



L'INSOSTENIBILE IMPREVEDIBILITA' DEI VACCINI COVID-19

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22 Agosto 2022

L'INSOSTENIBILE IMPREVEDIBILITA' DEI VACCINI COVID-19

1) Farmacologia dei vaccini a RNA

Ovvero: giocare a dadi con la spike

2) Analisi di laboratorio negli studi autorizzativi

Cosa comporta il fatto che non siano state svolte?

3) Curare il COVID-19

Ruolo delle cure nella valutazione dell'efficacia vaccinale

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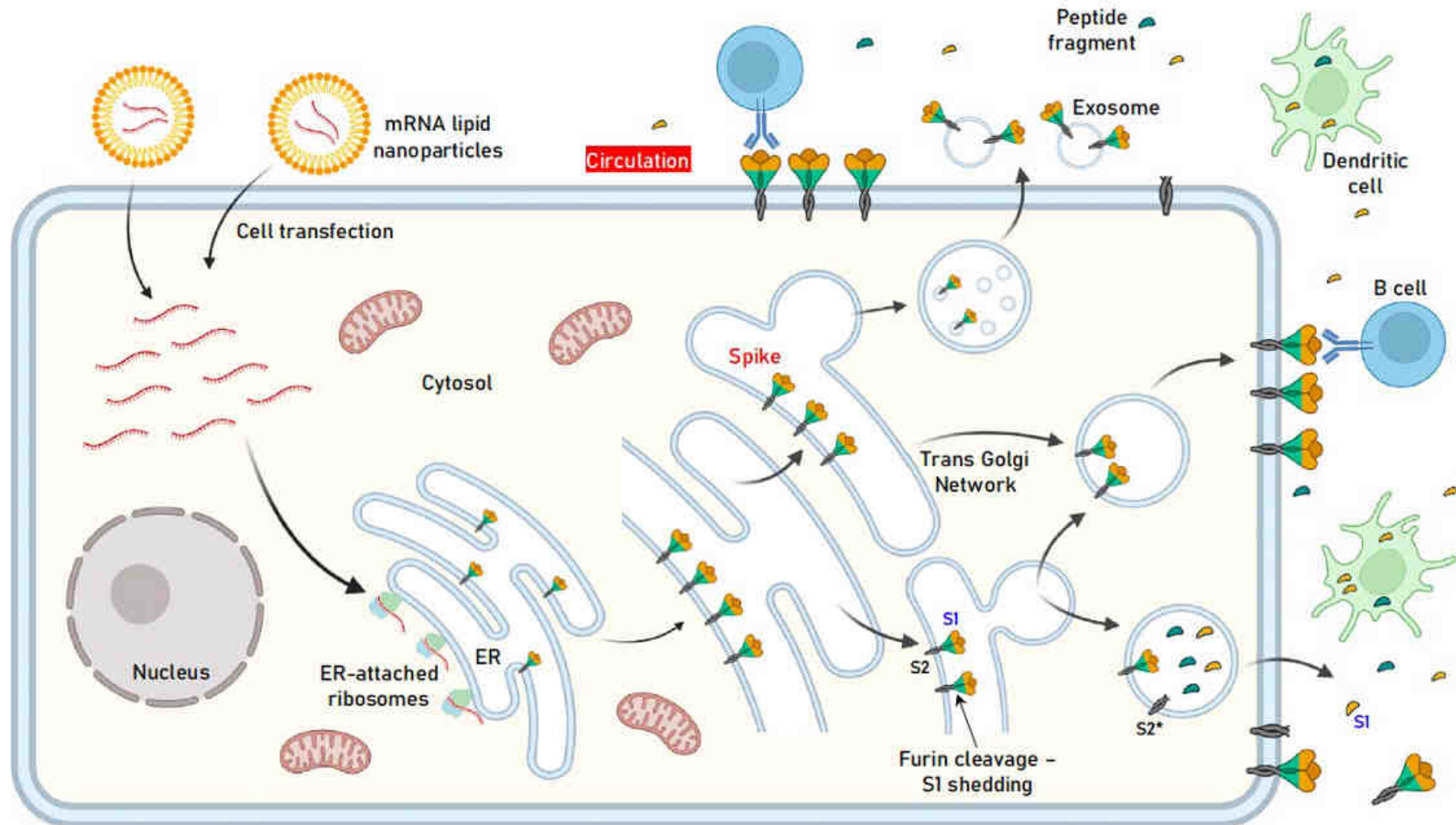
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Cosa comporta il fatto che non siano state svolte?

