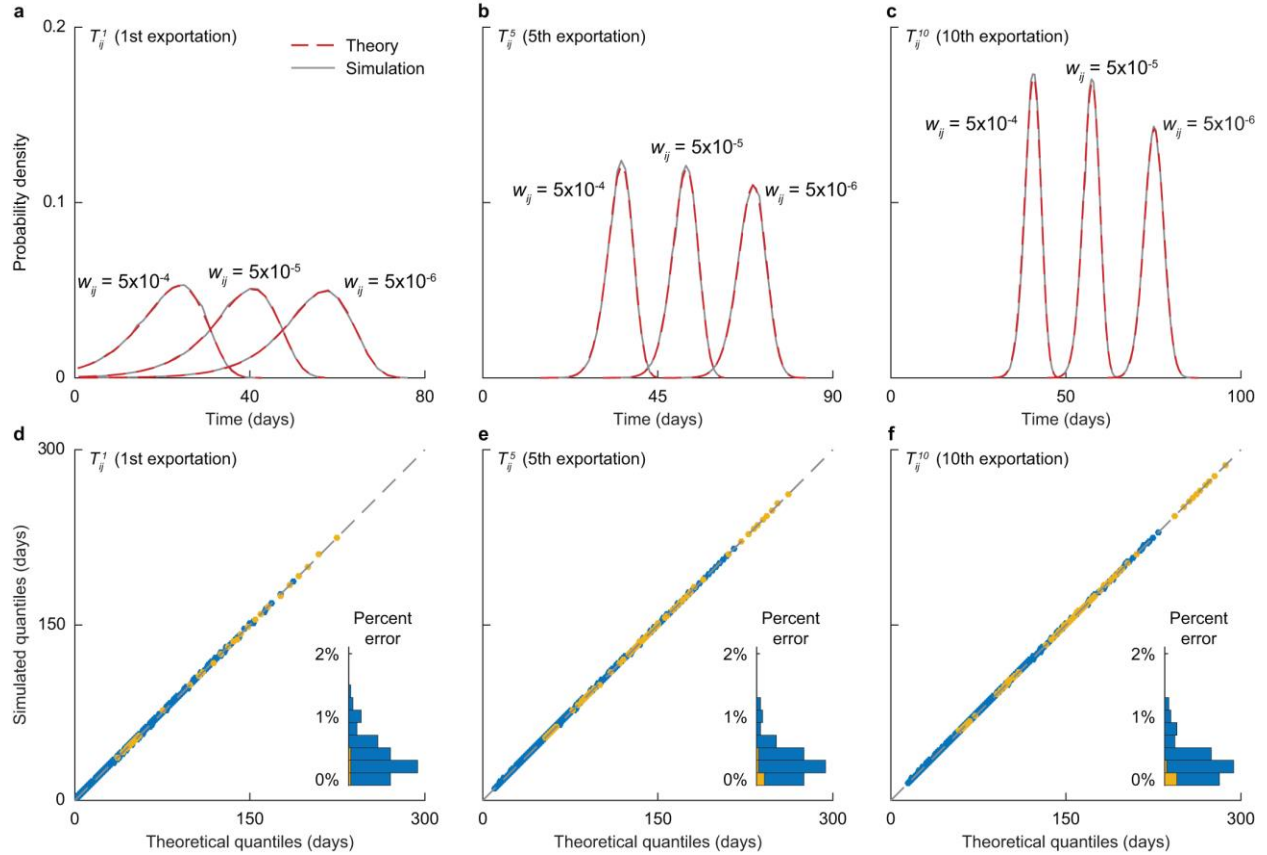
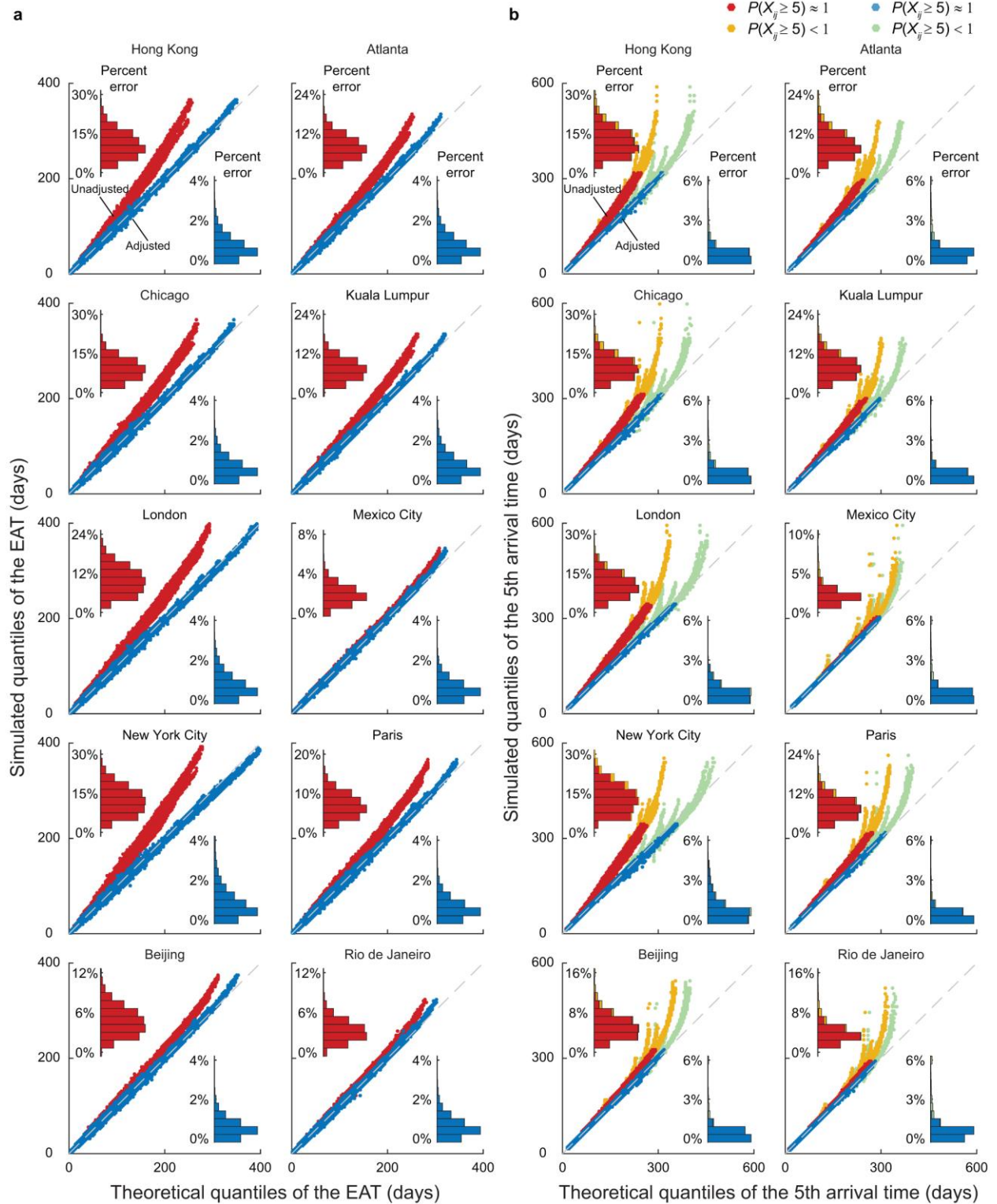


