

Supporting Table 8

			Lineage				Total
			EAI	EAm	EAs	IO	
local of infection	pumonary TB	Count	12 _a	43 _a	9 _a	15 _a	79
		% within Lineage	66,7%	67,2%	90,0%	48,4%	64,2%
	extra-pulmonary TB	Count	1 _a	10 _a	0 _{a, b}	13 _b	24
		% within Lineage	5,6%	15,6%	0,0%	41,9%	19,5%
	pulmonary+extra-pulmonary TB	Count	5 _a	11 _a	1 _a	3 _a	20
		% within Lineage	27,8%	17,2%	10,0%	9,7%	16,3%
Total	Count	18	64	10	31	123	
	% within Lineage	100,0%	100,0%	100,0%	100,0%	100,0%	

Each subscript letter denotes a subset of LSP_lineage categories whose column proportions do not differ significantly from each other at the 0.05 level (column proportions compared by the z-test with p-values adjusted by the Bonferroni method).