COVID-19 Vaccine Associated Cutaneous Lymphomatoid Reactions

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COVID-19 VACCINE

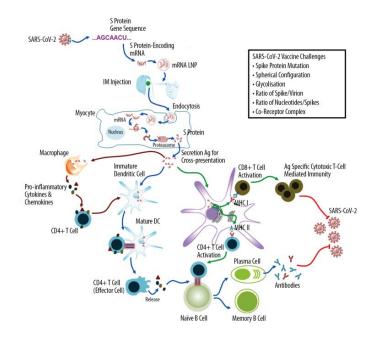
- Although there were nearly 50 candidate vaccines in clinical testing ultimately three became approved by the FDA with most patients receiving the Moderna or Pfizer vaccine
 - Moderna/NIAID Nucleoside-modified messenger RNA that encodes the full length of SARS-CoV-2 spike (S) protein (mRNA-1273)
 - Pfizer/BioNTech Nucleoside-modified messenger RNA that encodes the full of the SARS-CoV-2 spike protein (BNT162b2)
 - Janssen(Johnson and Johnson) modified adenovirus (Ad26) encodes full length S protein (Ad26.COV2.S)

Family: Coronaviridae Δ Subfamily Torovirinae Coronavirinae Genus Deltacoronavirus Alphacoronavirus Betacoronavirus Gammacoronavirus BuCoV-HKU11 Avian coronavirus Lineage Primarily infect birds HCoV-229E HCoV-OC43 SARS-CoV MERS-CoV Bt-HCoV-NL63 HCoV-HKU1 SARS-CoV-2 BtCoV-HKU-4 CoV-HKU-9 BtCoV-HKU-5 Primarily infect mammals в Spike protein (S) C Receptor binding motif Membrane S1 subunit glycoprotein (M) (Globular receptorbinding domain) - S2 subunit (Stalk fusion domain) . . . Viral membrane RNA and Intracellular tail Nucleocapsid protein (N) protein (E) n S'-UTR Orflab mannen 5 orf3a E M orf6 orf7a orf7b orf8 N orf10 3'-UTR -- S1 Subunit -----S 2 Subunit ----- RBD ------HRI HR NTD 53.52 CTD \$15.826

Fig 3. Classification and structure of coronavirus.

Mittal A, Manjunath K, Ranjan RK, Kaushik S, Kumar S, et al. (2020) COVID-19 pandemic: Insights into structure, function, and hACE2 receptor recognition by SARS-CoV-2. PLOS Pathogens 16(8): e1008762. https://doi.org/10.1371/journal.pbat.1008762 https://journals.plos.org/olospathogens/article/10.1371/journal.pbat.1008762.

PLOS PATHOGENS

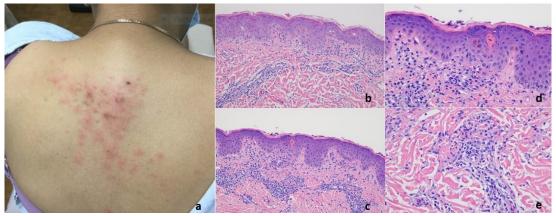




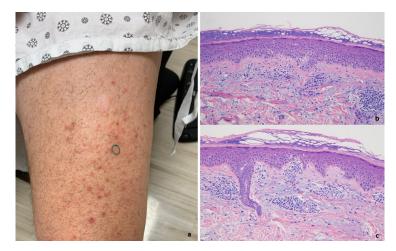
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Results

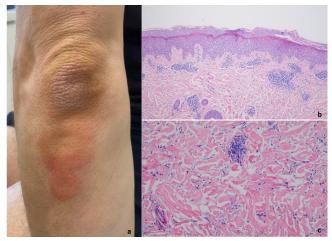
- Classic clinical and morphologic depictions of type IV cutaneous hypersensitivity with features of eczematous dermatitis, interface dermatitis, interstitial granulomatous inflammation, and/or granulomatous and or lymphocytic vasculitic component
- Clinical and/or histologic features of perniosis, pityriasis rosea, pityriasis rubra pilaris, and guttate psoriasis
- · Leukocytoclastic vasculitis, possibly reflective of an Arthus type III immune complex action
- Biopsy specimens of normal deltoid (standard site for assessing for systemic complement
 pathway activation) skin post vaccine and of skin affected by the post-vaccine eruption
 showed rare deep microvessels positive for spike glycoprotein with no complement
 deposition contrasting with greater vascular deposition of spike protein and complement in
 deltoid skin biopsies from patients experiencing severe COVID-19.
- These reactions developed within 48 hours up to 4 weeks after the first or second dose of Pfizer or Moderna vaccine.



