

A data-driven model for influenza transmission incorporating media effects

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Appendix

In Figure 1 we show the residuals from linear (blue circles) and quadratic (red crosses) least-square fits to the retweet-infected data from Figure 1 of the main text, for all seasons 2009/10–2014/15 considered in this study. Table 1 shows the relative Akaike weights for the linear and quadratic models for each year, where the relative weights p_i are given by

$$p_i = \exp\left(\frac{AIC_{\min} - AIC_i}{2}\right)$$

and AIC_i are the AIC values for each model, AIC_{\min} is the minimum value of AIC_i . These weights represent the probability that the i th model minimises the information loss in describing the data, relative to the best-fitting model [1]. Figure 2 shows relative Akaike weight of the quadratic model as a function of final outbreak size, suggesting that nonlinear effects will increasingly come into play for more severe outbreaks and pandemics.

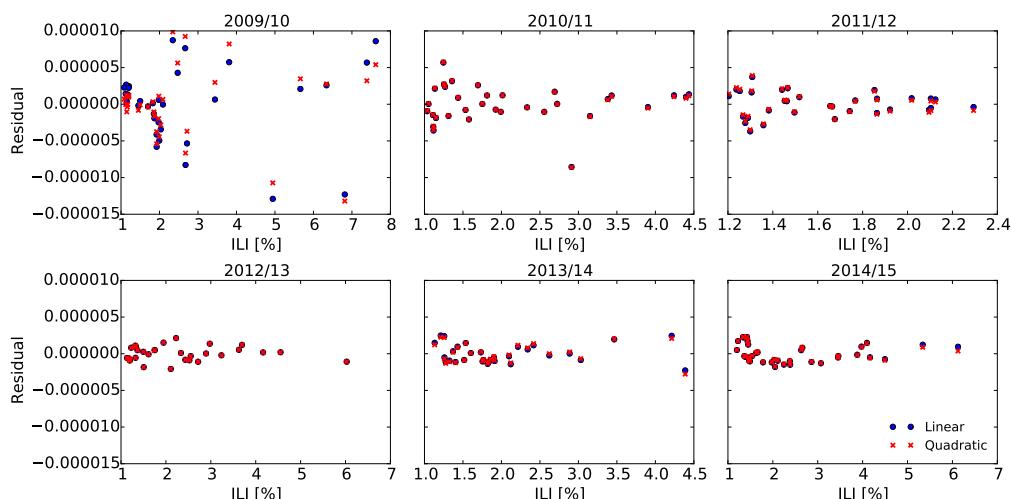


Figure 1. Residual plots for linear (blue dots) and quadratic (red crosses) media function models.

Table 1. Relative Akaike weights p_{lin} and p_{quad} for selecting the linear and quadratic models during each influenza season 1998/99–2014/15. Weights for each preferred model are shown in bold.

year	p_{lin}	p_{quad}
2009/10	0.4454	0.5546
2010/11	0.7769	0.2231
2011/12	0.8211	0.1789
2012/13	0.7781	0.2219
2013/14	0.6966	0.3034
2014/15	0.5811	0.4189

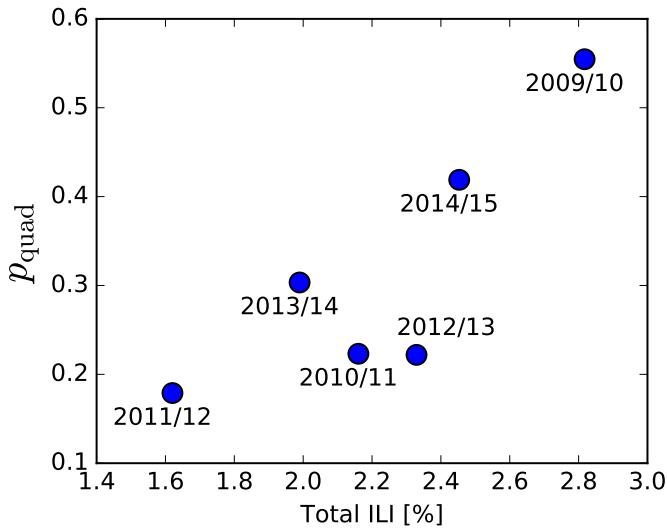


Figure 2. Relationship between relative Akaike weight for the quadratic model p_{quad} and the observed final outbreak size for each influenza season 2009/10–2014/15.

Table 2 shows the best-fitting parameters for the SEEIIR and SEEIIR-M models for all seasons 1998/99–2014/15, analogously to Table 1 in the main text. For completeness we also present results where we have used a standard SEIR model with only one exposed and infectious compartment each. Table 3 is an analog of Table 2 from the main text, showing the average conditional probability of selecting each model, and Figure 3 shows example fits to observations of the percentage of new laboratory-confirmed influenza cases per week for the SEIR and SEEIIR-M models, similarly to Figure 3 in the main text. In terms of model fitting, the confidence intervals in Table 3 show no significant difference in the likelihood of selecting each media model. Figure 3 suggests that in 2013/14 the media functions f_1 , f_2 and f_3 fit the data slightly worse using an SEIR model than with an SEEIIR model, while the $f(I) \equiv 1$ and f_m fit describe the data essentially as well using an SEIR model than with an SEEIIR model.

