

A Novel RNA In Situ Hybridization Method for the Detection of Mycobacteria in **Clinical FFPE Samples**

Background

Acid fast staining (AFS) for the detection of mycobacteria and other acid-fast organisms provides a rapid way of identifying pathogenic organisms in FFPE tissue in contrast to culture methods, which require weeks to perform. While rapid and straightforward, there are several issues with AFS. First, AFS can be challenging and time-consuming to interpret. Second, standard tissue fixatives and processing alters the lipid-rich cell wall of acid-fast organisms which reduces the detection sensitivity in FFPE samples. Last, AFS does not allow for discrimination between various acid-fast organisms, such as mycobacteria tuberculosis (MTB) vs. non-tuberculous mycobacteria (NTM). To address these issues with AFS, we have developed a highly sensitive and specific RNA in situ hybridization (RISH) assay using the RNAscope technology for the detection of mycobacteria rRNA in FFPE tissues and distinction of MTB and NTM.

Design

Serial sections from 50 FFPE lung samples from patients with a clinical diagnosis of pulmonary TB (not histologic) were collected from five hospitals in China. Samples were tested with AFS or using the RNAscope probes B-MTB-23s-rRNA-1-C1 and B-MTB-NTM-16srRNA-O1 targeting MTB only or both MTB and NTM, respectively, using a modified RNAscope protocol (Protocol I). A subset of the RISH positive samples were subsequently examined using an alternative protocol (Protocol II), which eliminates one pretreatment step (Figure 1). AFS and RISH slides were then analyzed for the presence of positively stained organisms.



Figure 1. RNAscope ISH assay overview



Results

MTB-NTM⁺ samples tested using Protocol II yielded the same results (Figure 2 and Table 1).

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Results



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Results

Sample number	Acid fast	RNAscope 2.5 Reagent Kit- RED. Protocol I	KNASCOPE 2.5 Reagent Kit-RED. Protocol I	KIT-RED. Protocol II	RNAscope 2.5 Reagent Ki RED. Protocol II
	(AFS)	B-MTB-NTM-16srRNA-O1	B-MTB-23s-rRNA-1-C1	B-MTB-NTM-16srRNA-	B-MTB-23s-rRNA-1-C1
				01	
1	-		-	N/A	N/A
2	-	_	-	N/A	N/A
3	_	_	_	N/A	N/A
4	-	+	+	N/A	N/A
5	_	_	-	N/A	N/A
6	+	+	+	+	+
7	+	+	+	+	+
8	-	+	+	+	+
9	-	_	-	N/A	N/A
10	+	+	+	+	+
11	+	+	-	N/A	N/A
12	+	+	+	N/A	N/A
13	+	+	-	N/A	N/A
14	-	-	-	N/A	N/A
15	-	-	-	N/A	N/A
16	-	-	-	N/A	N/A
17	-	_	-	N/A	N/A
18	-	-	-	N/A	N/A
19	-	-	-	N/A	N/A
20	+	+	+	+	+
21	+	+	+	+	+
22	+	+	+	+	+
23	+	+	+	+	+
24	+	+	+	+	+
25	+	+	+	+	+
26	+	+	-	N/A	N/A
27	+	+	+	+	+
28	+	+	+	+	+
29	+	+	+	+	+
30	+	+	+	+	+
31	+	+	+	+	+
32	+	+	+	+	+
33	+	+	+	+	+
34	+	+	+	+	+
35	+	+	+	+	+
36	+	+	+	+	+
37	+	+	+	+	+
38	+	+	+	+	+
39	-	-	_	N/A	N/A
40	+	+	+	+	+
41	+	+	+	+	+
42	+	+	+	+	+
43	+	+	+	+	+
44	+	+	+	+	+
45	+	+	+	N/A	N/A
46	+	+	+	N/A	N/A
4/	+	+	+	N/A	N/A
48	+	+	+	N/A	N/A
49	+	+	+	N/A	N/A

Conclusions

In the current study we have developed a novel RISH method for the detection of mycobacteria in FFPE tissue. In comparison to AFS, the RISH methodology appears to be more sensitive, detecting 2 additional positive cases. In addition, the RISH method was able to identify 3 cases of probable NTM, a distinction of high clinical relevance owing to the different treatments required for MTB vs. NTM infections. Finally, the RISH signals were easier to visualize and locate compared to AFS. Overall, our results indicate that the RNAscope assay is a sensitive method to detect mycobacteria in FFPE tissues that allows for differentiation of MTB vs. NTM and is easy to interpret.



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