

“SARS-CoV-2 vaccination induces immunological memory able to cross-recognize variants Alpha to Omicron”



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Article

SARS-CoV-2 vaccination induces immunological T cell memory able to cross-recognize variants from Alpha to Omicron

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Spike-specific T cell reactivity to SARS-CoV-2 variants in vaccinated individuals

SARS-CoV-2



Alpha
(B.1.1.7)



Beta
(B.1.351)



Gamma
(P.1)



B.1.1.519



Kappa
(B.1.617.1)



Delta
(B.1.617.2)



Lambda
(C.37)



R.1

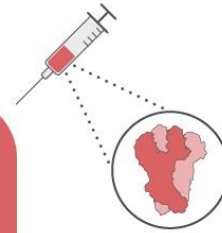


Mu
(B.1.621)



Omicron
(B.1.1.529)

Vaccinated



- mRNA-1273
- BNT162b2
- Ad26.COVS.S
- NVX-CoV2373

2 week post-vax
3-4 month post-vax
6 month post-vax

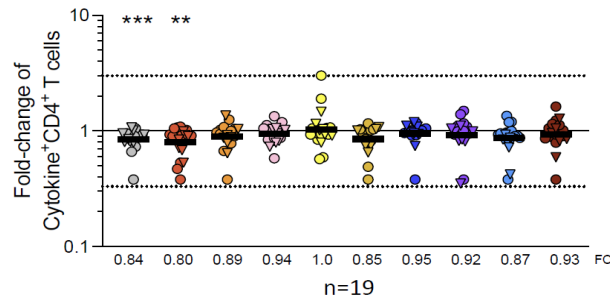
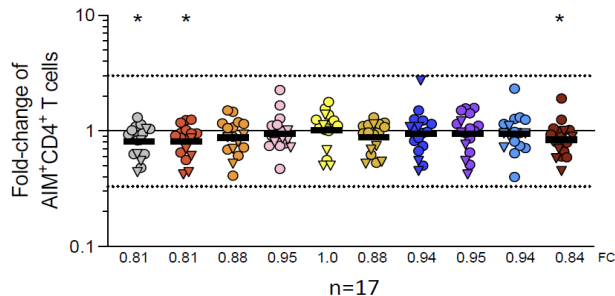


CD4 : Avg. 90%

CD8 : Avg. 87%

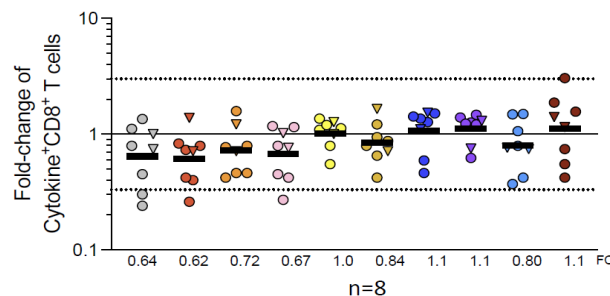
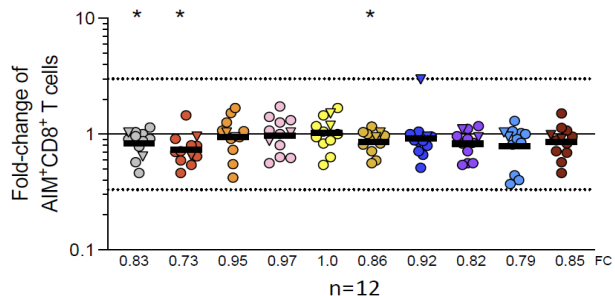
Cross-recognition of SARS-CoV-2 variants by memory T cells

Omicron 84%



SARS-CoV-2 variant:

- Alpha (B.1.1.7)
- Beta (B.1.351)
- Gamma (P.1)
- B.1.1.519
- Kappa (B.1.617.1)
- Delta (B.1.617.2)
- Lambda (C.37)
- R.1
- Mu (B.1.621)
- Omicron (B.1.1.529)

