

Figure S1:

Number of confirmed and suspected cases of Ebola up to 30 September 2014 in districts of Guinea (black), Liberia (red) and Sierra Leone (green), as a function of district population size.

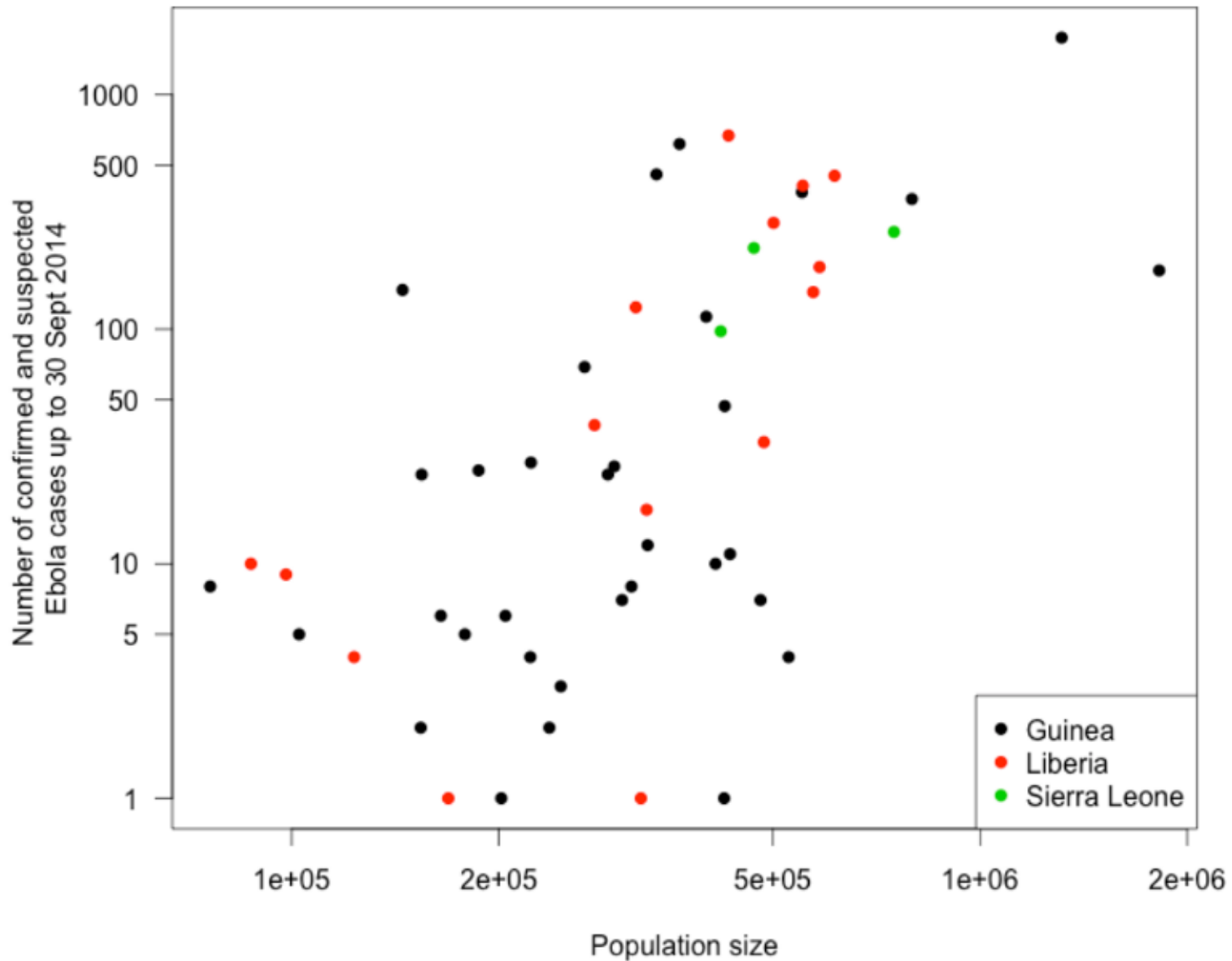


Figure S2:

Micro-scale fit of competing models. Predicted day of infection for infected administrative units based on each of the models (excluding best fit model, shown in Fig 2). The predicted day of infection for nodes based on simulations from April 24th. For each node, the 95% prediction interval is in light gray with the interquartile range specified as dark gray and the median infection as a break. Dots represent the observed day of infection for each node. Infection locations used in fitting are to the left of the dotted line and infection subsequent to October 1st are to the right of the dotted line.

