Global number of reported smallpox cases

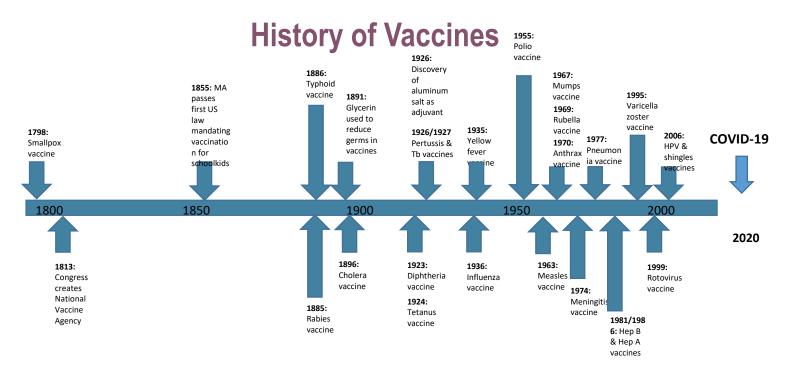


The number of reported cases is lower than the number of actual cases. This is due to limited testing and reporting and challenges in the attribution of the cause of death.



Source: World Health Organization (2011)

about 250 years ago life expectancy started to increase. In 1900, mankind had already reached a life expectancy of approximately 50 years (3). Nowadays, a child born in a high-income country can expect to live 85 years. The additional 35 years of life that we gained during the last century are substantially due to the conquest of infectious diseases, which used to kill 50% of people before the age of 20. These were viral diseases such as smallpox, rabies, measles,



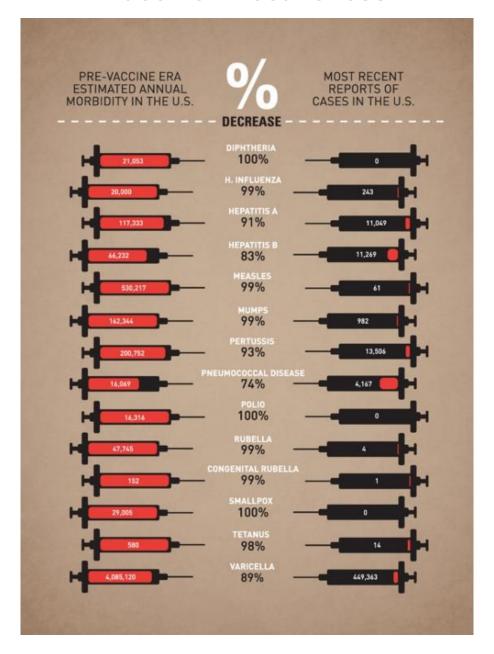
^{*}Note that this timeline is abbreviated to give an overview of vaccine development and the recent explosion of discovery

IL SISTEMA IMMUNITARIO 2048 IMMUNITA' INNATA IMMUNITA ADATTIVA Immunologic memory 1024 Mean antibody titer 512 256 128 64 32 16 Serum antibody Nature Reviews | Cancer 12 Time, months Vaccine Muscle Vaccine antigen Adjuvant (containing danger signals) Dendritic Memory B cell proliferation Maturation of the MHC class II Soluble antibody response vaccine Peptide of antigen vaccine antigen CD4+T cell BCR Activation and trafficking to draining lymph T cell help Proliferation Plasma cell differentiation and antibody production node TCR-B cell Tcell MHC help class II CD8 effector T cell Bone marrow MHC class I CD8 CD8° T cell memory Tcell Long-lived plasma cell

Sorveglianza

- Impatto della vaccinazione
- Effectiveness del vaccino
- Effetti della pressione selettiva prodotta dal vaccino, con eventuale emergenza di ceppi che sfuggono all'effetto del vaccino
- Sicurezza, eventi avversi, reazioni avverse

Vaccine Effectiveness



Eventi avversi e reazioni avverse

Evento avverso (Adverse Event Following Immunization, AEFI)

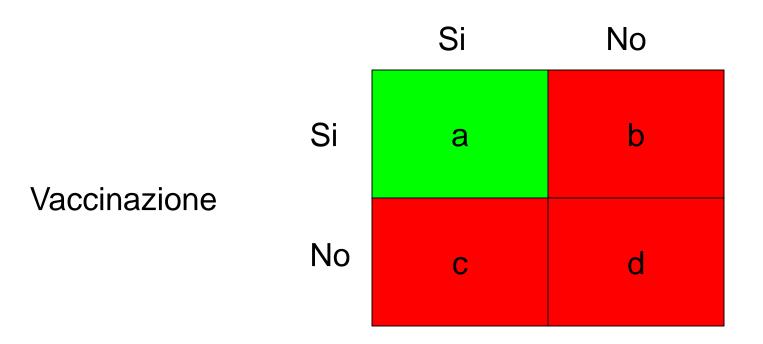
Evento indesiderato dopo vaccinazione (non necessariamente è presente una relazione causale)

Reazione avversa

Evento indesiderato di cui è dimostrata la relazione causale con la vaccinazione

Necessario distinguere tra relazione causale (= rapporto causa-effetto) e temporale (= dovuta al caso)

Evento



Frequenza dell'evento nei non vaccinati =
$$\frac{c}{c + d}$$

La pandemia di COVID-19 in Italia

Giampiero Carosi, Roberto Cauda, Andrea Pession, Guido Antonelli

Questo capitolo è stato scriito da autori italiani che hanno lavorato in autonomia rispetto agli elitor dell'edizione originale inglese. Pertanto McGraw-Hill e gli dotto dell'edizione originale non hanno nessuna responsabilità circa di contenuto del capitolo, la cui pubblicazione in formato elettronico e in esclusivus per l'edizione italiana è stato auto-rizzata da McGraw-Hill.