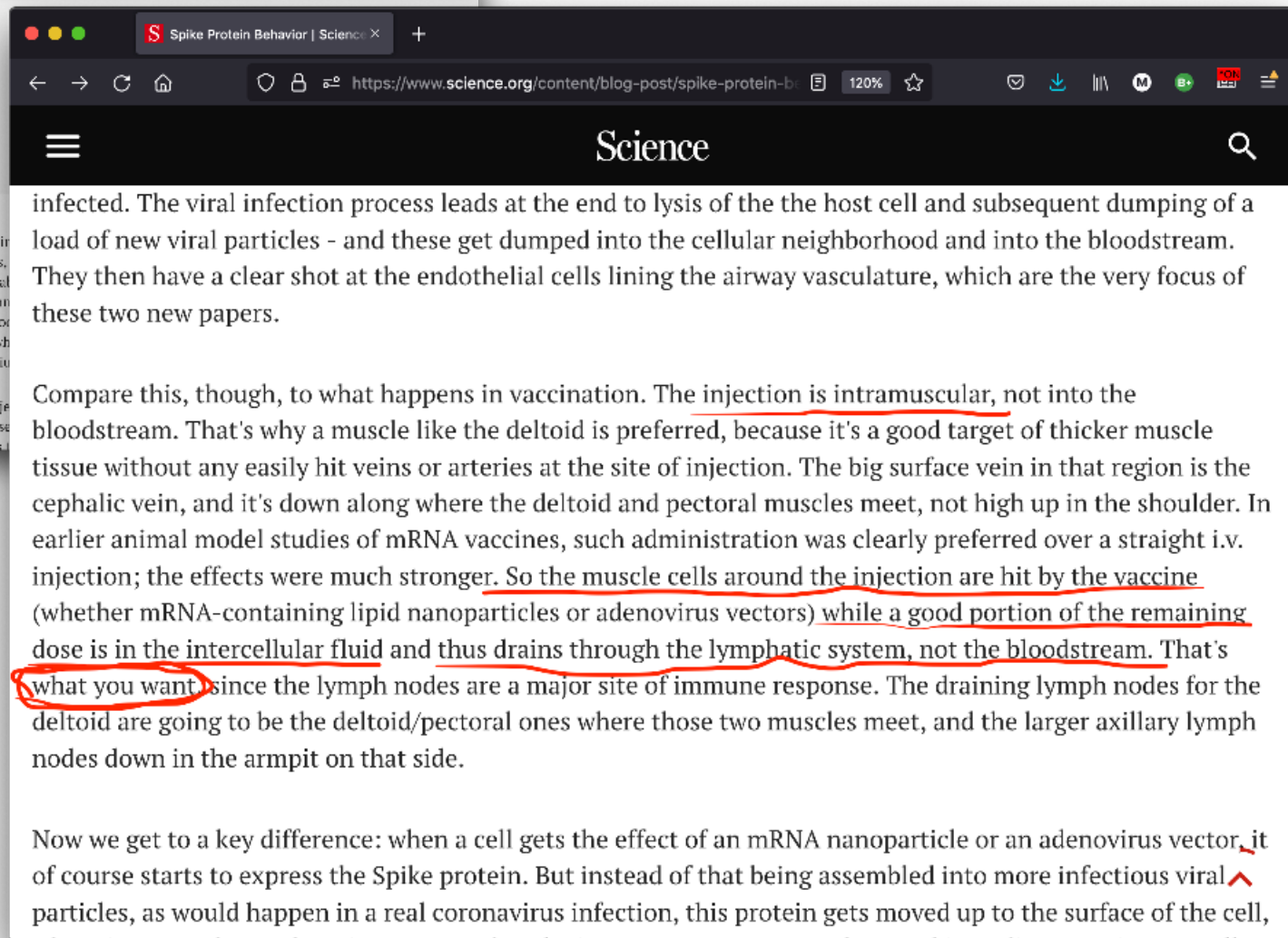
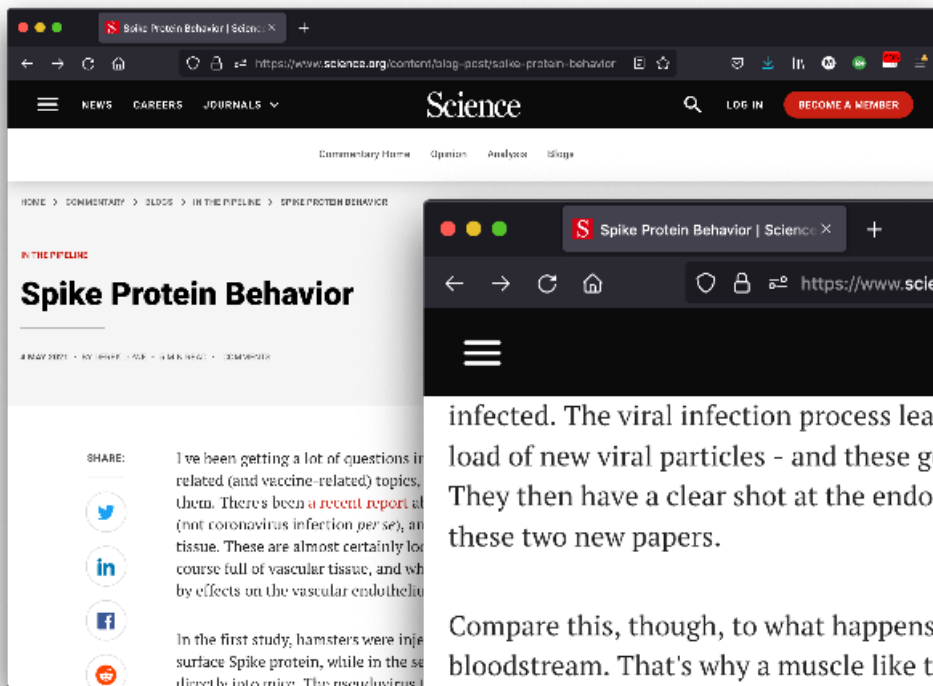
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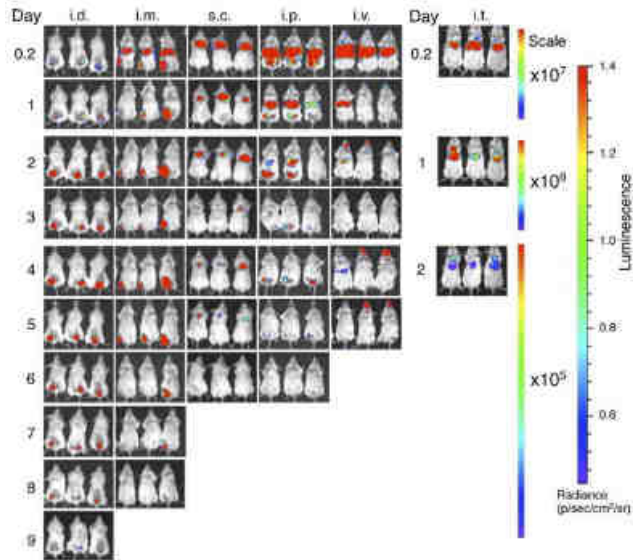
Antigen expression-localization following cell transfection with S protein mRNA-containing lipid nanoparticles (LNPs) used in anti-SARSCoV-2 mRNA vaccines



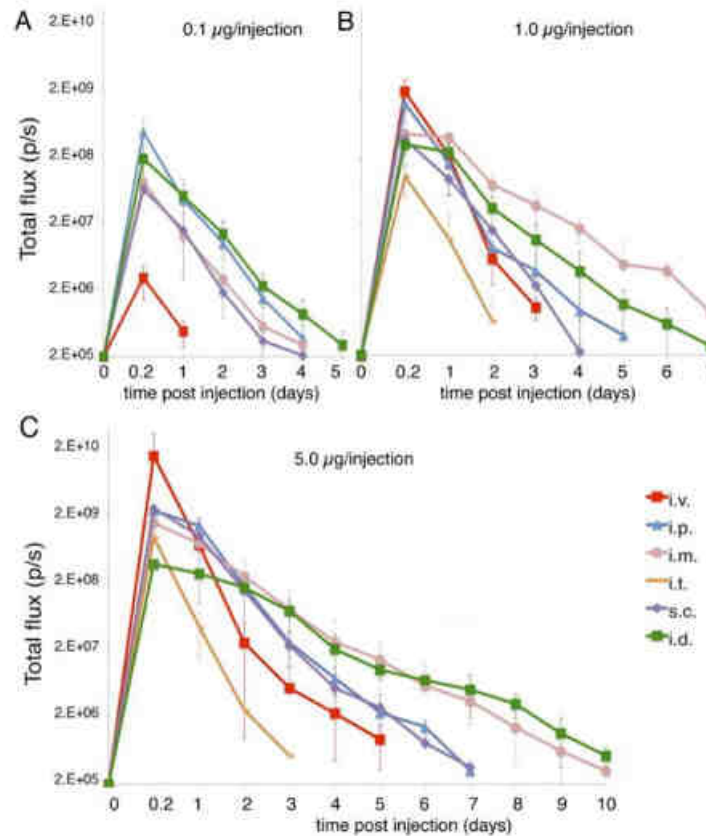
Trougakos, I.P. et al. (2022) **Adverse effects of COVID-19 mRNA vaccines: the spike hypothesis.** *Trends Mol. Med.* (in press)



