

A short memo for (10)2e about highlights of the proposed study

(10)(2e)  
16 July, 2020

### **Context**

Eyal, Lipsitch, and Smith (2020) *JID* proposed using human challenge studies for the acceleration of vaccine development, and Britton et al. (2020) *Science* insists on the influence of heterogeneity on the herd immunity. Beside, Gomes et al. (2020) *MedRxiv* also argues that individual variation in susceptibility lowers the herd immunity threshold.

### **The main message of this study**

With challenge studies, we can empirically infer the expected impact of preventive measures such as vaccination at the population-level. The evaluation of mitigation measures with a few healthy volunteers is much more safe, fast, and ethical than waiting until another epidemic runs its course.

### **Different points from our previous norovirus work**

1. This paper used only hypothetical human challenge data.
2. As we have multiple (hypothetical) data points, this study discusses dose-response curves for vaccinated a bit more generally.
3. Since the space is limited, I suggest not to include Tweedie models in this study. (I thought that a proposal of using Tweedie models is novel by itself, so this point should be emphasized in a different paper, intending different target readers.)

### **Important references**

Eyal, Lipsitch, and Smith (2020) *JID*

<https://academic.oup.com/jid/article/221/11/1752/5814216>

Britton et al. (2020) *Science*

<https://science.sciencemag.org/content/early/2020/06/22/science.abc6810>

Gomes et al. (2020) *MedRxiv*

<https://www.medrxiv.org/content/10.1101/2020.04.27.20081893v3>