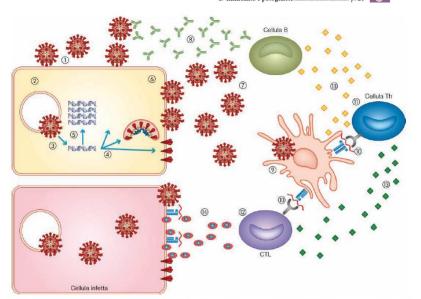
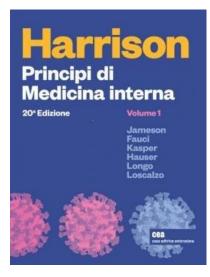
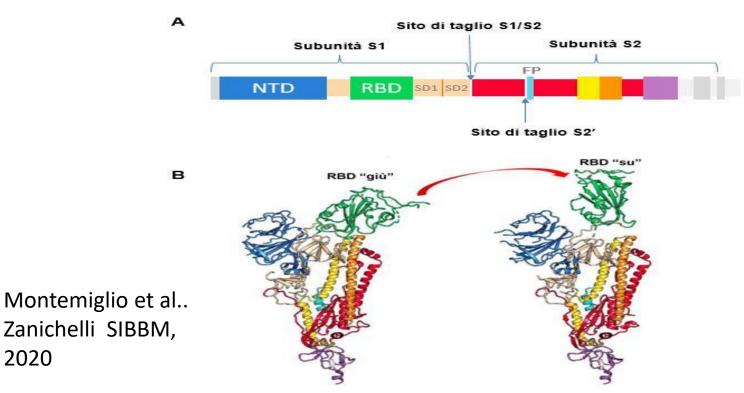


Figura 19 Principali passaggi del ciclo vitale virale e dello sviluppo della risposta immunitaria. (1) Adesione del virione di SARS-CoV-2 alla superficie cellulare tramite il recettore cellulare ACE2. (2) Ingresso nella cellula. Le proteine virali possone essere riconosciute dai recettori dell'immunità innata (per es. TIR3, TL4 e TL7), potando al rilascio dei pattern moleccian associati al pericolo, all'attivazione della reproteina e rilascio dell'RNA all'interno della cellula. (4) Traduzione dell'RNA per produrre le proteine virali. (5) L'RNA viene copiato e unito alle proteine del unicocapasio. (6) Assemblamento dei virioni figlii. (7) Riconoscimento della diplicoproteina del nucleocapasio. (6) Assemblamento dei leganti la glicoproteina spike e particori presentaria del nucleocapasio. (8) Assemblamento dei leganti la glicoproteina spike e particori presentaria del nucleocapasio. (8) Pasarone dell'RNA per si periodi della proteine stituturali en nu (11) Attivazione del CTL (13) Il linicota Tb produce citichine (principalmente IRN-y, L-2 e TNF-a). (14) Riconoscimento e Alling delle cellule infettate da parte dei CTL. (Modificata de Poland G.A. et al., SARS-CoV-2 immunity: review and applications to phase 3 vaccine candidates. Lancet. 396(10262): 1595-1609. (14) noembre 2020.)

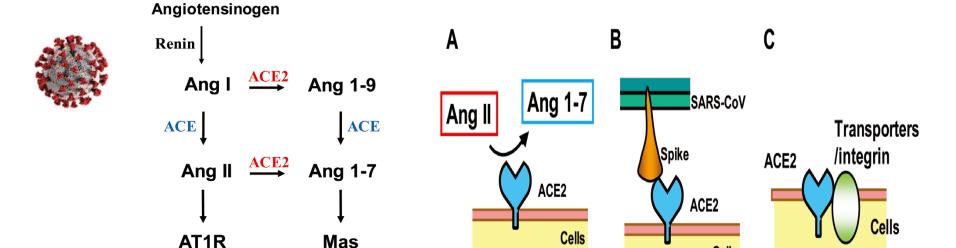


Spike e RBD di SARS-CoV-2





ACE, ACE 2 functions and SARS-CoV-2



Cells

Fig 1b

Fig1a

Cassone A, Gucciardo D, Cauda R.Pathog Glob Health.
2020 May 18;114(4):165-167.

Vasodilatation

Vasoprotection

Cardioprotection

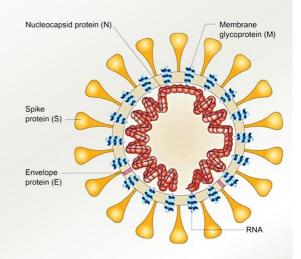
Vasoconstriction

Hypertension Cardiac hypertrophy

Infection-induced immunity to COVID-19

Infection-induced immunity to COVID-19:

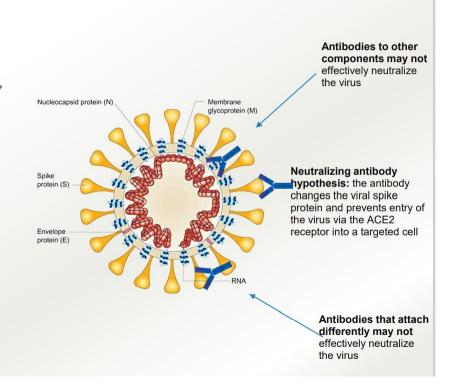
- lasts many months^{1,2}
- is multi-faceted and generates antibodies against the spike protein plus other non-structural proteins (Nucleoprotein (N), Matrix protein (M), Envelope protein (E))
- · induces systemic immunity and mucosal immunity

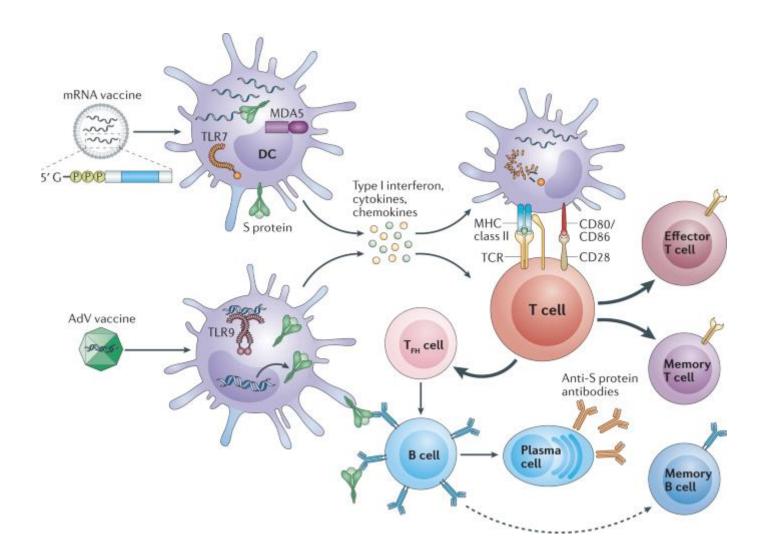


https://www.bmj.com/content/373/bmj.n1605 https://www.nature.com/articles/s41586-021-03696-9

