

The transmission dynamics of Norovirus in England: a genotype-specific modelling study

Juan F Vesga¹, Amy Douglas², Cristina Celma², Edward S. Knock³, Marc Baguelin^{3,4}, W. John Edmunds⁴

¹ Modelling & Economics Unit, UK Health Security Agency, London, UK; ² Gastrointestinal Infections, Food Safety and One Health Division, UK Health Security Agency, London, United Kingdom; ³ MRC Centre for Global Infectious Disease Analysis, School of Public Health, Imperial College London, London, UK; ⁴ Centre for Mathematical Modelling of Infectious Diseases, Department of Infectious Disease Epidemiology, London School of Hygiene & Tropical Medicine, London, UK

Background

- Seasonal acute gastroenteritis virus, mostly active in winter
- In the UK it causes ~3.7 million infections a year¹
- Severe cases affect older people and infants
- Costing the NHS over £100 million annually²
- Currently no vaccines are available

Aims

- Understand transmission dynamics of Norovirus in England and Wales
- Assess the impact of introducing a new vaccine formulation, and the optimal roll-out and targets

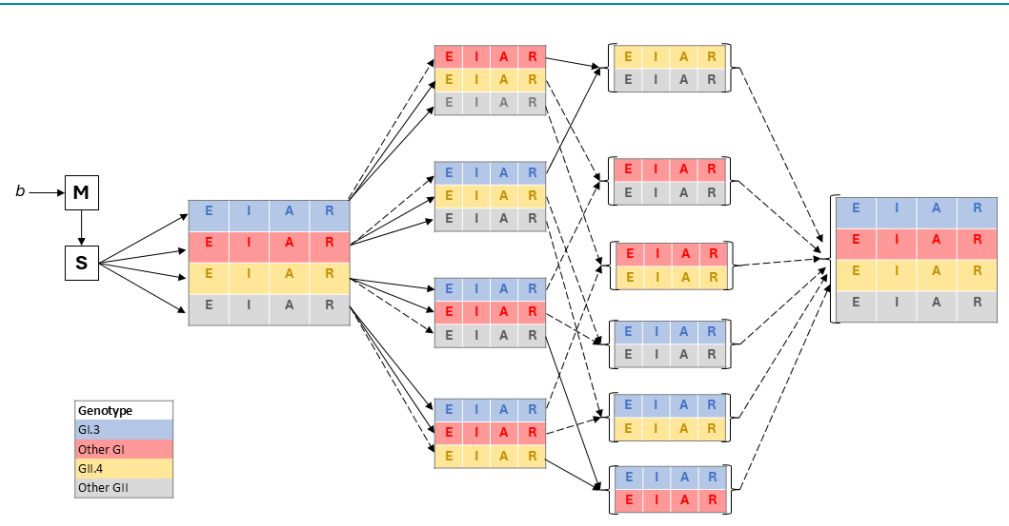
Methods

- Age-structured, multi-strain, stochastic compartmental transmission model
- Calibrated to
 - weekly surveillance time series (UKHSA)
 - Genogroup specific community incidence estimates
 - GII.4. Seroprevalence in Children
- Model selection to assess alternative mechanisms of immunity

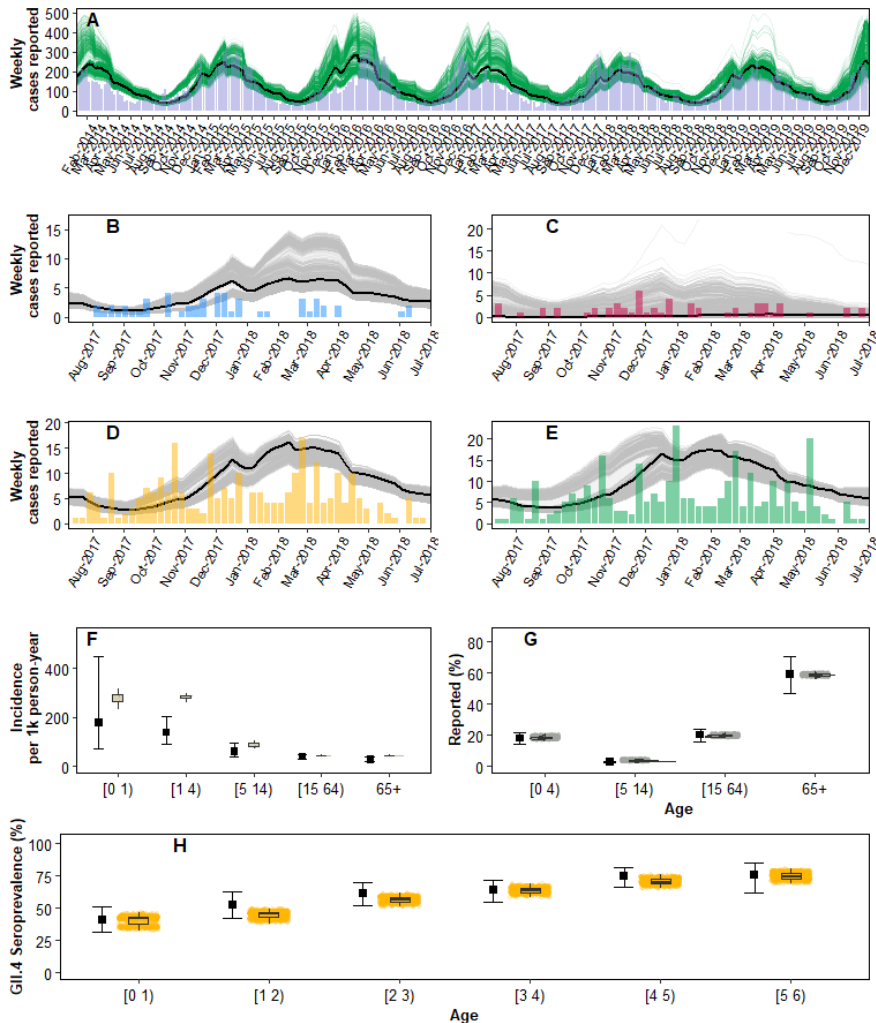
Target	Pre-clinical	Phase I	Phase IIA	Phase IIB	Phase III
Bivalent GI.1 GII.4			NVSI		Hillevar
GI.1 GI.3 GII.4 GII.17			Anhui		
Monovalent GI.1 or GII.4			Vaxart		
Trivalent GI.1 GII.3 GII.4					Moderna

