



HA												1	1 (20, 0.5–72)	5
CA	3				1								4 (80, 28–99)	
ST125						1						5	6 (75, 35–97)	8
HA													0	
CA														
ST152							4						4 (67, 22–96)	6
HA														
CA							2						2 (33, 4–78)	
ST228														
HA				1			5					11	1 (95, 74–100)	19
CA													0	
ST239														
HA			22	1	5		30			3			61 (94, 85–98)	65
CA			1										1 (2, 0.04–8)	
ST241														
HA			4		5					1		1	11 (92, 62–100)	12
CA													0	
Total HA:CA (% , 95% CI)														
HA	7 (39, 17–64)	31 (94, 80–94)	43 (47, 36–57)	11 (46, 26–67)	20 (40, 26–55)	15 (71, 48–89)	56 (62, 51–72)	1	9 (53, 28–77)	5 (42, 15–72)	31 (86, 71–95)	15 (83, 59–96)	244 (59, 54–64)	
CA	9 (50, 26–74)	2 (6, 1–20)	37 (40, 30–51)	9 (38, 19–59)	20 (40, 26–55)	5 (21, 7–42)	29 (32, 23–43)	0	3 (21, 5–51)	4 (33, 10–65)	0	1 (6, 0.1–27)	119 (29, 24–33)	
Total	18	33	92	24	50	24	90	1	14	12	36	18	414	

\*Shown are 414 independent cases (not belonging to the same transmission chain) of 444 cases acquired abroad. Cases are stratified as healthcare-acquired (HA) and community-acquired (CA) for each sequence type (ST) and region of acquisition. In some instances, numbers in rows and columns do not add up to the totals because of persons with STs other than the ones listed or because cases were indeterminable with regard to transmission setting. CI, confidence interval.