<https://www.science.org/content/blog-post/spike-protein-behavior>



Duration and translational pattern of mRNA-LNPs in mice injected by various routes. Representative IVIS images of groups of 3 BALB/c mice injected with 5.0 µg mRNA-LNP by the intradermal (i.d.), intramuscular (i.m.), subcutaneous (s.c.), intravenous (i.v.), intraperitoneal (i.p.) and intratracheal (i.t.) routes. Relative luminescence plot is shown and the scale of luminescence is indicated.



Translational kinetics of mRNA-LNP delivered by different routes in vivo. Quantification of the bioluminescent signal measured in BALB/c mice injected with (A) 0.1 µg, (B) 1.0 µg or (C) 5.0 µg mRNA-LNPs by intradermal (i.d.), intramuscular (i.m.), subcutaneous (s.c.), intravenous (i.v.), intraperitoneal (i.p.) and intratracheal (i.t.) routes. Error bars are standard error of the mean (SEM).

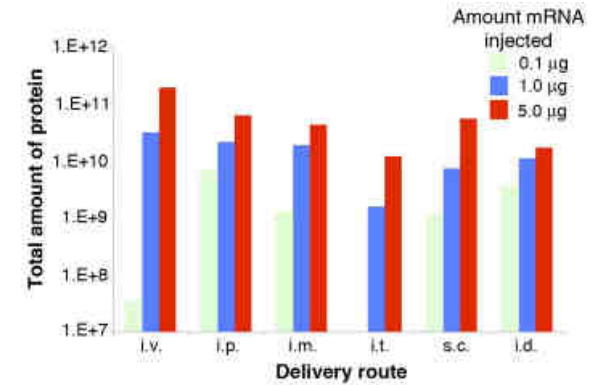


Table 1

Half-life of protein production

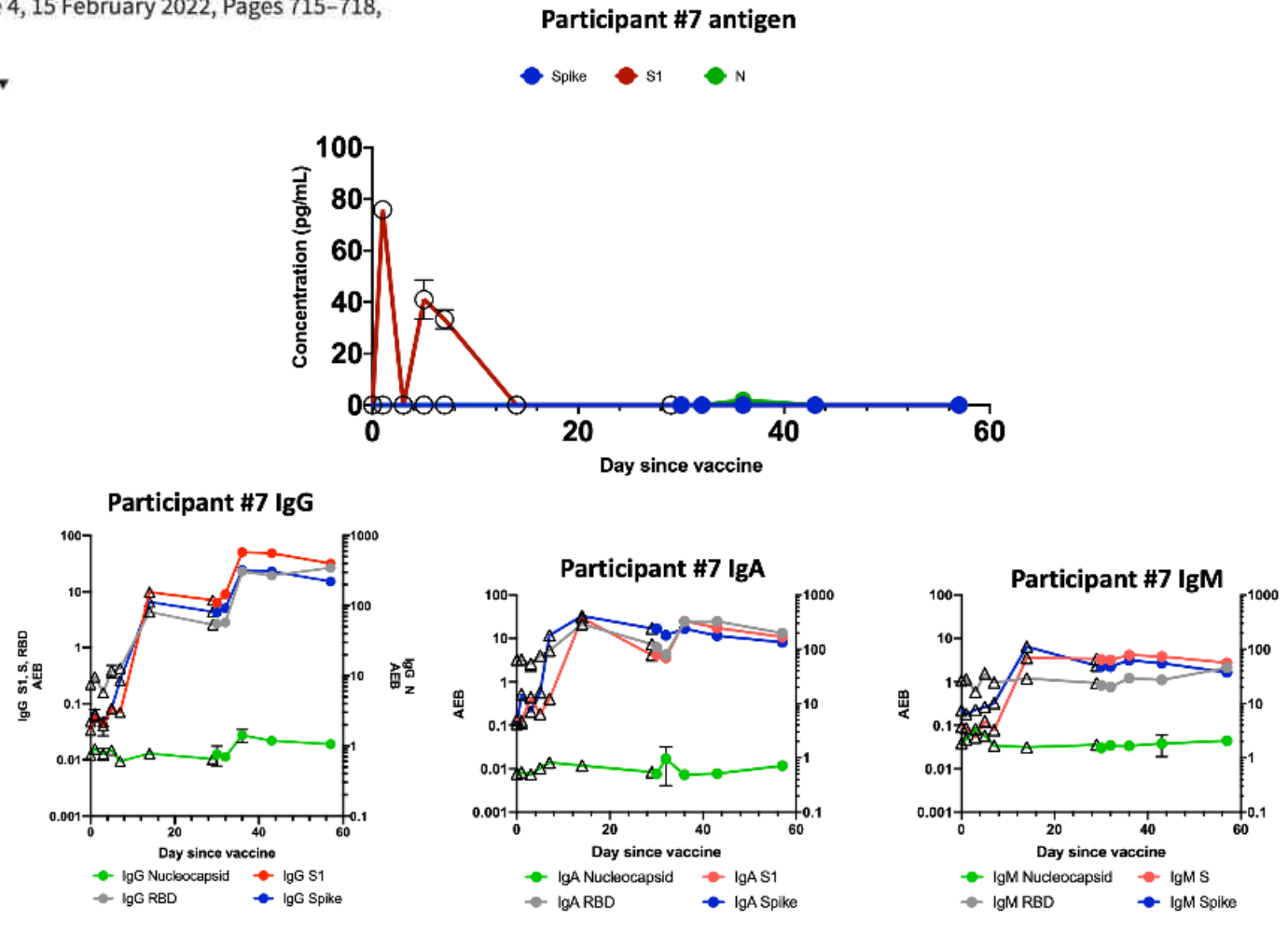
Delivery site	Half-life of mRNA translation (hours)
intradermal	29.6
intramuscular	20.6
subcutaneous	14.7
intravenous	6.8
intraperitoneal	14.8
intratracheal	7.5

