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Supplemental information

Systemic IL-15, IFN-γ, and IP-10/CXCL10 signature

associated with effective immune response

to SARS-CoV-2 in BNT162b2 mRNA vaccine recipients

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	COVID Symptoms	COVID PCR test	COVID Ab test
Recipient 1	March, 2020	Not done	Not done
Recipient 2		Not done	Not done
Recipient 3	Fever, fatigue, headache	Positive (08/13/2020)	Positive (September, 2020)
Recipient 4	none		Positive (May, 2020)
Recipient 5	Flu-like symptoms		Positive (June, 2020; December, 2020)
	(May 2020)		

Table S1. Relevant history of vaccine recipients with pre-existing CoV-2 immunity, Related to Figure 1.

Analytes with changes (n=19)	CRP (C-Reactive Protein)		
	Eotaxin (CCL11)		
	IFN-γ		
	IP-10 (CXCL10)		
	IL-1Ra		
	IL-3		
	IL-6		
	IL-7		
	IL-8		
	IL-12/IL-23p40		
	IL-15		
	IL-16		
	MCP-1 (CCL2)		
	MDC (CCL22)		
	MIP-1 α (CCL3)		
	MIP-1 β (CCL4)		
	SAA (serum amyloid A)		
	TNF-α		
	VEGF-A		
Analytes with minimal or no change (n=8)	Eotaxin-3		
	ICAM-1		
	IL-17C		
	IL-17B		
	MCP-4		
	TARC		
	TSLP		
	VCAM-1		
Analytes below detection threshold (n=14)	IL-1a		
	IL-1β		
	IL-2		
	IL-4		
	IL-5		
	IL-9		
	IL-10		
	IL-12p70		
	IL-13		
	IL-17A		
	IL-17A/F		
	IL-17D		
	GM-CSF		
	TNF-β		

Table S2. Cytokines and Chemokines tested, Related to Figures 2, 3, 4, and 5.

Adverse effects	Vaccination 1	Vaccination 2	
	(40 vaccinees, %)	(50 vaccinees, %)	
injection site pain	32 (80%)	38 (76%)	
fatigue	4 (10%)	9 (18%)	
headache	2 (5%)	7 (14%)	
myalgia		9 (18%)	
fever, shivers		9 (18%)	
diarrhea	1 (2.5%)		
axillar pain	1 (2.5%)		
dizziness	1 (2.5%)	2 (4%)	

Table S3. Adverse effects elicited by mRNA vaccination, Related to Figures 2, 3, 4, and 5.

	Spike-RBD Antibody		
	Spearman r		
	correlation	<i>p</i> value*	
	coefficient		
MIP-1a (CCL3)	0.432	0.001	
IL-1Ra	0.334	0.013	
MCP-1 (CCL2)	0.241	0.074	
ΜΙΡ-1β (CCL4)	0.322	0.015	
CRP (C-Reactive Protein)	0.167	0.219	
SAA (serum amyloid A)	0.271	0.043	
IP-10 (CXCL10)	0.145	0.285	
IL-6	0.21	0.121	
TNF-α	0.103	0.451	
IFN-γ	0.433	0.001	
IL-15	0.389	0.003	
IL-3	-0.145	0.29	
IL-7	0.24	0.075	
IL-8	0.114	0.402	
Eotaxin (CCL11)	0.120	0.377	
MDC (CCL22)	0.394	0.003	
IL-12/IL-23p40	0.353	0.008	
IL-16	0.248	0.066	
VEGF-A	0.303	0.023	

Table S4. Correlations of anti-Spike antibodies at day 36 and cytokines log2 fold changes upon 2nd vaccination (d23-d22), Related to Figure 6.

* These univariate correlations were determined using a cut-off for a Spearman correlation coefficient corresponding to p value <0.05 (non-adjusted for multiple comparisons).



Figure S1. Correlations of Spike and Spike-RBD antibodies, Related to Figure 1.

Correlations of antibodies against full length trimeric Spike (ELISA, endpoint titer in log) and Spike-RBD (ROCHE, log) at day 22 (A), day 36 (B) and day 50 (C). Spearman r and p values are given (GraphPad Prism).



Figure S2. Correlation of binding and neutralizing antibodies, Related to Figure 1.

NAb assays were performed in using a HIV- $1_{NL}\Delta Env$ -NanoLuc derived pseudotype neutralization assay in HEK-ACE2 cells using Wuhan-Hu-1 Spike. using sera from a subset of naïve vaccinees (day 36) selected to cover a range of low to high responses. Correlations of (A) Spike-RBD (ROCHE, log) and (B) antibodies against full length trimeric Spike (ELISA, endpoint titer in log) and pseudotype NAb measured at day 36 (samples from Figure 1). Spearman r and p values are given (GraphPad Prism).



Figure S3. Additional serum cytokine/chemokine levels affected by COVID-19 vaccination, Related to Figure 2. Cytokine and chemokine levels were measured over time using the MSD assay as described in Figure 2 for the 58 COVID-19 naïve vaccine recipients. Serum levels of 8 of 19 analytes showing changes upon vaccination are plotted over time.



Figure S4. Differential levels of cytokine in male and female vaccine recipients upon COVID-19 vaccination, Related to Figure 3.

Comparison of log2 fold changes after the 2^{nd} vaccination (d23_d22) in serum of (A) IFN- γ , (B) IL-15, (C) IL-6 and (D) IP-10/CXCL10 which are significantly higher in female vaccine recipients (N=33) to male vaccine recipients (N=23) at one day after the 2^{nd} vaccination (d23). P values are from unpaired non-parametric t test (Mann-Whitney).





Correlations of log2 fold changes of selected analytes after the 1st vaccination are shown. r and non-adjusted p value for the individual correlations are shown in the plots.



Figure S6. Coordinated vaccine response featuring inflammatory molecules and chemokines, Related to Figure 5.

Correlations of log2 fold changes of selected analytes after the 2^{nd} vaccination are shown derived from data depicted in heatmap in Figure 5A (grey box). r is shown in plots; all correlations are characterized by an adjusted p<0.05.