

# Six-month persistence and safety of mRNA-based influenza or multicomponent influenza/COVID-19 vaccines versus licensed comparators in adults aged ≥18 years

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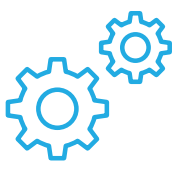
## BACKGROUND

- Seasonal influenza and SARS-CoV-2 viruses are global health concerns responsible for substantial disease burden<sup>1,2</sup>
- mRNA vaccine technology allows for quick updates in vaccine composition to match circulating strains due to the rapid manufacturing timeline<sup>3</sup>
- The investigational mRNA-1010 influenza vaccine contains mRNAs encoding haemagglutinin surface glycoproteins of seasonal influenza strains recommended by the World Health Organization<sup>4</sup>
  - In a phase 3 study among adults aged ≥18 years (P303; NCT05827978), a single dose of mRNA-1010 induced strong immune responses at Day 29 against seasonal influenza A and B strains that were superior to licensed standard-dose and high-dose influenza vaccines, with an acceptable safety profile<sup>4</sup>
- The investigational multicomponent vaccine mRNA-1083 combines the components of mRNA-1010 with a second-generation COVID-19 vaccine, mRNA-1283, which demonstrated noninferior efficacy than the first-generation COVID-19 vaccine (mRNA-1273) in adults aged ≥18 years in a phase 3 trial<sup>5</sup>
  - In a phase 3 study among adults aged ≥50 years (P301; NCT06097273), a single dose of mRNA-1083 induced strong immune responses at Day 29 against seasonal influenza A and B and SARS-CoV-2 strains, which were superior to licensed standard-dose influenza and COVID-19 vaccines in adults aged 50-64 years, and noninferior to licensed high-dose influenza and COVID-19 vaccines in adults aged ≥65 years, with an acceptable safety profile<sup>6</sup>



## OBJECTIVE

- To evaluate the 6-month immunogenicity and safety from two phase 3 studies of mRNA-based standalone vaccine against seasonal influenza (mRNA-1010) or multicomponent vaccine against seasonal influenza + COVID-19 (mRNA-1083)



## METHODS

### Study Design and Population

#### mRNA-1010 P303 Study (NCT05827978)

- The P303 trial was a 3-part, phase 3, randomised, observer-blind, active-controlled study conducted across 103 US sites
  - Part A: Participants aged ≥18 years were randomly assigned to receive either a single dose of mRNA-1010 or licensed standard-dose inactivated influenza vaccine, quadrivalent (SD-IIV4)
  - Part B: Participants aged 18-64 years were randomly assigned to receive either a single dose of mRNA-1010 or licensed SD-IIV4
  - Part C: Participants aged ≥65 years were randomly assigned to receive either a single dose of mRNA-1010 or licensed high-dose inactivated influenza vaccine, quadrivalent (HD-IIV4)

#### mRNA-1083 P301 Study (NCT06097273)

- The P301 trial was a phase 3, randomised, observer-blind, active-controlled study, including 2 age-cohort substudies, conducted across 146 US sites
  - Cohort A: Participants aged ≥65 years were randomly assigned to receive mRNA-1083 + placebo or licensed HD-IIV4 + COVID-19 vaccine
  - Cohort B: Participants aged 50-64 years were randomly assigned to receive mRNA-1083 + placebo or licensed SD-IIV4 + COVID-19 vaccine

### Immunogenicity Assessments

- Blood samples for immunogenicity assessments were collected from all participants at baseline (Day 1) and Day 29, and from a participant subset at Day 181
- Haemagglutination inhibition assay (HAI) assessed immune responses to vaccine-matched strains of influenza A/H1N1, A/H3N2, B/Victoria, and B/Yamagata, utilising cell-grown live viruses
- Pseudovirus neutralisation assay (PsVNA) assessed immune responses to a vaccine-matched strain of SARS-CoV-2 (XBB.1.5)

### Safety Assessments

- Solicited local or systemic adverse reactions (ARs) were recorded for 7 days after vaccination
- Unsolicited adverse events (AEs) were collected for 28 days after vaccination
- Serious AEs, AEs of special interest (AESIs), medically attended AEs, and AEs leading to study discontinuation were collected throughout study duration (to Day 181)