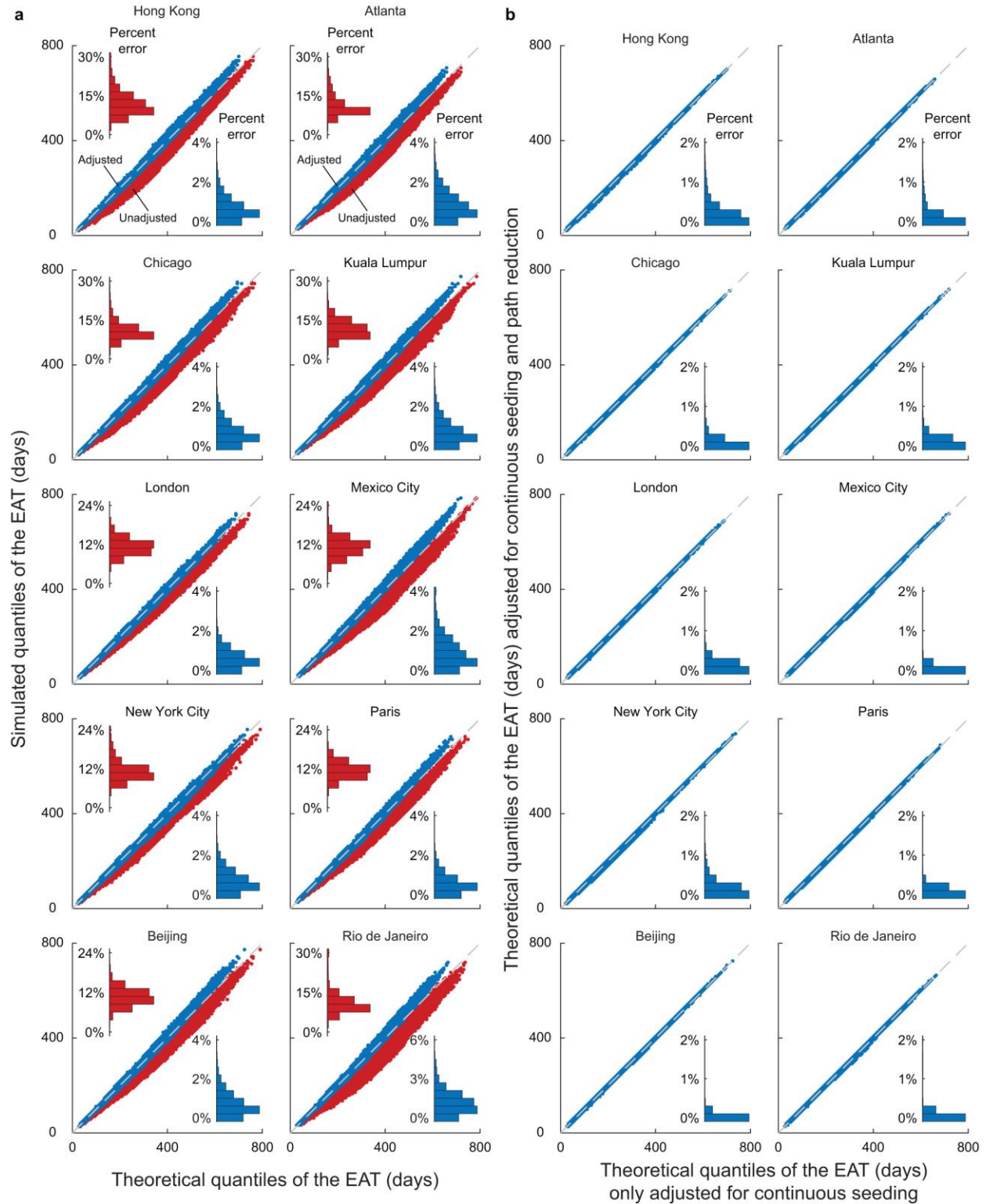
**Supplementary Figure 1. Mobility rates in the WAN. a,** Histograms of outbound mobility rates  $w_{jk}$  in the WAN stratified by population size. **b,** Histograms of total outbound mobility rates  $W_j = \sum_k w_{jk}$  in the WAN stratified by population size.



