

## Background

Hemagglutinin and neuraminidase antibodies are essential for preventing infection and reducing severe illness. Older adults are vulnerable to influenza-related complication. There is a lack of study regarding the relationship between these antibodies and protection against influenza in older adults in the community, particularly in an Asian population with low vaccination coverage.

## Objectives

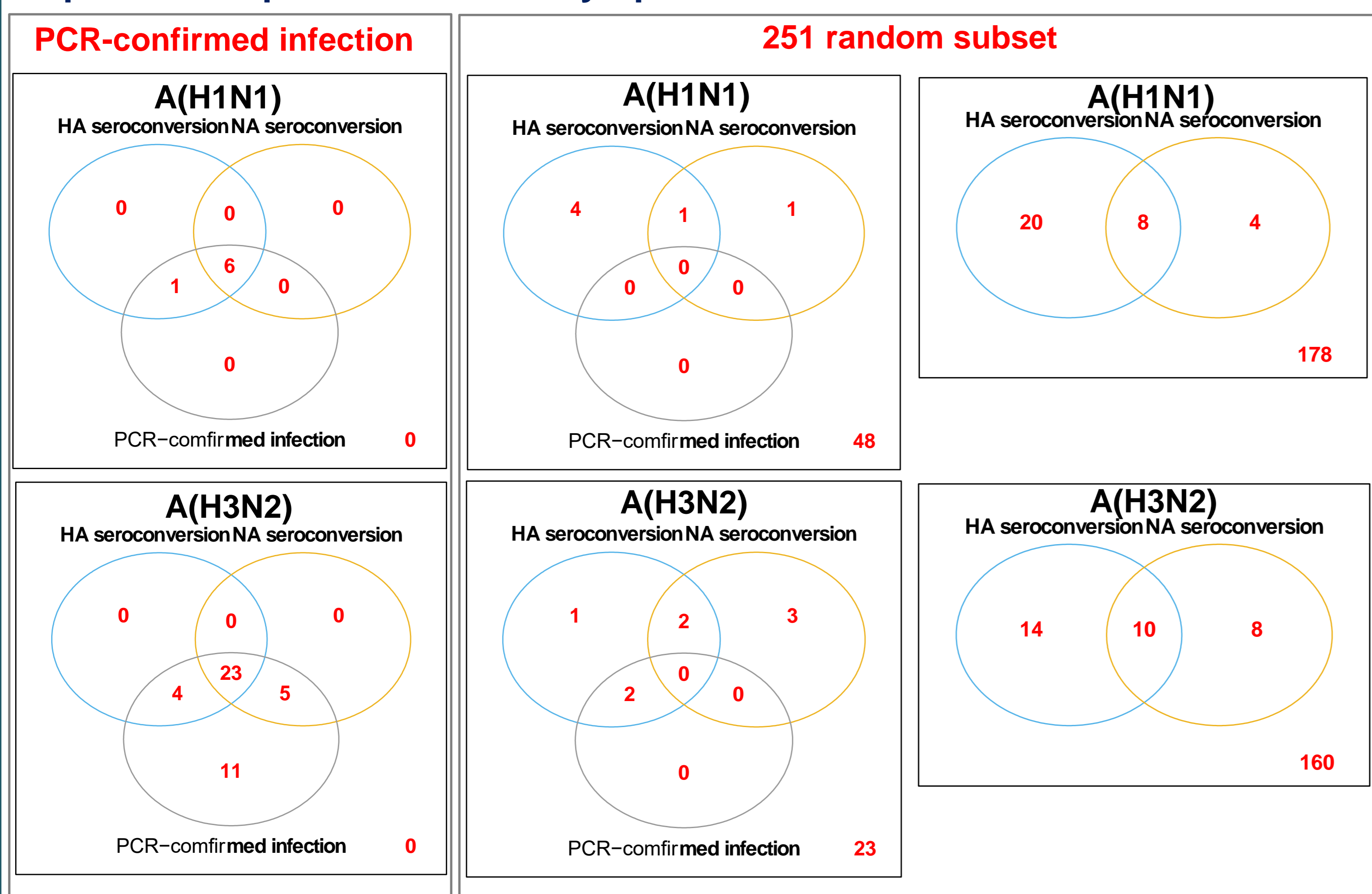
- To estimate the cumulative incidence of influenza virus infection using different methods;
- To compare the association between pre-existing HAI and NAI titers and risk of influenza infection, among those with symptomatic and asymptomatic/subclinical infections;
- To evaluate the relationship between pre-existing antibody titers and the magnitude of fold change in antibody titer after infection.