Circulating Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) Vaccine Antigen Detected in the Plasma of mRNA-1273 Vaccine Recipients FREE

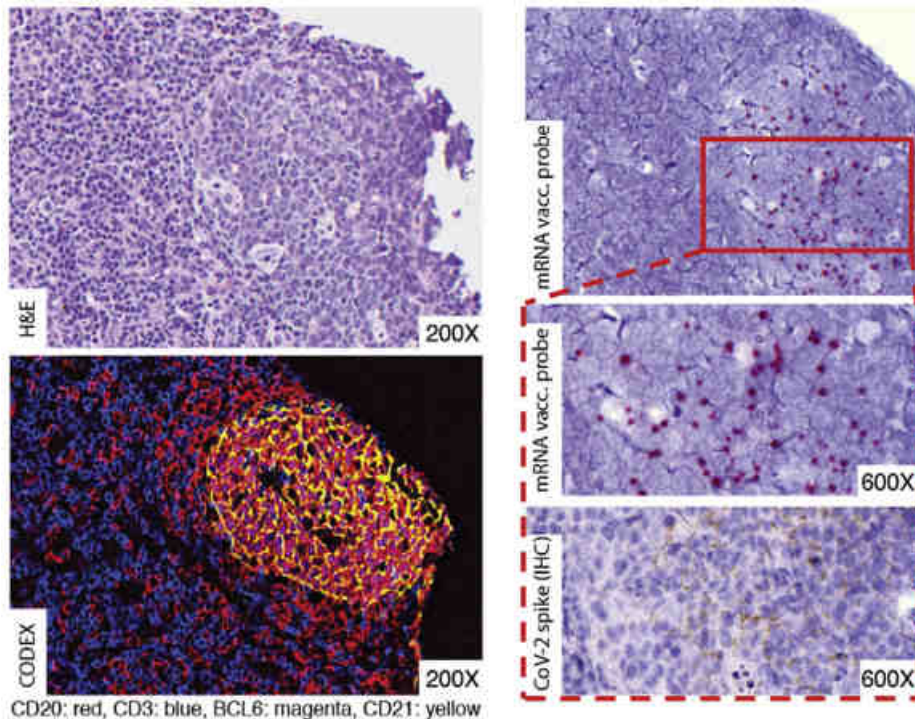
Alana F Ogata, Chi-An Cheng, Michaël Desjardins, Yasmeen Senussi, Amy C Sherman, Megan Powell, Lewis Novack, Salena Von, Xiaofang Li, Lindsey R Baden ... [Show more](#)
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Clinical Infectious Diseases, Volume 74, Issue 4, 15 February 2022, Pages 715–718,
<https://doi.org/10.1093/cid/ciab465>

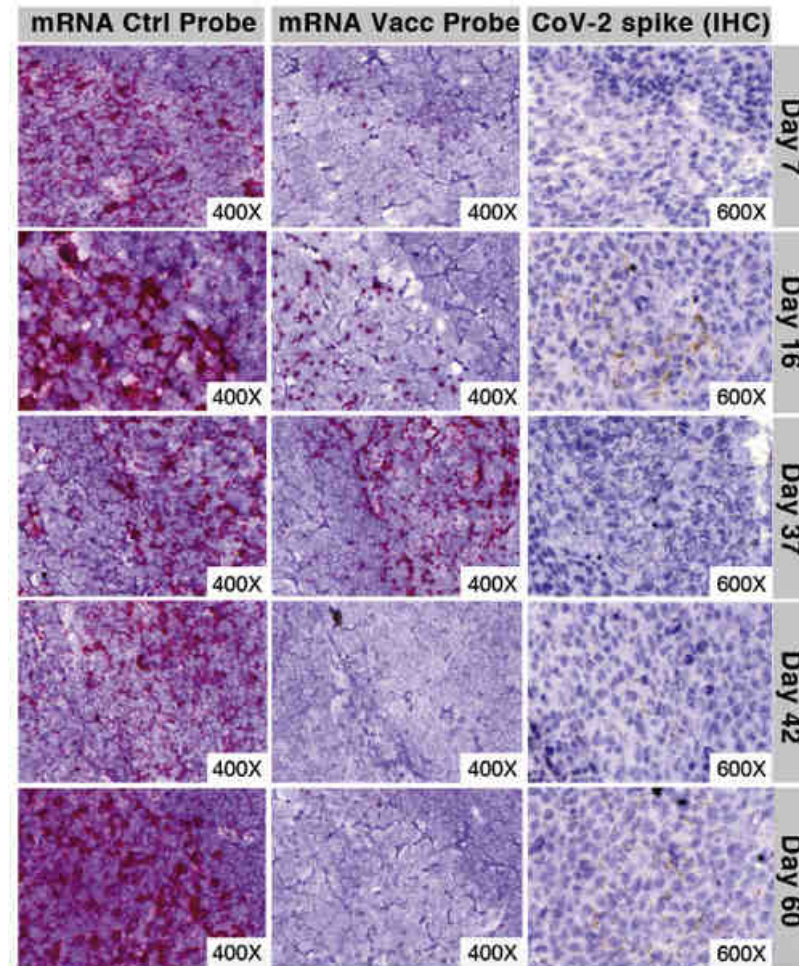
Published: 20 May 2021 [Article history](#) ▾



Ogata, A.F. et al. (2022) Circulating Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) Vaccine Antigen Detected in the Plasma of mRNA-1273 Vaccine Recipients. *Clin. Infect. Dis.* 74, 715-718



Representative LN GC after mRNA vaccination showing hematoxylin and eosin staining (upper left), four-color Codex staining (lower left), in situ hybridization of a SARS-CoV-2 mRNA vaccine-specific probe (upper right [lower magnification] and middle right [greater magnification]), and immunohistochemical (IHC) staining for spike antigen (lower right). Vaccine mRNA probe hybridization was visualized by colorimetric development with Fast Red chromogen, and positive IHC staining for spike antigen was visualized as granular brown color from 3,3'-diaminobenzidine (DAB) reagent.



Representative in situ hybridization of an RNAScope control probe (left panels) and SARS-CoV-2 mRNA vaccine-specific probe (middle panels) within ipsilateral axillary core needle LN biopsies of female patients 7–60 days after second mRNA-1273 or BNT162b2 dose. Probe hybridization is indicated by red chromogen spots. IHC signal for spike antigen (right panels) is detected as granular brown staining.