**Supplementary Figure 2. Validating Assumption 1 in the two-population model.** This figure is analogous to Fig. 1 but without the imposition of Assumption 2. That is, the analytical pdf of  $T_{ij}^n$  here is given by equation (S1). **a-c**, The analytical (red dashed lines) and simulated (grey lines) pdf of  $T_{ij}^1$ ,  $T_{ij}^5$  and  $T_{ij}^{10}$  for the same scenario of exemplary influenza pandemic as in Fig. 1**a-c**. **d-f**, Q-Q plots for the analytical and simulated quantiles of  $T_{ij}^1$ ,  $T_{ij}^5$  and  $T_{ij}^{10}$  across the same 100 scenarios (days) considered in Fig. 1**d-f**. In the Q-Q plots, deviations from the diagonal indicate discrepancies between the analytical and simulated quantiles. Q-Q plots are colored in blue if the number of exportations is  $n$  or above with probability 1 and yellow otherwise. Insets show the corresponding histograms of percent error in  $E[T_{ij}^n]$ .

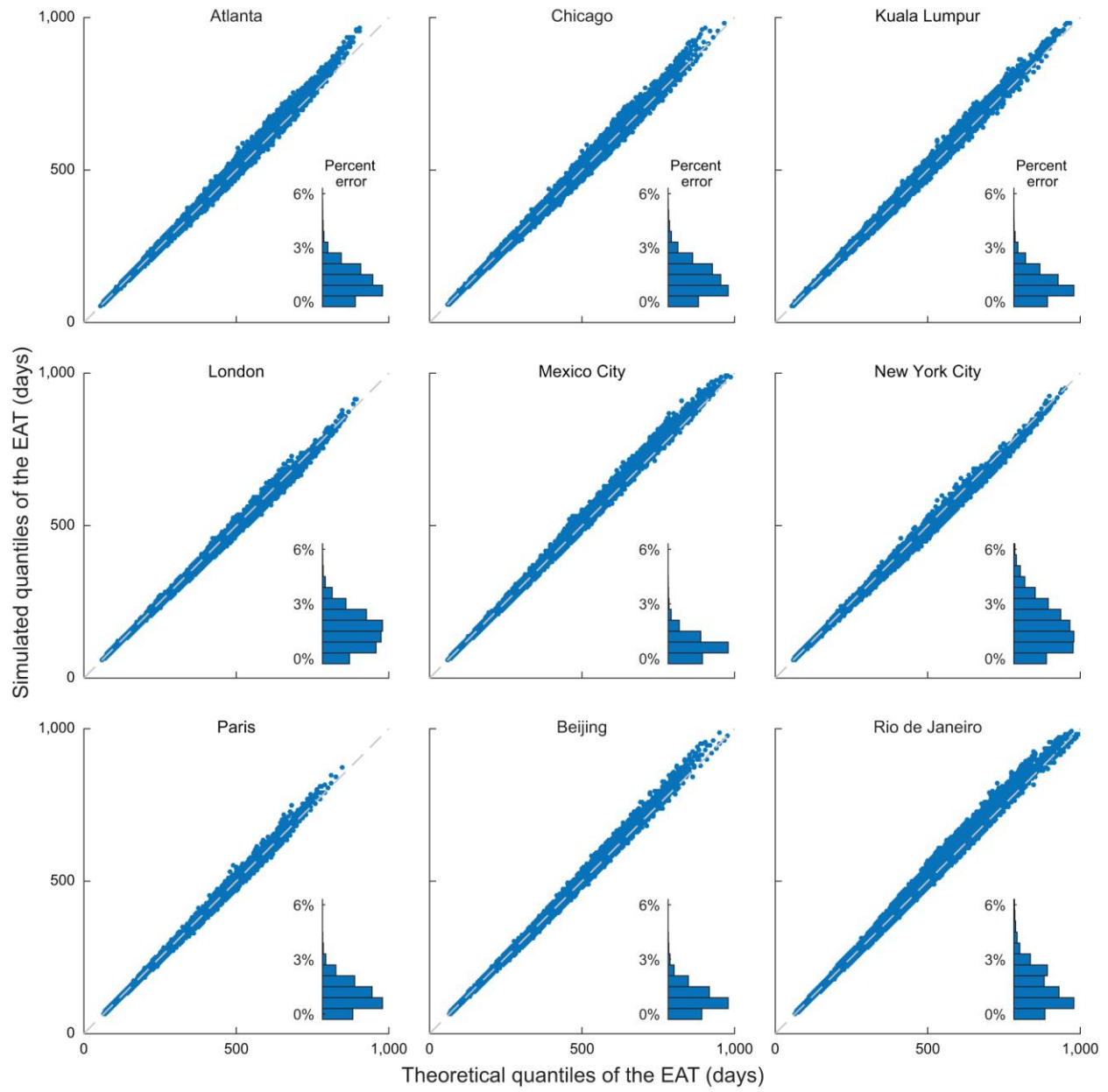


**Supplementary Figure 3. Hub-effect adjustments for  $D_{i,1}$  populations.** **a**, Analogous to Fig. 2d with different major hubs in the WAN as the epidemic origin. **b**, Same as panel **a** but for the 5<sup>th</sup> arrival time  $T_{ij}^5$ .

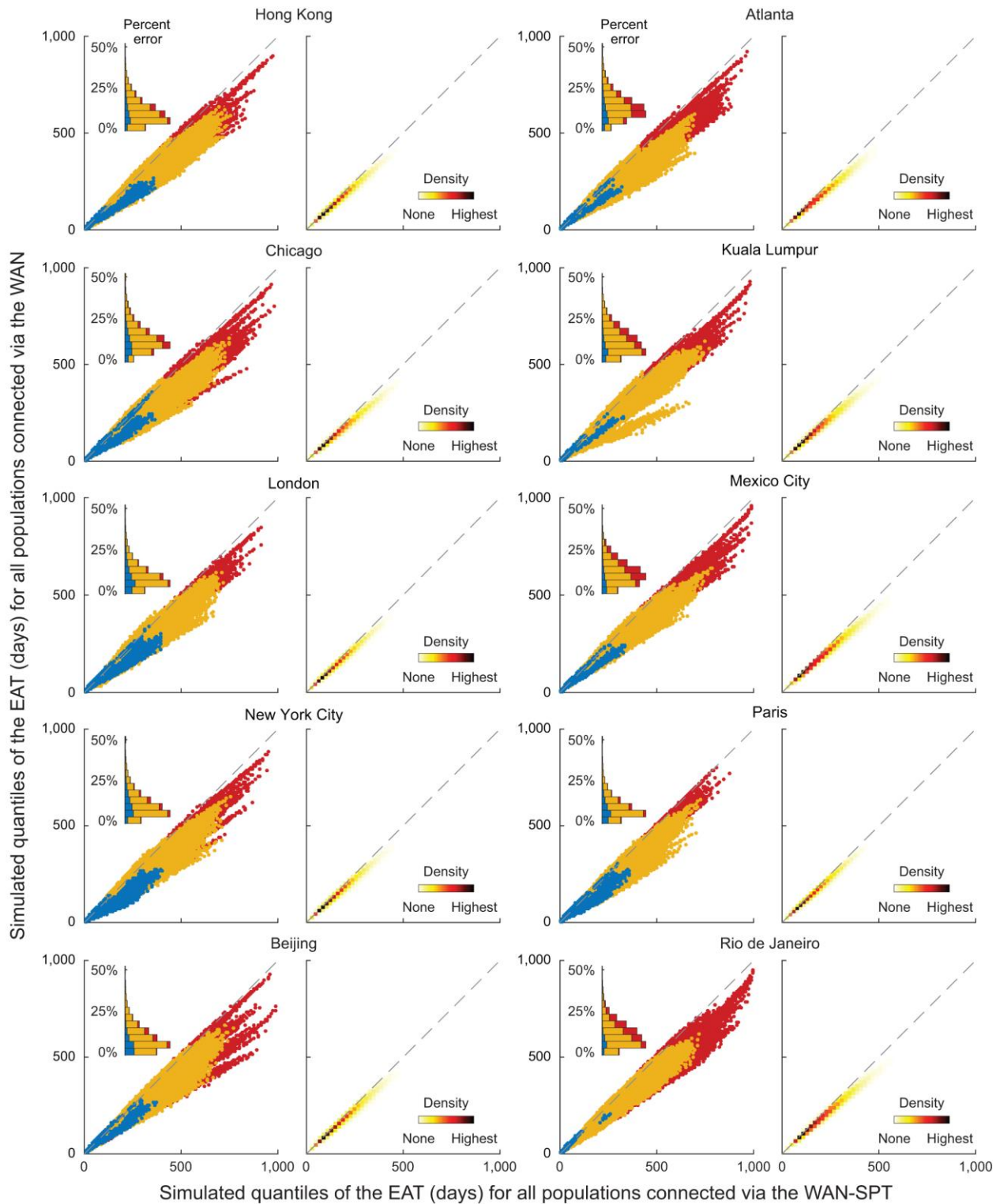


**Supplementary Figure 4. Analogous to Fig. 2e for  $D_{i,2}$  populations with different major hubs in the WAN as the epidemic origin. a, Q-Q plots for the analytical and simulated quantiles of EATs before (red) and after (blue) adjusting for continuous seeding. b, Q-Q plots for**