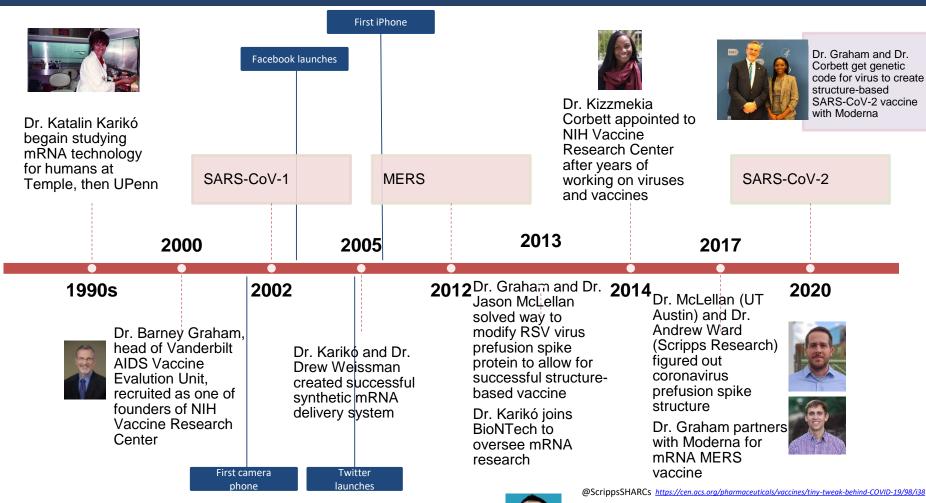
Vaccine-induced immunity to COVID-19

- COVID-19 vaccines induce neutralizing antibodies against the spike protein
- When SARS-CoV-2 is encountered naturally, neutralizing antibodies bind to the SARS-CoV-2 spike protein and block the virus from entering and multiplying in the cell
- A 'weaker' immune system (e.g. in elderly people or those with underlying health problems) may result in delayed and low stimulation of the antibody response after vaccination
- Current COVID 19 vaccines induce systemic immunity only and no mucosal immunity





Timeline of mRNA technology and key figures in vaccine development





I VACCINI E IL LORO RUOLO NEL MANTENIMENTO DELLO STATO DI SALUTE DELLA COMUNITA'

EVIDENZE SCIENTIFICHE E VACCINAZIONI

 IL RUOLO DEL SISTEMA IMMUNITARIO NELLA PROTEZIONE POST-VACCINALE

FERNANDO AIUTI

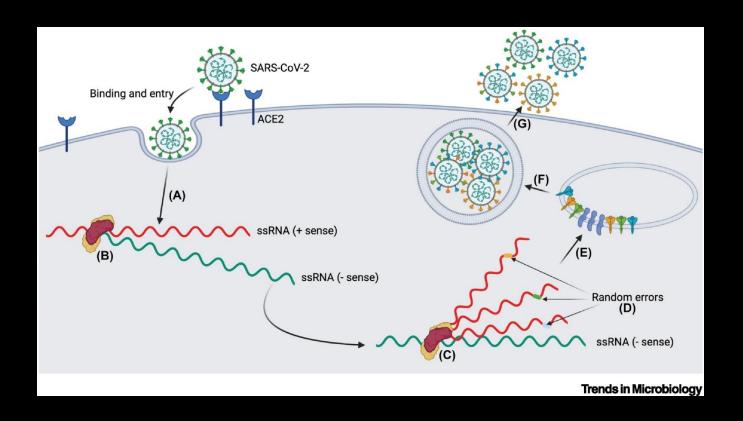
Professore Emerito Univ. "Sapienza" Roma

Corso organizzato da SIMEDET Vicariato di Roma Casa Bonus Pastor Roma 10 ottobre 2018

EVIDENZE IMMUNOLOGICHE

Vaccini di nuova concezione a RNA:
 vaccini che evocano immunità innata con
 stimolazione di cellule dendritiche o
 vaccini basati su vettori virali nel cui
 genoma viene inserito il gene che codifica
 per la proteina verso cui si vuole evocare
 risposta immune (es. produzione di
 anticorpi anti- virus influenza di recente
 produzione)

Figure 1







Its transmissibility

(relative to circulating variants)

Its virulence

(ability to cause severe disease)

Its ability to evade immune responses

(prior infection and vaccines & therapeutics)

DOI: 10.1002/jmv.26104

SHORT COMMUNICATION



. . .

MEDICAL VIROLOGY WILEY

Evidence for mutations in SARS-CoV-2 Italian isolates potentially affecting virus transmission

¹Department of Medical Statistics and Molecular Epidemiology, University Campus Bio-Medico of Rome, Rome, Italy

²Department of Medical Biology, Izmir University of Economics, Izmir, Turkey ³Deparment of Flavivirus, Instituto Oswaldo Cruz, Fundação Oswaldo Cruz, Rio de Janeiro - RJ, Brasii

⁴Department of Biochemical Sciences "A. Rossi Fanelli", University of Rome "La Sapienza", Rome, Italy

⁵Department of Clinical Laboratory Science, University Campus Bio-Medico of Rome, Rome Italy

⁶Department of Malattie Infettive – Fondazione Policlinico Universitario "A. Gemelli" IRCCS, Rome, Italy

⁷Department of Healthcare Surveillance and Bioethics, Catholic University of Sacred Heart, Rome, Italy

Biology, University of Siena, Siena, Italy

Correspondence

Domenico Benvenuto, Department of Medical Statistics and Molecular Epidemiology, University Campus Bio-Medico, Via Álvaro del Portillo, 21, 00128 Rome, Italy. Email: domenicobenvenuto95@gmail.com

Abstract

Italy is the first western country suffering heavy severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) transmission and disease impact after coronavirus disease-2019 pandemia started in China. Even though the presence of mutations on spike glycoprotein and nucleocapsid in Italian isolates has been reported, the potential impact of these mutations on viral transmission has not been evaluated. We have compared SARS-CoV-2 genome sequences from Italian patients with virus sequences from Chinese patients. We focussed upon three nonsynonymous mutations of genes coding for S(one) and N (two) viral proteins present in Italian isolates and absent in Chinese ones, using various bioinformatics tools. Amino acid analysis and changes in three-dimensional protein structure suggests the mutations reduce protein stability and, particularly for S1 mutation, the enhanced torsional ability of the molecule could favor virus binding to cell receptor(s). This theoretical interpretation awaits experimental and clinical confirmation.

