



Correlates of immunity in SARS-CoV-2 post-vaccine infections in the Prospective Assessment of **SARS-CoV-2 Seroconversion (PASS) Study**

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Background

We sought to determine pre-infection correlates of immunity against SARS-CoV-2 post-vaccine infections in a prospective cohort of healthcare workers.

Inclusion criteria:

- \geq 18 years old
- Generally healthy
- Works at Walter Reed National Military Medical Center (WRNMMC)

hjf

Exclusion criteria: • Severely immunocompromised

- History of COVID-19
- SARS-CoV-2 seropositive at time of study entry

Research Design

- Baseline peripheral blood mononuclear cells (PBMCs), serum, plasma, saliva
- Monthly serum antibody testing for SARS-CoV-2 (changed to quarterly in Fall 2021) • PBMCs drawn at scheduled intervals
- PCR testing at the WRNMMC COVID-19 center every time a participant has symptoms of viral respiratory infection
- Baseline and then serial symptom & risk exposure/PPE use/social distancing questionnaires

Methods

Single-center, observational cohort of 183 healthcare workers.

Testing for ancestral (WT) SARS-CoV-2 Spike (S)-specific IgG antibodies was performed on serum collected during the Fall 2021 visit and conducted using a microsphere-based multiplex immunoassay (research assay) interpolated against an internal standard curve for binding antibody units and then calibrated to the US National Standard (readout: WHO Binding Antibody Units/ml - BAU/ml)

Testing for WT SARS-CoV-2 S-specific IgG, IgA and secretory IgA was performed on saliva collected during the same visit and using the same microsphere-based multiplex immunoassay interpolated against an internal standard curve for binding antibody units/ml.

Serum samples were also tested using a commercial assay in collaboration with Quest Diagnostics (Elecsys[®] anti-SARS-CoV-2 S for use on the Roche Cobas analyzer). This electrochemiluminescence immunoassay is intended for qualitative detection of total antibodies to SARS-CoV-2 S protein receptor binding domain (RBD).

Neutralizing antibody titers against the wild-type virus (D614G) were determined by microneutralization assays, and against the Delta variant (B.1.617.2), and 2 Omicron subvariants (BA.1, BA.1.1) by lentiviral pseudovirus neutralization assays.

20 participants hadn't

STROBE flowchart of the study cohort.

Participants screened by		Participants enrolled in PASS study by March	doses of vax 2021	by Dec 1,	Participants that received at least 2 doses of vaccine by December 1, 2021 n = 251			
March 30, 2021 n = 284	13 individuals excluded	30, 2021 n = 271		F				
	 1 prior symptomatic COV 8 positive SARS-CoV-2 Ige 4 BMI > 40* * removed as an exclusion criteria Oct 19, 2021 	/ID G	48 participants were not seen between Oct 1 and Dec 15, 2021					
			Participants that were seen between Oct 1 and Dec 15, 2021 n = 203					
	ž	20 participants excluded from manuscript analyses (received a booster dose after the late fall visit and before Apr 1, 2022)						
			Participants with a post-vaccination infection n = 35 Participants with no post-vaccination infection n = 148 All participants included in the study n = 183					





A) Comparison of anti-Spike (S)-saliva IgG (AU/ml) between the uninfected group (n=147) and the post-vaccination infection group (n=35) using the research assay. P value determined using the Mann-Whitney U test (ns p = 0.1195). B) Comparison of anti-Spike (S)-saliva IgA (AU/ml) between the uninfected group (n=147) and the post-vaccination infection group (n=35) using the research assay. P value determined using the Mann-Whitney U test (ns p = 0.4731). C) Percentages of uninfected vs post-vaccination infected participants depending on the saliva levels of anti-S IgG (AU/ml). D) Percentages of uninfected vs post-vaccination infected participants depending on the saliva levels of anti-S IgA (AU/ml). E) Spearman correlation (p) to determine whether the anti-S serum IgG correlated with the anti-S saliva IgG. F) Spearman correlation (p) to determine whether the anti-S serum IgA correlated with the anti-S saliva IgA.

Pre-infection neutralization titers against Omicron subvariants provide robust correlation with protection against post-vaccine infections



Neutralization assays used lentiviral pseudoviruses bearing SARS-CoV-2 S proteins from WT D614G (ns p=0.0627), Delta variant (* p=0.0379) or Omicron subvariants BA.1 (* p=0.0114) and BA.1.1 (** p=0.0074)(n = 144 uninfected group; n= 34 post-vaccination infection group). P values determined using the Mann-Whitney U test.

