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**Increased Frequency of Nontuberculous Mycobacteria Detection at Potable Water Taps
within the United States**

Number of Pages: 2 (includes coversheet)

Table S1: Summary of sampling process used by each NTM occurrence study

	Study	von Reyn <i>et al.</i> , 1993	Glover <i>et al.</i> , 1994	Covert <i>et al.</i> , 1999	Falkinham <i>et al.</i> , 2001	Hilborn <i>et al.</i> , 2006	This study
Sample collection	Flush time	First pull	First pull	1 to 2 min	Not reported	First pull	15 sec
	Sample preservative	Sodium thiosulfate	Sodium thiosulfate	Sodium thiosulfate	Sodium thiosulfate	Sodium thiosulfate	Not added
Sample treatment	Decontamination	None	CPC* %0.04	CPC %0.04	None	CPC %0.04	CPC %0.04
	Decontamination time		24 hr.	30 min		30 min	30 min
Sample concentrating	Concentration method	Centrifugation@ 5,000 g, Spread plate 0.1 mL	Vacuum filtration 500 mL	Vacuum filtration 500 mL	Centrifugation@ 5,000 g, Spread plate 0.1 mL	Vacuum filtration 500 mL	Vacuum filtration 500 mL
	Volume of water represented in NTM recovery	2.5 mL/plate 3 plates 7.5 mL analyzed	500 mL	500 mL	30 mL/plate 5 plates 150 mL analyzed	500 mL	500 mL
NTM isolation	Media	Tsukamura minimal-Tween 80 cycloheximide agar media	Middlebrook 7 H10	Middlebrook 7 H10	Middlebrook 7H10	Middlebrook 7H10	Middlebrook 7H10
	Incubation Time	Eight weeks	Eight weeks,	Eight weeks	Three weeks	Eight weeks	Eight weeks
	Incubation Temperature	37 °C	37 °C	37 °C	37 °C	37 °C	37 °C
Identification	Presumptive NTM	Acid fast	Acid fast	Acid fast	Acid fast	Acid fast	
	MAC identification	DNA probes (SNAP or Gen-Probe)	DNA probes (Syngene)	16S partial sequencing	Multiplex PCR	Multi locus electrophoresis	16S partial and <i>hsp65</i> genes
	NTM identification excluding MAC isolates	Not done	Biochemical screening	16S partial sequencing	Endonuclease digest <i>hsp65</i> (BstEII and HaeIII)	HPLC mycolic acid	16S partial and <i>hsp65</i> genes

*CPC, cetylpyridinium chloride

Table S1: Summary of sampling approach used by each NTM occurrence study.