KEYWORDS

bioinformatics, COVID-19, molecular evolution, mutation, SARS coronavirus

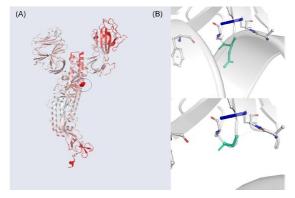


FIGURE 1 A, A model of spike glycoprotein monomer displaying the amino acids colored according to the vibrational entropy change upon mutation, red regions are those gaining in flexibility. The amino acidic mutation is blue circled; (B) the top image shows the molecular interaction between the side chain of the wild-type amino acid and the side chains of the surrounding amino acid; the bottom image shows the molecular interaction between the side chain of the mutated amino acid and the side chains of the surrounding amino acid.

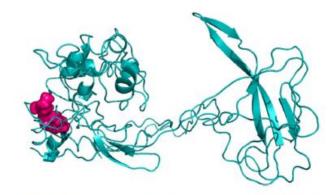


FIGURE 2 Cartoon model of the nucleocapsid of the SARS-CoV-2 where the mutated amino acids have been shown in purple

Mutazioni in RBD, S1 ed S2 della proteina Spike di SARS-CoV-2



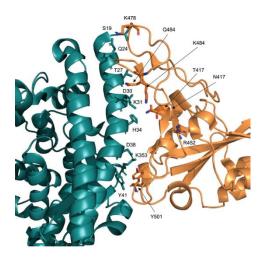


Garcia-Beltra, et al. MedRxiv. https://doi.org/10.1101/202 1.02.14.21251704 | Received: 21 May 2021 | Revised: 11 June 2021 | Accepted: 8 July 2021 | DOI: 10.1002/jmv.27210

RESEARCH ARTICLE

MEDICAL VIROLOGY WILEY

SARS-CoV-2 B.1.617 Indian variants: Are electrostatic potential changes responsible for a higher transmission rate?



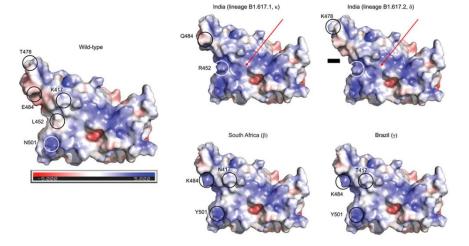


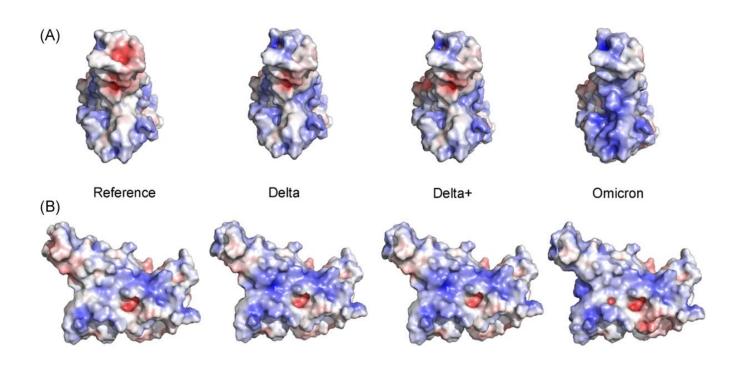
TABLE 1 Omicron mutations occurring in the Spike RBD

Mutationsa	Structural contextb	Interactions
G339D	α-helix	Possible H-bond to NAG
S371L	α-helix	VdW interaction with ACE2 V367 e F374
S373P	Loop	
S375F	N-terminal side of a β-strand	
K417N	C-terminal side of an α-helix	Removes the salt bridge with ACE2 D30
N440K	α-helix	
G446S	Loop within the RBM	
S477N	Loop within the RBM	
T478K	Loop within the RBM	
E484A	Loop	Removes the salt bridge between E484 and ACE2 K31
Q493R	short β-strand	Forms a salt bridge with ACE2 E35
G496S	Loop	H-bond with ACE2 D38
Q498R	Loop	Forms a salt bridge with ACE2 D38
N501Y		H-bond with ACE K353; Aromatic interaction with ACE2 Y41
Y505H	α-helix	Removes the H-bond with ACE2 E37

Note: Structural context and interaction changes (added or removed) compared to those found in the reference Spike. Boldfaced mutations are in the RBD.

Abbreviation: RBM, receptor-binding motif.

The electrostatic potential of the Omicron variant spike is higher than in Delta and Delta-plus variants: A hint to higher transmissibility?