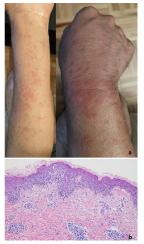
Type IV T-cell response with features of interface dermatitis and interstitial granulomatous inflammation (case 1). The patient was a 38-year-old woman who presented with blanchable erythematous papules (reproduced with permission from Dr. Henry J. Lee, New York, NY). The patient had received a COVID-19 vaccine, either the Moderna or the Pfizer, on February 15, 2021. (B, C) The biopsy showed a lymphocyte-mediated interface dermatitis associated with focal areas of epidermal attenuation. Lymphocyte satellitosis is visible around injured keratinocytes. (D) Focal areas of interstitial granulomatous inflammation accompanied by some degree of mesenchymal mucin deposition were observed. COVID-19, coronavirus disease 2019.(E)



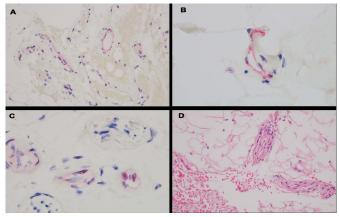
T-cell–mediated cytotoxic interface dermatitis (case 5). A 67-year-old man presented with a 2-week history of pruritic eruption on the trunk and extremities. The patient received the first and second doses of the Moderna vaccine on January 15, 2021 and February 16, 2021, respectively. (A) He developed an itchy eruption after the second dose (reproduced with permission from Dr. Silvia Mancebo, New York, NY). (B, C) The histologic findings are those of a classic morbilliform type IV hypersensitivity reaction combining delayed dermal hypersensitivity with a very mild cytotoxic interface dermatitis



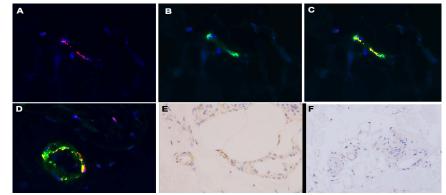
T-cell response predominated by eczematous features (case 16). (A) The patient is a 37-year-old man who developed an eruption that began on the elbows in February of 2021 and then subsequently spread to the knees (*reproduced with permission from Dr. Andrew Avarbock, New York, NY*). The eruption on the elbows and knees improved, but subsequently spread to the lower legs. (B) The epidermal changes are predominated by eczematoid alterations, but a subtle interface dermatitis is also noted. (C) The dermal component exhibits features of delayed dermal hypersensitivity characterized by vasocentric lymphocytic and eosinophilic infiltrates along with an interstitial granulomatous component.



T-cell response predominated by eczematous features (case 6). (A) A 34-year-old man presented with papulovesicular eruptions on the extremities, hands, and palms 1 week after receiving the Moderna vaccine (reproduced with permission from Dr. Paul Dantzig, New York, NY). The patient was treated with prednisone 60 mg daily, and the eruption cleared 3 days later. The biopsy showed intercellular edema within the epidermis along with lymphocytic exocytosis. A concomitant interface dermatitis was identified as evidenced by basilar vacuolar change with a few lymphocytes present along the dermal-epidermal junction. (B) Scattered eosinophils are noted.



Normal deltoid skin in patients with fatal COVID-19 versus healthy patients after vaccination. (A) In both deltoid skin samples from patients with fatal COVID-19 versus healthy patients after vaccination, there are a relatively greater number of positive staining vessels for ACE2 in the deeper dermis and in subcutaneous fat compared with the microvessels present superficially (red chromagen, 200 ×). The ACE2 distribution pattern mirrors spike glycoprotein endothelial cell localization. (B) Granular deposition within the endothelium for spike glycoprotein was present in the setting of fatal COVID-19 (red chromagen, 1000 ×). (C) A similar pattern of endothelial cell staining for spike glycoprotein was noted in the post-vaccine biopsy (red chromagen, 1000 ×). (D) A microvessel after vaccine in which a mononuclear cell response is evident (hematoxylin and eosin, 1000 ×). ACE2, angiotensin converting enzyme 2; COVID-19, coronavirus disease 2019.