TO THE EDITOR:

SARS-CoV-2 spike-dependent platelet activation in COVID-19 vaccine-induced thrombocytopenia

Jacob Appelbaum,¹ Donald M. Arnold,² John G. Kelton,² Terry Gernsheimer,¹ Stefan D. Jevtic,² Nikola Ivetic,² James W. Smith,² and Ishac Nazy²

¹Division of Hematology, Department of Medicine, University of Washington School of Medicine, Seattle, WA; and ²Department of Medicine, Michael G. DeGroote School of Medicine, McMaster University, Hamilton, ON, Canada

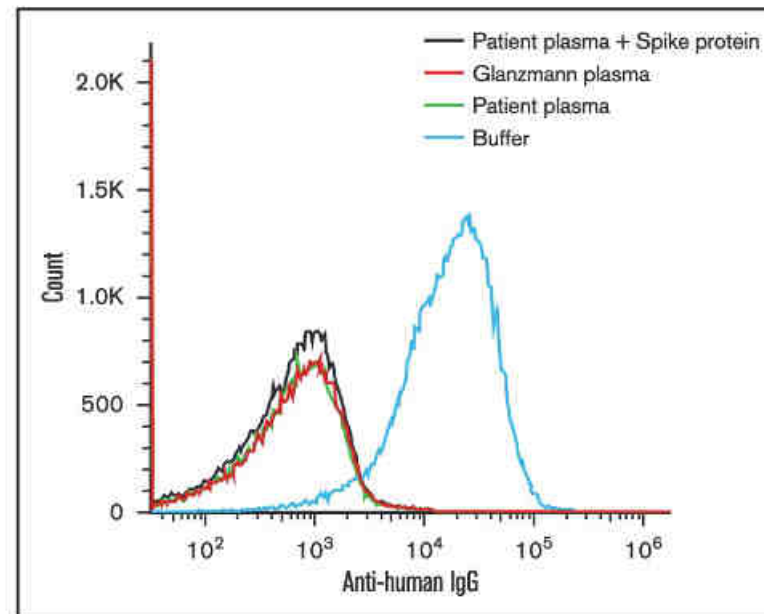


Figure 2. Flow cytometry of IgG binding to the platelet surface. Platelets incubated with plasma sample from our patient (green) failed to show IgG binding with exogenous spike protein administration (red). Negative buffer control (black) and positive Glanzmann thrombasthenia (blue) are shown for comparison.



Article

Intramyocardial Inflammation after COVID-19 Vaccination: An Endomyocardial Biopsy-Proven Case Series

Christian Baumeier ^{1,*}, Ganna Aleshcheva ¹, Dominik Harms ¹, Ulrich Gross ¹, Christian Hamm ^{2,3}, Birgit Assmus ³, Ralf Westenfeld ⁴, Malte Kelm ⁴, Spyros Rammos ⁵, Philip Wenzel ⁶, Thomas Münzel ⁶, Albrecht Elsässer ⁷, Mudather Gailani ⁸, Christian Perings ⁹, Alae Bourakkadi ¹⁰, Markus Flesch ¹¹, Tibor Kempf ¹², Johann Bauersachs ¹², Felicitas Escher ^{1,13,14} and Heinz-Peter Schultheiss ¹



Citation: Baumeier, C.; Aleshcheva, G.; Harms, D.; Gross, U.; Hamm, C.; Assmus, B.; Westenfeld, R.; Kelm, M.; Rammos, S.; Wenzel, P.; et al. Intramyocardial Inflammation after COVID-19 Vaccination: An Endomyocardial Biopsy-Proven Case Series. *Int. J. Mol. Sci.* **2022**, *23*, 6940. <https://doi.org/10.3390/ijms23136940>

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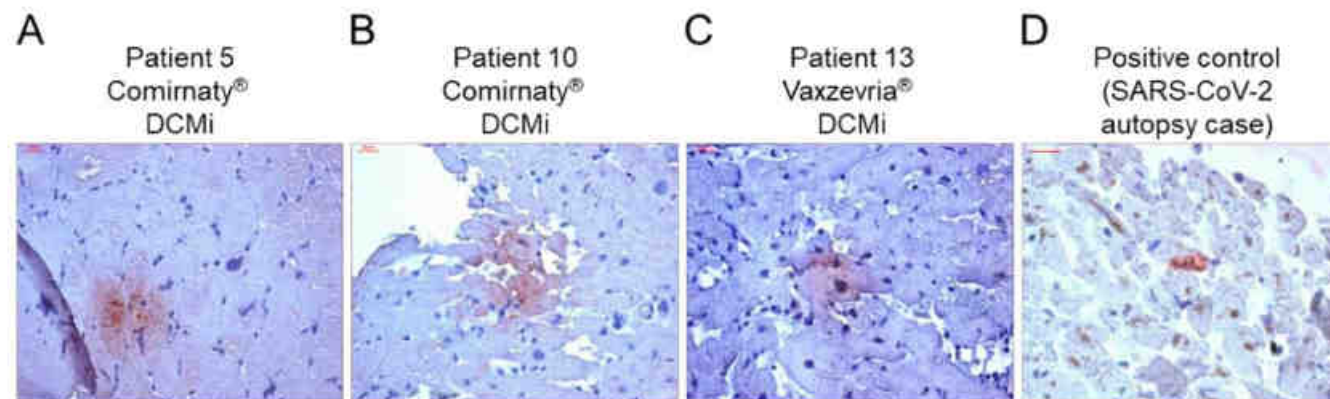


Figure 2. Evidence of SARS-CoV-2 spike protein in cardiac tissue after COVID-19 vaccination. (A–C) Representative immunohistochemical stainings of SARS-CoV-2 spike protein in EMBs from patients diagnosed with DCMi after receiving Comirnaty[®] (panel A and B, patients 5 and 10) or Vaxzevria[®] (panel C, patient 13). (D) SARS-CoV-2-positive cardiac tissue served as positive control. Magnification 400 \times . Scale bars 20 μ m.