- Stefano Pascarella¹
- Massimo Ciccozzi²
 - Martina Bianchi¹
- Domenico Benvenuto²
 - Roberto Cauda³
 - Antonio Cassone⁴

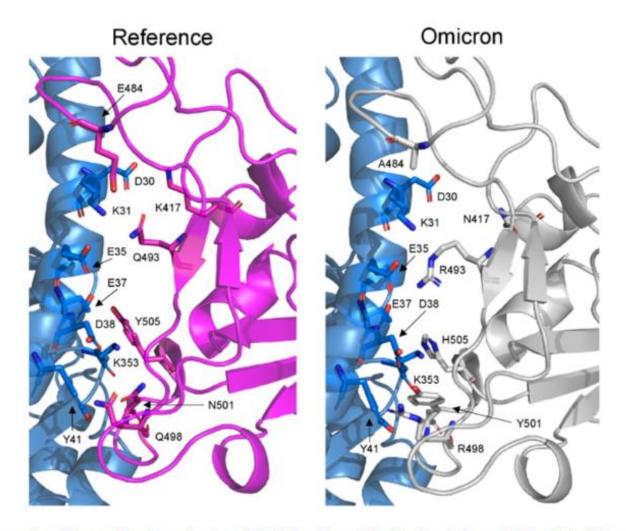


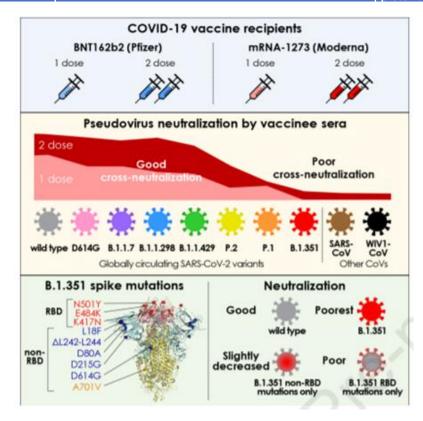
FIGURE 1 Comparison between the observed and predicted interactions at the interface between ACE2 and the Reference and Omicron RBDs, respectively. Spike Reference RBD ad Spike Omicron RBD are displayed as magenta and light gray ribbons, respectively. Blue cartoons indicate ACE2. Relevant Reference and Omicron side chains are displayed as sticks and labeled

Garcia-Beltram WF et al

Cell 2020

https://www.cell.com/cell/fulltext/S0092-8674(21)00298-

Multiple SARS-CoV-2 variants escape neutralization by vaccineinduced humoral immunity Il siero di 99 individui vaccinati con vaccino a mRNA Pfizer o MODERNA contro SARS-CoV-2 neutralizza il virus wildtype e alcune varianti, ma non la B.1.351 ("sudafricana") e la P.2 ("brasiliana"), per cui andrebbe supportata secondo gli autori la sintesi di ulteriori dosi di vaccino adeguate alle mutazioni della proteina Spike.



Check for updates

https://doi.org/10.1038/s41467-021-27649-y

OPEN

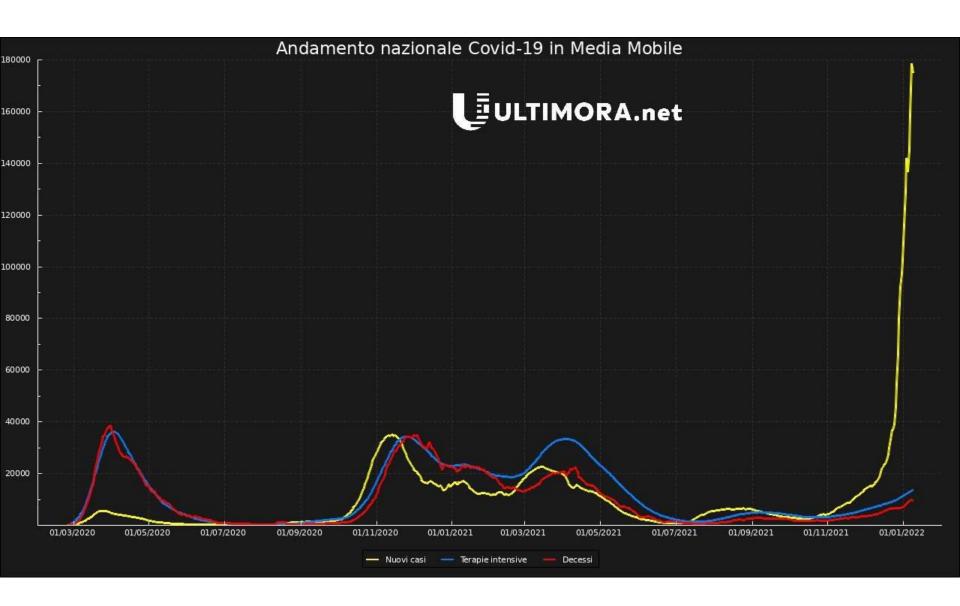
Dynamics of spike-and nucleocapsid specific immunity during long-term follow-up and vaccination of SARS-CoV-2 convalescents

Nina Koerber 1,12, Alina Priller 2,12, Sarah Yazici 2,12, Tanja Bauer 1,3,12, Cho-Chin Cheng 1, Hrvoje Mijočević 1, Hannah Wintersteller 2, Samuel Jeske 1, Emanuel Vogel 1, Martin Feuerherd 1, Kathrin Tinnefeld 1, Christof Winter 4, Jürgen Ruland 4, Markus Gerhard 5, Bernhard Haller 6, Catharina Christa 1, Otto Zelger 1, Hedwig Roggendorf 2, Martin Halle 7, Johanna Erber 8, Paul Lingor 9, Oliver Keppler 10,3, Dietmar Zehn 11, Ulrike Protzer 11,3,12 & Percy A. Knolle 2,3,12

Anti-viral immunity continuously declines over time after SARS-CoV-2 infection. Here, we characterize the dynamics of anti-viral immunity during long-term follow-up and after BNT162b2 mRNA-vaccination in convalescents after asymptomatic or mild SARS-CoV-2 infection. Virus-specific and virus-neutralizing antibody titers rapidly declined in convalescents over 9 months after infection, whereas virus-specific cytokine-producing polyfunctional T cells persisted, among which IL-2-producing T cells correlated with virus-neutralizing antibody titers. Among convalescents, 5% of individuals failed to mount long-lasting immunity after infection and showed a delayed response to vaccination compared to 1% of naïve vaccinees, but successfully responded to prime/boost vaccination. During the follow-up period, 8% of convalescents showed a selective increase in virus-neutralizing antibody titers without accompanying increased frequencies of circulating SARS-CoV-2-

specific T cells. The same convalescents, however, responded to vaccination with simultaneous increase in antibody and T cell immunity revealing the strength of mRNA-vaccination to increase virus-specific immunity in convalescents.

