

Spatio-temporal patterns of proportions of influenza B cases

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Supplementary Material

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S1 Effects of excluding data from 2010 and 2011

Table S1: Effects of setting the start date to 2011-1 and 2012-1.

factor	ref country	time interval	n	θ	ρ	95%CI	p-value	signif
effd1	Mexico	2011-1 to 2015-40	31	2000	0.633	(0.359, 0.806)	1.324e-04	***
effd2	Mexico	2011-1 to 2015-40	31	2000	0.652	(0.388, 0.818)	6.957e-05	****
effd1	China	2011-1 to 2015-40	31	2000	-0.167	(-0.492, 0.199)	0.3683	
longitude	-	2011-1 to 2015-40	31	2000	0.558	(0.255, 0.762)	0.0011	**
latitude	-	2011-1 to 2015-40	31	2000	0.116	(-0.248, 0.452)	0.5341	
effd1	Mexico	2011-1 to 2016-8	31	2000	0.450	(0.114, 0.694)	0.0110	*
effd2	Mexico	2011-1 to 2016-8	31	2000	0.513	(0.194, 0.734)	0.0032	**
effd1	China	2011-1 to 2016-8	31	2000	-0.234	(-0.543, 0.131)	0.2046	
longitude	-	2011-1 to 2016-8	31	2000	0.533	(0.221, 0.747)	0.0020	**
latitude	-	2011-1 to 2016-8	31	2000	-0.018	(-0.370, 0.339)	0.9251	
effd1	Mexico	2012-1 to 2015-40	25	2000	0.563	(0.216, 0.784)	0.0034	**
effd2	Mexico	2012-1 to 2015-40	25	2000	0.599	(0.268, 0.804)	0.0015	**
effd1	China	2012-1 to 2015-40	25	2000	-0.320	(-0.635, 0.086)	0.1187	
longitude	-	2012-1 to 2015-40	25	2000	0.591	(0.256, 0.799)	0.0019	**
latitude	-	2012-1 to 2015-40	25	2000	0.087	(-0.319, 0.466)	0.6785	
effd1	Mexico	2012-1 to 2016-8	25	2000	0.412	(0.020, 0.694)	0.0408	*
effd2	Mexico	2012-1 to 2016-8	25	2000	0.447	(0.063, 0.716)	0.0250	*
effd1	China	2012-1 to 2016-8	25	2000	-0.313	(-0.630, 0.094)	0.1278	
longitude	-	2012-1 to 2016-8	25	2000	0.499	(0.129, 0.747)	0.0112	*
latitude	-	2012-1 to 2016-8	25	2000	-0.064	(-0.448, 0.340)	0.7611	

S2 Proportions of B virus v.s. country median age

Table S2: Summary of correlation tests of proportions of B virus against country median age. A marginally negative correlation is detected when a threshold of 500 cases is used.

factor	time interval	n	θ	ρ	95%CI	p-value	signif
median.age	2006-1 to 2016-8	74	500	-0.295	(-0.491, -0.072)	0.0106	*
median.age	2006-1 to 2008-52	48	500	-0.354	(-0.580, -0.078)	0.0135	*
median.age	2010-1 to 2016-8	74	500	-0.294	(-0.490, -0.071)	0.0109	*
median.age	2010-1 to 2015-40	74	500	-0.283	(-0.480, -0.058)	0.0147	*
median.age	2006-1 to 2016-8	34	2000	-0.025	(-0.361, 0.315)	0.8864	
median.age	2006-1 to 2008-52	30	2000	-0.260	(-0.567, 0.110)	0.1646	
median.age	2010-1 to 2016-8	34	2000	-0.054	(-0.385, 0.289)	0.7610	
median.age	2010-1 to 2015-40	34	2000	-0.120	(-0.440, 0.228)	0.5005	

We downloaded country median age from https://en.wikipedia.org/wiki/List_of_countries_by_median_age (checked on March 22, 2016).

