

Supporting Table 3

<i>mclx#2</i>			multivariate ORs (95% CI)
			Model 2
patient-related	age		0.958 (0.912-1.007) <i>p</i> =0.089 <i>B</i> =-0.043; <i>S.E.</i> =0.025 <i>Wald</i> =2.897
	ethnicity	native dutch	3.702 (0.811-16.898) <i>p</i> =0.091 <i>B</i> =1.309; <i>S.E.</i> =0.775 <i>Wald</i> =2.854
		foreign-born	1 (ref)
microorganism-related	transmissibility	no	1.754 (0.439-7.000) <i>p</i> =0.426 <i>B</i> =0.562; <i>S.E.</i> =0.706 <i>Wald</i> =0.632
		yes	1 (ref)
	lineage		<i>p</i> =0.357 <i>Wald</i> = 3.237
		EAI	0.136 (0.015-1.196) <i>p</i> =0.072 <i>B</i> =-1.998; <i>S.E.</i> =1.111 <i>Wald</i> =3.237
		EAm	1 (ref)
		EAs	<i>p</i> =0.999 0.000 (0.000-.) <i>B</i> =-20.230; <i>S.E.</i> =12299.315 <i>Wald</i> =0.000
		IO	<i>p</i> =0.998 0.000 (0.000-.) <i>B</i> =-20.251; <i>S.E.</i> =7120.660 <i>Wald</i> =0.000
Omnibus Test (chi-square/<i>p</i>)			40.620/ <i>p</i> <0.001
Cox & Snell R²			0.285
Nagelkerke R²			0.459
Hosmer and Lemeshow (chi-square/<i>p</i>)			0.792/ <i>p</i> =0.999
<i>n</i>			121