Potential of Cell-Free DNA in Plasma and Urine for Rapid Detection of *Mycobacterium tuberculosis*

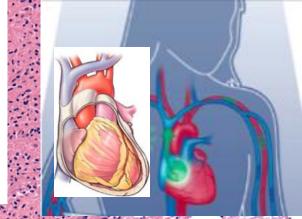
> Niaz Banaei MD Director, Clinical Microbiology Laboratory Associate Professor of Pathology and Medicine Stanford University

Disclosures

- Banaei:
 - IP interest in GWiS PCR
 - Industry links or funding related to this talk
 - Research support form KariusDx

60 y/o Vietnamese with restrictive pericarditis Pericardial biopsy

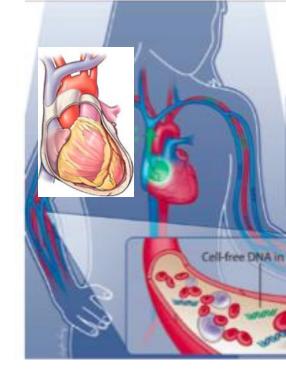
Necrotizing granulomas, no AFB TB PCR and culture: Positive for MTC Sputum culture x3: Negative



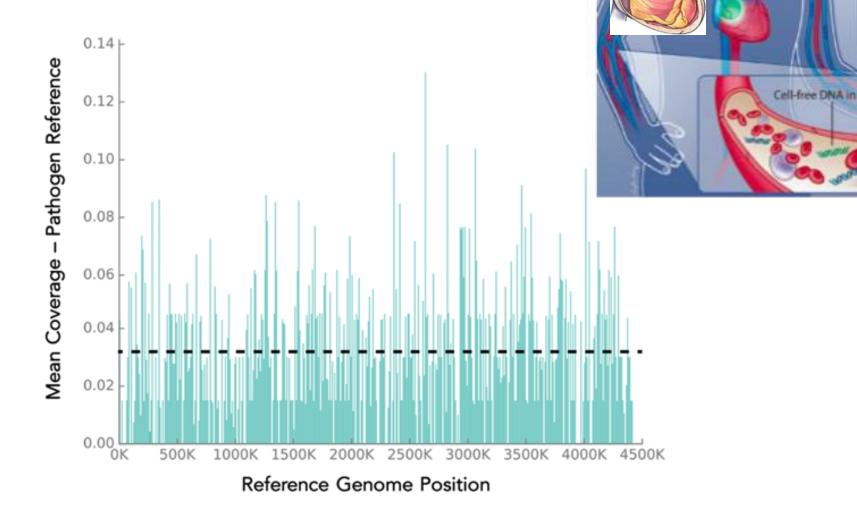


60 y/o Vietnamese with restrictive pericarditis EDTA Plasma

Deep sequencing of plasma cell-free DNA on Illumina NextSeq500, 75 million reads



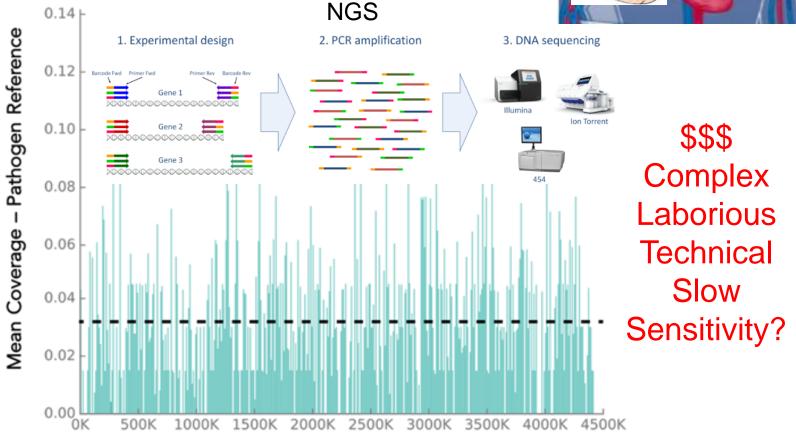
60 y/o Vietnamese with restrictive pericarditis EDTA Plasma Deep sequencing of plasma cell-free DNA on Illumina NextSeq500



KARIUS

Hong et al ASM Microbe 2016 Hong et al In preparation 60 y/o Vietnamese with restrictive pericarditis EDTA Plasma Deep sequencing of plasma cell-free DNA on Illumina NextSeq500





Reference Genome Position

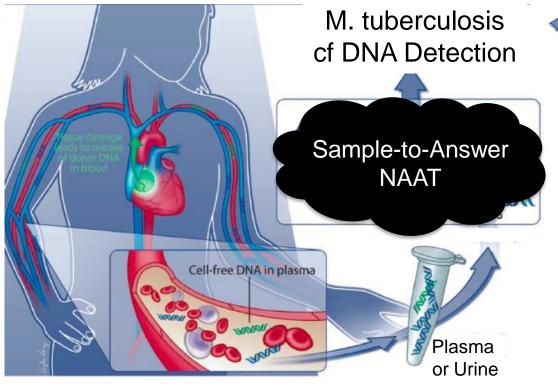
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Application of cf DNA for Diagnosis of TB