S3 Patterns of total confirmations

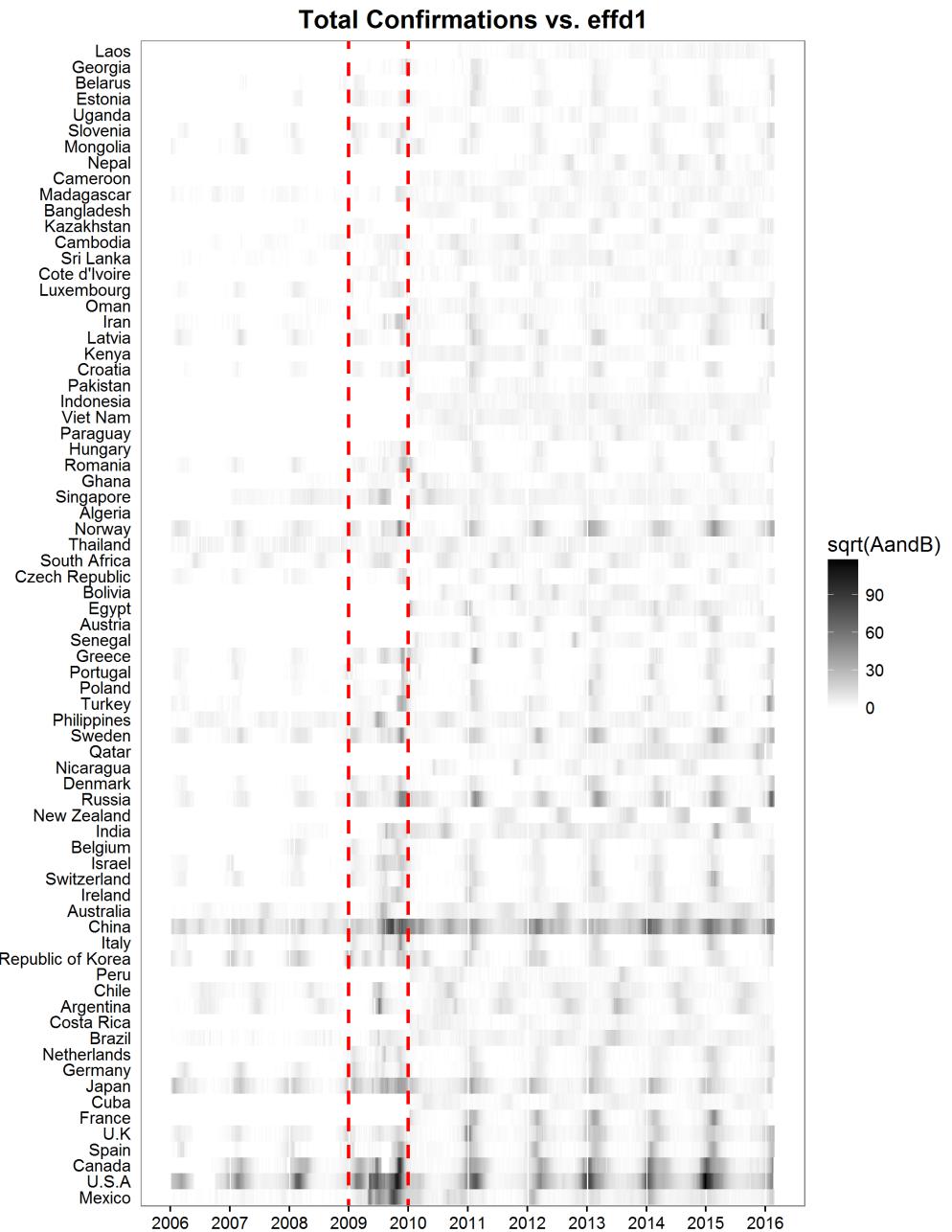


Figure S1: Patterns of total confirmations. Countries are ordered according to their effective distances to Mexico (definition 1).

