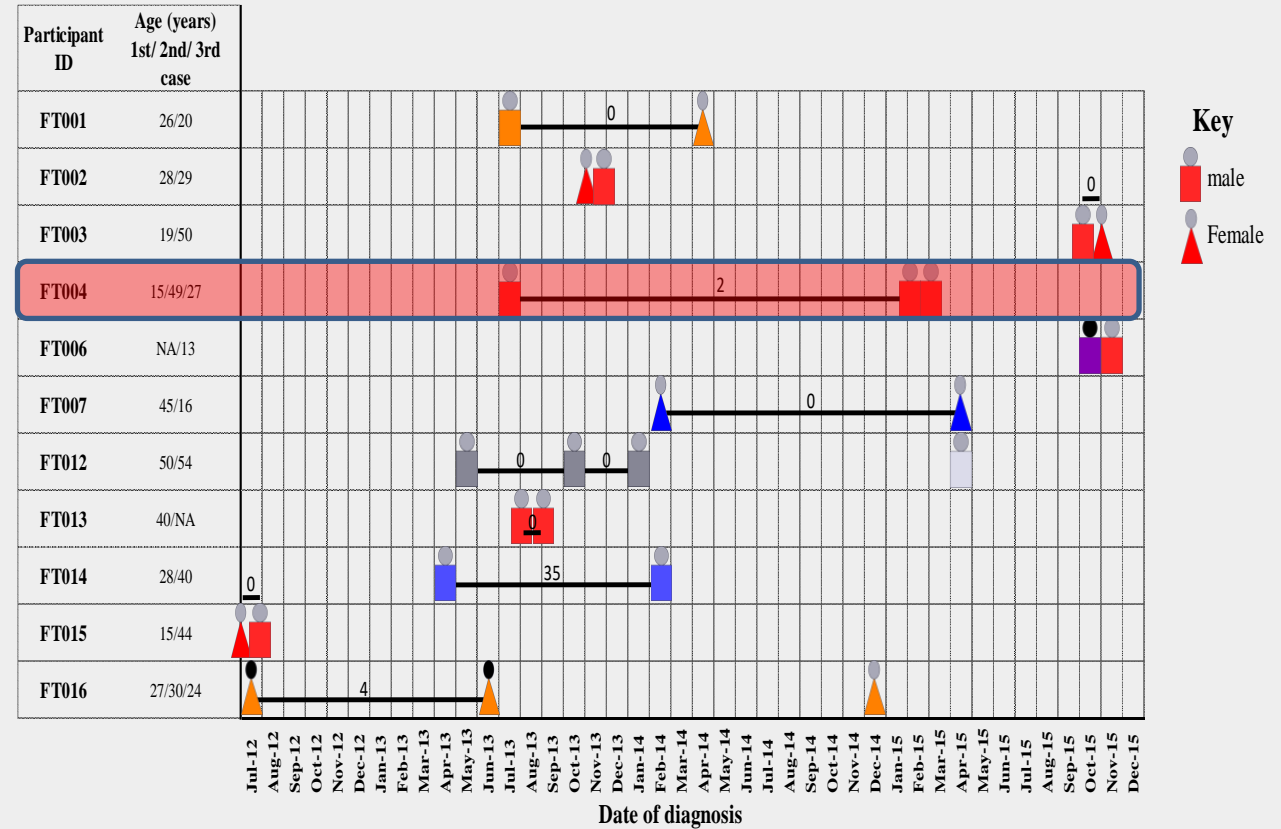
Using genomics to support both diagnosis and epidemiology