## Methods

We compared all PCR-confirmed influenza infections and a random subset of 251 participants from a cohort of 1527 adults aged ≥60 years in eastern China in 2015-2017. Nose and throat swabs were collected during acute respiratory illness for influenza type/subtype identifying using polymerase chain reaction (PCR). Sera were collected every 6-12 months using hemagglutination inhibition (HAI) assay and neuraminidase inhibition (NAI) titer using enzyme-linked lectin assay (ELLA).

## Results

- High serological response rates in PCR-confirmed infections;**
- Only 2/9 PCR-confirmed infection with data available antibody titer, indicating **potential representation of asymptomatic or subclinical infections**



**Figure 1.** Overlap testing using PCR, HAI and NAI assay among participants with PCR-confirmed influenza A subtype and in a random subset of 251 individuals.

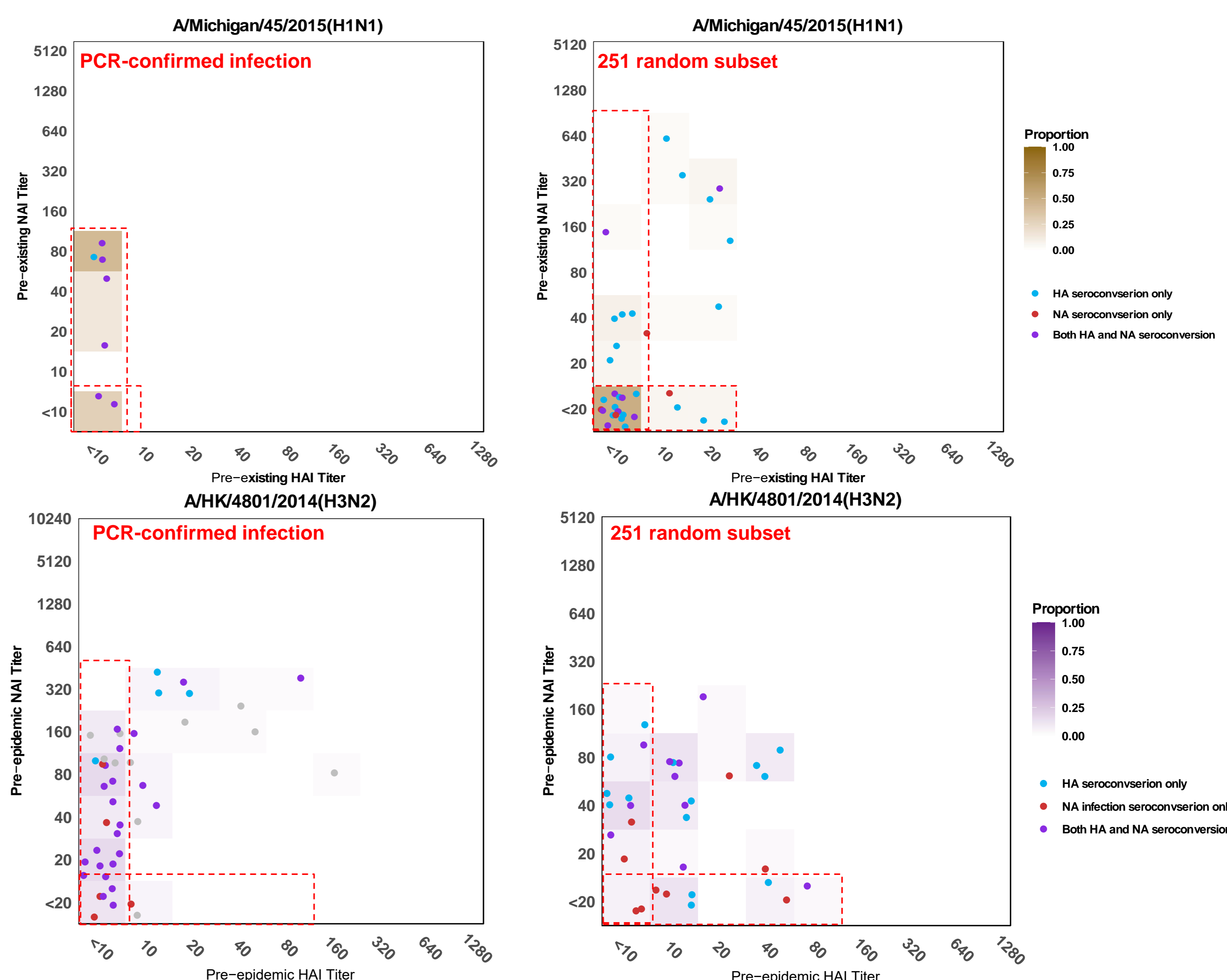
- HAI method achieves higher** cumulative incidence coverage compared to NAI;
- Maximum cumulative incidence** can be estimated using different methods