	IFNγ	IL-2	TNF	GzmB
convalescents	23	85	30	0
11 months	(±116)	(±116)	(±748)	(±10)
naive	0	15	290	0
11 months	(±5)	(±78)	(±1906)	(±13)
convalescents	255	625	260	0
vacc	(±285)	(±353)	(±418)	(±8)
naive	38	328	228	0
vacc	(±285)	(±266)	(±974)	(±10)
convalescents	225	415	165	0
boost	(±299)	(±213)	(±305)	(±13)
naive	165	435	275	0
boost	(±228)	(±376)	(±894)	(±22)



Effectiveness of mRNA-1273 against SARS-CoV-2 omicron and delta variants

Authors: Hung Fu Tseng, PhD,^{1,2,*} Bradley K. Ackerson, MD,¹ Yi Luo, PhD,¹ Lina S. Sy, MPH,¹ Carla A. Talarico, PhD,³ Yun Tian, MS,¹ Katia J. Bruxvoort, PhD,⁴ Julia E. Tubert, MPH,¹ Ana Florea, PhD,¹ Jennifer H. Ku, PhD,¹ Gina S. Lee, MPH,¹ Soon Kyu Choi, MPH,¹ Harpreet S. Takhar, MPH,¹ Michael Aragones, MD,¹ Lei Qian, PhD¹

Figure 1. Vaccine effectiveness of 2-dose mRNA-1273 against omicron and delta variants by time since vaccination

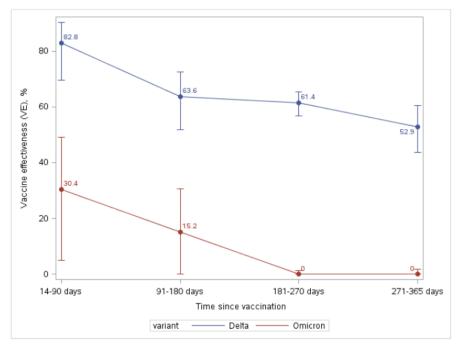
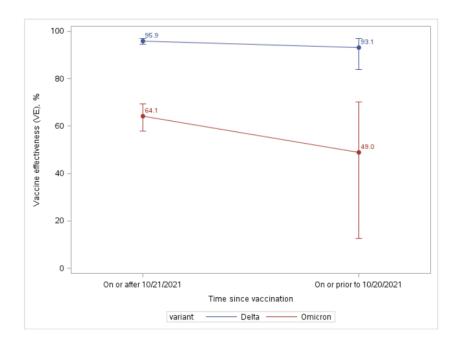
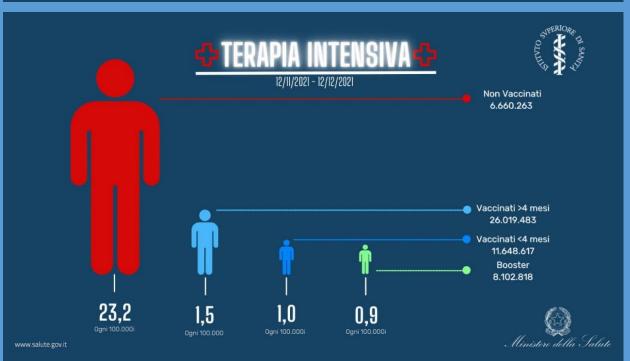


Figure 2. Vaccine effectiveness of 3-dose mRNA-1273 against omicron and delta variants by time since vaccination among immunocompetent population

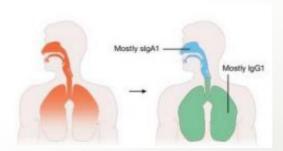




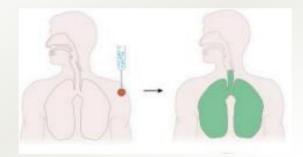


Mucosal immunity may work as a barrier to infection

- Current intramuscular COVID-19 vaccines do not induce mucosal immunity. They do not induce the same multifaceted immune response as a natural infection but do protect from severe disease
- Nasal COVID-19 vaccines are being investigated to protect from infection as well as from severe disease



Infection-induced immunity induces systemic immunity but also mucosal immunity because SARS-CoV-2 infection starts in the upper respiratory tract



Vaccine-induced immunity induces systemic immunity only and no mucosal immunity



Vaccine

journal homepage: www.elsevier.com/locate/vaccine



Commentary

Multicomponent vaccines to fight SARS-CoV-2 variants of concern



Antonio Cassone a,*, Roberto Cauda b,1

interacts with the host cell membrane and allows the viral fusion with, and entry, the cell so initiating infection [1]. The choice of nucleic acid technology and S protein as antigen for vaccine formulation was pathogenically logical and allowed generation of safe and highly efficacious vaccines at an unprecedented speed (around 1 year). These vaccines elicit the production of specific antibodies that inhibit S protein binding to ACE2 receptor and neutralize virus infection. They also activate cell-mediated, B and T memory immunity, and are highly effective in protecting against severe disease, hence abating COVID-19 mortality [2].

2. Vaccines challenged by SARS-CoV-2 variants

Unfortunately, though expected, viral mutations have soon appeared among spreading SARS-CoV-2 lineages, the so-called Variants of Interest (VOI) or Concern (VOC), which seriously threaten vaccine effectiveness. The most threatening mutations are those leading to amino acid substitutions or deletions in the S protein, particularly in its RBD sequence. Some of them make SARS-CoV-2 more contagious because they confer to the S protein increased affinity for ACE2 receptor but do not appear to markedly undermine vaccine effectiveness. Some others, however, appear to substantially lower antibody protection as shown by in vitro

3. Proposal

Scientists, public health administrators and stakeholders appear to be well aware of the above threatening situation, and some vaccine manufacturers have started production and validation of a second generation vaccines for VOC better fighting. To do this, the high efficiency and flexibility of nucleic acid-based technologies, as witnessed by the success of current vaccines, is being rightly advocated, and a logical option would seem to be just replacing, or adding to, the present nucleic acid sequences with those of coronavirus variants [1,2].