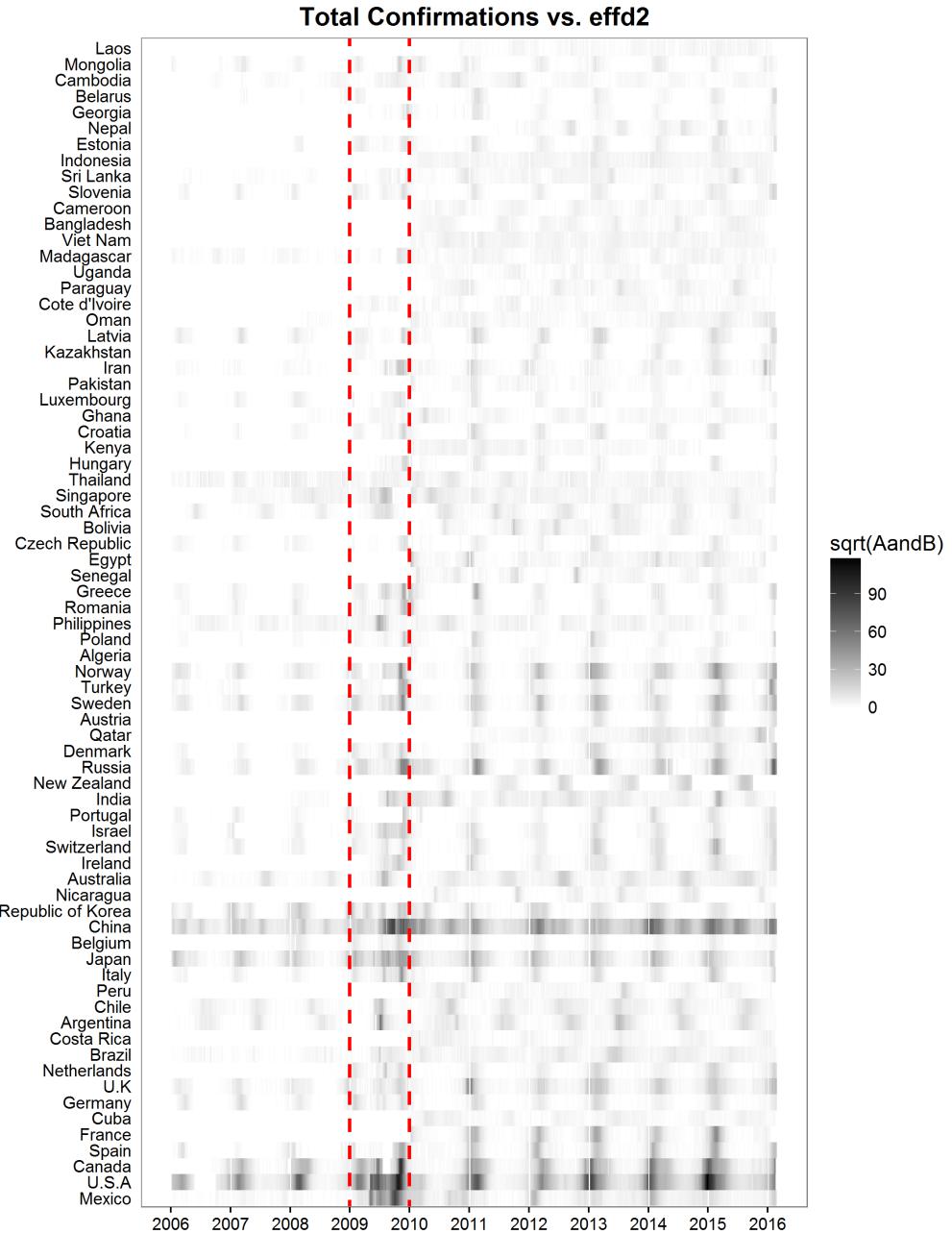


Figure S2: Patterns of total confirmations. Countries are ordered according to their effective distances to Mexico (definition 2).