## RESULTS

### Study Populations

- Participant demographics in the mRNA-1010 and mRNA-1083 phase 3 studies were generally reflective of the US population and were well balanced across vaccine groups within each study part/cohort (**Table 1**)

**Table 1. Participant Demographics**

|                                            | mRNA-1010 study <sup>a,b</sup> |                                 |                       |                                 |                       |                     |
|--------------------------------------------|--------------------------------|---------------------------------|-----------------------|---------------------------------|-----------------------|---------------------|
|                                            | Part A: ≥18 years              |                                 | Part B: 18-64 years   |                                 | Part C: ≥65 years     |                     |
|                                            | mRNA-1010<br>N = 1220          | SD-IIV4<br>N = 1180             | mRNA-1010<br>N = 1492 | SD-IIV4<br>N = 1488             | mRNA-1010<br>N = 1502 | HD-IIV4<br>N = 1490 |
| Median age (min, max), years               | 50 (18, 86)                    | 50 (18, 91)                     | 49 (18, 64)           | 50 (18, 64)                     | 70 (65, 93)           | 70 (64, 93)         |
| Female                                     | 660 (54.1)                     | 623 (52.8)                      | 877 (58.8)            | 904 (60.8)                      | 878 (58.5)            | 852 (57.2)          |
| White                                      | 919 (75.3)                     | 851 (72.1)                      | 1012 (67.8)           | 966 (64.9)                      | 1255 (83.6)           | 1220 (81.9)         |
| Black/African American                     | 248 (20.3)                     | 270 (22.9)                      | 420 (28.2)            | 444 (29.8)                      | 224 (14.9)            | 235 (15.8)          |
| Not Hispanic/Latino ethnicity              | 937 (76.8)                     | 940 (79.7)                      | 1073 (71.9)           | 1079 (72.5)                     | 1037 (69.0)           | 1021 (68.5)         |
| Received influenza vaccine in prior season | 484 (39.7)                     | 457 (38.7)                      | 512 (34.3)            | 496 (33.3)                      | 488 (32.5)            | 496 (33.3)          |
|                                            | mRNA-1083 study <sup>a,c</sup> |                                 |                       |                                 |                       |                     |
|                                            | Cohort A: ≥65 years            |                                 | Cohort B: 50-64 years |                                 |                       |                     |
|                                            | mRNA-1083<br>N = 2011          | HD-IIV4 + mRNA-1273<br>N = 2006 | mRNA-1083<br>N = 1993 | SD-IIV4 + mRNA-1273<br>N = 2005 |                       |                     |
| Median age (min, max), years               | 70 (67, 74)                    | 70 (67, 74)                     | 58 (54, 61)           | 58 (54, 61)                     |                       |                     |
| Female                                     | 1078 (53.6)                    | 1098 (54.7)                     | 1156 (58.0)           | 1194 (59.6)                     |                       |                     |
| White                                      | 1577 (78.4)                    | 1565 (78.0)                     | 1374 (68.9)           | 1344 (67.0)                     |                       |                     |
| Black/African American                     | 370 (18.4)                     | 370 (18.4)                      | 516 (25.9)            | 551 (27.5)                      |                       |                     |
| Not Hispanic/Latino ethnicity              | 1688 (83.9)                    | 1689 (84.2)                     | 1576 (79.1)           | 1603 (80.0)                     |                       |                     |
| Received influenza vaccine in prior season | 1019 (50.7)                    | 1016 (50.6)                     | 784 (39.3)            | 784 (39.1)                      |                       |                     |
| Received COVID-19 vaccine in prior season  | 854 (42.5)                     | 853 (42.5)                      | 631 (31.7)            | 611 (30.5)                      |                       |                     |

HD, high dose; IIV4, inactivated influenza vaccine, quadrivalent; max, maximum; min, minimum; SD, standard dose. Data are presented as n (%) unless otherwise specified.

<sup>a</sup>The safety population consisted of all participants who were randomly assigned and received any study vaccine.