Patients with normal deltoid skin after vaccination shows endothelial cells activated by the spike protein. (A) Using the Nuance software, a microvessel positive for spike glycoprotein using a red chromagen gives a red signal localized to endothelium (red chromagen, 1000 ×). (B) The same microvessels show a green signal in a similar distribution highlighting interleukin 6, using a diaminobenzidine stain (diaminobenzidine, 1000 ×). (C) The combined signal fluoresces yellow, proving colocalization of spike glycoprotein and interleukin 6 in the biopsy of a person post-vaccination (1000 ×). (D) Panel demonstrates a similar strong coexpression between the spike protein and caspase 3 in a post-vaccine biopsy specimen (1000 ×). (C) An extensive microvascular deposition of C5b-9 is visible in a case of severe COVID-19 (diaminobenzidine, 400 ×). (F) In contrast, the post-vaccine sample does not show significant complement deposition (diaminobenzidine, 400 ×). COVID-19, coronavirus disease 2019.

What is the basis of these vaccine triggered reactions in the skin ? Pathophysiology of the vaccine associated reactions

- An adaptive T or B cell immune response to spike glycoprotein although as a generalized immune response given the small amount of human synthesized spike in skin samples
- A type IV immunologic response to a component of the vaccine vehicle
- Unmasking of subclinical cutaneous eruptions with a genetic component (psoriasis/atopic eczema) or subclinical hypersensitivity due to immune boosting properties of the vaccine

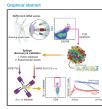
Pathophysiology of the type IV immune reactions: The role of superantigens and the preferential follicular helper T cell response

- The pattern of interface and interstitial granulomatous dermatitis with granulomatous/lymphocytic vasculitis is among the commonest histologic COVID-19 vaccine reaction patterns and is also typical hypersensitivity reactions triggered by microbial pathogens with superantigen properties(Magro CM et al, 1998).
- Spike glycoprotein has potent superantigen properties and therefore can stimulate a significant component of the T cell repertoire(Cheng MH et al, 2020).
- Spike glycoprotein induces a robust follicular helper T cell response(Mudd PA et al, 2022)

Cell

Article

SARS-CoV-2 mRNA vaccination elicits a robust and persistent T follicular helper cell response in humans



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In brief

Analysis of draining lymph nodes of individuals vaccinated with BNT162b2 mRNA vaccine against SARS-CoV-2 identifies viral-spike-specific follcular helper CD4* T cells that persist for months and contribute to long-term immunity.

Highlights

- SARS-CoV-2 vaccines induce robust human T_{FH} cell responses in draining lymph nodes
- HLA-DPB1*04 restricts the immunodominant SARS-CoV-2 S₁₄₇₋₁₈₀ epitope
- S_{nar-tao} is recognized by T cell receptors with a public *x*-chain motif
- S-specific T_{FH} cells are maintained in draining lymph nodes 6 months after vaccination



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Superantigenic character of an insert unique to SARS-CoV-2 spike supported by skewed TCR repertoire in patients with hyperinflammation

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Contributed by lvet Bahar, June 29, 2020 (sent for review May 26, 2020; reviewed by Talal A. Chatila, Ruth Nussinov, and Celia A. Schiffer)