S4 Patterns of proportions of B Lineages

S4.1 Proportions of B Lineages v.s. median age

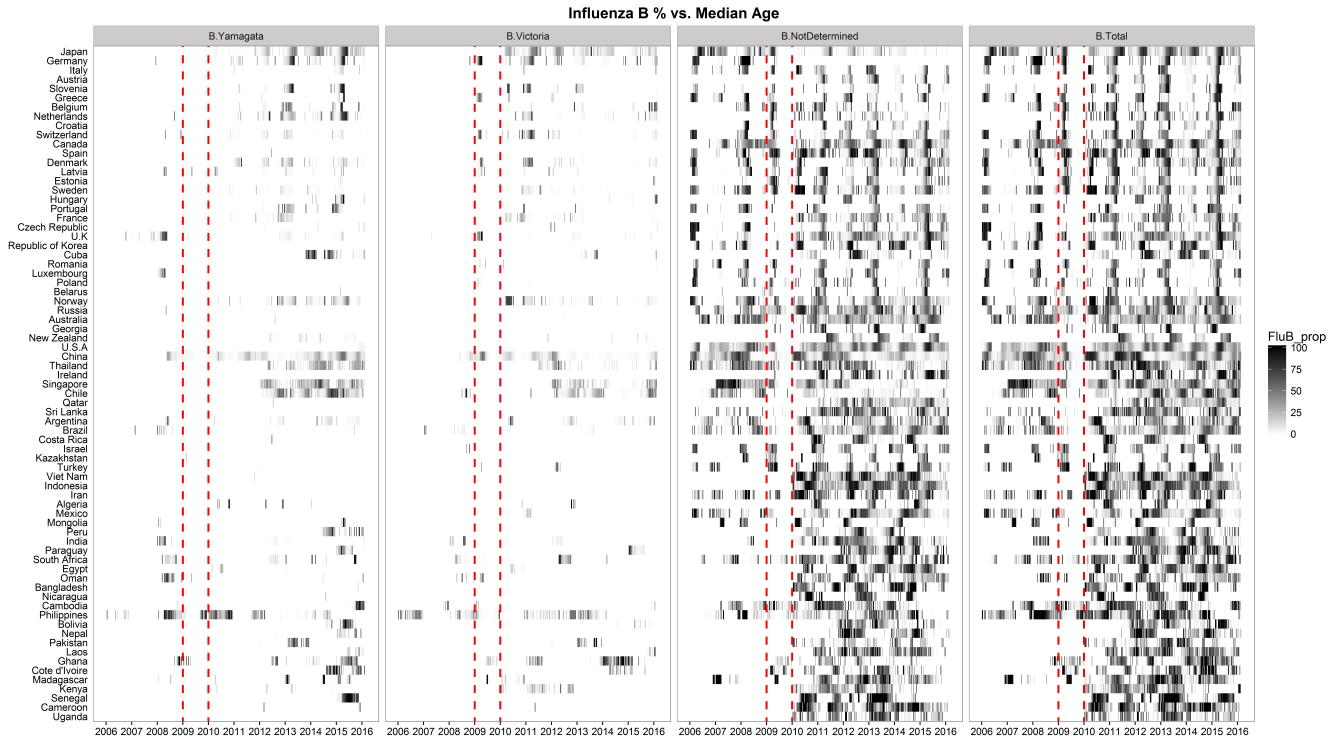


Figure S3: Patterns of proportions of different B lineage influenza strains out of total of A and B. Countries are ordered according to the population's median ages.