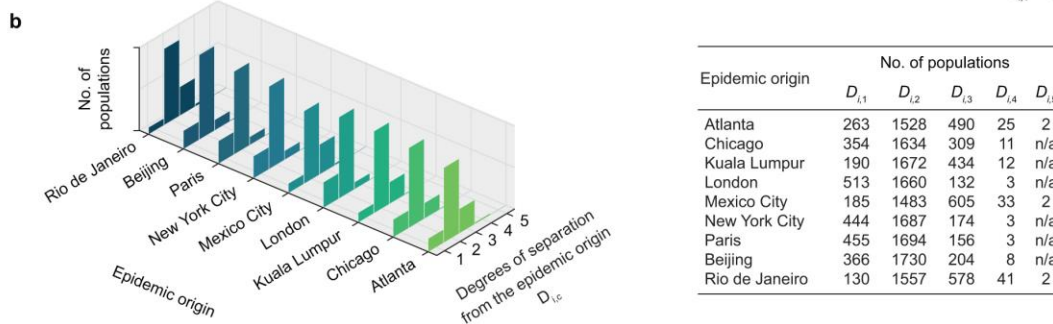
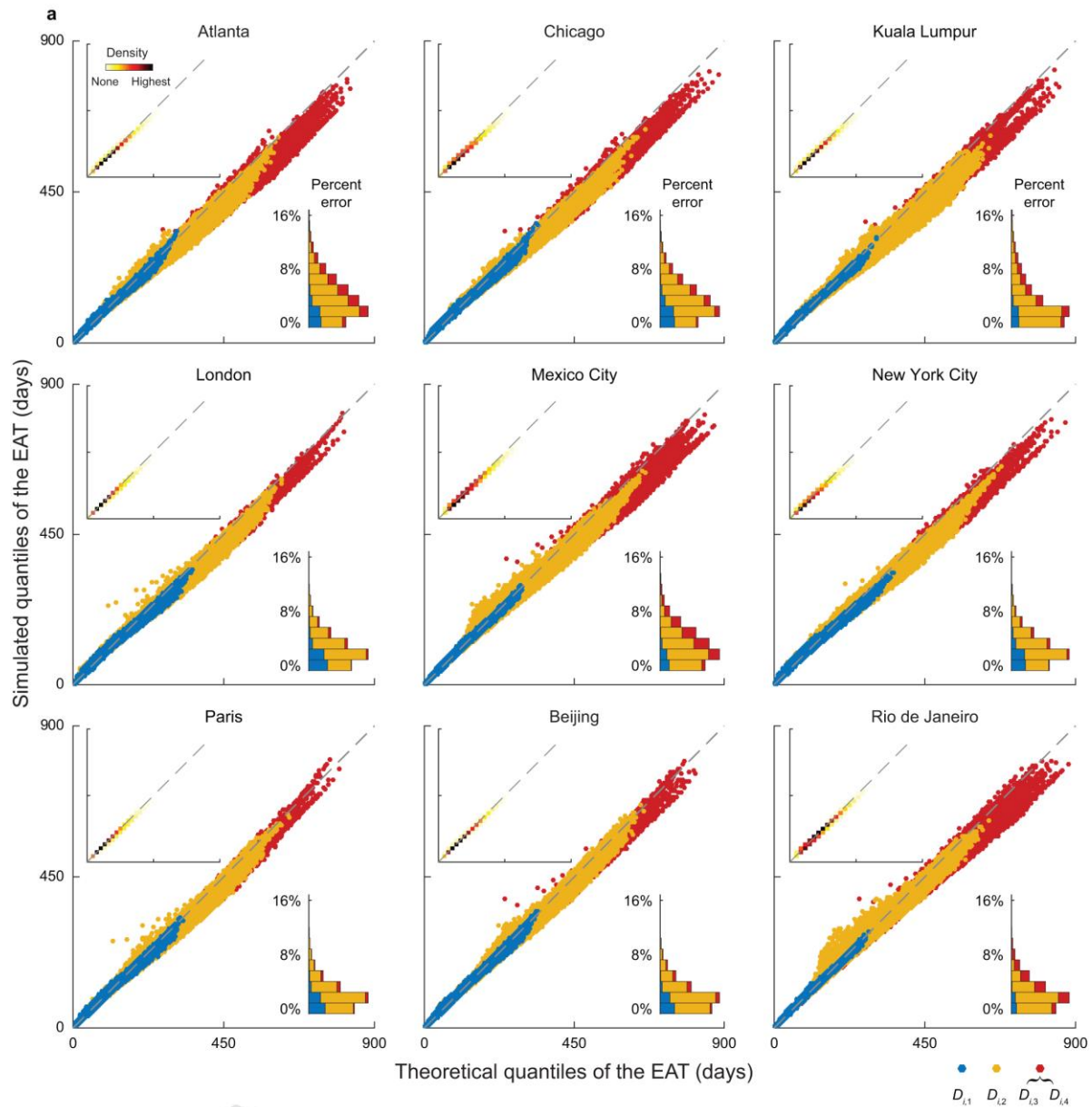
the analytical quantiles of EATs with and without path reduction after adjusting for continuous seeding.