We believe there is time to consider an integrative option. We posit the necessity of broadening antigen composition of the second generation, SARS-CoV-2 vaccines by including sequences of genes encoding non RBD, non S constituents of the coronavirus genome. We invite vaccine companies to seriously consider the potential of a multicomponent vaccine to generate protective antibodies and cell-mediated immunity to which SARS-CoV-2 could much less likely escape by mutations than with the current, single component vaccines. A number of structural and non-structural viral proteins of critical importance for the reproduction cycle and architecture of the beta-coronaviruses, as well as for their interaction with human cells, could in principle be considered. Past studies have indeed documented that SARS-CoV-2 -infected subjects raise strong immune responses to antigenic components other than S protein, some of which are likely to contribute to the immune protection against the virus, as discussed below. Most of these immune responses are targeted to the nucleoprotein (N) constituent of the viral particle.

^a Polo d'innovazione della genomica, genetica e biologia, c/o Toscana Life Sciences, Strada del Petriccio e Belriguardo, 53100 Siena, Italy

^b Dipartimento di Sicurezza e Bioetica, Università Cattolica del Sacro Cuore, Rome, Italy

^{*} Corresponding author.

E-mail addresses: a.cassone@pologgb.com (A. Cassone), roberto.cauda@unicatt.it (R. Cauda).

¹ Istituto di Clinica della Malattie Infettive, Università Cattolica del Sacro Cuore, Largo Francesco Vito, 1, 00168-Roma, Italy.



https://doi.org/10.1038/s41467-021-27674-x

OPEN

Cross-reactive memory T cells associate with protection against SARS-CoV-2 infection in COVID-19 contacts

Rhia Kundu 12 → Janakan Sam Narean 1,2, Lulu Wang 1,2, Joseph Fenn 1,2, Timesh Pillay 1,2, Nieves Derqui Fernandez 1,2, Emily Conibear 1,2, Aleksandra Koycheva 1,2, Megan Davies 1,2, Mica Tolosa-Wright 1,2, Seran Hakki 1,2, Robert Varro 1,2, Eimear McDermott 1,2, Sarah Hammett 1,2, Jessica Cutajar 1,2, Ryan S. Thwaites 2, Eleanor Parker 3, Carolina Rosadas 3, Myra McClure 3, Richard Tedder 3, Graham P. Taylor 3, Jake Dunning 4,5 & Ajit Lalvani 1,2

Cross-reactive immune responses to SARS-CoV-2 have been observed in pre-pandemic cohorts and proposed to contribute to host protection. Here we assess 52 COVID-19 household contacts to capture immune responses at the earliest timepoints after SARS-CoV-2 exposure. Using a dual cytokine FLISpot assay on peripheral blood mononuclear cells, we enumerate the frequency of T cells specific for spike, nucleocapsid, membrane, envelope and ORF1 SARS-CoV-2 epitopes that cross-react with human endemic coronaviruses. We observe higher frequencies of cross-reactive (p = 0.0139), and nucleocapsid-specific (p = 0.0355) IL-2-secreting memory T cells in contacts who remained PCR-negative despite exposure (n = 26), when compared with those who convert to PCR-positive (n = 26); no significant difference in the frequency of responses to spike is observed, hinting at a limited protective function of spike-cross-reactive T cells. Our results are thus consistent with pre-existing non-spike cross-reactive memory T cells protecting SARS-CoV-2-naïve contacts from infection, thereby supporting the inclusion of non-spike antigens in second-generation vaccines.

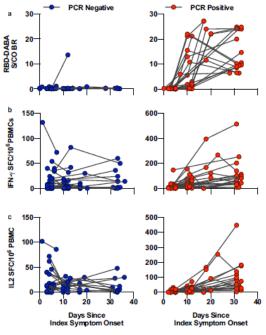


Fig. 3 Dynamics of cross-reactive T cells and RBD-specific antibody in PCR-positive and negative contacts. Serum sampled from COVID-19 contacts at the baseline, D7 and D28 visit were assayed for RBD-specific antibody, represented as sample/control ratios (a). PBMCs from these visits were rested overnight at high density prior to stimulation with 1 µg/ml cross-reactive peptide pool cultured for 20 h in a FLISpot assay to detect IL-2- (b) and IFNy (c) secreting T cells. Serum from these visits were assayed for RBD-specific antibody, represented as sample/control ratios (c). Left-hand panels and blue circles represent PCR-negative contacts whilst right-hand panels and red circle represent PCR-positive contacts.



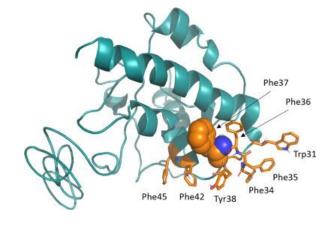
Journal of Infection

Volume 81, Issue 1, July 2020, Pages e24-e27

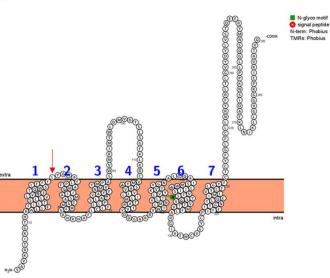


Evolutionary analysis of SARS-CoV-2: how mutation of Non-Structural Protein 6 (NSP6) could affect viral autophagy

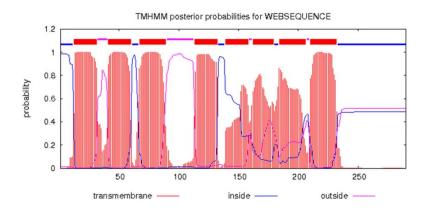
Domenico Benvenuto * 🌣 🖾 , Silvia Angeletti ⁶, Marta Giovanetti ^c, Martina Bianchi ^d, Stefano Pascarella ^d, Roberto Cauda * ^f, Massimo Ciccozzi ^e, Antonio Cassone ^g