Case Report

Clinical and Molecular Characterization of a Rare Case of BNT162b2 mRNA COVID-19 Vaccine-Associated Myositis

Eli Magen ^{1,2,*}, Sumit Mukherjee ^{3,4,†}, Mahua Bhattacharya ³, Rajesh Detroja ³, Eugene Merzon ^{2,5}, Idan Blum ¹, Alejandro Livoff ⁶, Mark Shlapobersky ⁶, Gideon Baum ³, Ran Talisman ⁷, Evgenia Cherniavsky ⁸, Amir Dori ^{9,10} and Milana Frenkel-Morgenstern ^{3,*}



Citation: Magen, E.; Mukherjee, S.; Bhattacharya, M.; Detroja, R.; Merzon, E.; Blum, I.; Livoff, A.; Shlapobersky, M.; Baum, G.; Talisman, R.; et al. Clinical and Molecular Characterization of a Rare Case of BNT162b2 mRNA COVID-19 Vaccine-Associated Myositis. *Vaccines* 2022, 10, 1135. <https://doi.org/10.3390/vaccines10071135>

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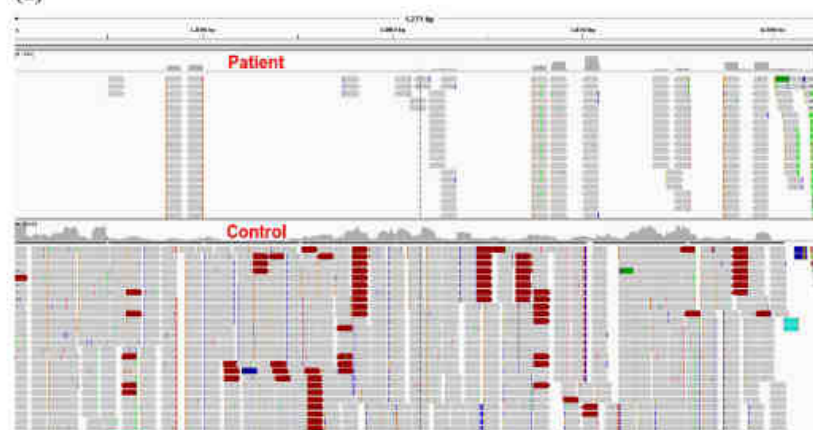
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(I)



(II)

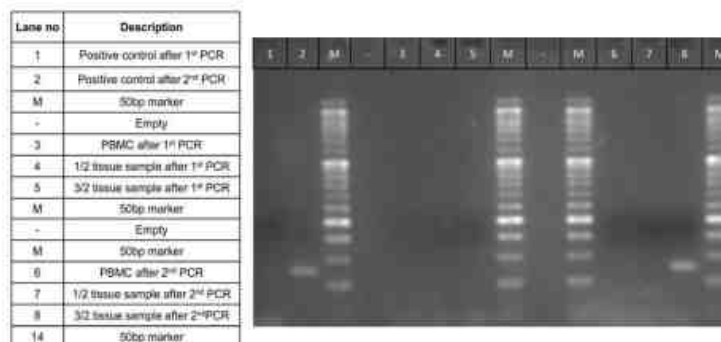


Figure 3. (I) Distribution of mapped RNA-seq reads across spike mRNA. (II) PCR validation of vaccine spike mRNA expression in the patient's muscle tissue biopsy samples one-month post-vaccination.

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Short Report:

SARS-CoV-2 S1 Protein Persistence in SARS-CoV-2 Negative Post-Vaccination Individuals with Long COVID/PASC-Like Symptoms

Bruce K. Patterson, Edgar B. Francisco, Ram Yogendra, Emily Long, and 8 more

This is a preprint; it has not been peer reviewed by a journal.

<https://doi.org/10.21203/rs.3.rs-1844677/v1>
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Abstract

Background

We sought to determine the immunologic abnormalities in patients following SARS-CoV-2 vaccines who experience post-acute sequelae of COVID-19 (PASC)-like symptoms > 4 weeks post vaccination. In addition, we investigated whether the potential etiology was similar to PASC.

Design:

We enrolled 50 post-vaccination individuals who experience PASC-like symptoms, 10 healthy individuals, and 35 individuals post-vaccination without symptoms. We performed multiplex cytokine/chemokine profiling with machine learning as well as SARS-CoV-2 S1 protein detection on monocyte subsets using flow cytometry and mass spectrometry.

Results

We determined that post-vaccination individuals with PASC-like symptoms had similar symptoms to PASC patients. When analyzing their immune profile, post-vaccination individuals had statistically significant elevations of sCD40L, CCL5, IL-6, and IL-8. SARS-CoV-2 S1 and S2 protein were detected in CD16 + monocytes using flow cytometry and mass spectrometry on sorted cells.

Conclusions

Post-vaccination individuals with PASC-like symptoms exhibit markers of platelet activation and pro-inflammatory cytokine production which may be driven by the persistence of SARS-CoV-2 S1 protein persistence in intermediate and non-classical monocytes.

Immunology

COVID-19

PASC

long COVID

long haulers

SARS CoV-2 S1 protein

non-classical monocytes

CCR5

fractalkine

<https://www.researchsquare.com/article/rs-1844677/v1>

OPINION | VOLUME 28, ISSUE 7, P542-554, JULY 01, 2022

PDF [2 MB]

Figures

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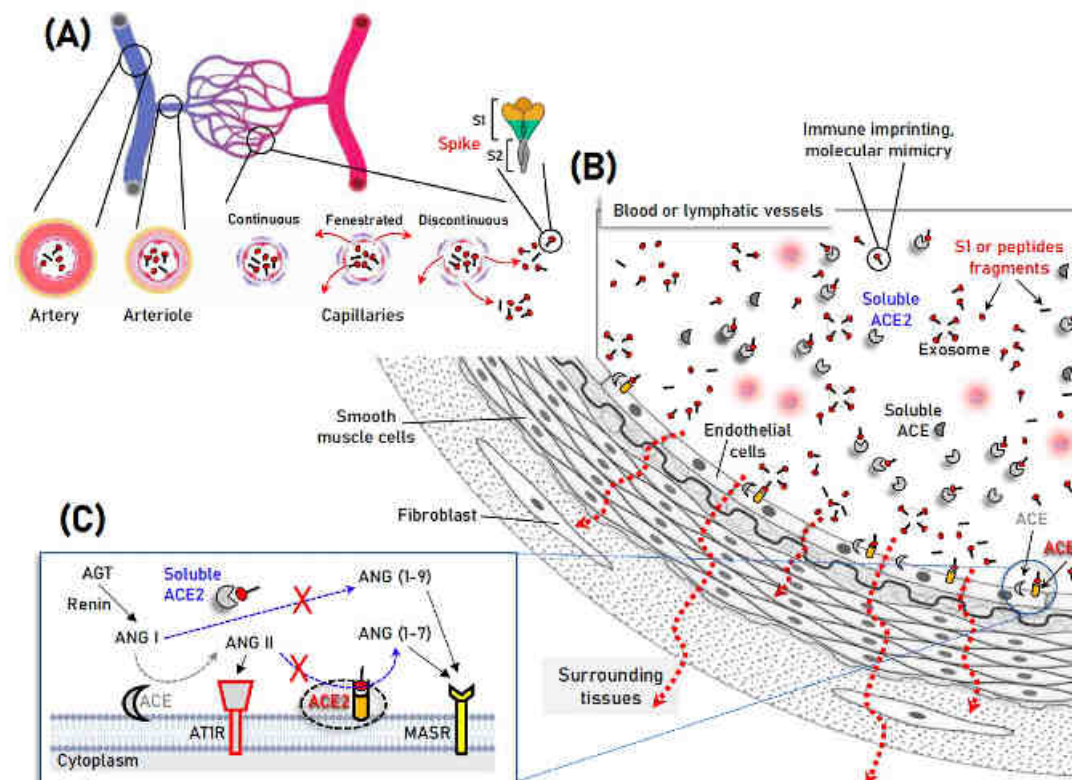
Reprints