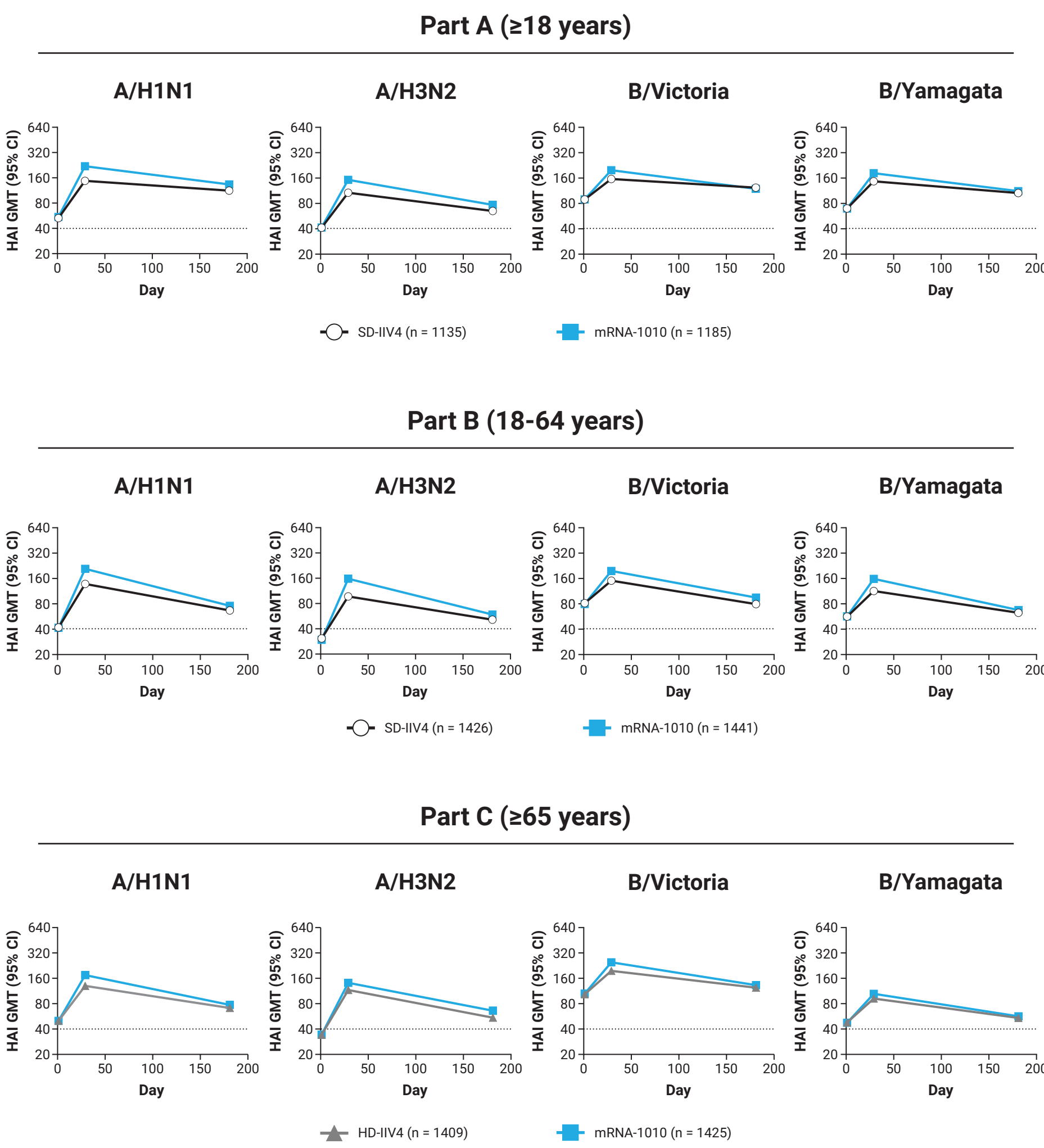
<sup>b</sup>Random assignment to vaccine groups in Part A was stratified by age group (18-49 years, 50-64 years, or ≥65 years) and by receipt of influenza vaccine in the prior 12 months (received or not received). Random assignment in Parts B and C was stratified by receipt of influenza vaccine since September 2022 (received or not received and, if received, whether or not it was from participation in the mRNA-1010-P302 study [NCT05566639]).

<sup>c</sup>In both age cohort substudies of the mRNA-1083 P301 study, random assignment to vaccine groups was stratified by receipt of influenza vaccine since September 2022 (received or not received). Additionally, random assignment in the ≥65 years substudy was stratified into 65-74 or ≥75 years age groups.