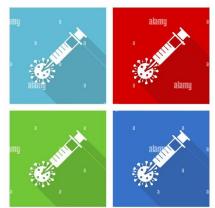


В



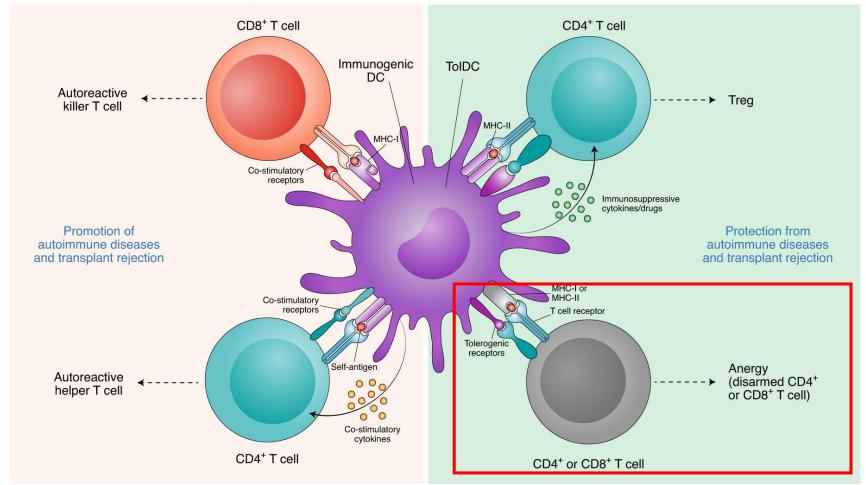


- 1) Il Sars-Cov2 muta molto in fretta e alcune mutazioni conferiscono vantaggi al virus
- 2) I vaccini attuali nonostante non siano aggiornati sulle varianti forniscono un'ottima protezione dalla malattia
- 3) I ricercatori del centro Walter Reed hanno messo a punto un vaccino pan-coronavirus
- 4) Il vaccino pan-coronavirus dovrebbe proteggere dalle varianti attuali e dalle future
- 5) Il vaccino pan-coronavirus accoppiato al vaccino spray potrebbe segnare una svolta nella pandemia



Autoimmunity

Tolerance



IDEAL+ Student Training Course - Paris 2021

Monday March Truckdy March Wednesday 10 March Truckdy 11 March Friday 1 March Privally 11 March Privally 12 March						
Domains		Monday 8 March	Tuesday 9 March	Wednesday 10 March	Thursday 11 March	Friday 12 March
09:30 Welcome	Rooms		Online	Pavillon 3		Pavillon 3
Welcome		Click here	Click here	Click here	Click here	Click here
09:30	08:30					
Sepsis B Infectious Diseases Discol - 10.50 Lung infections Emergencies (Annexery) Emergencies (Name) E	09:00					
10:30	09:30	Sepsis &				
11:00 11:00 11:00 11:00	10:00		(Rome)	(Rome)	(Antwerp & Hamburg)	
11:30 11:30 11:30 11:30 Tuberculosis (Rome) 11:00-12:50	10:30					(Antwerp & Athens)
11:30 11:00-12:50 Tuberculosis (Name) 11:00-12:50 Tuberculosis (Name) 12:00-12:50 12:00-12:50 12:00-12:50 Hiv infection (Rome) 12:00-12:50 Cher tropical diseases (Edinburgh) 13:30 14:30 Lunch break	11:00					
12:00 12:00	44.20		11:00-12:20	Animal bites & scratches	Malaria	
12:30 Lunch break 13:00-14:10 Congenital infections (Athens & Paris) (Antwerp) Mey rules in management of Infectious Diseases (Antwerp) (Antwerp) 15:00 15:00-15:50 Hepatitis (Edinburgh) 15:00-16:20 Skin rash (Edinburgh) 16:00 16:00-16:50 Sm (Paris) 16:30-17:20 Diarrhea (Idinburgh) 17:00 Wrap-up session	11.30		(Rome)	(Parts)	(Edinourgh)	
13:30	12:00	(Paris)		HIV infection	Other tropical diseases	
13:30 Lunch break Lunch	12:30					Lunch break
Lunch break 14:00-14:50 Roundtable on COVID-19 (Rame & Edinburgh) Itico-15:00 Skin rath Infection (Antheria) Listo-16:30 Lymphadenopathy (Edinburgh) Listo-16:30 Lymphadenopathy (Edinburgh) Vrap-up session	13:00	-	Lunch break			
14:00	13.55	Lunch break		Lunch break	Lunch break	13:00-14:10
14:00	13:30					
Reyrules in management of Infectious Diseases (Antwerp)	14:00	14:00-14:50				
14:30 (Rome) (Rome) (Rome & Edinburgh) 14:20-15:40 Infection control (Antwerp) 15:00 - 16:50 Skin rash (Edinburgh) Skin and soft tissue infections & Arthritis (Hamburg & Paris) (Athens) (Edinburgh)			(Paris)			
15:00	14:30					14:20-15:40
15:30 Hepatitis 15:00-16:20 Skin rash (Edinburgh) Skin and soft tissue infections & Arthritis (Hamburg & Paris) 16:30 (Pana) 16:30-17:20 Dierrhea (Edinburgh) 16:30-17:20 Dierrhea (Edinburgh) 17:00 Wrap-up session Wrap-up session Wrap-up session (Antwerp) (An	15:00					
15:30 (Hamburg) Skin rash (Edinburgh) Skin and soft tissue infections & 15:00-16:50 Bronchiolitis, whooping cough and other miscellaneous (Athens) (Edinburgh) (Edinburgh) Total Cough and other miscellaneous (Athens) (Edinburgh) (Edinburgh) Wrap-up session	25.00	Hepatitis	45:00-46:20	45-00-46-50		(Antwerp)
16:00 16:00 16:00 Arthritis (Athens) (Edinburgh) 16:30 (Pana) 16:30 -17:20 (Diarrhea (Edinburgh)) 17:00 (Value of the content of the conten	15:30	16:00-16:50	Skin rash	Skin and soft tissue infections & Arthritis	Bronchiolitis, whooping cough and other miscellaneous	15:40-45:20
16:30 - 17:20 Diarrhea (Fdinburgh) Wrap-up session	16:00					Lymphadenopathy
17:00 (Edinburgh) Wrap-up session	16:30					
17:30	17:00					Wrap-up session
	17:30					