NVSI National Vaccine and Serum Institute, China - Recombinant polysaccharide conjugate; Anhui Zhifei Biologic Pharmacy Ltd, China - Recombinant polysaccharide conjugate; Vaxart, USA - Viral vector vaccine
Moderna USA (UK Trial) - mRNA

- 39 sites across the UK
- 2,500 participants in the UK between October 2024 and early 2025
- Part of the Moderna-UK Strategic partnership

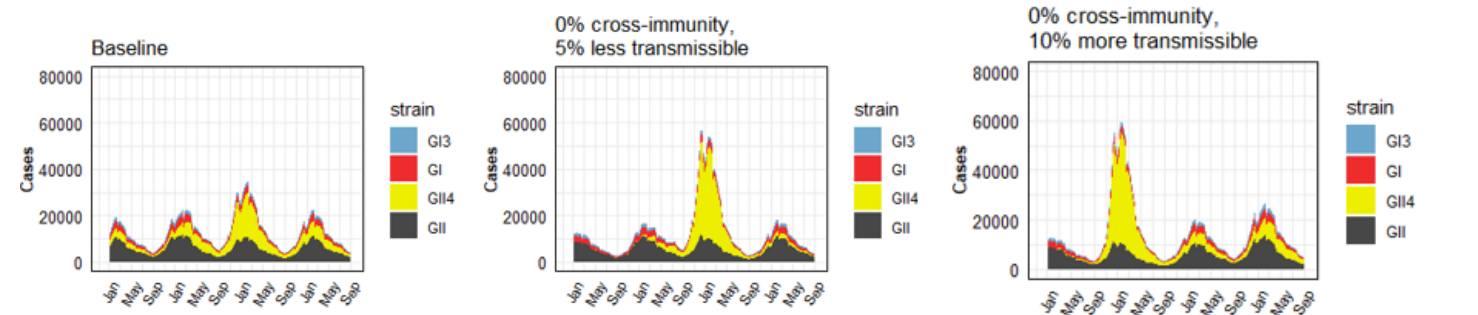


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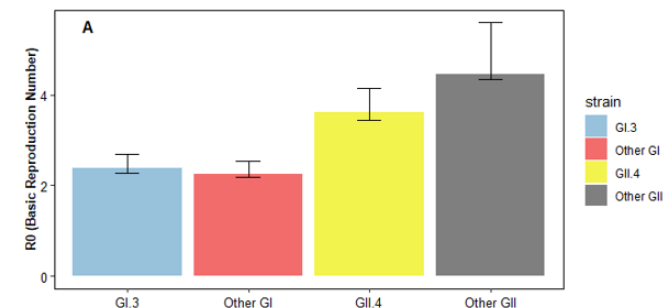
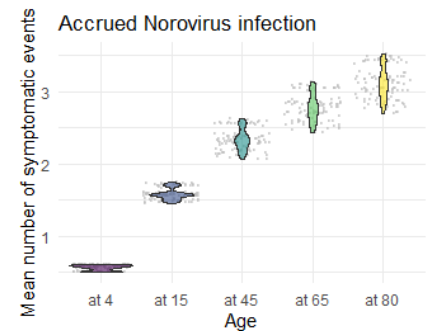
Model selection favors a calibration with a model involving 2 immunity parameters

- Maternal antibodies have an estimated mean protection of 239 days
- Waning immunity after a first symptomatic episode lasts a mean of 16 years
- Genogroup specific cross-protection estimates (e.g., protection against G.II.4 after a single episode of other GII strains) indicates low cross-protection between GI strains (5%, and negligible protection within the GII group (1%).



Emergence of a novel GII.4 variant will cause a large season of Noro cases, and the peak is determined by the potential relative transmissibility.

On average in the UK an individual will have ~3 symptomatic events over a lifetime



R0 ~ 3-4 for GII genogroups and ~ 2.2 for GI