Multisystem Inflammatory Syndrome in Children (MIS-C) associated with COVID-19 is a newly recognized condition in children with recent severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) infection. These children and adult patients with severe hyperinflammation present with a constellation of symptoms that strongly resemble toxic shock syndrome, an escalation of the cytotoxic adaptive immune response triggered upon the binding of pathogenic superantigens to T cell receptors (TCRs) and/or major histocompatibility complex class II (MHCII) molecules. Here, using structure-based computational models, we demonstrate that the SARS-CoV-2 spike (S) alvcoprotein exhibits a high-affinity motif for binding TCRs, and may form a ternary complex with MHCI. The binding epitope on S harbors a seguence motif unique to SARS-CoV-2 (not present in other SARS-related coronaviruses), which is highly similar in both sequence and structure. to the bacterial superantigen staphylococcal enterotoxin B. This interaction between the virus and human T cells could be strengthened by a rare mutation (D839Y/N/E) from a European strain of SARS-CoV-2. Furthermore, the interfacial region includes selected residues from an intercellular adhesion molecule (ICAM)-like motif shared between the SARS viruses from the 2003 and 2019 pandemics. A neurotoxin-like sequence motif on the receptor-binding domain also exhibits a high tendency to bind TCRs. Analysis of the TCR repertoire in adult COVID-19 natients demonstrates that those with severe hyperinflammatory disease exhibit TCR skewing consistent with superantigen activation. These data suggest that SARS-CoV-2 S may act as a superantigen to trigger the development of MIS-C as well as cytokine storm in adult COVID-19 patients, with important implications for the development of theraneutic annroaches

COVID-19 | superantigen | SARS-CoV-2 spike | toxic shock syndrome | TCR binding

Service acute respiratory syndrome consorting 2 (5488-CoV-52), the virtual factors COVLD-19 is a heatcorrestoring to the syndrome international syndrome in the syndrome targe Syndrome (MESS)-CoV (1), COVLD-19 can marifest in adds as a server interial in potentiaria that heperializations, while severe meginatory manifestations are rate in differen (2-4). (MISC) has been observed in patients that chief the total opation (MISC) has been observed in patients that chief total opation links to COVID-19 (5-7). Alter of the syndrome in the syndrome in the syndrome COVID-19 (5-7). Alter of the syndrome in the syndrome in the syndrome (MISC) has been observed in patients that chief total position links to COVID-19 (5-7). Alter of the syndrome in the syndrome in the syndrome syndrome in the syndrome in the syndrome in the syndrome in the syndrome syndrome in the syndrome in the

shock syndrome (TSS) (8, 9) (Table 1), rather than Kawasaki disease (KD), due to marked demographic, clinical, and laboratory differences (6). Indeed, a recent uncontrolled retrospective case study concluded that MIS-C is distinct from KD and KD shock syndrome (10). The similarities to TSS and the association of MIS-C with COVID-19 led us to hypothesize that SARS-CoV-2 may possess superantigenic fragments that induce an inflammatory cascade and may also contribute to the hyperinflammation and cytokine storm observed in severe adult COVID-19 patients (3, 4). The question we raised is, does SARS-CoV-2 spike (S) possess superantigenic fragments that could elicit such reactions upon binding proteins involved in the host cell cytotoxic adaptive immune response? Such a reaction was not observed in the SARS-CoV pandemic of 2003 (SARS1 hereafter). What is unique to SARS-CoV-2, and how might recent mutations in SARS-CoV-2 S have promoted such an increased virulence?

TSS can be caused by two types of superantigens (SAgs): bacterial or viral. Bacterial SAgs have been broadly studied. They include proteins secreted by *Staphylococcus aureus* and *Streptococcus pyogenes* that stimulate massive production of

Significance

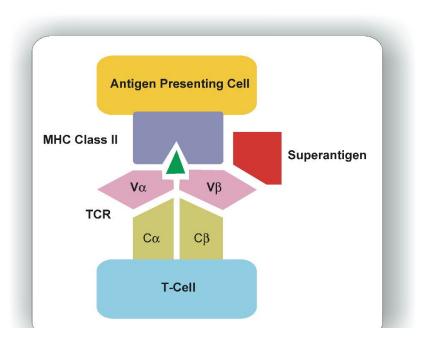
A hyperinflumnatory syndrome neministent of task shock syndrome (SS) is descered in severe COPD patterts, Indular, ability, why Multipaten influmnatory Syndrome in Collient with the syndrome influence of the syndrome in Collient and the syndrome influence of the short syndrome in the short with SMS-COV-2 spike contains sequence and structure motifs hyperinfluence in the short single sequence and structure motifs hyperinfluence in the syndrome influence of the short single sequences sequence in a syndrome in the syndrome influence of the sequence of the syndrome in the syndrome influence of the late motif and parsent in other SMS family concentrates, which differs and the syndrome is motioned and do CODD-13.