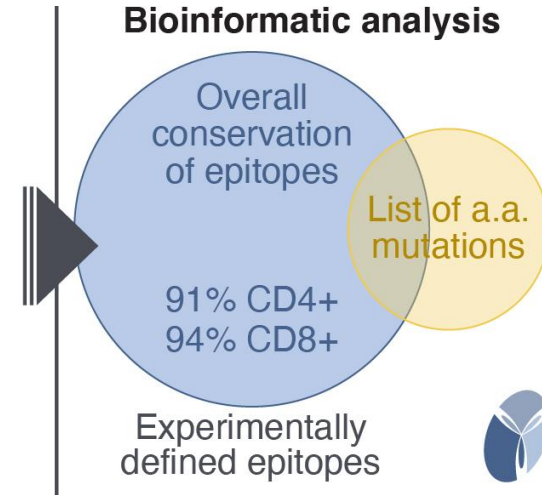
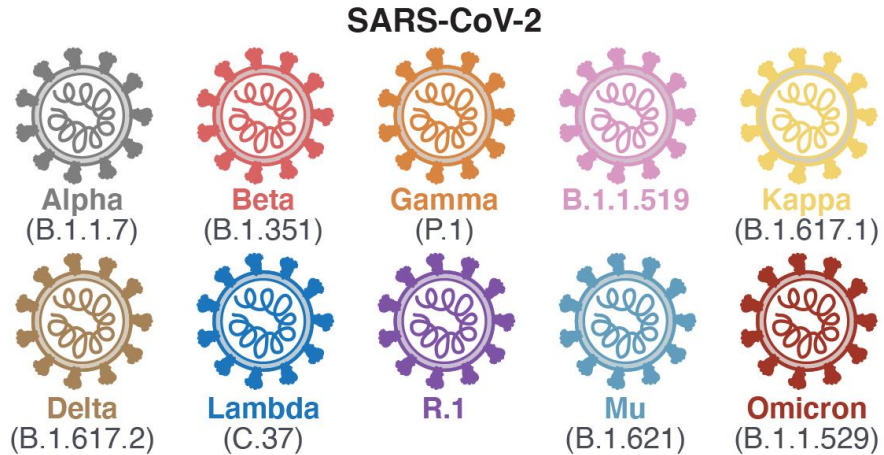


COVID-19 vaccine:

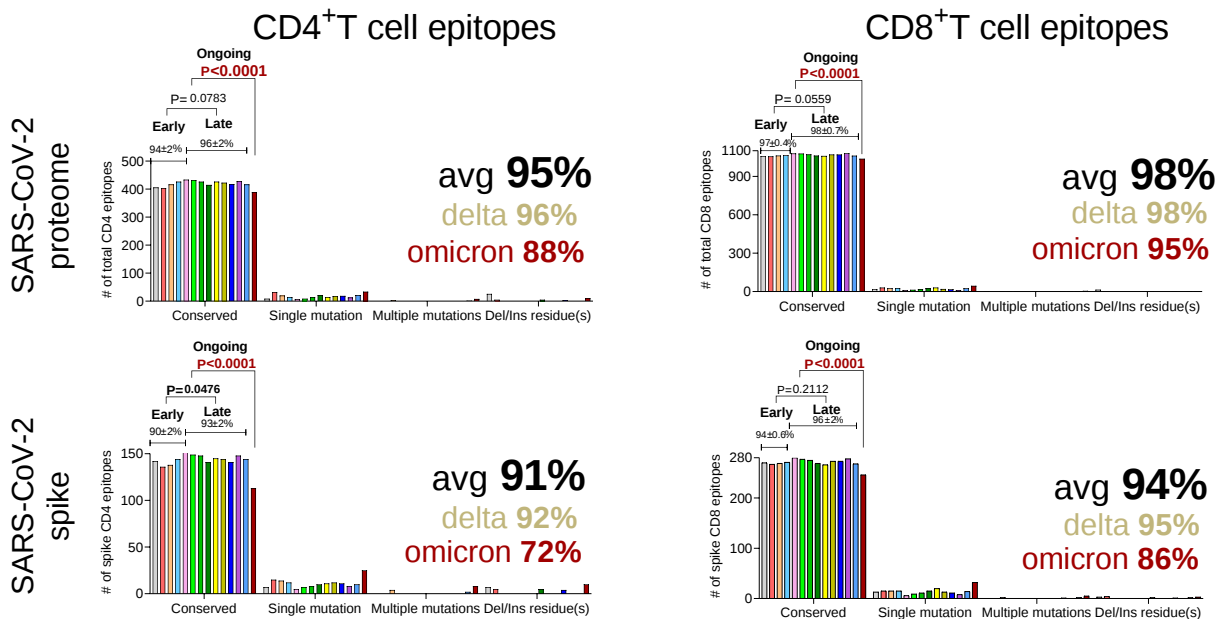
- mRNA-1273
- ▽ BNT162b2

Omicron 85%

Predicted impact of SARS-CoV-2 variants on T cell epitopes

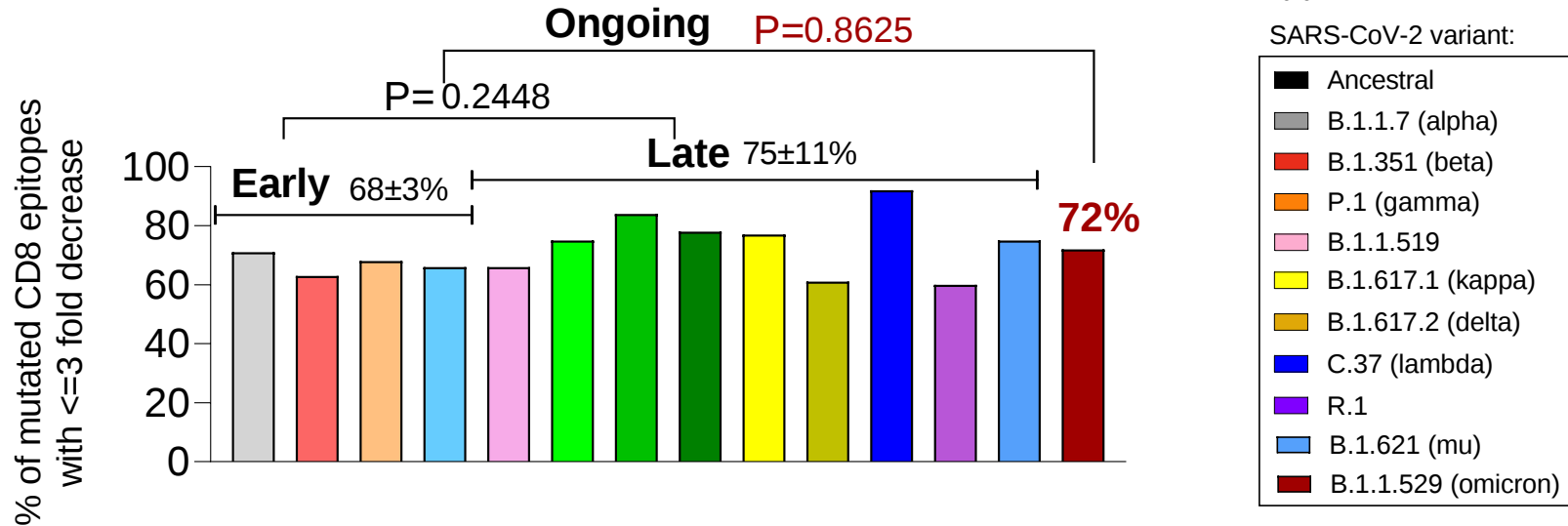


Conservation of T cell epitopes in Omicron

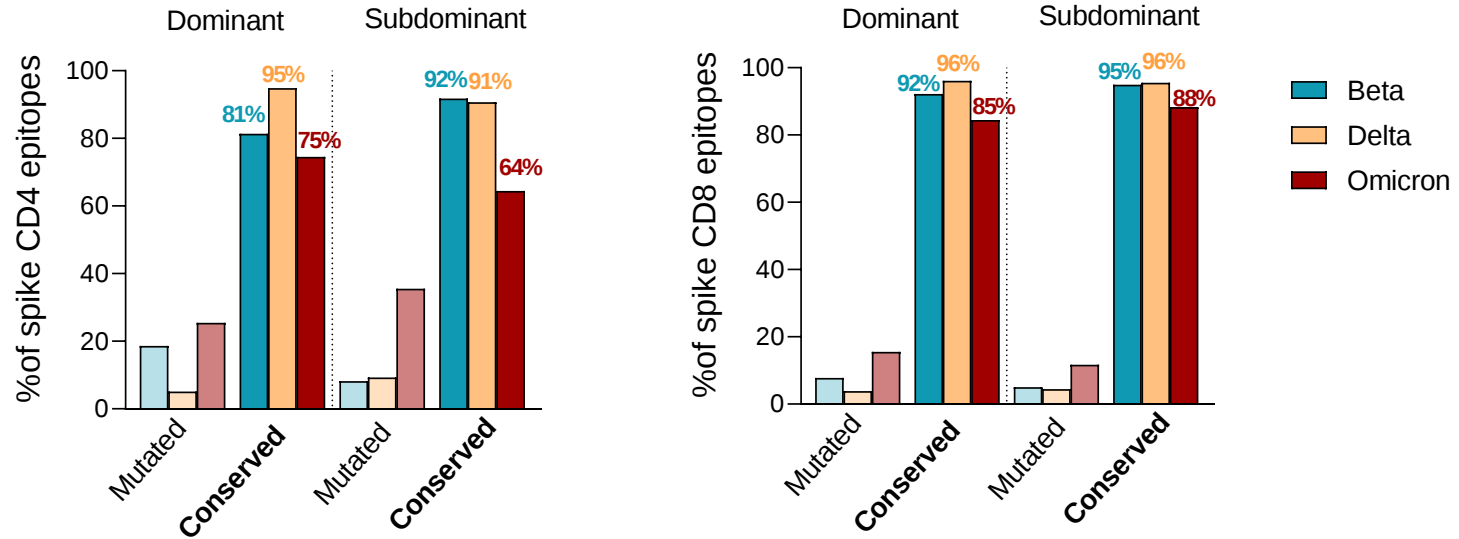


B.1.1.7 (alpha)	B.1.351 (beta)	P.1. (gamma)	B.1.427/429 (epsilon)	Early
B.1.1.519	B.1.525 (eta)	B.1.526 (iota)	B.1.526.1	Late
B.1.617.1 (kappa)	B.1.617.2 (delta)	C37 (lambda)	R1	
B.1.1529 (Omicron)				Ongoing

The majority of mutated CD8+T cell epitopes in Omicron are still able to bind HLA class I molecules

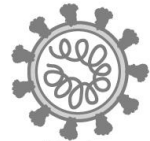


Dominant CD4 T cell epitopes are more conserved in Omicron



Breadth of T cell epitope repertoire and its conservation in variants

SARS-CoV-2



Alpha
(B.1.1.7)



Beta
(B.1.351)



Gamma
(P.1)



B.1.1.519



Kappa
(B.1.617.1)



Delta
(B.1.617.2)



Lambda
(C.37)



R.1



Mu
(B.1.621)



Omicron
(B.1.1.529)