**Table 1.** Cumulative incidence by PCR or serology with corresponding 95% CI.

|  | Suzhou<br>(N=108) <sup>a</sup> | Yancheng<br>(N=143) <sup>a</sup> | Overall<br>(N=251)                       |
|--|--------------------------------|----------------------------------|--|
| <b>A(H1N1)</b>                                       |                                |                                  |  |
| PCR confirmed infection no. (%) <sup>b</sup> 2015-16 | 0 (0)                          | 1 (0.70)                         | 1 (0.40)                                 |
| Seroconversion rate % (95% CI) <sup>c</sup> 2015-16  | (N=108)                        | (N=139)                          | (N=247)                                  |
| Anti-H1 % (95% CI)                                   | 9.3 (4.5, 16)                  | 16 (10, 23)                      | 13 (9, 18)                               |
| Anti-N1 % (95% CI)                                   | 7.4 (3.2, 14)                  | 2.9 (0.79, 7.2)                  | 4.9 (2.5, 8.3)                           |
| <b>Cumulative incidence by PCR or seroconversion</b> |                                |                                  | 15 (10, 20)<br>17 (14, 19) <sup>a</sup>  |
| <b>A(H3N2)</b>                                       |                                |                                  |  |
| PCR confirmed infection no. (%) <sup>b</sup> 2016-17 | 0 (0)                          | 5 (3.5)                          | 5 (2.0)                                  |
| Seroconversion rate % (95% CI) <sup>c</sup> 2016-17  | (N=100)                        | (N=126)                          | (N=226)                                  |
| Anti-H3 % (95% CI)                                   | 12 (6.4, 20)                   | 13 (7.4, 20)                     | 12 (8.4, 17)                             |
| Anti-N2 % (95% CI)                                   | 9.0 (4.2, 16)                  | 8.7 (4.4, 15)                    | 8.8 (5.5, 13)                            |
| <b>Cumulative incidence by PCR or seroconversion</b> |                                |                                  | 19 (14, 24)<br>17 (14, 20) <sup>a</sup>  |
| PCR confirmed infection no. (%) <sup>b</sup> 2017-18 | 1 (0.93)                       | 2 (1.4)                          | 3 (1.2)                                  |
| Seroconversion rate % (95% CI) <sup>c</sup> 2017-18  | (N=107)                        | (N=124)                          | (N=231)                                  |
| Anti-H3 % (95% CI)                                   | 5.6 (2.1, 12)                  | 6.5 (2.8, 12)                    | 6.1 (3.4, 10)                            |
| Anti-N2 % (95% CI)                                   | 6.5 (2.7, 13)                  | 0.8 (0.02, 4.4)                  | 3.5 (1.5, 6.7)                           |
| <b>Cumulative incidence by PCR or seroconversion</b> |                                |                                  | 12 (7.8, 17)<br>12 (10, 13) <sup>a</sup> |