Author contributions: M.H.C., M.A., and I.B. designed research; M.H.C., S.Z., R.A.P., L.P., E.W., M.B., M.A., and I.B. performed research; M.H.C., S.Z., R.A.P., M.N.R., M.B., M.A., and I.B. analyzed data; and M.H.C., R.A.P., M.A., and I.B. wrote the paper.

Reviewers: T.A.C., Children's Hospital Bostory, R.N., Frederick National Laboratory, and C.A.S., University of Massachusetts Medical School.

Competing interest statement: Patent filing process has been started for short peptide sequences to neutralize the superantigenic fragment.

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The T Cell lymphomatoid reactions in the setting of the COVID-19 Vaccine Three cases have been encountered in the routine and consultative practice at Weill Cornell Medicine where these apparent vaccine reactions appeared atypical cytomorphologically, architecturally and phenotypically.

A 35 year old male who developed primary cutaneous CD4+ small medium sized T cell lymphoproliferative disease 1 week after the first dose of the Pfizer vaccine.

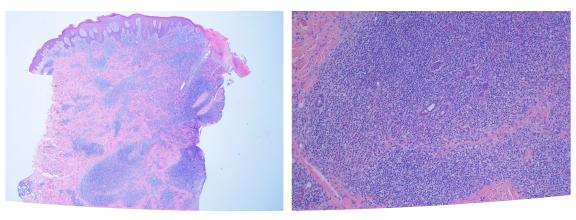
A 18 year old male who developed LYP within 1 month following the second dose of the Pfizer vaccine, responding to MTX and then recurring when he had a COVID-19 break through infection.

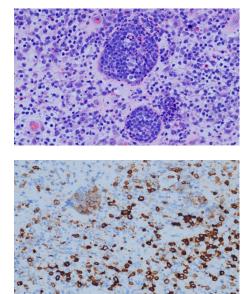
A 63 year old female who developed CD8+ CD25+ peripheral T cell lymphoma NOS after the second dose of the Pfizer vaccine

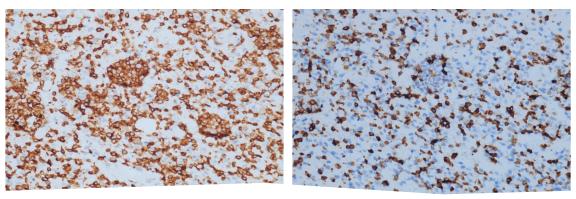
Prior reported cases include recurrent CD30+ primary cutaneous LPD(2 days after first dose of Pfizer in a 79 year old male)(Brumfiel et al,2021), the progression of AILD(Goldman S et al,2021)(66 year old male developing AILD within 5 months of receiving the Pfizer and then worsening of the AILD with the booster

Clinical History

- 18-year-old healthy male who abruptly developed a papular nodular skin rash in his right axilla after he received the second SARS-CoV-2
- The lesions began as a singular nodule, which then diminished with the development of multiple others within the same location
- He later developed an oval pink papule on his left thigh
- Both were biopsied as LYP overlapping type A and type C
- Began methotrexate with improvement, and stopped due a mononucleosis infection while at college
- Was doing well without lesions, until a recent COVID-19 infection where he then developed new pink crops of papules on the shaft of his penis
- He was restarted on methotrexate