Request

Adverse effects of COVID-19 mRNA vaccines: the spike hypothesis

Ioannis P. Trougakos   • Evangelos Terpos • Harry Alexopoulos • ... Efstathios Kastritis •
Evangelos Andreakos • Meletios A. Dimopoulos • Show all authors

Published: April 20, 2022 • DOI: <https://doi.org/10.1016/j.molmed.2022.04.007> •  Check for updates

 PlumX Metrics

Schematic of the vasculature components showing vaccination-produced S protein/subunits/peptide fragments in the circulation

Letter

The spike hypothesis
in vaccine-induced
adverse effects:
questions and
answers

Marco Cosentino ^{1,*} and
Franca Marino¹

- 1) Evidence for systemic biodistribution of COVID-19 vaccine induced S protein**
- 2) Adverse effects following COVID-19 vaccination: too much S protein, for too long and/or in the wrong place?**
- 3) COVID-19 mRNA vaccines: pharmaceutical drugs rather than conventional vaccines**

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Phase I/II study of COVID-19 RNA vaccine BNT162b1 in adults

[Mark J. Mulligan](#), [Kirsten E. Lyke](#), [Nicholas Kitchin](#), [Judith Absalon](#) , [Alejandra Gurtman](#), [Stephen Lockhart](#), [Kathleen Neuzil](#), [Vanessa Raabe](#), [Ruth Bailey](#), [Kena A. Swanson](#), [Ping Li](#), [Kenneth Koury](#), [Warren Kalina](#), [David Cooper](#), [Camila Fontes-Garfias](#), [Pei-Yong Shi](#), [Özlem Türeçli](#), [Kristin R. Tompkins](#), [Edward E. Walsh](#), [Robert Frenck](#), [Ann R. Falasay](#), [Philip R. Dormitzer](#), [William C. Gruber](#), [Uğur Şahin](#) & [Kathrin U. Jansen](#)

[Nature](#) **586**, 589–593 (2020) | [Cite this article](#)

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No grade 1 or greater change in routine clinical laboratory values or laboratory abnormalities were observed for most participants after either of the BNT162b1 vaccinations. Of those with laboratory changes, the largest changes were decreases in the lymphocyte count after the first dose in 8.3% (1 out of 12), 45.5% (5 out of 11) and 50.0% (6 out of 12) of participants who received 10 µg, 30 µg and 100 µg BNT162b1, respectively. One participant each in the 10-µg (8.3% (1 out of 12)) and 30-µg (9.1% (1 out of 11)) groups and 4 participants in the 100-µg group (33.3% (4 out of 12)) had grade 3 decreases in the lymphocyte count. These decreases in lymphocyte count after the first dose were transient and returned to normal 6–8 days after vaccination (Extended Data Fig. 1). In addition, grade-2 neutropenia was noted 6–8 days after the second dose in 1 participant each in the 10-µg and 30-µg BNT162b1 groups. These two participants continue to be followed in the study, and no adverse events or clinical manifestations of neutropenia have been reported to date. None of the post-vaccination abnormalities observed were associated with clinical findings.

ICH HARMONISED GUIDELINE

REVISION OF M4E GUIDELINE ON ENHANCING THE FORMAT AND
STRUCTURE OF BENEFIT-RISK INFORMATION IN ICH

EFFICACY - M4E(R2)

Current Step 4 version
dated 15 June 2016

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STRUCTURE OF BENEFIT-RISK INFORMATION IN ICH

EFFICACY - M4E(R2)

Current Step 4 version
dated 15 June 2016

elaborate analysis.

- examination of which subjects experience extreme laboratory value abnormalities ("outliers") may be useful in identifying subgroups of individuals who are at particular risk for certain adverse events.

Groups of studies that could be used in pooled safety analyses include:

https://database.ich.org/sites/default/files/M4E_R2__Guideline.pdf

There is an urgent need to assess any patterns of clinical laboratory findings occurring in subjects before and following vaccination, as it could be of major help:

- in the assessment, prediction and prevention of individual risk, allowing the identification of subgroups of individuals who are at particular risk for AE as well as the better understanding of the pathophysiological mechanisms underlying specific AE;**
- in the evaluation of suspect AE following vaccination, contributing to the definition of any eventual relationship between AE and the vaccines and encouraging the spontaneous reporting of the AE to the pharmacovigilance systems;**
- in general, in reassuring the public about the safety assessment process of these COVID-19 vaccines, consequently leading to a more safe and effective use of these vaccines and to the overcoming of vaccine hesitancy.**

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REPORT ESTESO ISS

COVID-19: SORVEGLIANZA, IMPATTO DELLE INFEZIONI ED EFFICACIA VACCINALE

Agglomerato nazionale
17/08/2022 - ore 12:00

DATA PUBBLICAZIONE: 19/08/2022

Fonte: Istituto Superiore di Sanità (ISS), Roma, 17 agosto 2022

TABELLA 5 - TASSO DI INCIDENZA DI DIAGNOSI DI INFEZIONE DA SARS-CoV-2, DI OSPEDALIZZAZIONE, DI RICOVERO IN TERAPIA INTENSIVA E DI DECESSO PER 100.000 E RISCHIO RELATIVO PER STATO VACCINALE E FASCIA DI ETÀ