Affordable Sensitive Specific User-friendly Robust, rapid Equipment min Deliverable

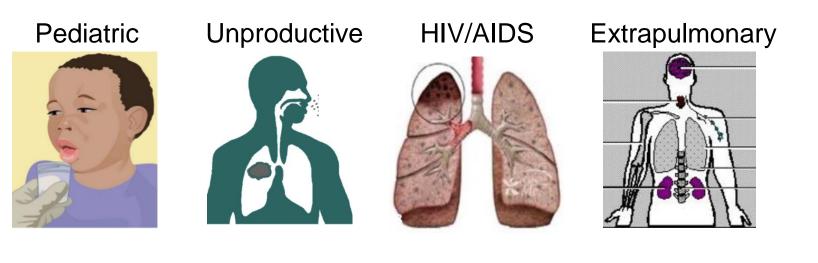
Application of cf DNA for Diagnosis of TB



adapted from De Vlaminck Sci Transl Med 2014

Affordable Sensitive Specific User-friendly Robust, rapid Equipment min Deliverable

Target Population for cf DNA TB Diagnosis

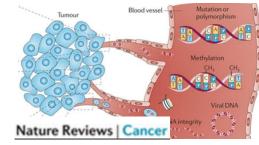


Application of cf DNA in Diagnostics

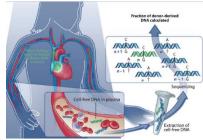
Fetal aneuploidy



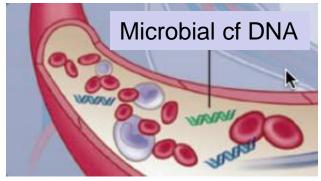
Cancer mutations



Organ rejection



Infectious diseases



- EBV→nasopharyngeal CA (Cancer Res 1999)
- Invasive fungal infection (CID 2013)

www.thelancet.com/infection Vol 9 August 2009

Rapid diagnosis of tuberculosis through the detection of mycobacterial DNA in urine by nucleic acid amplification methods

Clare Green, Jim F Huggett, Elizabeth Talbot, Peter Mwaba, Klaus Reither, Alimuddin I Zumla

	Sensitivity by tu	berculosis present	ation	Effect of HIV coir	nfection* on sensitivity	Method of tuberculosis confirmation	Target size bp†
	Pulmonary	Extrapulmonary	Other	HIV positive	HIV negative	-	
Sechi et al ³⁴			13% ND (77/602)	16% (65/412)	6% (12/190)	Suspected—3% (18/602) urine culture positive	182 (566)
Aceti et al ³⁵	100% (13/13)			100% (13/13)		Sputum smear or culture	309 (566)
Kafwabulula et al ³³	56% (35/63)			64% (32/50)	23% (3/13)	Sputum smear or culture	181 (556)
Torrea et al ³⁶	44% (108/247)	57% (48/84)		59% (86/145)	38% (70/186)	Pulmonary tuberculosis diagnosed by sputum smear or culture; extrapulmonary tuberculosis diagnosed by clinical criteria	309 (566)
Rebollo et al ³⁷ ‡	7% (2/27)	14% (2/14)	16% D (5/31)	28% (7/25)	6% (2/32)	Culture from any clinical sample and response to treatment	123
Cannas et al ³⁸ §	79% (34/43)					Sputum smear or culture	67 (129)
Gopinath and Singh ³⁹	52% (24/46)					Sputum culture	786

All studies used IS6110 as a target except Gopinath³⁹ who amplified from cfp32. The tabulated data present the most relevant comparisons between studies relating to transrenal DNA detection. For all studies, except the initial study by Sechi and colleagues³⁴ who relied on empirical observations, the gold standard for pulmonary diagnosis was sputum smear or culture positives. Data have been divided to distinguish between different presentations where appropriate. D=disseminated. ND=not disclosed. *Studies in which fewer than ten HIV-positive cases included in the study are not detailed. †Size of external product shown in brackets where amplification was nested. \$0–1 month after presentation and initiation of treatment. SOnly data relating to the transrenal DNA are detailed. The authors considered the urine pellet separately, which produced significantly lower detection rates.