Figure S3:

Coverage vs. correlation of unit-infection dates. The correlation between the actual infection date and simulated median date of infection for a representative subset of compared models is plotted versus the coverage for the same model. Coverage is defined as percent of actual infection dates occurring in the 95% prediction intervals (Fig S2). The best-fit and other gravity models have the best combination of accuracy and precision by these measures.

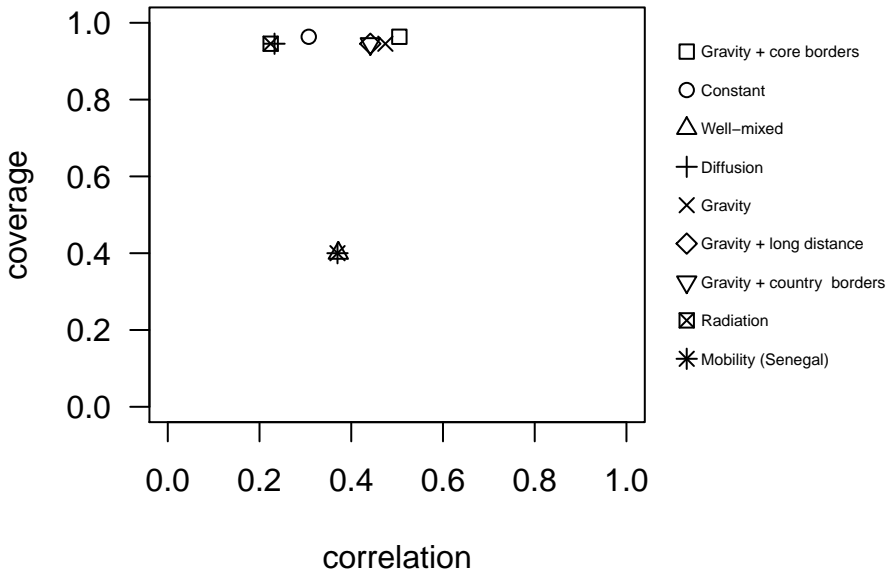
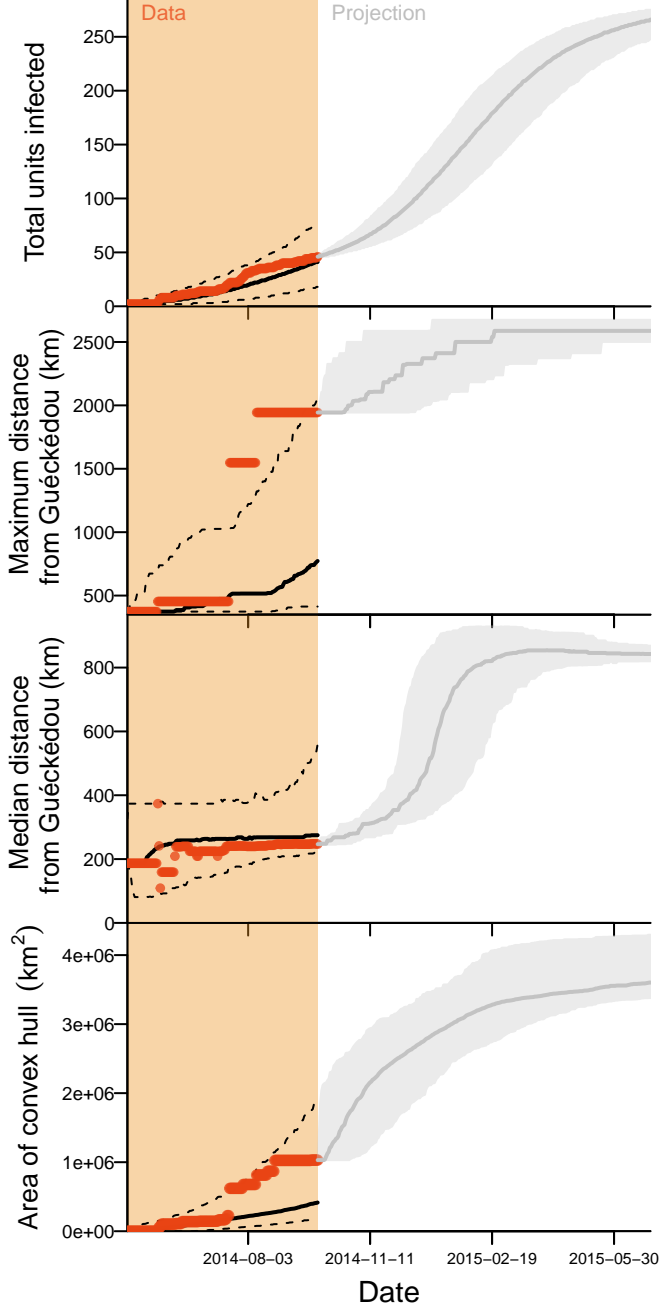