Gruppo	Fascia di età	Tasso (per 100,000)				Rischio relativo		
		Non vaccinati	Vaccinati con ciclo completo >120 giorni	Vaccinati con ciclo completo >120 giorni	Vaccinati con ciclo completo + dose aggiuntiva/booster	Non vaccinati rispetto a vaccinati con ciclo completo >120 giorni	Non vaccinati rispetto a vaccinati con ciclo completo = 120 giorni	Non vaccinati rispetto a vaccinati con ciclo completo + dose aggiuntiva/booster
Diagnosi tra 15/07/2022-14/08/2022	12-39	2.947,1	1.653,5	2.572,1	2.426,6	1,8	1,1	1,2
	40-59	2.654,5	2.173,4	3.702,5	2.710,6	1,2	0,7	0,9
	60-79	3.448,3	4.233,5	3.329,5	2.754,5	0,8	1,0	1,3
	80+	5.100,0	2.355,2	2.637,2	2.395,6	2,2	1,9	2,1
	Totale	3.124,9	2.545,2	3.152,0	2.603,4	1,2	1,0	1,2
Diagnosi tra 01/07/2022-31/07/2022 con ospedalizzazione	12-39	31,9	13,4	16,2	12,5	2,4	2,0	2,6
	40-59	25,8	16,1	30,0	15,0	1,6	0,9	1,7
	60-79	153,6	95,8	97,9	51,3	1,6	1,5	3,0
	80+	1.124,4	301,5	382,4	224,4	3,7	2,9	5,0
	Totale	153,5	59,7	72,7	41,2	2,6	2,1	3,7
Diagnosi tra 01/07/2022-31/07/2022 con ricovero in TI	12-39	0,9	0,4	0,7	0,4	2,2	1,3	2,2
	40-59	1,7	0,7	1,1	0,8	2,4	0,4	2,1
	60-79	13,0	5,7	1,7	2,9	2,3	7,6	4,5
	80+	28,5	4,0	14,9	5,4	7,1	1,9	5,3
	Totale	6,6	2,2	3,3	1,5	3,0	2,0	4,1
Diagnosi tra 24/06/2022-24/07/2022 con decesso	12-39	0,2	0,1	0,0	0,1	2,0	Inf	2,0
	40-59	2,2	0,7	0,0	0,5	3,1	Inf	4,4
	60-79	19,1	11,6	16,2	5,3	1,6	1,2	3,6
	80+	327,2	75,2	90,1	48,2	4,4	3,6	6,5
	Totale	33,4	9,6	11,8	6,6	3,6	2,8	6,0

Note: Per maggiori dettagli vedere Nota metodologica paragrafo 4.

- Il tasso relativo alla popolazione complessiva ("Totale") equivale al tasso standardizzato rispetto alla popolazione 2022 (<https://demo.istat.it/>).

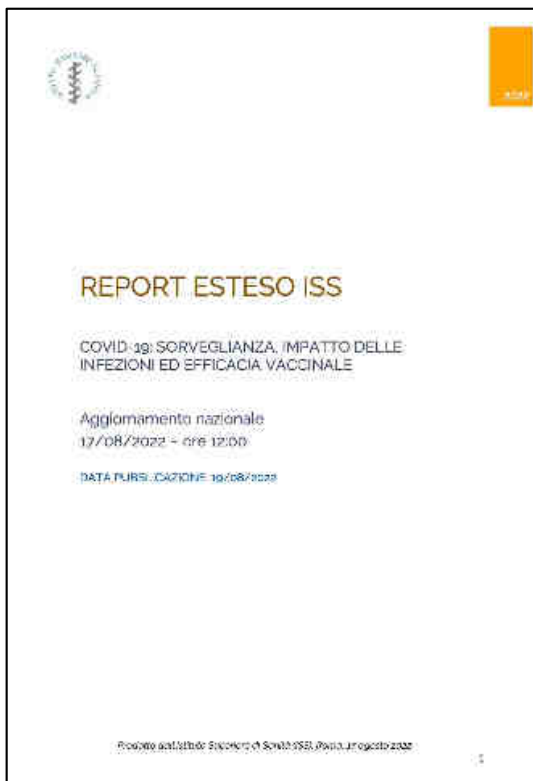
- Il tasso di incidenza degli ultimi 30 giorni potrebbe essere sottovalutato in quanto considero a rischio tutte le persone, tranne quelle che sono state diagnosticate e riportate alla sorveglianza negli ultimi 3 mesi. A causa dell'elevato numero di nuove infezioni, spesso non diagnosticate o autodiagnosticate e quindi non riportate alla sorveglianza, il numero delle persone a rischio considerate per il calcolo del tasso di incidenza è quindi verosimilmente sovrastimato, in particolare nelle fasce 12-39 e 40-59. E inoltre verosimile la presenza di una più elevata sottorapportata delle diagnosi nella popolazione non vaccinata e vaccinata da oltre 120 giorni. (Per maggiori dettagli vedere Nota metodologica paragrafo 4).

Prodotto dall'Istituto Superiore di Sanità (ISS), Roma, 17 agosto 2022



Gruppo	Fascia di età	Non vaccinati
Diagnosi tra 15/07/2022-14/08/2022	12-39	2.947
	40-59	2.554
	60-79	3.448
	80+	5.100
	Totale	3.124
Diagnosi tra 01/07/2022-31/07/2022 con ospedalizzazione	12-39	31
	40-59	25
	60-79	153
	80+	1.124
	Totale	153
Diagnosi tra 01/07/2022-31/07/2022 con ricovero in TI	12-39	0
	40-59	1
	60-79	13
	80+	28
	Totale	6
Diagnosi tra 24/06/2022-24/07/2022 con decesso	12-39	0
	40-59	2
	60-79	19
	80+	327
	Totale	33

Note: Per maggiori dettagli vedere Nota n. 1



???

???

???

Gruppo	Fascia di età	Non vaccinati
Diagnosi tra 15/07/2022-14/08/2022	12-39	2.947
	40-59	2.554
	60-79	3.448
	80+	5.100
	Totale	3.124
Diagnosi tra 01/07/2022-31/07/2022 con ospedalizzazione	12-39	31
	40-59	25
	60-79	153
	80+	1.124
	Totale	153
Diagnosi tra 01/07/2022-31/07/2022 con ricovero in TI	12-39	0
	40-59	1
	60-79	13
	80+	28
	Totale	6
Diagnosi tra 24/06/2022-24/07/2022 con decesso	12-39	0
	40-59	2
	60-79	19
	80+	327
	Totale	33

Note: Per maggiori dettagli vedere Nota n. 1

Remote management of COVID-19 patients at home: the neglected role of drugs

Marco Cosentino^{1*}, Veronica Vernocchi^{2*}, Franca Marino¹, Mauro Rango²

¹Center for Research in Medical Pharmacology, University of Insubria, Varese, Italy, and

²IppocrateOrg Association, Lugano, Switzerland

*: these authors provided equal contributions to the study.

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L'INSOSTENIBILE IMPREVEDIBILITA' DEI VACCINI COVID-19

1) Farmacologia dei vaccini a RNA

Ovvero: giocare a dadi con la spike

2) Analisi di laboratorio negli studi autorizzativi

Cosa comporta il fatto che non siano state svolte?

3) Curare il COVID-19

Ruolo delle cure nella valutazione dell'efficacia vaccinale

Umanità



Ragione