COVID-19 risk by immunologic marker. The pre-Omicron wave covariates were adjusted for age, sex and home risk score. Odds ratios (OR) represent the relative change in the odds of infection corresponding to an increase in the independent variable of 1/10 of its range. CI, confidence interval; AUC, area under the curve; ID50, 50% inhibitory dilution. ROC analyses (not shown) showed that a 90% protection was associated with anti-S IgG serum levels ≥ 7200 BAU/ml, B.1.617.2 ID50 > 1:1200 and BA.1.1 ID50 > 1:1400. An 80% protection was associated with anti-S IgG serum levels ≥ 4900 BAU/ml, B.1.617.2 ID50 > 1:1100 and BA.1.1 ID50 > 1:900.

vaccination infection group); Home precautionary score, ns p=0.5705 (n=144 uninfected group; n=35 post-vaccination infection group). P values determined using the Mann-Whitney U test.

Pre-Omicron epidemic serum anti-S IgG antibody levels, and neutralization titers against the Delta variant B.1.617.2 and Omicron subvariant BA.1.1 are strong covariates of protection against post-vaccine infections

Immunologic Marker	OR	Lower Cl	Upper Cl	AUC	P-value
Serum Binding IgG to Spike (BAU/ml)	0.655	0.467	0.918	0.919	0.014
Serum Binding IgA to Spike (AU/ml)	0.346	0.079	1.513	0.908	0.159
Roche Serum anti-S total Igs (BAU/ml)	0.720	0.497	1.041	0.903	0.081
ID50 neutralizing titer BA.1.1	0.476	0.239	0.945	0.914	0.034
ID50 neutralizing titer D614G	0.308	0.095	1.000	0.907	0.050
ID50 neutralizing titer B.1.617.2	0.365	0.134	0.995	0.907	0.049
ID50 neutralizing titer BA.1	0.663	0.418	1.051	0.908	0.081
Saliva Binding IgG to Spike (AU/ml)	0.972	0.770	1.227	0.887	0.811
Saliva Binding IgA to Spike (AU/ml)	0.790	0.329	1.898	0.891	0.597

Conclusions

Anti-spike IgG binding antibody (bAb) serum levels correlate with protection against symptomatic infection during the 1st Omicron wave, but complete protection was only observed with very high levels (> 15,000 BAU/ml).

There is a high correlation between the research assay and the commercial Roche assay, raising the prospect of translational clinical implementation.

Spike IgG bAb serum levels correlate with reduced duration of symptoms and fewer symptoms in the nasal, eyes, and body/systemic domains.

Pre-Omicron epidemic saliva levels of anti-spike IgG and IgA antibodies do not show statistically significant correlation with protection against post-vaccine infection.

Neutralization titers against the Omicron subvariants provide robust correlation of protection against post-vaccine infection, but are not markedly better correlates than binding antibody levels.

The Home risk score provides a robust correlate of protection against post-vaccine

Regression analysis for immunologic markers after adjustment for non-immunologic markers (sex, age, and home risk score) shows that pre-Omicron wave Spike IgG bAb serum levels (research assay) and neutralization titers against the Delta variant B.1.617.2 and Omicron subvariant BA.1.1 are strongly correlated with protection against post-vaccine infection (p=0.014, p=0.049, and p=0.034, respectively)

Acknowledgments

The investigators gratefully acknowledge all research participants for their many contributions to the PASS study.

The views expressed in this presentation are the sole responsibility of the presenter and do not necessarily reflect the views, opinions, or policies of the Uniformed Services University of the Health Sciences, the Department of Defense, the Department of the Navy, Army, Air Force, the United States Government, or the Henry M. Jackson Foundation for the Advancement of Military Medicine, Inc. (HJF). Mention of trade names, commercial products, or organizations does not imply endorsement by the U.S. Government. The study protocol was approved by the USUHS Institutional Review Board in compliance with all applicable federal regulations governing the protection of human subjects. The HJF, in support of the USU IDCRP, was funded to conduct or augment unrelated Phase III Mab and vaccine trials as part of US Govt. COVID19 response. The authors declare no competing

The protocol was executed by the Infectious Disease Clinical Research Program (IDCRP), a Department of Defense (DoD) program executed by the Uniformed Services University of the Health Sciences (USUHS) through a cooperative agreement by the Henry M. Jackson Foundation for the Advancement of Military Medicine, Inc. (HJF). This work was supported in whole, or in part, with federal funds from the Defense Health Program (HU00012020067, HU00012020094) and the Immunization Healthcare Branch (HU00012120104) of the Defense Health Agency, United States Department of Defense. The sponsors had no involvement in the study design, the collection of data, the analysis of data, the interpretation of data, the writing of the report, or in the decision to submit the article for publication.

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