Figure S4:

Macro-scale spread statistics of gravity + core border model. Orange shading represents simulated spread from April 24th to Oct 1st, 2014 and unshaded represents forecasts obtained by forward simulations until June 30th 2015. Simulations forward were based on epidemic transmission remaining unchanged from the fitting period and therefore does not account for the realized success of control. Dots are observed data, solid lines are median values, and dotted lines/shaded areas represent 95% prediction intervals. a) The number of newly infected administrative units is consistent with observation and continues to accelerate if spread continues uncontrolled. The maximum b) and median c) distance from Guéckédou; the maximum distance increased faster than predicted due to early spread to Nigeria. d) The convex hull of infection area is the total area within the minimum convex polygon of infected unit centroids.



Gravity + core borders

Figure S5:

Predicted day of infection for infected administrative units based on the best fit model. Predicted intervals for the counties without observed infections are included for assessment of lack of spread. The predicted day of infection for nodes is based on simulations from April 24th. For each node, the 95% prediction interval is in light blue with the interquartile range specified as dark blue and the median infection as a break. Dots represent the observed day of infection for each node colored by country. Infection locations used in fitting are to the left of the dotted line and infection subsequent to Oct 1 are to the right of the first dotted line. After the second dotted line are the counties in the dataset that were not infected. Most have median infection dates much later than predicted for the infected counties and the actual infections also fall below the interquartile range of these counties. Given that interventions and changes in behavior began to slow the outbreak in late 2015 it is unreasonable to expect many additional infections after this point. However, it is interesting that much of the remainder of uninfected West African locations have similar, wide ranges of the predicted date of infection, this illustrates the importance of stochasticity in determining the realized path of spread. We have further illustrated the difference between core-countries and the rest of the region by coloring core-countries with a lighter color.

Supplementary Information Table 1: Infection data and sources.

The following details the sources of information used to determine the initial dates of infection for each administrative unit used in model fitting. Information was accumulated and updated as the epidemic progressed and dates used for fitting were finalized Oct 10th, 2014. Recently WHO has released new data showing confirmed infections in counties earlier (and in a few cases later) than the data below. These new data also differed in being aggregated weekly. We used these recent patient databases to fit the same models to weekly data and found qualitatively similar results with the same best fit model and the same support for gravity models overall with AIC.