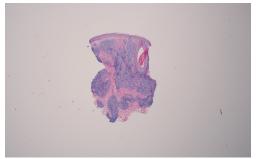


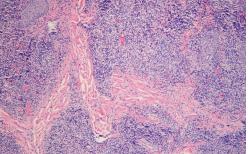


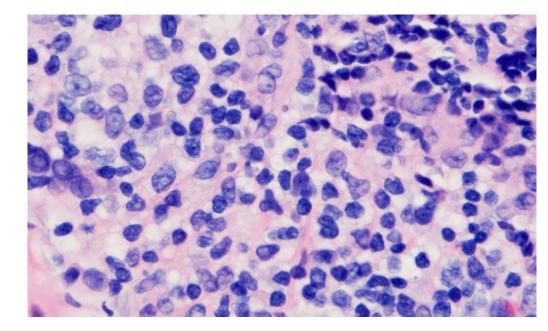


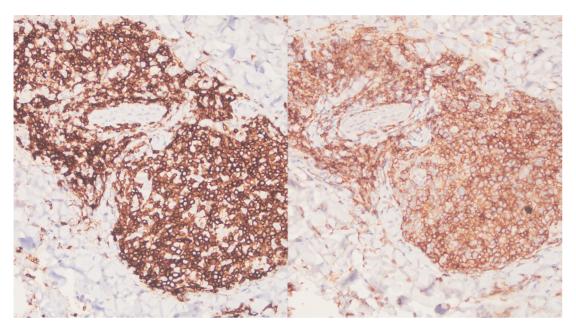
Clinical history

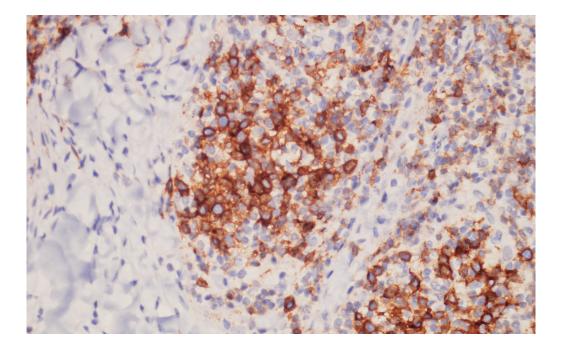
- 35-year-old male with no significant past medical history developed a nodule on his forehead 1 week after his first dose of the Pfizer-BioNTech COVID-19 vaccine
- The lesion was stable and otherwise asymptomatic for 3 months, and the patient had no systemic symptoms
- Physical examination revealed a 0.5 x 1.0 cm non-tender, nonulcerated nodule on the mid forehead
- A punch biopsy was performed
- ٠











Case of post vaccine PTCL, NOS

63-year-old female with hypertension developed a non-itching rash on her bilateral arms and legs after her $2^{\rm nd}$ Pfizer COVID-19 vaccine.

The initial biopsies in April/May were interpreted as a delayed hypersensitivity reaction followed by a biopsy in June 2021 interpreted as probable hypersensitivity but with atypical lymphomatoid features versus an evolving T cell LPD.

The rash resolved after systemic corticosteroids

The patient developed erythematous patches on her face, extremities and trunks after she received the vaccine booster

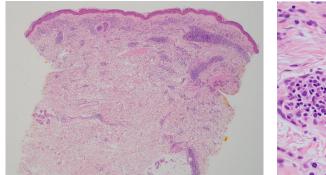
Again, systemic corticosteroids succeeded in resolving the rash

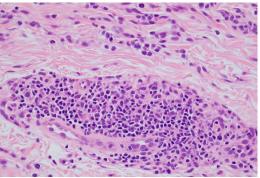
The patient then had breakthrough COVID and quickly recovered, however she developed new violaceous patches, plaques as well as nodules(November 2021, biopsies were performed on patches and nodules)

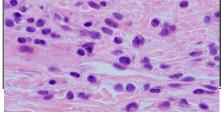
She has continued to develop new skin lesions and now has extracutaneous lymph node involvement.