S4.2 Confirmations of B Lineages v.s. effective distances

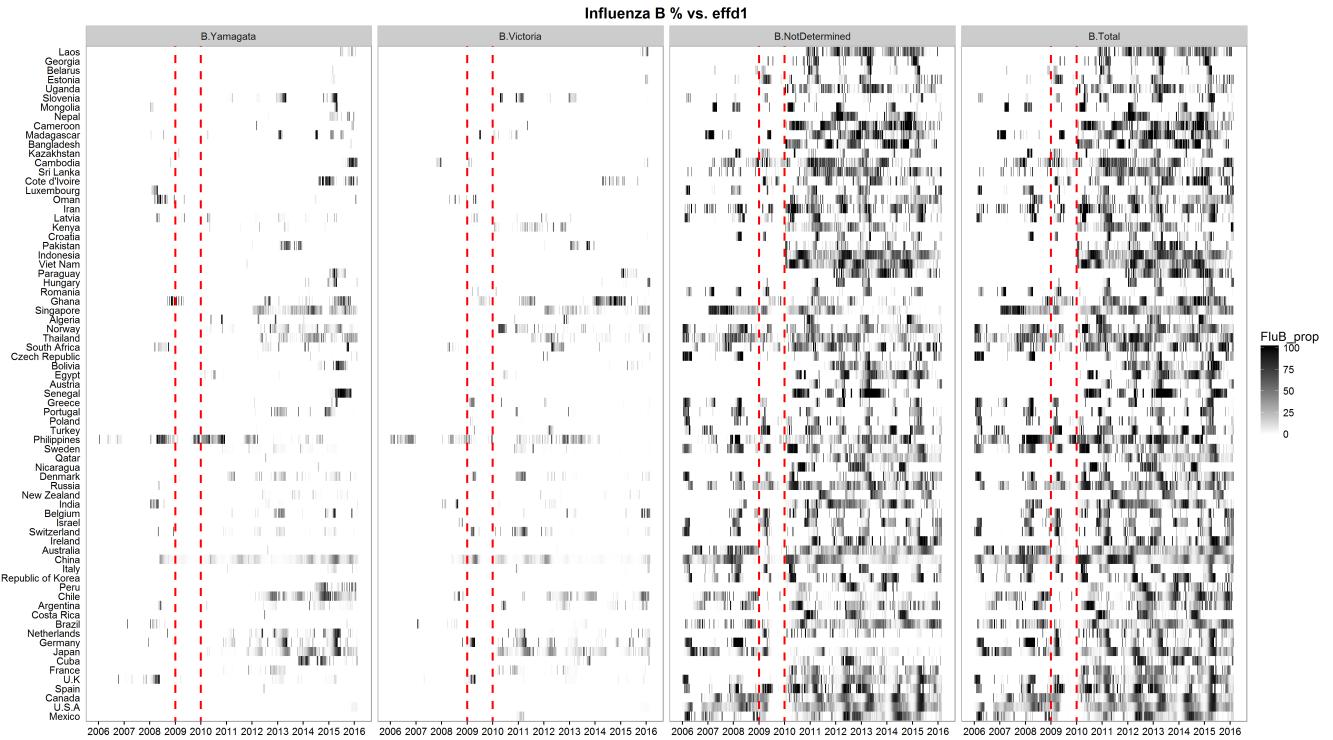


Figure S4: Patterns of proportions of different B lineages out of total of A and B. Countries are ordered according to the effective distance (definition 1) from Mexico.