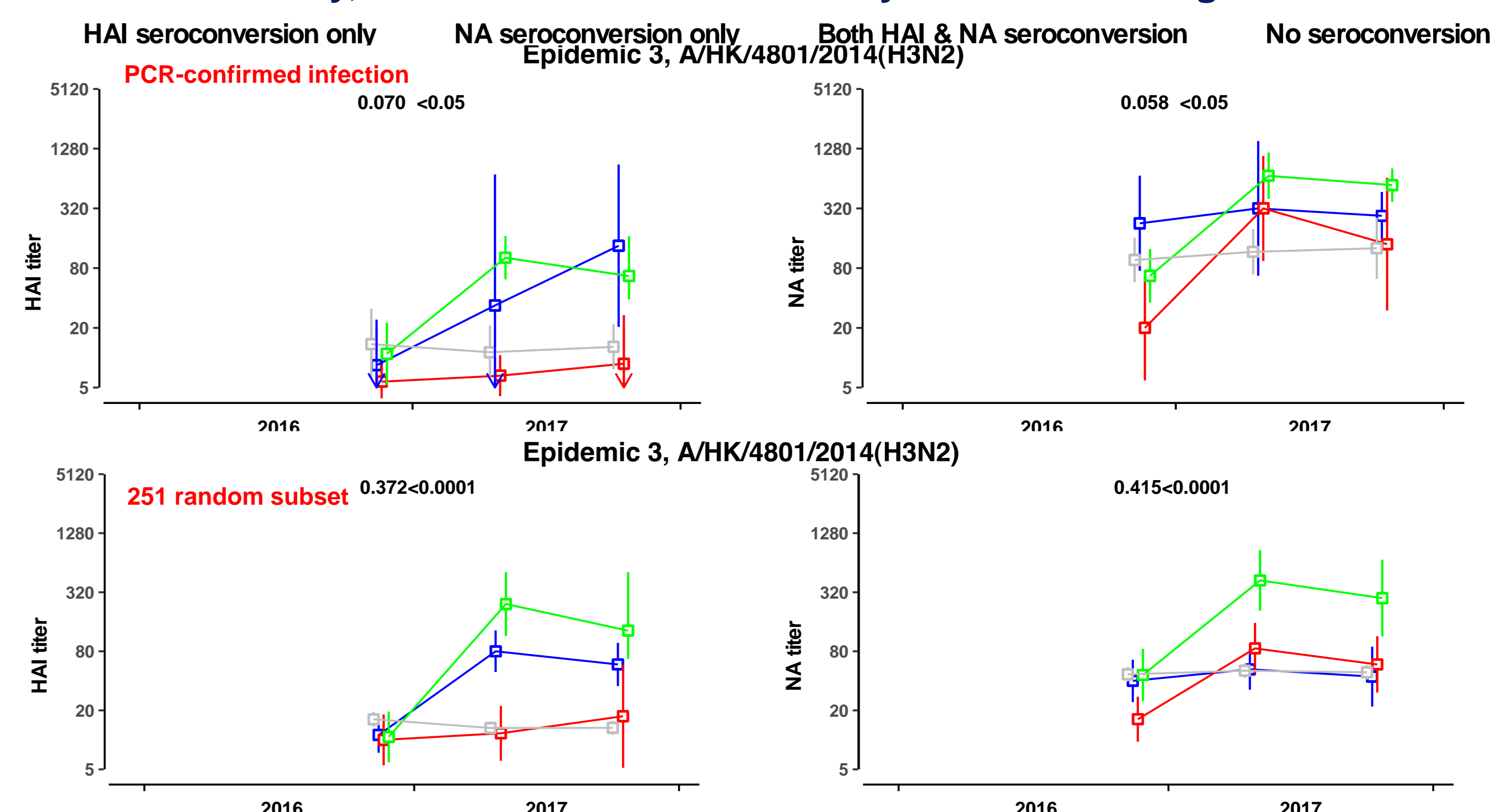
<sup>a</sup> Weighted by standardizing to the population age and sex of Jiangsu province, China in 2010.

- Lower pre-existing HAI and NAI titers** are associated with an increased risk of infection, particularly in relation to **symptomatic infection risk**;
- HAI** demonstrates **more conservative** in identifying PCR-confirmed infections through the **titer limitation (≤10)**, while in **wide range NAI titers**;
- Broader protection against strains in anti-NA antibody** with various levels, while **strain-specific in anti-HA antibody** mainly in lower levels



**Figure 2.** Correlation of pre-existing HAI and NAI titer against influenza A virus infection. Points indicated pre-existing/pre-epidemic antibody titer among different seroconversion, and arrows indicated pre- and post- HAI and NAI antibody trends and fold change.

- Pre-epidemic HAI and NAI titer were slightly higher in **no seroconversion** group but without statistically significant;
- More pronounced increase** in antibody levels in seroconversion in **both HAI and NAI assay**, and **NAI seroconversion may contribute to higher HAI titers**



**Figure 3.** Dynamics of antibody titer over time categorized by seroconversion in the random subset. Rectangles indicate geometric mean titer with arrow segments representing 95% confidence intervals at each timepoint.

## Conclusion

HAI testing is more effective in capturing a broader range of infections, and NAI can take as a supplementary method. Lower pre-existing HAI and NAI titers are associated with an increased risk of infection, particular for symptomatic infections. Our findings emphasize the potential value of incorporating pre-existing HAI testing into surveillance and risk assessment strategies, as well as considering the impact of NAI seroconversion on HAI antibody levels in vaccine development.

## References

[1] Cowling, BJ., et al., Cohort profile: the China Ageing REspiratory infection Study (CARES), a prospective cohort study in older adults in Eastern China. *BMJ Open* 2017; 7(10): e017503.

## Acknowledgements

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