Country	County	Date	Official Confirmation	Additional Source
Sierra Leone	Kailahun	5/24/2014	WHO Outbreak News ¹ , WHO Essay ²	
Sierra Leone	Kenema	6/17/2014	24 June GoSL SitRep ³	Media ⁴
Sierra Leone	Kono	7/29/2014	NGO (Wellbody Alliance) ⁵	
Sierra Leone	Bombali	7/20/2014	30 July IFRC-SL SitRep ⁶	Media ^{7,8}
Sierra Leone	Kambia	6/5/2014	6 June GoSL SitRep ⁹	
Sierra Leone	Port Loko	6/6/2014	11 June GoSL SitRep ¹⁰	Media ¹¹
Sierra Leone	Tonkolili	8/3/2014	4 August GoSL SitRep ¹²	
Sierra Leone	Bo	6/23/2014	24 June GoSL SitRep ³	
Sierra Leone	Bonthe	7/17/2014	20 July US Embassy ¹³	Blog (text of SL Facebook post) ¹⁴
Sierra Leone	Moyamba	7/29/2014	4 August GoSL SitRep ¹²	Media ¹⁵
Sierra Leone	Pujehun	8/1/2014	4 August GoSL SitRep ¹²	Media ¹⁶
Sierra Leone	Western Rural	8/7/2014	8 August GoSL SitRep ¹⁷	
Sierra Leone	Western Urban	7/11/2014	20 July US Embassy ¹³	Media ¹⁸
Guinea	Boffa	5/22/2014	6 June GoG SitRep ¹⁹	
Guinea	Conakry	4/26/2014		
Guinea	Kissidougou	9/14/2014	16 September GoG SitRep ²⁰	
Guinea	Krouan	8/31/2014	31 August GoG SitRep ²¹	
Guinea	Kouroussa	5/25/2014	5 June GoG SitRep ¹⁹	
Guinea	Siguir	7/18/2014	22 July GoG SitRep ²²	
Guinea	Coyah	9/1/2014	2 September GoG SitRep ²³	
Guinea	Dubrka	8/13/2014	14 August GoG SitRep ²¹	
Guinea	Forcariah	8/27/2014	29 August GoG SitRep ²⁴	
Guinea	Kindia	9/20/2014	20 September GoG SitRep ²⁵	
Guinea	Tliml	5/23/2014	WHO Outbreak News ²⁶ , 27 May GoG SitRep ²⁷	
Guinea	Dalaba	9/11/2014	12 September GoG SitRep ²⁸	
Guinea	Pita	7/27/2014	4 August GoG SitRep ²⁹	Media ³⁰
Guinea	Beyla	9/20/2014	25 September GoG SitRep ³¹	
Guinea	Guckdou	4/26/2014		
Guinea	Lola	9/30/2014	1 October GoG SitRep ³²	
Guinea	Macenta	5/23/2014	WHO Outbreak News ²⁶ , 27 May GoG SitRep ²⁷	
Guinea	Nzrkor	8/2/2014	4 August GoG SitRep ²⁹	
Guinea	Yamou	8/7/2014	7 August GoG SitRep ³³	
Liberia	Nimba	7/17/2014	20 July GoL SitRep ³⁴ , 16 July UNICEF SitRep ³⁵	
Liberia	River Cess	7/31/2014	31 July GoL SitRep ³⁶ , 15 August UNICEF SitRep ³⁷	
Liberia	River Gee	9/25/2014	25 September GoL SitRep ³⁸	
Liberia	Sinoe	8/20/2014	20 August GoL SitRep ³⁹	
Liberia	Bomi	7/10/2014	10 July GoL SitRep	

Liberia	Bong	7/15/2014	16 July UNICEF SitRep ³⁵	
Liberia	Grand Cape Mount	7/30/2014	31 July GoL SitRep ³⁶ , 2 August GoL SitRep ⁴⁰	
Liberia	Grand Bassa	8/2/2014	2 August GoL SitRep ⁴⁰ Probable, WHO Patient-level: there are earlier cases ⁴¹ , GoL Ministry of Information ⁴²	Media ^{43,44}
Liberia	Lofa	5/23/2014	6 June GoL SitRep ⁴⁵	
Liberia	Margibi	6/25/2014	25 June GoL SitRep ⁴⁶	
Liberia	Montserrado	6/11/2014	11 June GoL SitRep ⁴⁷ , 12 June GoL SitRep ⁴⁸	
Nigeria	Lagos	7/20/2014	WHO Outbreak News ⁴⁹	
Nigeria	Rivers	8/11/2014	WHO Situation Assessment ⁵⁰	
Senegal	Dakar	8/26/2014	WHO Outbreak News ⁵¹	