**Supplementary Figure 5. Analogous to Fig. 2f for  $D_{i,3}$  and  $D_{i,4}$  (and  $D_{i,5}$  if any) populations with different major hubs in the WAN as the epidemic origin.**

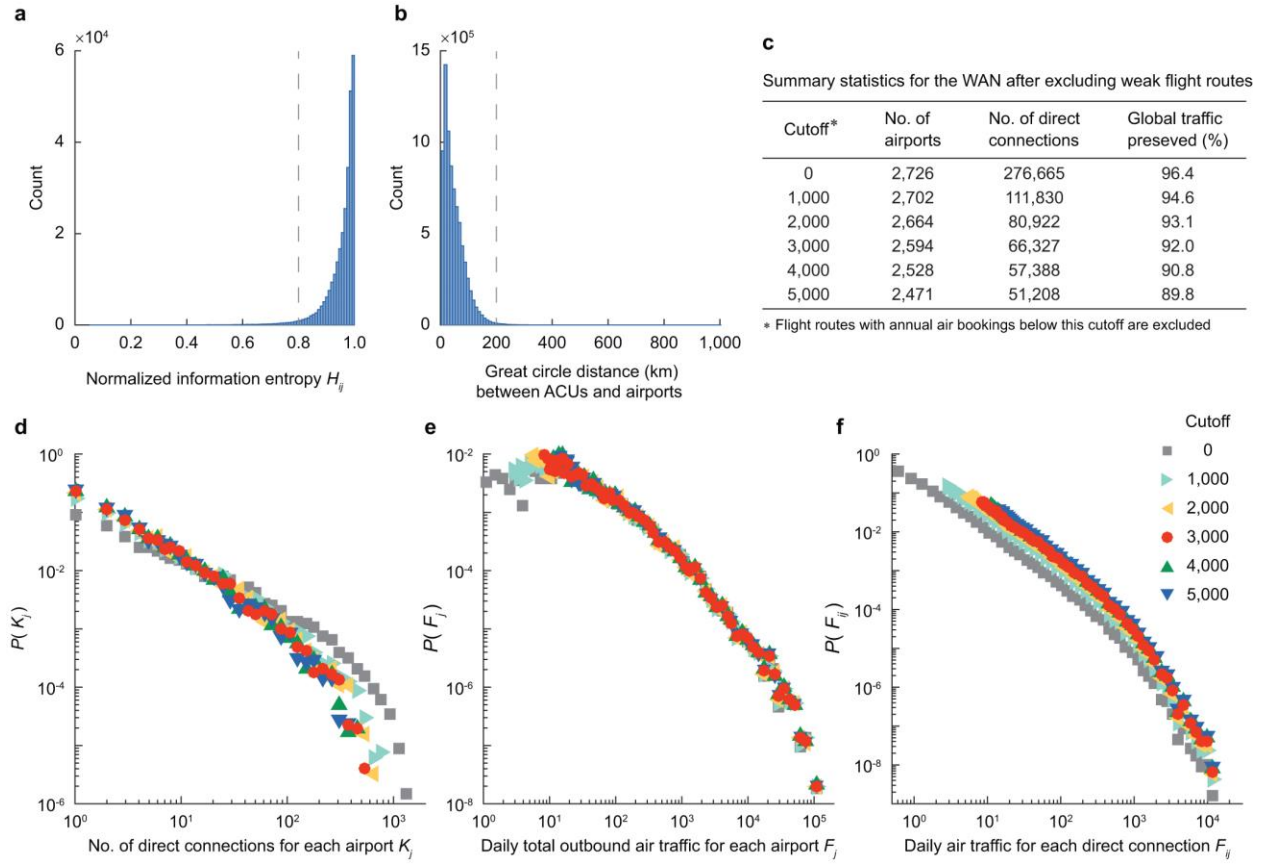


**Supplementary Figure 6. Q-Q plots for the simulated EATs in the WAN-SPT and WAN and the corresponding density plots with different major hubs in the WAN as the epidemic origin.**

