

**S2 Appendix Spline model.** The log-log transformation of the survival function is modelled as a natural cubic spline function of  $\log(t)$  [13, 14].

$$\log(-\log(S(t))) = \gamma_0 + \gamma_1 \log(t) + \gamma_2 v_1(\log(t)) + \dots + \gamma_{m+1} v_m(\log(t))$$

where  $v_j(x)$  is the  $j$ -th basis function

$$v_j(x) = (x - k_j)_+^3 - \lambda_j (x - k_{min})_+^3 - (1 - \lambda_j) (x - k_{max})_+^3,$$

where  $\lambda_j = (k_{max} - k_j)/(k_{max} - k_{min})$  and  $(x - a)_+ = \max(0, x - a)$ .

Two separate spline models with 4 internal knots are fitted for the two cause-specific hazards for recovery and death based on the complete data of all confirmed and hospitalized cases. The location of the knots and the values of  $\gamma_j$  are given in Table A.

Table A: **Spline models.**

Model	Parameter	Index j					
		1	2	3	4	5	6
1	$k_j$	-1.39	2.57	2.77	3.00	3.22	5.38
	$\gamma_j$	-7.85	0.67	-1.94	-6.47	16.67	-8.27
2	$k_j$	-1.39	1.79	2.08	2.40	2.77	5.57
	$\gamma_j$	-5.52	0.97	-1.56	0.34	2.81	-1.63

Knots ( $k_j$ ) on the log-scale and coefficients ( $\gamma_j$ ) of the two spline models ( $j = 1, \dots, 6$ ).