Evidence of Household Recent TB Transmission



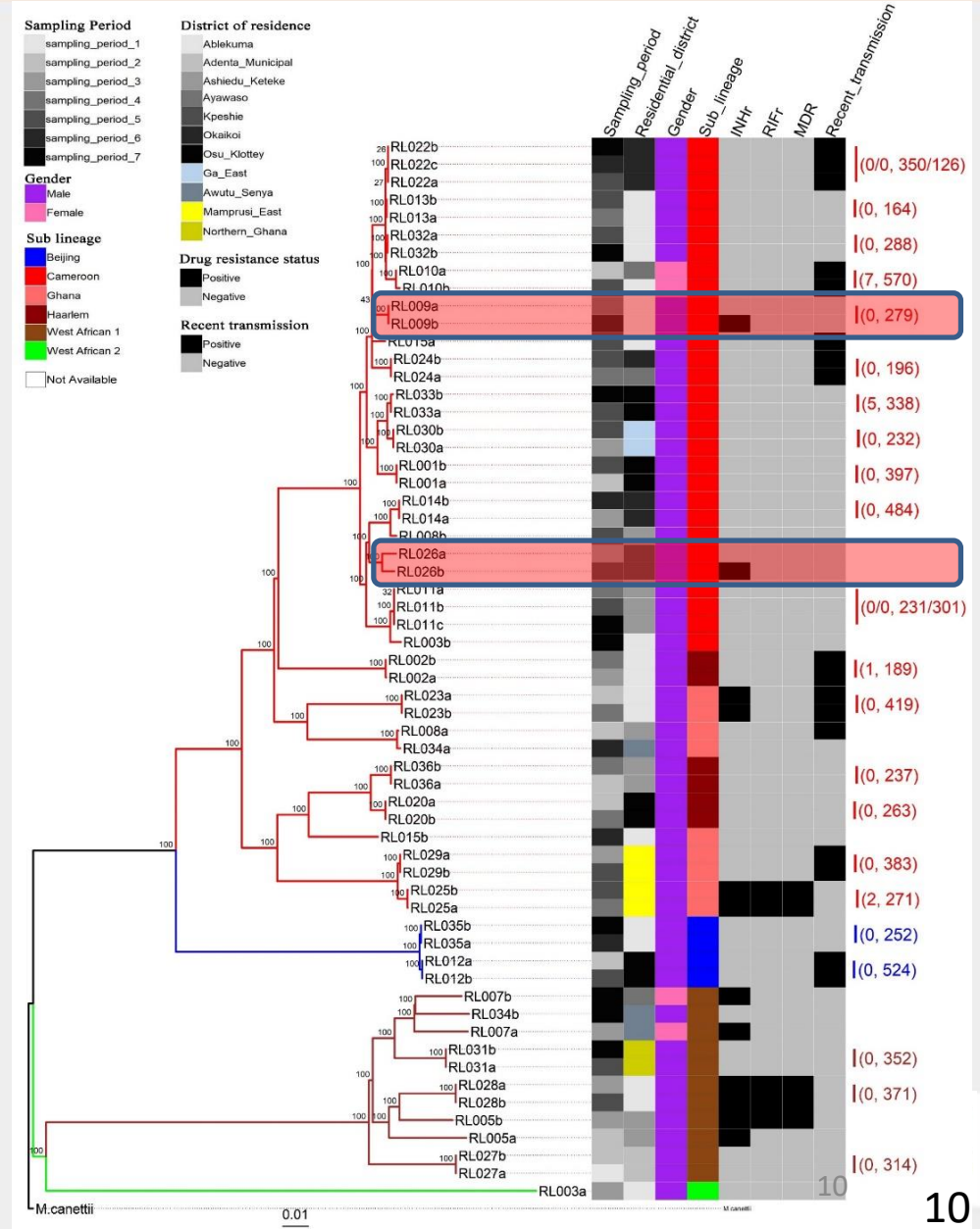
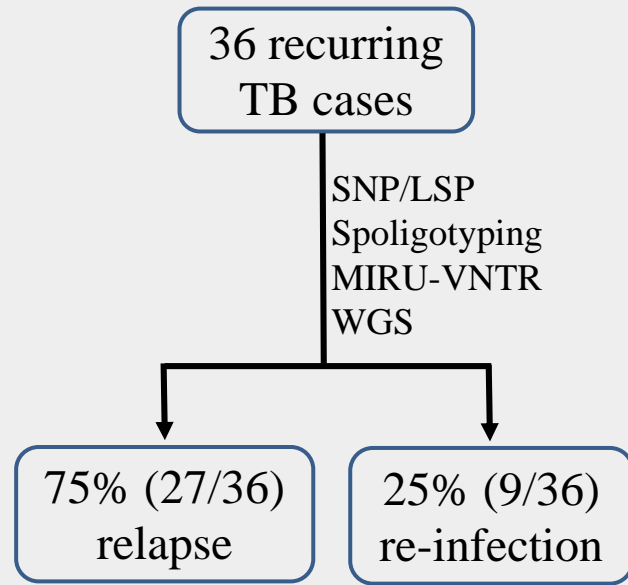
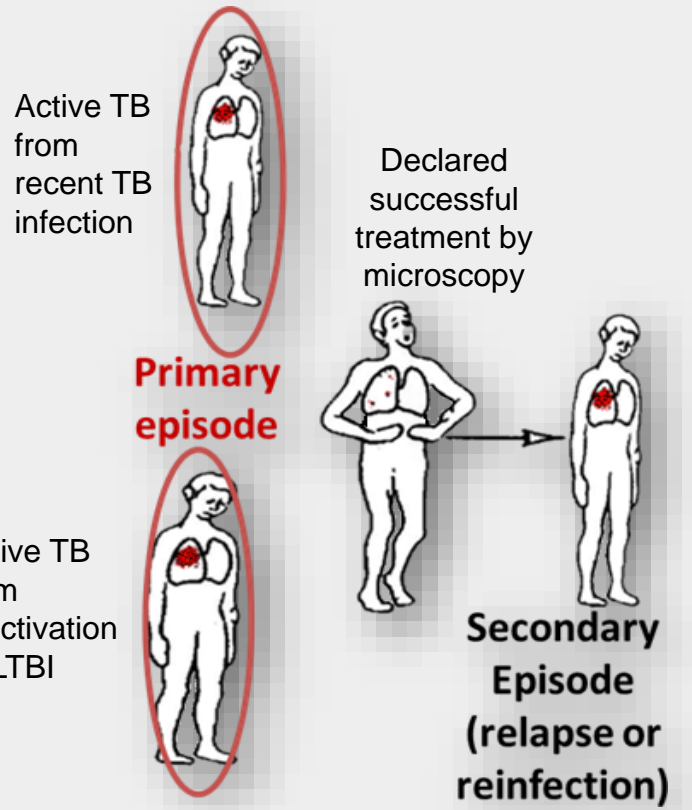
Phylogenetic reconstruction showing the genomic relationship between 19 MTBC isolates from 9 households



Time till event of household related tuberculosis cases

Need for contact screening

Delineating the Occurrence of Recurrent TB in Ghana



Patients with recurring TB are largely due to relapse (inadequate treatment) of previous infection

Community Engagement

Educating Market women on TB



Student Engagement on TB/DR-TB



Explaining TB in Local Language (Twi) on National TV



Free Health Screening at identified hotspots of transmission



Education on TB during the free health screening



MDR Survivor shared her story to sensitize others of TB



Take Home Message

- Lineage diversity is very important ==>Geography
- Evidence of Community and household spread
- Intensify public education to improve early case reporting
- Observed relapse rate calls for measures to improve compliance and treatment monitoring

THANK YOU