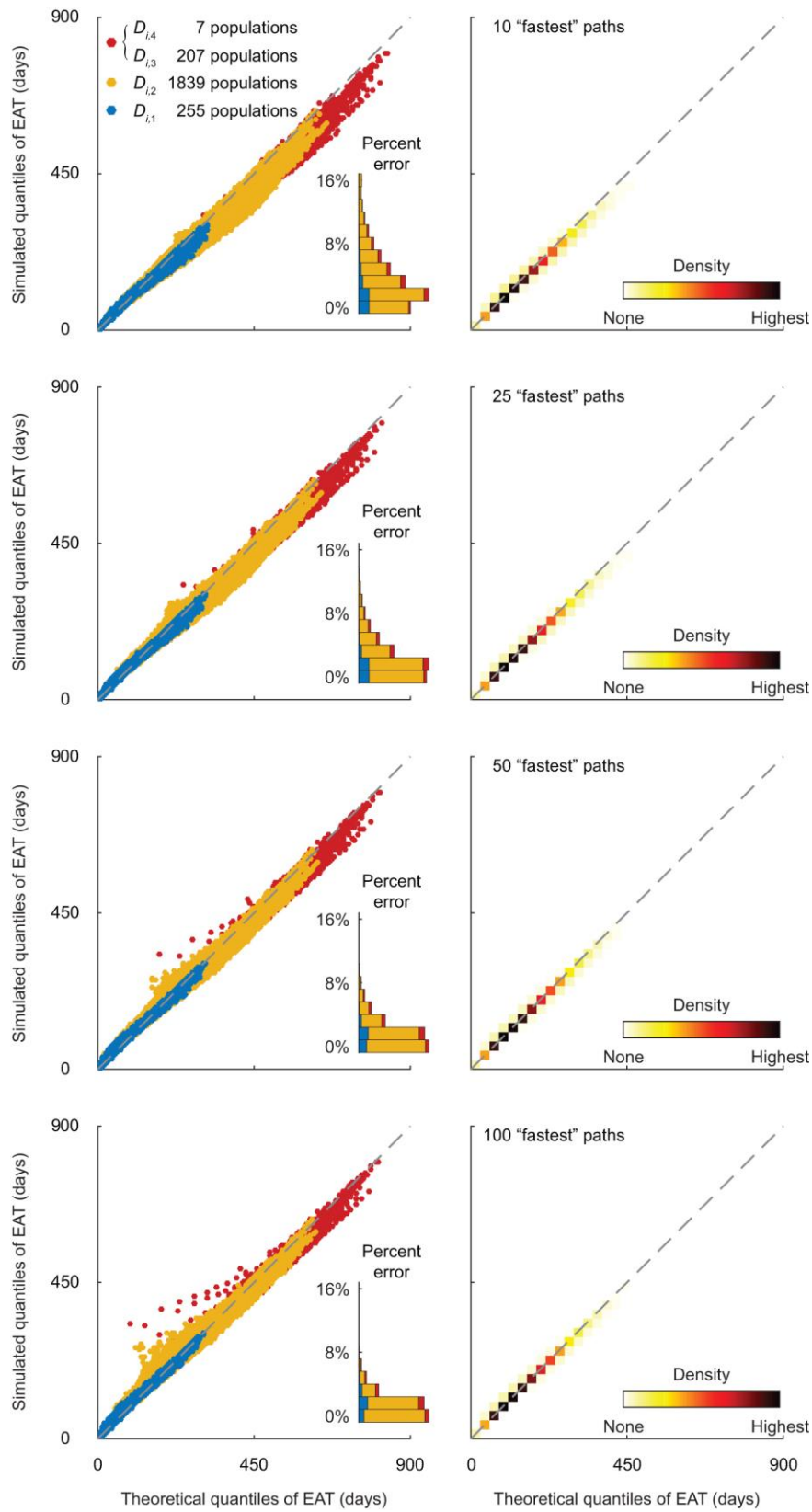


**Supplementary Figure 7. Analogous to Fig. 3 with other major hubs in the WAN as the epidemic origin.**

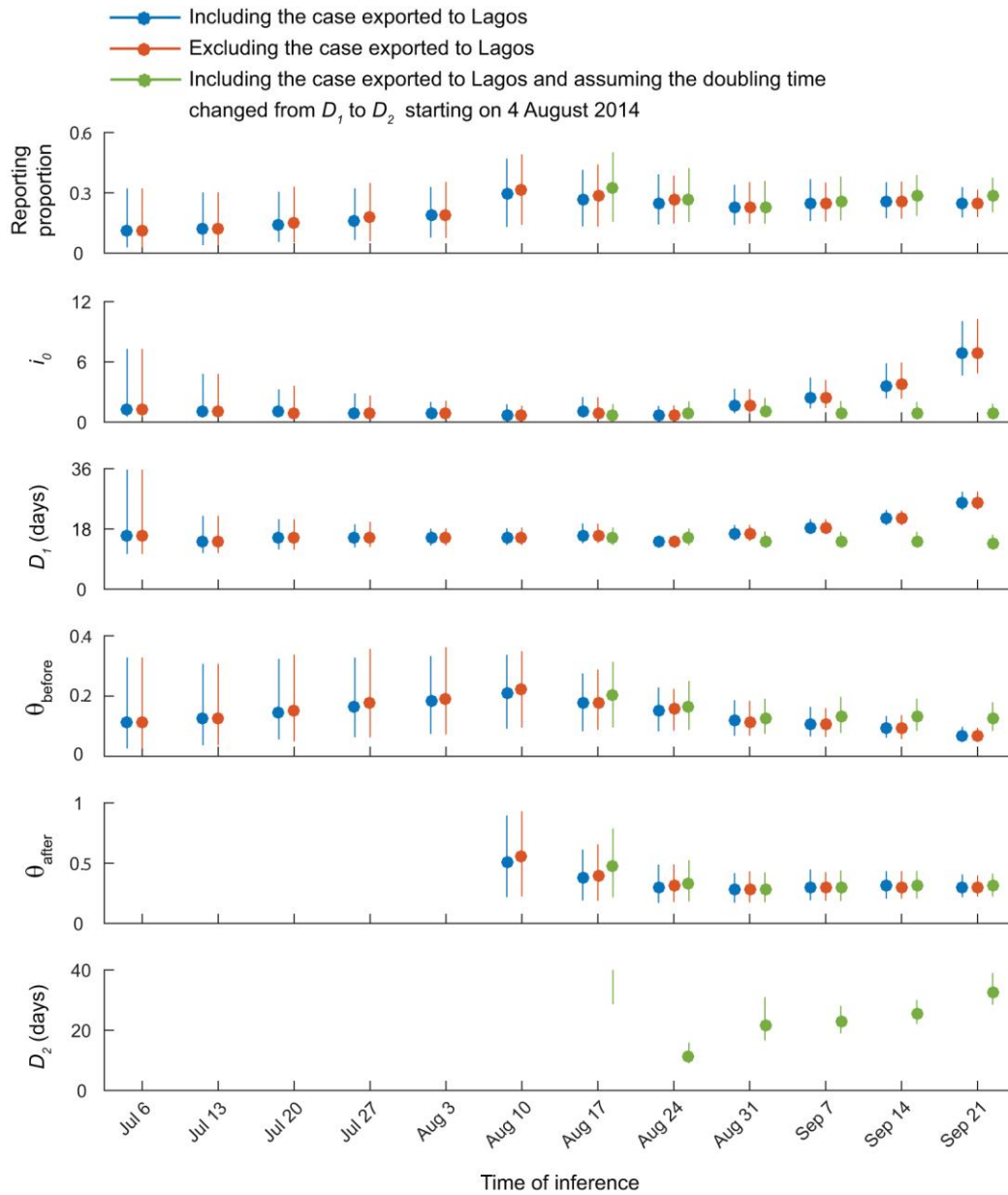




**Supplementary Figure 8. Building the global epidemic simulator.** **a**, Histogram of the seasonality measure  $H_{ij}$  (based on normalized information entropy) calculated with the raw data in the OAG flight booking dataset. **b**, Distribution of great circle distance for all combinations of ACUs in the original GPWv4 dataset and airports in the OAG dataset after flight routes with  $H_{ij} < 0.8$  have been excluded. **c**, Summary network statistics with different cutoff for eliminating flight routes with weak traffic. **d–f**, Distributions of key network statistics with different cutoff for eliminating flight routes with weak traffic.



**Supplementary Figure 9. Superposition of paths in the WAN.** Analogous to Fig. 3 with Hong Kong as the epidemic origin, this figure shows the effect of increasing the number of "fastest" paths for superposition on estimating the epidemic arrival times in the WAN. From top to bottom, the epidemic arrival time for each population in the WAN is computed with the superposition of the 10, 25, 50, and 100 "fastest" paths, respectively.



**Supplementary Figure 10. Parameter estimates in the case study of the 2014 West African Ebola epidemic in Montserrado and Margibi, Liberia.** Circles and bars indicate the medians and 95% credible intervals of the posterior distributions. Under the assumption that the doubling time changed from  $D_1$  to  $D_2$  starting on 4 August 2014,  $D_2$  is statistically non-identifiable or unreliable until the week of 18-24 August 2014.

**Supplementary Table 1. Raw data used in the case study of the 2009 influenza A/H1N1 pandemic in greater Mexico City.**

Country*	Onset of symptoms <sup>†</sup>	Flight arrival <sup>†</sup>	Confirmation <sup>†</sup>	Air traffic during March-April 2009 <sup>‡</sup>
US	28-Mar-09	n/a	21-Apr-09	545,368
Canada	11-Apr-09	8-Apr-09	23-Apr-09	52,702
El Salvador	n/a	19-Apr-09	3-May-09	15,090
UK	24-Apr-09	21-Apr-09	27-Apr-09	8,970
Spain	25-Apr-09	22-Apr-09	27-Apr-09	52,390