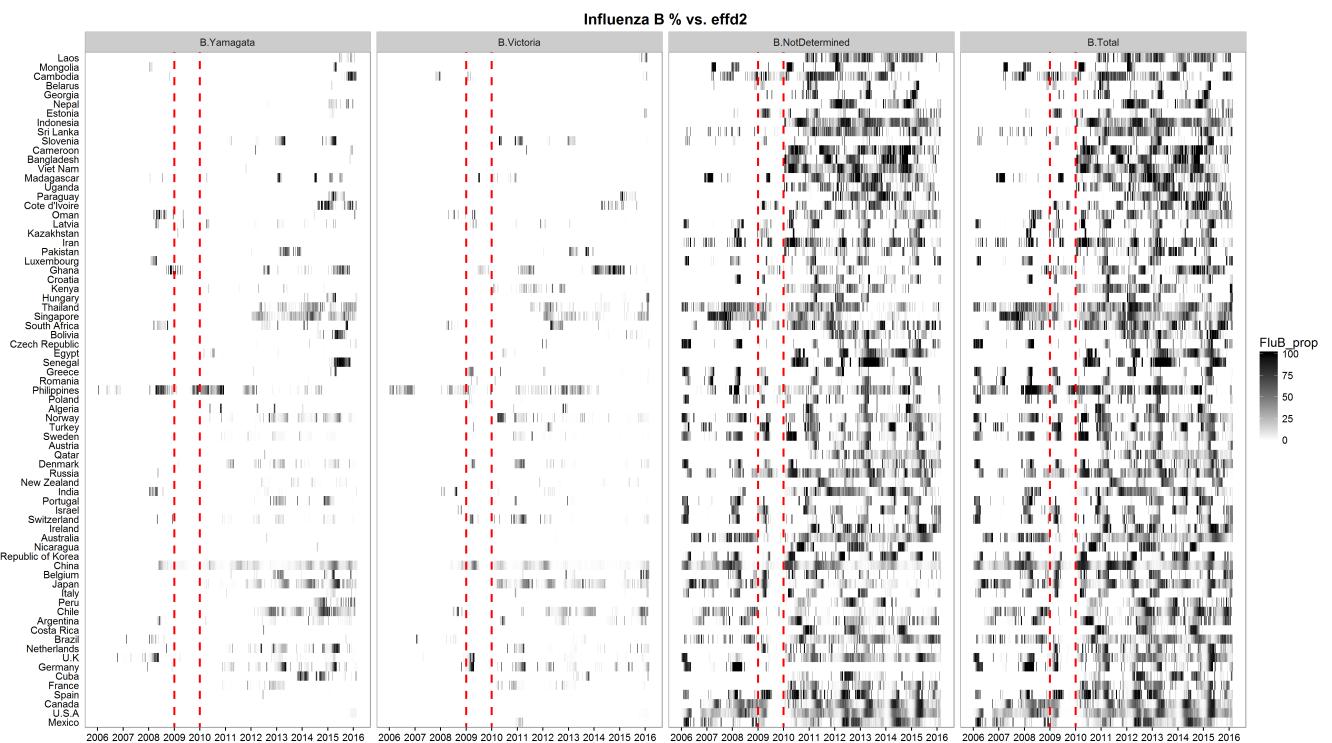


Figure S5: Patterns of proportions of different B lineages out of total of A and B. Countries are ordered according to the effective distance (definition 2) from Mexico.

S4.3 Proportions of B Lineages v.s. longitude and latitude

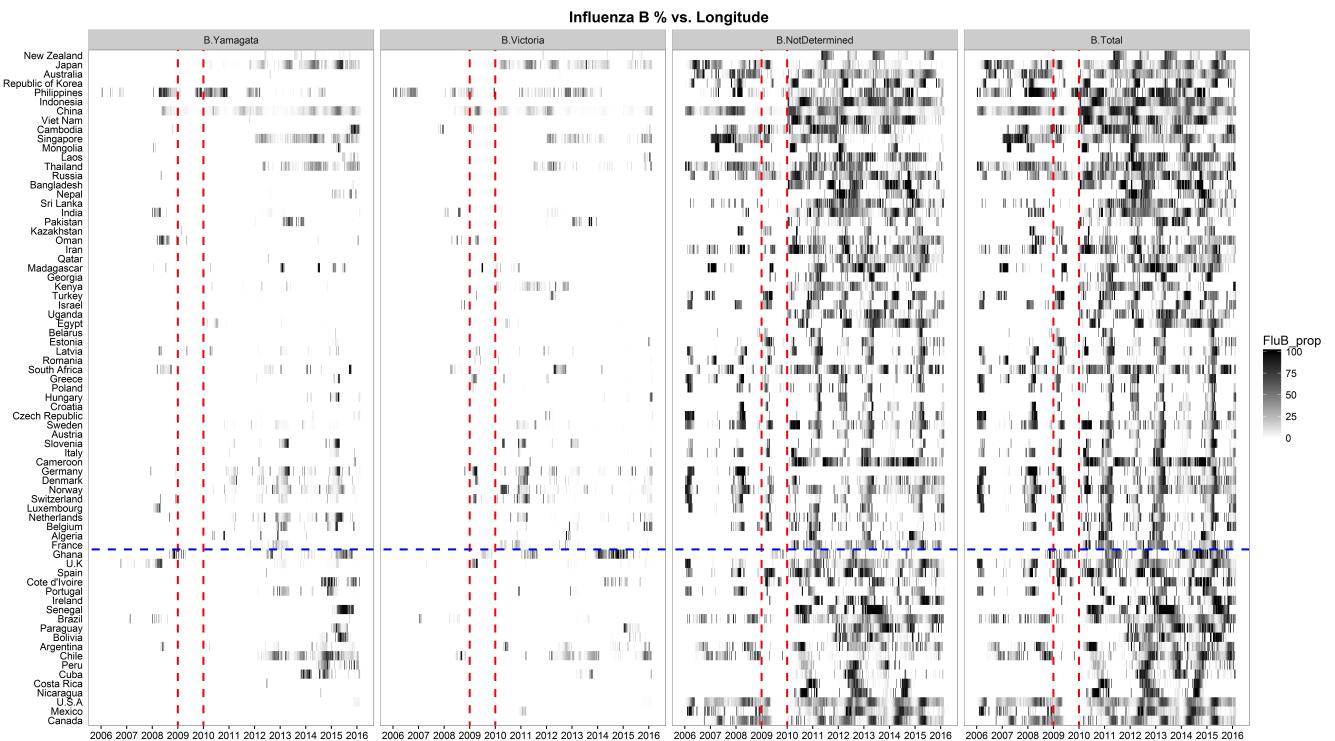


Figure S6: Patterns of proportions of different B lineages out of total of A and B. Countries are ordered according to longitude.