### mRNA-1010 Immunogenicity

- In Parts A-C of the mRNA-1010 study, HAI titres to all 4 vaccine-matched influenza strains remained above baseline through 6 months (Day 181) post-vaccination with mRNA-1010 or licensed standard-dose or high-dose influenza vaccines (**Figure 1**)
  - mRNA-1010 elicited HAI titres at Day 29 that were higher than those elicited by licensed SD-IIV4 or HD-IIV4 comparators for all vaccine-matched influenza strains; titres were numerically similar to or higher than comparators for all age groups at 6 months post-vaccination

**Figure 1. Influenza HAI GMTs Through 6 Months in the mRNA-1010 Study**

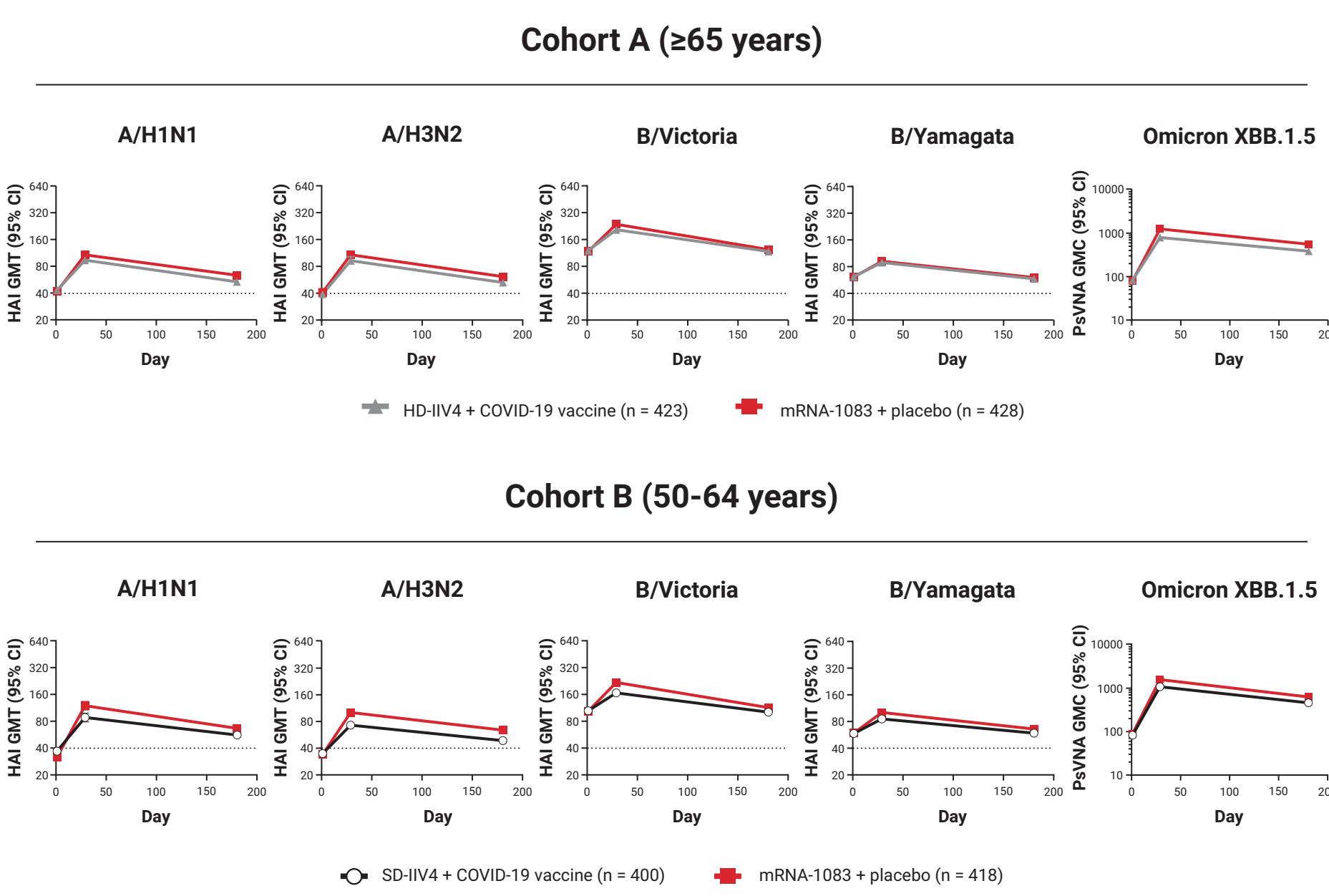


GMT, geometric mean titre; HAI, haemagglutination inhibition assay; HD, high dose; IIV4, inactivated influenza vaccine, quadrivalent; SD, standard dose. Per-protocol immunogenicity population. Horizontal dotted line indicates the HAI GMT associated with a 50% reduction in risk of influenza infection in healthy adults.