Cuba	n/a	25-Apr-09	13-May-09	23,100
Costa Rica	25-Apr-09	25-Apr-09	2-May-09	16,950
Netherlands	n/a	27-Apr-09	30-Apr-09	16,680
Germany	28-Apr-09	n/a	29-Apr-05	21,350
France	n/a	n/a	1-May-09	39,690
Guatemala	1-May-09	n/a	5-May-09	36,758
Colombia	n/a	n/a	3-May-09	24,535

\* The 12 countries directly seeded by Mexico during 2009 influenza A/H1N1 pandemic <sup>1</sup>.

<sup>†</sup> The epidemic arrival times for these 12 countries are based on the Supplementary Table 5 in Balcan et al. <sup>1</sup>. Given the first confirmed case imported in each country, its epidemic arrival time (EAT) is assumed to be (i) the onset date of symptoms or the date of flight arrival (whichever was earlier); or (ii) left-censored at the date of confirmation if neither the onset date nor the flight arrival date was reported.

<sup>‡</sup> Number of air passengers departing from airports in greater Mexico City during March-April 2009, based on the Supplementary Table S3 in Fraser et al. <sup>2</sup>.

**Supplementary Table 2. Raw data used in the case study of the 2014 West African Ebola epidemic in Montserrado and Margibi, Liberia.**

Month	No. of monthly total outbound international flight bookings*	Total outbound mobility rate in each month <sup>†</sup>
May	12,146	$2.142 \times 10^{-4}$
Jun	12,760	$2.326 \times 10^{-4}$
Jul	13,017	$2.296 \times 10^{-4}$
Aug	12,417	$2.190 \times 10^{-4}$
Sep	5,939	$1.082 \times 10^{-4}$

\* Monthly international air traffic departing from airports in Montserrado and Margibi, Liberia, between May and September 2014 is obtained from the OAG database. Flights to Guinea and Sierra Leone are excluded from analysis.

<sup>†</sup> The total population size of Montserrado and Margibi in Liberia is 1,828,846 in year 2015 according to the GPWv4 database.

**Supplementary References**

- 1 Balcan, D. *et al.* Seasonal transmission potential and activity peaks of the new influenza A(H1N1): a Monte Carlo likelihood analysis based on human mobility. *BMC Med.* **7**, 45, doi:10.1186/1741-7015-7-45 (2009).
- 2 Fraser, C. *et al.* Pandemic potential of a strain of Influenza A (H1N1): Early findings. *Science* **324**, 1557-1561, doi:10.1126/science.1176062 (2009).