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Table S2: Model comparison

Model name	Infection probability	NLL	AIC
Constant	β_0 (no network)	346.1	348
Well-mixed	β_0	350.7	353
Diffusion	$\frac{1}{1 + e^{\beta_0 + \beta_1 d_{ij}}}$	281.9	353
Diffusion + long distance	$\frac{1}{1 + e^{\beta_0 + \beta_1 d_{ij}^{\beta_2}}}$	275.1	281
Gravity	$\frac{1}{1 + e^{\beta_0 + \beta_1 \frac{d_{ij}}{(p_i p_j)^{\beta_2}}}}$	273.2	279
Gravity + long distance	$\frac{1}{1 + e^{\beta_0 + \beta_1 \frac{d_{ij}^{\beta_2}}{(p_i p_j)^{\beta_3}}}}$	272.2	280
Mobility (Senegal)	$\frac{1}{1 + e^{\beta_0 + \beta_1 \text{mob}}}$	350.4	354
Mobility (Senegal) + diffusion	$\frac{1}{1 + e^{\beta_0 + \beta_1 \text{mob} + \beta_2 d_{ij}}}$	281.8	288
Diffusion + country borders	$\begin{cases} \beta_2 \frac{1}{1 + e^{\beta_0 + \beta_1 d_{ij}}} & \text{if crossing border} \\ \frac{1}{1 + e^{\beta_0 + \beta_1 d_{ij}}} & \text{if not crossing border} \end{cases}$	280.0	286
Diffusion + core borders	$\begin{cases} \beta_2 \frac{1}{1 + e^{\beta_0 + \beta_1 d_{ij}}} & \text{if crossing core border} \\ \frac{1}{1 + e^{\beta_0 + \beta_1 d_{ij}}} & \text{if not crossing core border} \end{cases}$	274.9	281
Gravity + country borders	$\begin{cases} \beta_3 \frac{1}{1 + e^{\beta_0 + \beta_1 \frac{d_{ij}}{(p_i p_j)^{\beta_2}}}} & \text{if crossing border} \\ \frac{1}{1 + e^{\beta_0 + \beta_1 \frac{d_{ij}}{(p_i p_j)^{\beta_2}}}} & \text{if not crossing border} \end{cases}$	272.5	281
Gravity + core borders	$\begin{cases} \beta_3 \frac{1}{1 + e^{\beta_0 + \beta_1 \frac{d_{ij}}{(p_i p_j)^{\beta_2}}}} & \text{if crossing core border} \\ \frac{1}{1 + e^{\beta_0 + \beta_1 \frac{d_{ij}}{(p_i p_j)^{\beta_2}}}} & \text{if not crossing core border} \end{cases}$	270.0	278
Radiation	$\frac{1}{1 + e^{\beta_0 + \beta_1 d_{ij} + \beta_2 \text{rad}}}$	281.8	288

Population weighted opportunity	$\frac{1}{1 + e^{\beta_0 + \beta_1 d_{ij} + \beta_2 \text{pwo}}}$	281.9	288
Force of infection (unnormalized)	$1 - e^{\left(\beta_0 p_i^{\beta_1} \sum_j \frac{p_j^{\beta_2}}{d_{ij}^{\beta_3}} \right)}$ (no network)	275.4	283
Force of infection (normalized)	$1 - e^{\left(\frac{\beta_0 p_i^{\beta_1} \sum_j \frac{p_j^{\beta_2}}{d_{ij}^{\beta_3}}}{\left(\sum_{k \neq i} \frac{p_k^{\beta_2}}{d_{ik}^{\beta_3}} \right)^{\beta_4}} \right)}$ (no network)	274.0	284

Table S3: Estimated parameter values

Model name	Infection probability	β_0	β_1	β_2	β_3	β_4
Constant	β_0 (no network)	0.0010				
Well-mixed	β_0	0.00005				
Diffusion	$\frac{1}{1 + e^{\beta_0 + \beta_1 d_{ij}}}$	5.630	1.032			
Diffusion + long distance	$\frac{1}{1 + e^{\beta_0 + \beta_1 d_{ij}^{\beta_2}}}$	0.876	5.748	0.341		
Gravity	$\frac{1}{1 + e^{\beta_0 + \beta_1 \frac{d_{ij}}{(p_i p_j)^{\beta_2}}}}$	5.246	155.8	0.186		
Gravity + long distance	$\frac{1}{1 + e^{\beta_0 + \beta_1 \frac{d_{ij}^{\beta_2}}{(p_i p_j)^{\beta_3}}}}$	3.659	22.63	0.593	0.080	
Mobility (Senegal)	$\frac{1}{1 + e^{\beta_0 + \beta_1 \text{mob}}}$	9.886	0.000007			
Mobility (Senegal) + diffusion	$\frac{1}{1 + e^{\beta_0 + \beta_1 \text{mob} + \beta_2 d_{ij}}}$	5.594	0.000003	1.046		
Diffusion + country borders	$\beta_2 \frac{1}{1 + e^{\beta_0 + \beta_1 d_{ij}}}$ if crossing border	5.561	0.806	0.257		
	$\frac{1}{1 + e^{\beta_0 + \beta_1 d_{ij}}}$ if not crossing border					
Diffusion + core borders	$\beta_2 \frac{1}{1 + e^{\beta_0 + \beta_1 d_{ij}}}$ if crossing core border	6.484	0.510	0.070		
	$\frac{1}{1 + e^{\beta_0 + \beta_1 d_{ij}}}$ if not crossing core border					