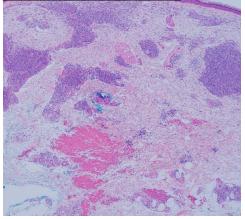


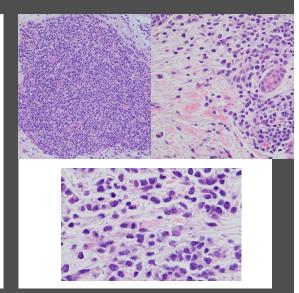


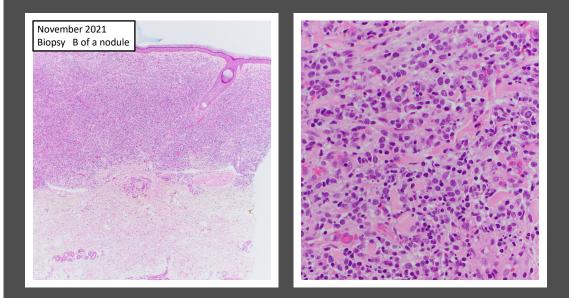


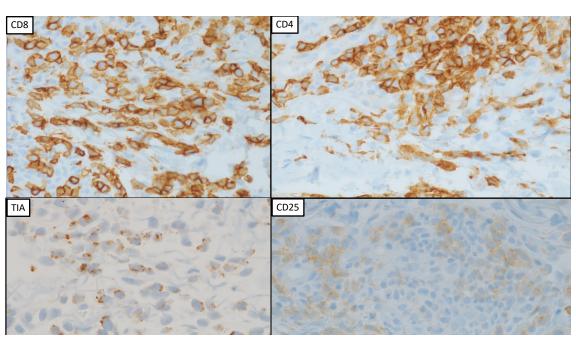
Initial Set of Biopsies in June 2021











Summary of Light Microscopic Findings

- Progression of an initial type IV immune response to the vaccine to a progressive atypical lymphomatoid process whereby throughout the biopsies there was evidence of an identical T cell clone and a rare unique T cell was implicated namely a CD8+ or double CD25+ positive non-regulatory cytotoxic CD8+ memory T cell.
- Earlier biopsies could be interpreted as a vaccine triggered CD8+ T cell lymphoproliferative disorder of non-regulatory CD8+ memory T cell origin.
- Subsequent biopsies showed progression to overt lymphoma consistent with a CD8+ positive peripheral T cell lymphoma of non-regulatory CD8+ memory T cell origin as a unique form of immunosenescent lymphoma, NOS.

Is there a benign T cell Counterpart that could be preferentially expanded in response to spike glycoprotein in the middle aged to older age setting? IL2/IL4 producing CD8+CD25+ non-regulatory memory T cells are seen in older patients with depleted naïve T cells

- A very rare T cell subset is represented by an IL-2/IL-4-producing CD8+CD25+ nonregulatory memory T cell population that occurs in a subgroup of healthy elderly persons who have depleted naïve T cells; these CD8 positive T cell can show a double positive phenotype(Herndler-Brandstetter et al 2005).
- This cell population represents a critical reservoir for launching an intact immune response to viruses and to a vaccine; the presence of double positive memory T cells in fact has been correlated with a better outcome in patients with COVID-19(Kalpakci et al, 2020,(Nascimbeni et al 2004).

How do we link the spike glycoprotein mediated type IV immune reactions to COVID-19 vaccine associated T cell Lymphoproliferative disease?

A multistep process involved in lymphomagenesis

Initial antigenic stimulus leads to a clonally restricted T or B cell response.

In the setting of underlying iatrogenic or endogenous immune dysregulation (underlying autoimmune disease or a genetic predisposition), an overzealous clonal immune response occurs.

Subsequent oncogenic transformation of clonally restricted T or B cells unrelated to any properties of the vaccine or virus eventuates in endogenous T/B cell lymphoproliferative disease.

The nature of the adaptive T cell response to spike glycoprotein could determine the type of vaccine associated LPD

- 1. In healthy patients especially younger patients a robust type IV immune response mediated by activated antigen specific CD4+ or cytotoxic CD8 T cells is characteristic (benign reaction is the typical type IV hypersensitivity vaccine reaction; atypical counterpart would be CD4+ CD30+ LYP).
- 2.Follicular helper T cells are also preferentially expanded by spike glycoprotein(atypical counterpart is primary cutaneous follicular helper T cell dyscrasia (i.e. primary cutaneous CD4+ small medium sized pleomorphic T cell LPD) and angioimmunoblastic lymphoma).
- 3. In older individuals where the naïve T cell repertoire is significantly diminished, expansion of CD25+CD8+ T cells occur in response to SARS CoV-2 and is protective against severe disease (atypical counterpart would be CD25+CD8+ PTCL NOS)