

Supplementary Table S3

Hospital-acquired infections, according to the ECDC case-source definition

Case source COVID-19	Total number (% of total HAI)	Number sequenced (% of category)	% belonging to a genome cluster
<i>Indeterminate HAI</i>	93 (34.6)	36 (38.7)	70.3
<i>Probable HAI</i>	86 (32.0)	40 (46.5)	87.8
<i>Definite HAI</i>	90 (33.5)	47 (52.2)	90.0