

Proteins that contaminate influenza vaccines have high homology to SARS-CoV-2 proteins thus increasing risk of severe COVID-19 disease and mortality

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Abstract

Influenza vaccines are manufactured using chicken eggs, canine kidney cells or insect cells. Chicken and dogs can be infected with numerous viruses including coronaviruses. Therefore influenza vaccines can be contaminated with coronavirus proteins. These coronavirus proteins of course have high homology to SARS-CoV-2 proteins. Further, we show that even chicken egg proteins have high homology to SARS-CoV-2 proteins. Influenza vaccine administration is known to create IgE mediated sensitization (allergy) to all proteins in the vaccine. In fact, vaccine-induced egg allergy is required for the vaccine to work and protect against influenza. This is because most influenza vaccines lack an adjuvant and they depend on the allergic reaction at the injection site to provide an adjuvant effect.

Upon COVID-19 infection, patients will suffer an allergic reaction due to cross reaction against SARS-CoV-2 proteins. This predictably produces anaphylaxis symptoms. Since the viral load increases over a few days, we have slow rolling anaphylaxis. The result is damage to multiple organs. Acute respiratory distress syndrome results from lung damage. Cardiac injury due to Kounis syndrome. Coagulation dysfunction, hypotension and shock are other possible outcomes. As in anaphylaxis treatment, epinephrine, histamine H1 and H2 blockers, etc. help prevent or treat these conditions.

Introduction

Most influenza vaccines are manufactured using specific pathogen free (SPF) chicken eggs. The SPF specification covers 18 pathogens but that list (1) does not include coronaviruses. So chicken and the SPF eggs they produce can be infected with coronaviruses. Madin-Darby Canine Kidney (MDCK) cells are also used to manufacture some influenza vaccines (2).

MDCK cells are susceptible to coronavirus infections (3). So MDCK cells derived from infected dogs can be contaminated with coronavirus proteins. Contamination of vaccines with viruses from infected animal tissues used in vaccine manufacturing, is not uncommon. The rotavirus vaccine contains viable porcine circoviruses, for example (4,5).

Influenza vaccine protection depends on vaccine-induced egg allergy

Influenza vaccines create long term persistent IgE mediated sensitization (allergy) to all proteins in the vaccine, including the influenza viral proteins (6–10). Upon subsequent influenza infection, people can suffer an allergic reaction to the influenza virus itself, which similar to COVID-19, can lead to a cytokine storm and influenza shock syndrome (11).

Jacob et al. (12). identified 293 chicken proteins in the influenza vaccine. Injecting such chicken egg proteins into humans induces long term persistent IgE mediated sensitization to these proteins (10). Subsequently administered egg protein-containing influenza vaccines elicit an allergic reaction at the injection site. This reaction provides the innate immune system costimulation required (adjuvant effect) to produce the protective immune response against influenza proteins in the vaccine. This is the reason why when the influenza vaccine is administered for the very first time, it does **not** produce an effective protective response. The CDC recommends a second dose of the vaccine to address this issue when a child receives an influenza vaccine for the first time (13). In other words, the influenza vaccine depends on sickening people with egg allergy, to produce its protective effect.

The Flublok influenza vaccine is egg-free (14). So as expected, it would elicit only a weak protective response. The solution? They increased the antigen content to 300% compared to a regular egg derived influenza vaccine. So the Flublok vaccine contains 45 mcg of each hemagglutinin (HA) protein compared to 15 mcg in the regular vaccine.

Vaccine-induced allergy causes COVID-19 severity

Therefore, as above, coronavirus proteins that contaminate the influenza vaccine as well as chicken proteins that have similarity to SARS-CoV-2 proteins, cause allergic sensitization to all those proteins. Upon subsequent COVID-19 infection, due to the cross reaction, people will suffer a severe allergic reaction which explains COVID-19 severity in a section of the population. Detailed BLASTP protein sequence analysis results showing homology between chicken and SARS-CoV-2 proteins are provided in a later section. Nursing home residents are likely to have higher influenza vaccine coverage. So predictably, they have a very high risk of suffering severe COVID-19.

This is the reason why allergy medications such as antihistamines (cetirizine, famotidine) and mast cell stabilizers help in COVID-19 (15–17). In general, since COVID-19 is simply a slow rolling version of anaphylaxis, proven anaphylaxis treatments will work (18).

Cardiac injury that occurs due to an allergic reaction is known as Kounis syndrome. The same injury occurs in influenza shock syndrome, COVID-19 and dengue because they all involve an allergic reaction. The details were previously described (19).