Table 1: Summary of studies on urine-based detection of mycobacterial DNA by PCR amplification

	Age	Country	TB Type	Cases	Controls	Method	Target	Sensitivity	Specificity
Cannas et al IJTLD 2008	>18 yo	Italy	РТВ	43	23	Nested PCR	IS6110	79% (34/43)	100% (23/23)
Fortún et al IJTLD 2014	>18 yo	Spain	EPTB PTB	82 25	0	ТМА	16S rRNA	70% (57/82) 18% (5/25)	Not Done
Labugger et al Infection 2017	>18 yo	Germany	РТВ	11	8	PCR	IS6110	64%* (7/11)	100% (8/8)

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Labugger et al Infection 2017	РТВ	11	8	PCR	IS6110	64%* (7/11)	100% (8/8)

Miliary	Multifocal	LAN	Pleural	Joint
90%	67%	72%	33%	45%
(9/10)	(16/24)	(18/25)	(1/3)	(5/11)

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Labugger et al Infection 2017	РТВ	11	8	PCR	IS6110	64%* (7/11)	100% (8/8)	vs Radiology ↑cfDNA	vs Smear+ None	vs TTCxP None	Wk1 ↑cfDNA	Wk12 9/11 Neg
						*100% with retesting					•	, -



Accuracy of Plasma cf DNA for TB Diagnosis

	Age	Country	TB Type	Cases	Controls	Method	Target	Sensitivity	Specificity
Ushio et al Tuberculosis	>18 vo	Japan	РТВ	33	19	Digital	IS6110	65% (21/33)	93% (18/19)
2016	>18 y0	Japan	110	55	19	PCR	gyrB	29% (10/33)	100% (19/19)

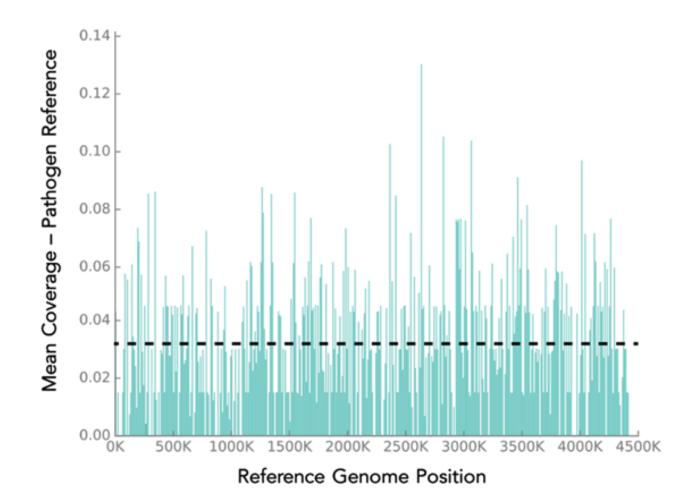


Accuracy of Plasma cf DNA for TB Diagnosis

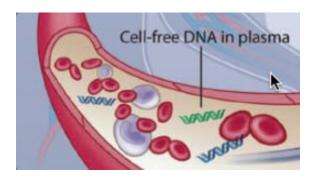
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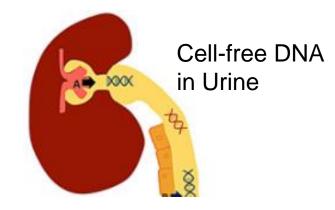
Bilateral vs	PTB+EPTB
Unilateral PTB	vs. PTB
个cfDNA	个cfDNA

Genome-Wide Sensitive PCR (GWiS PCR)



Potential of cf DNA in Diagnosis of TB





Accuracy
Pediatric
HIV/AIDS
Unproductive
Extrapulmonary

Sensitivity >70%
Specificity ≈100%
Image: Construction of the sense of the sens

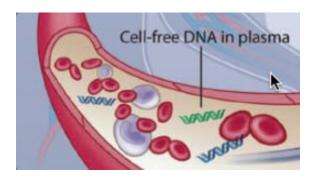


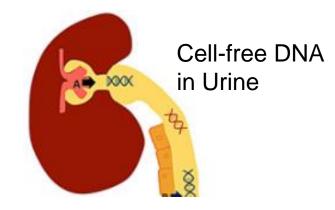
n mark X																			
	Age	Country	TB Type	Cases	Controls	HIV+	Smear+	Reference	Sample	Tx Naïve	Preserve	Fresh/F rozen		Extraction	A 2	Method	Target	Sensitivity	Specificity
Cannas et al IJTLD 2008	>18 yo	Italy	РТВ	43	23	5%	95%	Culture	Urine	No	EDTA	Frozen	SmL	Mannual/ Resin	LDT	nested PCR	IS6110	79% (34/43)	100% (23/23)
Fortún et al IJTLD 2014	>18 yo >18 yo	Spain Spain	ЕРТВ РТВ	82 25	0 0	? ?	NA ?	Culture Culture	Urine Urine	Yes Yes	? ?	? ?	? ?	? ?	MTD (Hologic) MTD	ТМА ТМА	16S rRNA 16S rRNA	70% (57/82) 18% (5/25)	Not Done Not Done
Labugger et al Infection 2017	>18 yo	Germany	РТВ	11	8	0%	60%	Culture	*Urine	Yes	EDTA	?Fresh	4	Mannual/ Resin	LDT	PCR	IS6110	64%* (7/11)	100% (8/8)



Ushio et al	>18 vo	Japan	РТВ	33	19	0%	100%	Culture	Plasma	2	EDTA	?Fresh	0.2	Qiagen	LDT	digital PCR	IS6110	65% (21/33)	93% (18/19)
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Acknowledgements

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FIND

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