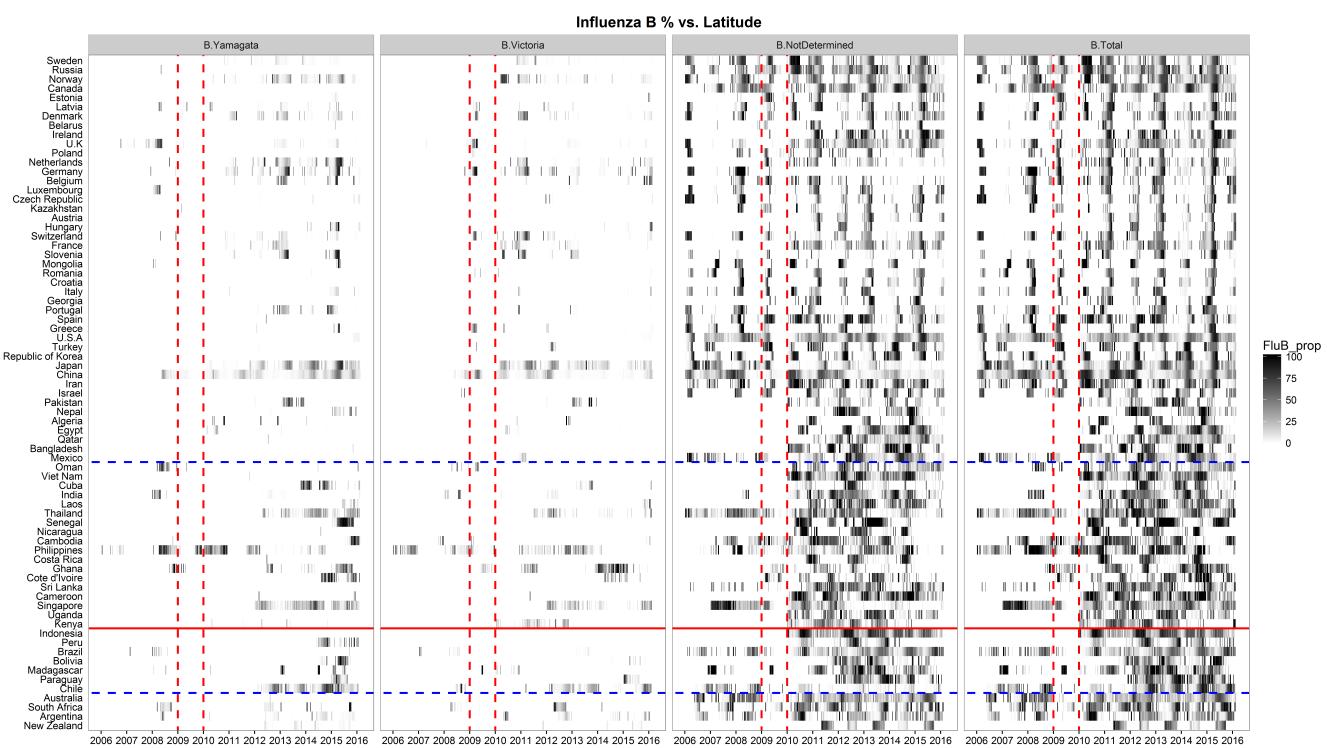


Figure S7: Patterns of proportions of different B lineages out of total of A and B. Countries are ordered according to latitude.