Conclusion

Contaminated vaccines, including influenza vaccines, contribute to COVID-19 severity and mortality. While this article focused on influenza vaccines, all vaccines have a similar problem (5). COVID-19 severity is the latest to be added to the huge list of diseases caused by contaminated vaccines (20).

Epidemiological studies are unreliable. One study will claim influenza vaccine is associated with COVID-19 severity and another one will claim it is not associated. The Institute of Medicine (IOM) now known as the National Academy of Medicine (NAM) found that vaccine safety epidemiological studies are useless, 93% of the time (21). We present reliable, mechanistic evidence of causation.

Detailed BLASTP result examples

QHD43415.1 orf1ab polyprotein [Severe acute respiratory syndrome coronavirus 2] vs.

[XP_025008461.1](#) poly [ADP-ribose] polymerase 14 isoform X3 [Gallus gallus]

Alignment statistics for match #1

Score Expect Method Identities Positives Gaps
60.1 bits(144) 1e-07 Compositional matrix adjust. 57/183(31%) 86/183(46%) 22/183(12%)

```
Query 1056 VVVNAANVYLKHGGVAGALNKATNNAMQVESDDYIATNGPLKVGGSVLSGHNLA-KHC 1114
          VVVNA+N LKH GG+A AL +A +Q E D + +G L+ G + + L K
Sbjct 864 VVVNASNEDLKHIGGLAWALLQAAGPELQAECDGVVRMSGSLQAGDAVITGAGKLPCKQV 923

Query 1115 LHVVGPNVNGED---IQLLKSA-----ENFNQHEVLLAPLLSAGIFGADPIHSLRV 1164
          +H VGP + + + LLK E +N H + P +S GIFG P+H
Sbjct 924 IHAVGPRWKEQDAEKCVYLLKKTIKKSLQLAETYN-HRSIAFPSVSGGIFGF-PLHK--- 978

Query 1165 CVDTVRTNVYLVAFDKNLYDKLVSSFLEMKSEKQVEQKIAEIPKEEV-KPFITESKPSVE 1223
          CV N ++ K L + S L+ V+++ + +E V K F +S SV
Sbjct 979 CV-----NAIVSAIKKTLEEFKRDSLKEIHLVAVDEETVRVLRRETQKEFTAKSSSVL 1033

Query 1224 QRK 1226
          Q++
Sbjct 1034 QQQ 1036
```

QHD43415.1 orf1ab polyprotein [Severe acute respiratory syndrome coronavirus 2] vs.

[XP_015131826.1](#) protein ZGRF1 [Gallus gallus]

Alignment statistics for match #1

Score Expect Method Identities Positives Gaps
55.5 bits(132) 3e-06 Compositional matrix adjust. 68/258(26%) 112/258(43%) 52/258(20%)

```
Query 5694 IVVFDEISMATNYDLSVVNARLRAKHVYVIGDPAQLP----APRTLLTKGTLEPEYFNSV 5749
          +VV DE S T + AR + + V +GDP QLP ++ KG LE F+ +
Sbjct 1488 VVVLDECSQMTEPTSLLPiARFQCEKLVLVGDPKQLPPTIQGSESIHEKG-LEQTLFDRL 1546

Query 5750 CRL-MKTIGPDMFLGTCRRCPAEIVDTVSALVYDNKL---KAHKDKS-----AQCF-- 5796
          C + KTI L T RC I + L Y+ L + K++S CF
Sbjct 1547 CLMGHKTi----LLRTQYRCHPAISAIANELFYEGNLIDGVSEKERSPLLDWLP TLCFYS 1602

Query 5797 -----KMFYKGVITH-----DVSSAINRPQIGVVREFLTRNPAWRKAVFISP 5838
          FY H ++S I+ +GV+ + ++ ++ +
Sbjct 1603 VNGLEQIERDNSFYNMAEVHFTVKFIQALIASGIDGSAVGVITFY--KSQMYKLQNL LRS 1660

Query 5839 YNSQNAVASKILGLPTQTVDSQSEYDYVIFT--QTTETAHSCNVNRFNVAITRAK--- 5893
          +S+ A + + TVD+ QG+E + V+ + +T +T + + R NVA+TRAK
Sbjct 1661 IHSE---AFPVKAVQVSTVDAFQGAKEIIVLSCVRTRQTGFTDSEKRMNVALTRAKRHL 1717

Query 5894 --VGILCIMS DRDLYDKL 5909
          VG L +S LY+++
Sbjct 1718 LIVGNLACLSKNRLYERV 1735
```

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