### mRNA-1083 Immunogenicity

- In Cohorts A and B of the mRNA-1083 study, HAI titres to all 4 vaccine-matched influenza strains remained above or similar to comparator through 6 months (Day 181) post-vaccination with mRNA-1083 (**Figure 2**)
- PsVNA concentrations to the vaccine-matched SARS-CoV-2 strain (XBB.1.5) remained higher than comparator through 6 months (Day 181) post-vaccination with mRNA-1083 in both cohorts

**Figure 2. Influenza HAI GMTs and SARS-CoV-2 Omicron XBB.1.5 PsVNA GMCs Through 6 Months in the mRNA-1083 Study**



GMC, geometric mean concentration; GMT, geometric mean titre; HAI, haemagglutination inhibition assay; HD, high dose; IIV4, inactivated influenza vaccine, quadrivalent; PsVNA, pseudovirus neutralisation assay; SD, standard dose. Per-protocol immunogenicity population subset who provided immunogenicity blood sampling at Day 181. Horizontal dotted line indicates the HAI GMT associated with a 50% reduction in risk of influenza infection in healthy adults.

### Safety

- Most solicited local or systemic ARs within 7 days after mRNA-1010 or mRNA-1083 vaccination were grade 1 or 2 in severity
  - Median duration of local or systemic ARs was 2-3 days across mRNA vaccine and active comparator groups
  - Injection site pain was the most commonly reported local AR; headache, fatigue, and myalgia were the most commonly reported systemic ARs
- In the mRNA-1010 study, the proportion of participants with any unsolicited AE of any cause occurring within 6 months post-vaccination was 23.0% (A), 18.5% (B), and 20.6% (C) for mRNA-1010, and 21.5% (A), 18.7% (B), and 19.5% (C) for active comparator
  - Few participants had serious AEs (2.2% [A], 1.7% [B], and 2.7% [C] for mRNA-1010; 1.8% [A], 1.5% [B], and 2.6% [C] for comparator)
  - AESIs classified by the investigator as having a relationship to the study vaccination were Bell's palsy in 1 (<0.1%) mRNA-1010 recipient (B) and face swelling in 1 (<0.1%) mRNA-1010 recipient (C); there were no related AESIs of myocarditis or pericarditis
- In the mRNA-1083 study, the proportion of participants with any unsolicited AE of any cause occurring within 6 months post-vaccination was 25.0% (A) and 18.9% (B) for mRNA-1083, and 24.9% (A) and 18.8% (B) for active comparator
  - Few participants had serious AEs (3.5% [A] and 1.5% [B] for mRNA-1083; 2.6% [A] and 1.3% [B] for comparator)
  - An AESI of confusional state in 1 (<0.1%) mRNA-1083 recipient (A) was classified by the investigator as having a relationship to the study vaccination; however, this was not a protocol-defined AESI and there were several confounders (ie, advanced age, history of post-traumatic stress disorder and hypertension, and concurrent SARS-CoV-2 infection)
  - There were no related AESIs of myocarditis or pericarditis



## CONCLUSIONS

- In these two phase 3 studies, mRNA-1010 (seasonal influenza) and mRNA-1083 (seasonal influenza + COVID-19) elicited robust humoral immune responses against vaccine-matched strains that persisted through 6 months post-vaccination, remaining comparable to or higher than licensed vaccine comparators
- For both mRNA-1010 and mRNA-1083 vaccines (N = 4214 and N = 4004, respectively), acceptable tolerability and safety profiles were observed over the 6-month period of follow-up
- These findings support the potential of mRNA-based standalone vaccination against seasonal influenza or multicomponent vaccination against seasonal influenza + COVID-19 and could represent a meaningful benefit for global public health

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### Disclosures

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