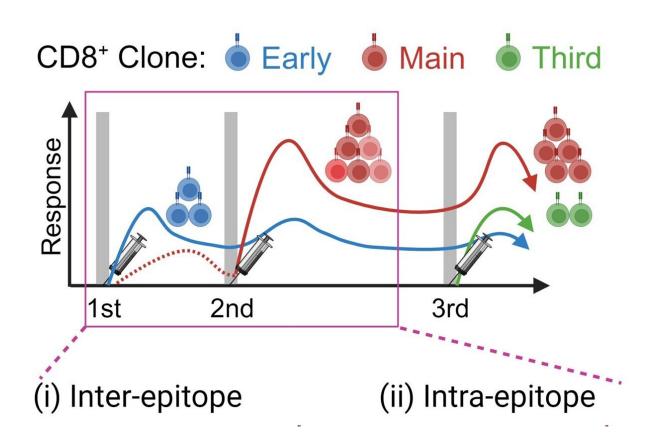


Optimizing boosters: How COVID mRNA vaccines reshape immune memory after each dose

March 8 2024



Changes in the clonal populations of CD8⁺ T- cells after successive vaccinations, an ability that can be tuned for developing more effective mRNA vaccines. Credit: Satoshi Ueha/Tokyo University of Science



mRNA vaccines developed against the spike glycoprotein of severe acute respiratory syndrome type 2 coronavirus (SARS-CoV-2), displayed remarkable efficiency in combating coronavirus 19 (COVID-19). These vaccines work by triggering both cellular and humoral immune responses against the spike protein of the virus.

Cellular immunity may play a more protective role than humoral immunity to variants of concern (VOC) against SARS-CoV-2, as it targets the conserved regions of spike protein and possibly cross-reacts with other variants.

Since a single spike epitope is recognized by multiple T-cell clones, the mRNA vaccination-induced T-cell response may consist of multiple spike-reactive clones. Thus, it is important to understand the mechanism of mRNA vaccination-induced cellular immune response. However, to address this clonal-resolution analysis on T-cell responses to mRNA vaccination has not been performed yet.

To bridge this gap, a team of researchers, led by Associate Professor Satoshi Ueha, including Professor Kouji Matsushima from the Tokyo University of Science (TUS), Japan, Mr. Hiroyasu Aoki from the University of Tokyo, and Professor Toshihiro Ito from Nara Medical University, aimed to develop a kinetic profile of spike-reactive T-cell clones during repetitive mRNA vaccination.

For this, they performed a longitudinal TCR sequencing on peripheral T cells of 38 participants who had received the Pfizer vaccine from before the vaccine to after the third vaccination and then analyzed the single-cell gene expression and epitope specificity of the clonotypes.

Their findings, <u>published in *Cell Reports*</u>, revealed that while the primary T-cell response of naïve T cells generally peaked 10–18 days after the first shot, expansion of "early responders" was detected on day 7 after



the first shot, suggesting that these early responders contain memory T cells against common cold coronaviruses.

They also found a "main responder" that expanded after the second shot and did not expand early after the first shot and a "third responder" that appeared and expanded only after the third shot.

By longitudinally tracking the total frequency of each response pattern, it was observed that, after the second shot, a shift among the clonotypes occurred, wherein the major population changed from early responders to main responders, suggestive of a shift in clonal dominance. A similar shift of responding clones was also observed in CD4⁺ T cells.

Prof. Ueha says, "We next analyzed the phenotype of main responders after the second and the third vaccination. The results showed that the main responders after the second and third shots mostly consist of effector-memory T cells (T_{EM}), with more terminally differentiated effector memory-like phenotype after the third shot."

The researchers then examined the repertoire changes of main responders, revealing that the expansion of main responders, which occurred after the second shot, diminished following the third shot, and the clonal diversity decreased and was partially replaced by the third responders. This may potentially mean that the third vaccination selected better-responding clones.

Due to the vaccination-induced shift in immunodominance of spike epitopes, the study supports the inter-epitope shift model. In addition, there were intra-epitope shifts of vaccine-responding clonotypes within spike epitopes.

Prof. Ueha explains, "Our analysis suggests that T cells can 're-write' themselves and reshape their memory populations after successive



vaccinations. This re-writability not only maintains the number of memory T cells but also maintains diversity that can respond to different variants of pathogens. Moreover, by tuning the replacement of memory cells, more effective vaccines can be developed that can also be tailored to an individual's unique immune response."

Overall, this study provides important insights into mRNA vaccineinduced T-cell responses, which will be crucial for developing nextgeneration vaccines for more effective and broad protection against viruses.

More information: Hiroyasu Aoki et al, CD8+ T cell memory induced by successive SARS-CoV-2 mRNA vaccinations is characterized by shifts in clonal dominance, *Cell Reports* (2024). <u>DOI:</u> <u>10.1016/j.celrep.2024.113887</u>

Provided by Tokyo University of Science

Citation: Optimizing boosters: How COVID mRNA vaccines reshape immune memory after each dose (2024, March 8) retrieved 11 May 2024 from <u>https://medicalxpress.com/news/2024-03-optimizing-boosters-covid-mrna-vaccines.html</u>

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