CD4+ T cells

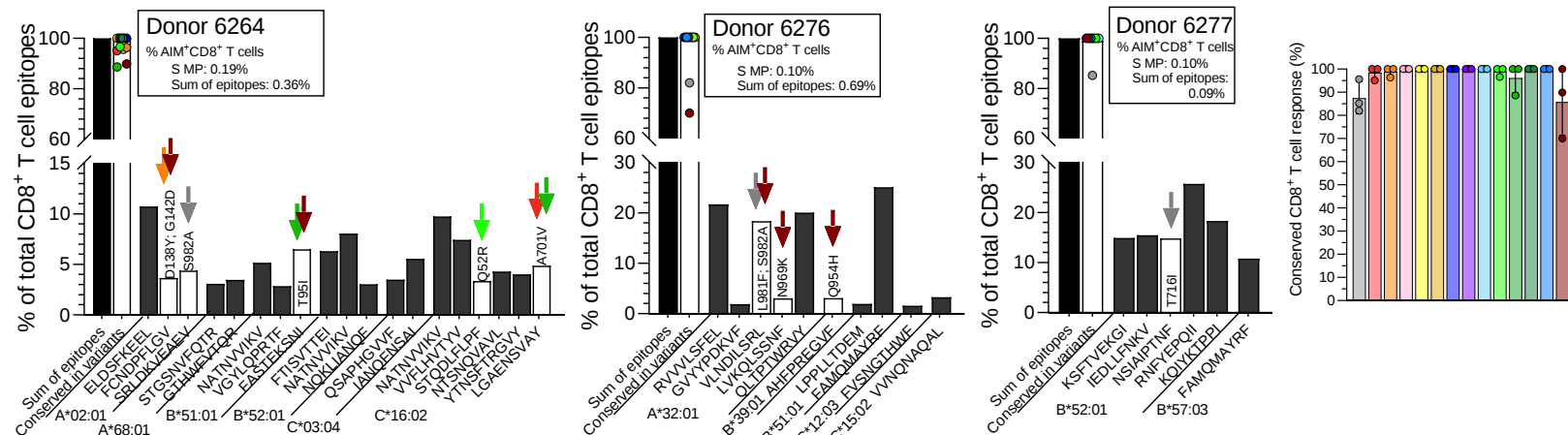


CD8+ T cells

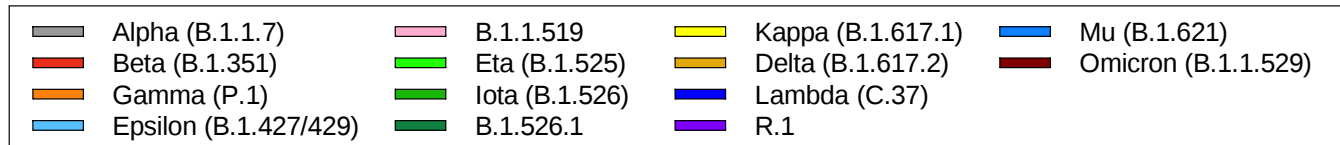


Broad epitope repertoire

CD8+ T cell epitope repertoire



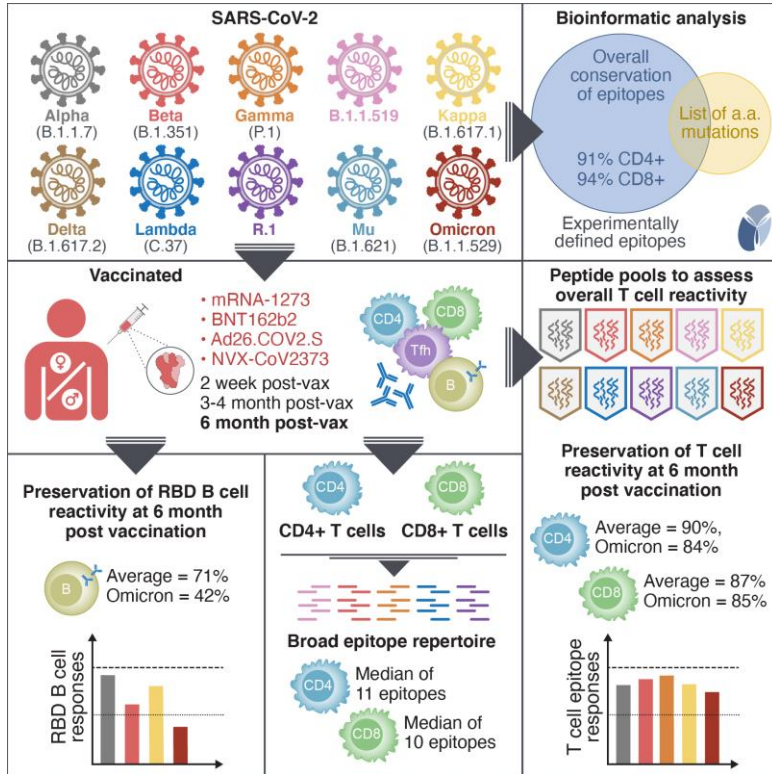
SARS-CoV-2 variant:



COVID-19 vaccine:



Cross-recognition of SARS-CoV-2 variants



➤ **Bioinformatic analysis show the majority of spike epitopes are fully conserved**

- CD4 : Avg. 91% Omicron 72%
- CD8 : Avg. 94% Omicron 86%
- 74% of mut CD8 epitopes are predicted to still bind HLA

➤ **B cells recognition is significantly reduced**

- (Avg. 71%, Omicron 42%)

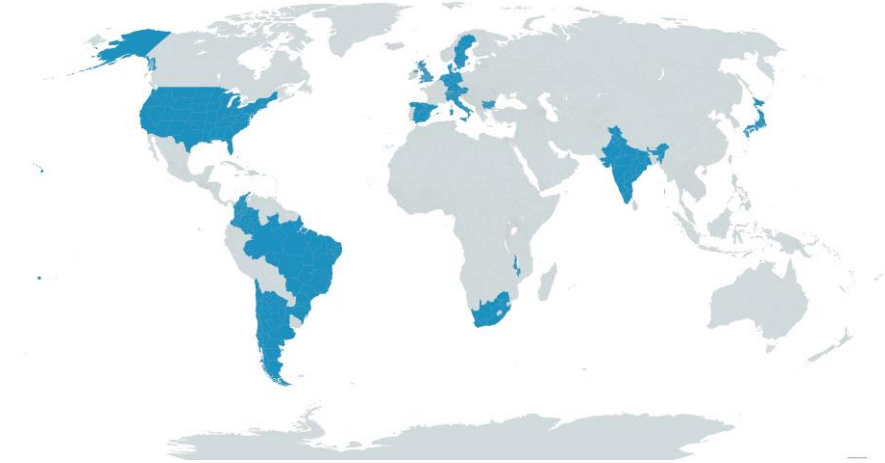
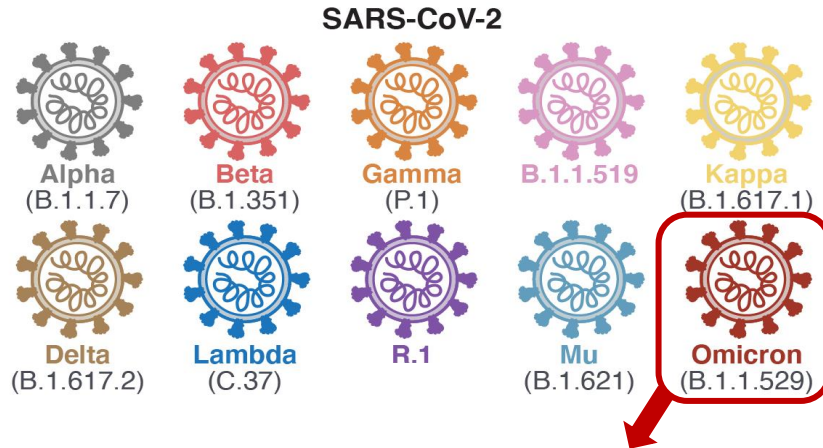
➤ **T cells of vaccinees cross-recognize SARS-CoV-2 variants**

- Responses are also retained at the ICS level
- Irrespective of vaccine platform
- Effective recognition also after a first shot
- Memory recognition:
 - CD4 : Avg. 90% Omicron 84%
 - CD8 : Avg. 87% Omicron 85%

➤ **T cell epitope repertoire**

- A median of 11 CD4 and 10 CD8 spike epitopes are
- Avg preservation > 80% for Omicron at the epitope level.

Worldwide collaboration sharing spike variant pools



❑ Madelon et al., medrxiv 2021; [Switzerland](#)

Christiane Eberhardt

❑ Keeton et al., Nature 2022; [South Africa](#)

Wendy Burgers

❑ GeurtsvanKessel et al., 2022 Sci Imm; [Netherlands](#)

Rory DeVries/ Bart Haagmans

❑ Gao et al., Nat Med 2022; [Sweden](#)

Marcus Buggert