Model name	Infection probability	β_0	β_1	β_2	β_3	β_4
Gravity + country borders	$\begin{cases} \frac{\beta_3}{1+e} \frac{1}{\beta_0+\beta_1 \frac{d_{ij}}{(p_i p_j)^{\beta_2}}} & \text{if crossing border} \\ \frac{1}{1+e} \frac{1}{\beta_0+\beta_1 \frac{d_{ij}}{(p_i p_j)^{\beta_2}}} & \text{if not crossing border} \end{cases}$	5.166	157.1	0.189	0.507	
Gravity + core borders	$\begin{cases} \frac{\beta_3}{1+e} \frac{1}{\beta_0+\beta_1 \frac{d_{ij}}{(p_i p_j)^{\beta_2}}} & \text{if crossing core border} \\ \frac{1}{1+e} \frac{1}{\beta_0+\beta_1 \frac{d_{ij}}{(p_i p_j)^{\beta_2}}} & \text{if not crossing core border} \end{cases}$	5.792	105.7	0.186	0.150	
Radiation	$\frac{1}{1+e^{\beta_0+\beta_1 d_{ij}+\beta_2 \text{rad}}}$	5.588	1.046	0.000002		
Population weighted opportunity	$\frac{1}{1+e^{\beta_0+\beta_1 d_{ij}+\beta_2 \text{pwo}}}$	5.606	1.041	0.0000006		
Force of infection (unnormalized)	$1 - e^{\left(\beta_0 p_i^{\beta_1} \sum_j \frac{p_j^{\beta_2}}{d_{ij}^{\beta_3}} \right)}$ (no network)	-12.14	0.544	-1.267	2.399	
Force of infection (normalized)	$1 - e^{\left(\frac{\beta_0 p_i^{\beta_1} \sum_j \frac{p_j^{\beta_2}}{d_{ij}^{\beta_3}}}{\left(\sum_{k \neq i} \frac{p_k^{\beta_2}}{d_{ik}^{\beta_3}} \right)^{\beta_4}} \right)}$ (no network)	-0.113	0.395	-1.075	2.621	0.404

Table S4: Model parameter confidence ranges

Model name	Infection probability	Δ_0	Δ_1	Δ_2
Constant	β_0 (no network)	0.0000002		
Well-mixed	β_0	0.0000000004		
Diffusion	$\frac{1}{1 + e^{\beta_0 + \beta_1 d_{ij}}}$	1.079	0.502	
Diffusion + long distance	$\frac{1}{1 + e^{\beta_0 + \beta_1 d_{ij}^{\beta_2}}}$	8.841	9.049	0.450
Diffusion + country borders	$\left\{ \begin{array}{ll} \beta_2 \frac{1}{1 + e^{\beta_0 + \beta_1 d_{ij}}} & \text{if crossing border} \\ \frac{1}{1 + e^{\beta_0 + \beta_1 d_{ij}}} & \text{if not crossing border} \end{array} \right.$	0.669	0.411	0.397
Diffusion + core borders	$\left\{ \begin{array}{ll} \beta_2 \frac{1}{1 + e^{\beta_0 + \beta_1 d_{ij}}} & \text{if crossing core border} \\ \frac{1}{1 + e^{\beta_0 + \beta_1 d_{ij}}} & \text{if not crossing core border} \end{array} \right.$	6.484	0.432	0.146

Δ values represent the confidence interval on either side of the estimated β values (see Table S2).
The remainder of the confidence intervals could not be estimated with this approach (see Methods).

Table S5A: Spearman correlation of selected transmission links

	Gravity + core borders	Gravity + long distance	Gravity	Gravity + country borders	Diffusion	Radiation
Gravity + long distance	0.968					
Gravity	0.981	0.991				
Gravity + country borders	0.982	0.990	0.999			
Diffusion	0.852	0.938	0.886	0.883		
Radiation	0.852	0.938	0.886	0.883	0.999	
Mobility (Senegal)	0.974	0.954	0.984	0.985	0.797	0.797

Table S5B: Pearson correlation of selected transmission links

	Gravity + core borders	Gravity + long distancel	Gravity	Gravity + country borders	Diffusion	Radiation
Gravity + long distancel	0.959					
Gravity	0.983	0.972				
Gravity + country borders	0.983	0.973	0.999			
Diffusion	0.809	0.908	0.834	0.830		
Radiation	0.809	0.908	0.834	0.830	0.999	
Mobility (Senegal)	0.151	0.188	0.146	0.150	0.135	0.133