S5 A variety of models with or without region as a random-effect factor

A variety of models with or without region as a random-effect factor

Linear mixed model fit by REML [‘lmerMod’]

Formula: r ~ pop + long + abslat + effd + (1 | region)

Data: rr

REML criterion at convergence: -162.7

Scaled residuals:

Min	1Q	Median	3Q	Max
-2.41305	-0.52911	0.05728	0.69284	2.02150

Random effects:

Groups	Name	Variance	Std.Dev.
region	(Intercept)	0.0004447	0.02109
Residual		0.0042312	0.06505

Number of obs: 77, groups: region, 19

Fixed effects:

	Estimate	Std. Error	t value
(Intercept)	0.274755	0.009241	29.731
pop	-0.002384	0.008135	-0.293
long	0.022551	0.009811	2.299
abslat	-0.004060	0.008993	-0.451
effd	0.022670	0.009025	2.512

Correlation of Fixed Effects:

	(Intr)	pop	long	abslat
pop	-0.008			
long	0.019	-0.238		
abslat	0.058	-0.009	0.050	
effd	0.010	0.295	-0.418	-0.012

Linear mixed model fit by REML [‘lmerMod’]

Formula: r ~ specimens + long + abslat + effd + (1 | region)

Data: rr

REML criterion at convergence: -161.5

Scaled residuals:

Min	1Q	Median	3Q	Max
-2.41554	-0.48051	-0.00582	0.73237	2.03358

Random effects:

Groups	Name	Variance	Std.Dev.
region	(Intercept)	0.0006186	0.02487
Residual		0.0039121	0.06255

Number of obs: 75, groups: region, 19

Fixed effects:

	Estimate	Std. Error	t value
(Intercept)	0.277405	0.009716	28.550
specimens	0.007829	0.008800	0.890
long	0.027741	0.010224	2.713
abslat	-0.003936	0.009331	-0.422
effd	0.021356	0.009560	2.234

Correlation of Fixed Effects:

	(Intr)	spcmns	long	abslat
specimens	-0.055			
long	0.066	0.020		
abslat	0.092	-0.126	0.073	
effd	-0.045	0.381	-0.381	-0.075

Call:

```
lm(formula = r ~ pop + long + abslat + effd, data = rr)
```

Residuals:

Min	1Q	Median	3Q	Max
-0.171101	-0.042221	0.005935	0.045810	0.153983

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	0.273880	0.007732	35.420	< 2e-16 ***
pop	-0.002944	0.008209	-0.359	0.72089
long	0.026653	0.008857	3.009	0.00361 **
abslat	-0.003699	0.007762	-0.477	0.63510
effd	0.020509	0.008941	2.294	0.02472 *

Signif. codes: 0 *** 0.001 ** 0.01 * 0.05 . 0.1 1

Residual standard error: 0.06785 on 72 degrees of freedom
(1 observation deleted due to missingness)
Multiple R-squared: 0.2703, Adjusted R-squared: 0.2298
F-statistic: 6.668 on 4 and 72 DF, p-value: 0.000127

Call:

```
lm(formula = r ~ specimens + long + abslat + effd, data = rr)
```

Residuals:

	Min	1Q	Median	3Q	Max
	-0.16828	-0.03902	0.00469	0.04140	0.15632

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	0.276333	0.007702	35.877	< 2e-16 ***
specimens	0.009057	0.008577	1.056	0.294610
long	0.032161	0.008998	3.574	0.000641 ***
abslat	-0.004274	0.007685	-0.556	0.579894
effd	0.020039	0.009602	2.087	0.040523 *

Signif. codes: 0 *** 0.001 ** 0.01 * 0.05 . 0.1 1

Residual standard error: 0.06656 on 70 degrees of freedom

(3 observations deleted due to missingness)

Multiple R-squared: 0.3043, Adjusted R-squared: 0.2646

F-statistic: 7.656 on 4 and 70 DF, p-value: 3.552e-05

Single term deletions

Model:

```
r ~ pop + long + abslat + effd + (region)
```

	Df	Sum of Sq	RSS	AIC	F value	Pr(>F)
<none>		0.22646	-402.83			
pop	1	0.001002	0.22747	-404.49	0.2388	0.62703
long	1	0.006700	0.23316	-402.59	1.5977	0.21166
abslat	1	0.000100	0.22656	-404.80	0.0238	0.87799
effd	1	0.019464	0.24593	-398.48	4.6411	0.03569 *
region	18	0.104956	0.33142	-409.51	1.3904	0.17457

Signif. codes: 0 *** 0.001 ** 0.01 * 0.05 . 0.1 1