Table 2: Parameters of best fit for SEEIIR and SEEIIR-M models for all influenza seasons.

	model	R_0	$1/\sigma$ (days)	$1/\gamma$ (days)	p_i	p_{AIC}
1998	$f(I) \equiv 1$	1.17	1.00	2.41	–	0.80
	f_m	2.00	1.92	1.41	0.22	0.02
	f_1	1.30	1.00	2.30	0.02	0.15
	f_2	1.35	1.21	2.06	0.06	0.03
	f_3	1.58	1.98	2.02	5.71	0.00
1999	$f(I) \equiv 1$	1.23	1.00	1.63	–	0.00
	f_m	1.47	1.28	1.58	0.11	0.99
	f_1	1.63	1.71	1.38	0.03	0.01
	f_2	2.00	1.95	1.92	0.62	0.00
	f_3	1.76	1.90	1.30	4.14	0.01
2000	$f(I) \equiv 1$	1.14	1.00	1.26	–	0.00
	f_m	1.30	1.00	1.23	0.21	1.00

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	model	R_0	$1/\sigma$ (days)	$1/\gamma$ (days)	p_i	p_{AIC}
2000	f_1	1.35	1.13	1.13	0.04	0.00
	f_2	1.33	1.99	1.97	1.05	0.00
	f_3	1.59	1.99	1.99	1.06	0.00
2001	$f(I) \equiv 1$	1.10	1.00	1.98	–	0.00
	f_m	1.45	1.14	1.33	0.24	0.89
	f_1	1.34	1.37	1.32	0.03	0.00
	f_2	1.38	1.29	1.37	0.26	0.00
	f_3	1.46	1.07	1.38	3.15	0.11
2002	$f(I) \equiv 1$	1.13	1.00	1.00	–	0.00
	f_m	1.28	1.00	1.00	0.29	1.00
	f_1	1.56	1.61	1.55	0.19	0.00
	f_2	1.28	1.35	1.00	0.70	0.00
	f_3	1.71	1.83	1.74	0.54	0.00
2003	$f(I) \equiv 1$	1.17	1.00	1.46	–	1.00
	f_m	1.17	1.00	1.46	0.00	0.00
	f_1	1.17	1.00	1.46	0.00	0.00
	f_2	1.17	1.00	1.46	0.00	0.00
	f_3	1.19	1.00	1.46	56.78	0.00
2004	$f(I) \equiv 1$	1.11	1.00	1.30	–	0.00
	f_m	1.36	1.02	1.00	0.30	1.00
	f_1	2.00	2.00	2.07	0.68	0.00
	f_2	1.23	1.16	1.10	0.41	0.00
	f_3	1.14	1.09	1.20	33.94	0.00
2005	$f(I) \equiv 1$	1.10	1.00	1.09	–	0.00
	f_m	1.24	1.00	1.01	0.28	1.00
	f_1	1.10	1.78	2.95	0.14	0.00
	f_2	1.33	1.00	1.00	2.58	0.00
	f_3	1.19	1.00	1.29	4.23	0.00
2006	$f(I) \equiv 1$	1.12	1.00	1.22	–	0.00
	f_m	1.40	1.00	1.00	0.36	1.00
	f_1	1.32	1.04	1.16	0.05	0.00
	f_2	1.62	1.04	1.00	2.51	0.00
	f_3	1.54	1.78	1.00	1.24	0.00
2007	$f(I) \equiv 1$	1.12	1.00	1.64	–	0.01
	f_m	1.26	1.00	1.42	0.13	0.99
	f_1	1.45	1.49	1.09	0.04	0.00
	f_2	1.72	1.27	1.00	0.66	0.00
	f_3	1.42	1.84	1.84	2.11	0.00
2008	$f(I) \equiv 1$	1.14	2.83	1.06	–	0.00
	f_m	1.28	1.00	1.07	0.27	0.96
	f_1	1.40	1.13	1.00	0.06	0.00
	f_2	1.25	1.02	1.02	0.57	0.04
	f_3	2.00	1.18	1.13	0.48	0.00
2009	$f(I) \equiv 1$	1.16	1.00	1.74	–	0.00
	f_m	1.10	1.10	3.39	0.06	0.00
	f_1	1.39	1.28	1.46	0.03	0.91
	f_2	1.41	1.64	1.19	0.11	0.00
	f_3	1.53	1.57	1.28	4.83	0.09
	$f(I) \equiv 1$	1.11	1.00	1.31	–	0.35

