Longitudinal trajectories of influenza immune response after repeat vaccination

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Supported with funding from UGA Graduate School and CIVIC - CIVR-HRP (contract: 75N93019C00052)

What could trajectories look like?



Expected patterns



Expected patterns

This is what we want!



When we consider boosting/waning as at least a 4-fold change, we see all patterns in real data.

- Data are from an ongoing cohort study of flu vaccination by Ted Ross' group at UGA. (Started 2016.)
- Participants give serum samples before vaccination and approx. 28 days after. Some repeated in consecutive years, so we can use these prevaccination titers as a "follow-up" measurement.
- Serum samples are tested against a panel of current and historical influenza samples.
- Figure shows: 200 randomly sampled HAI titer against B-Phuket-2013 strain (total sample size for this strain = 640 trajectories) across all five years of the study.



Most trajectories fall into the expected categories—others could be measurement error or biological.



No 4-fold change was, by far, the most common outcome across all strains and seasons.



Future research

- Quantify patterns with formal statistical analyses.
- Are there noticeable patterns when we stratify trajectories by host factors, like sex or age?
- How do patterns of homologous and heterologous trajectories differ? We have an entire panel of heterologous strains we can check!

Example of stratification:

Pattern	Age >= 65	Age < 65	Overall
boosting waning	3%	17%	16%
boosting no waning	8%	14%	13%
no boosting waning	29%	10%	11%
no boosting no waning	60%	58 %	58%
other 5 patterns	0%	1%	2%
For trajectories of B-Phuket-2013 response (n = 640 representing 376 unique individuals).			