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	model	R_0	$1/\sigma$ (days)	$1/\gamma$ (days)	p_i	p_{AIC}
2010	f_m	1.16	1.28	1.04	0.06	0.00
	f_1	1.30	1.86	2.32	0.22	0.00
	f_2	1.28	1.02	1.01	0.33	0.56
	f_3	1.21	1.00	1.16	7.44	0.09
2011	$f(I) \equiv 1$	1.10	2.75	1.09	–	0.00
	f_m	1.24	1.00	1.00	0.48	1.00
	f_1	1.57	1.00	1.33	0.38	0.00
	f_2	1.10	1.03	1.42	0.35	0.00
	f_3	1.25	1.09	1.56	1.24	0.00
2012	$f(I) \equiv 1$	1.15	1.00	1.94	–	0.00
	f_m	1.44	1.30	1.70	0.21	0.96
	f_1	1.57	1.63	1.51	0.07	0.01
	f_2	1.41	1.51	1.57	0.22	0.00
	f_3	1.54	1.44	1.61	2.96	0.03
2013	$f(I) \equiv 1$	1.16	1.00	1.25	–	0.00
	f_m	1.53	1.78	1.00	0.34	1.00
	f_1	1.68	2.13	1.01	0.07	0.00
	f_2	1.33	1.33	1.19	0.35	0.00
	f_3	1.75	1.23	1.13	1.19	0.00
2014	$f(I) \equiv 1$	1.19	1.00	2.46	–	0.00
	f_m	1.70	1.67	2.00	0.18	0.81
	f_1	1.65	2.00	2.02	0.06	0.00
	f_2	1.57	1.68	2.02	0.12	0.02
	f_3	1.96	1.93	1.85	2.64	0.17

Table 3. Average probability of selecting each SEIR-type model over the 1998/99-2014/15 seasons. (cf. Table 2 of the main text.) The 2009/10 pandemic influenza season has been excluded.

	p_{AIC}	95% CI
$f(I) \equiv 1$	0.0635	[0,0.1714]
$f_m(I) = 1 - p_m I$	0.9122	[0.7974,1]
$f_1(I) = \exp(-\gamma p_1 I)$	0.0010	[0,0.0026]
$f_2(I) = \frac{1}{1+p_2 I^2}$	0.0134	[0,0.0388]
$f_3(I) = \frac{I}{1+p_3 I}$	0.0100	[0,0.0255]

References

- 1 Anderson DR. Model Based Inference in the Life Sciences. New York: Springer; 2008.

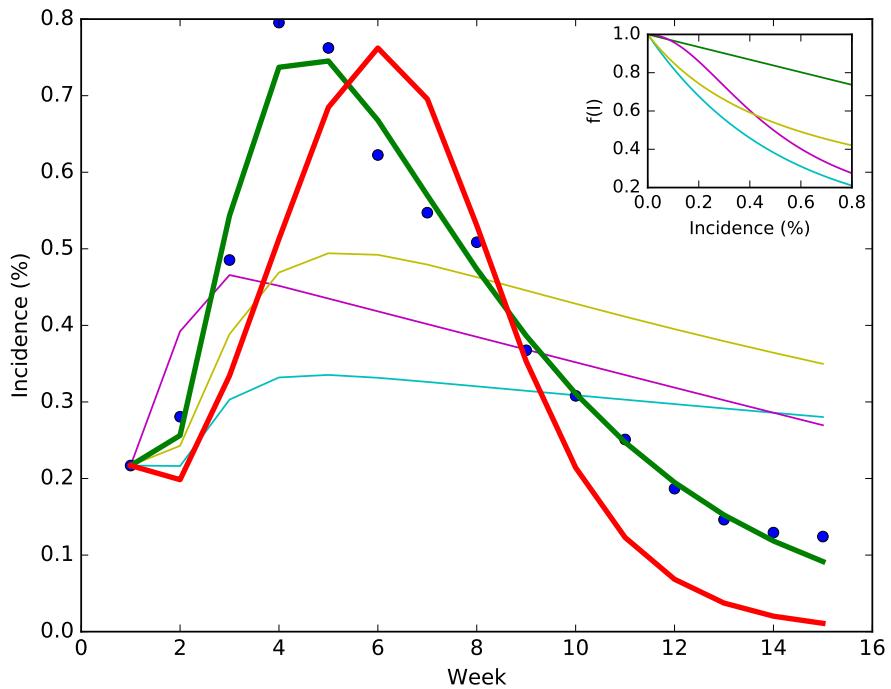


Figure 3. Best model fits to data from 2013/14 flu season. Weekly influenza data (blue dots), and model fits for SEIR model without media function (red), with media function (green), and variations f_1 (cyan), f_2 (magenta) and f_3 (yellow), for the 2013/14 influenza season (USA). (cf. Figure 3 from the main text.)