

P9185 - Project 5: protocol design and analysis for COVID-19

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- Vaccine efficacy protocol
- Adverse effect analysis for Vaccine v.s. Control
- Survival analysis COVID contraction after vaccine shot

- Coronavirus, the severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), has had devastating consequences globally.
- Control measures, such as the use of masks, have been variably implemented and have proved insufficient in impeding the spread of coronavirus disease 2019 (Covid-19), the disease caused by SARS-CoV-2.
- Vaccines are urgently needed to reduce the morbidity and mortality associated with Covid-19.

Vaccine efficacy protocol

- A pharmaceutical company therefore would like to conduct a phase III randomized (**1-to-1 ratio**), **stratified**, observer-blinded, placebo-controlled trial at **100 U.S. sites** to demonstrate the efficacy for their developing vaccine.

Define primary outcome

$$VE = 1 - \frac{p_1}{p_2}$$

- p_i : the number of new cases during 12 months over the total number at risk during 12 months in group i
- x_i cases in group i with n_i samples
- Goal: test the null hypothesis that the vaccine efficacy is 30% or less and provide 80% power to detect a 60% vaccine efficacy without planned interim analyses

Vaccine efficacy protocol – randomization procedure

- The study consists of **2 periods** :
 - Vaccine period for 2 injections
 - Follow up period:
 - Second injection - 14 days: if the subjects have symptoms/being positive at this period, regard as not at risk and will not contribute to the efficacy calculation
 - 12 months: follow up period,

Vaccine efficacy protocol – randomization procedure

- Collect study subjects with seronegative at baseline N_0
- Take 2 covid shots
- Collect status at day 14 after second shot
- Remove those becoming positive during the 14 days
- Count new cases during 12 months

Vaccine efficacy protocol – randomization procedure

- Blinding and Randomization procedure
 - The Phase III clinical trial will be conducted in multicenter, randomized, stratified, observer-blinded, and placebo-controlled design
 - There are 100 participating centers in the US
 - Eligible participants are those aged 16 years or older who tested seronegative for SARS-CoV-2 at the recruitment time
 - We stratify the participants by age and gender, and randomly assigned persons in each stratification in a 1:1 ratio to receive either two doses of vaccines or placebo
 - They will be monitored for 12 months by active surveillance of COVID-19.
 - The trial is observer-blinded to avoid introducing bias, i.e, the participants and those responsible for the evaluation are blinded to the treatment group

Vaccine efficacy protocol – analysis approach

- $H_0 : VE \leq 30\%$, $H_1 : VE > 30\%$
- Test stat:

$$Z = \frac{\phi_0 \hat{p}_2 - \hat{p}_1}{\sqrt{\frac{\tilde{p}_1(1-\tilde{p}_1)}{n_1} + \phi_0^2 \frac{\tilde{p}_2(1-\tilde{p}_2)}{n_2}}},$$

where $\phi_0 = 1 - VE_0$, $\hat{p}_1 = \frac{x_1}{n_1}$, $\hat{p}_2 = \frac{x_2}{n_2}$, and \tilde{p}_1 and \tilde{p}_2 are the maximum likelihood estimates under the null hypothesis, calculated by

$$\tilde{p}_1 = \phi_0 \tilde{p}_2, \quad \tilde{p}_2 = \frac{-B - \sqrt{B^2 - 4AC}}{2A},$$

where

$$A = (n_1 + n_2)\phi_0, \quad B = -(n_1\phi_0 + x_1 + n_2 + x_2\phi_0), \quad C = x_1 + x_2,$$

and we reject the null hypothesis if $Z > Z_\alpha$, the upper α -th percentile of a standard normal distribution.

Vaccine efficacy protocol – sample size calculation

To detect a vaccine efficacy of $VE_1 = 0.6$ (or vaccine event probability of $p_1 = 0.4p_2$) with $1 - \beta = 80\%$ power, we assume that $n_1 = n_2$, and the number of subjects needed will be determined by the following formula:

$$N = \frac{(Z_\alpha[\phi_0 p_2(1 - \phi_0 p_2)/0.5 + \phi_0^2 p_2(1 - p_2)/0.5]^{1/2} + Z_\beta[p_1(1 - p_1)/0.5 + \phi_0^2 p_2(1 - p_2)/0.5]^{1/2})^2}{(\phi_0 p_2 - p_1)^2}.$$

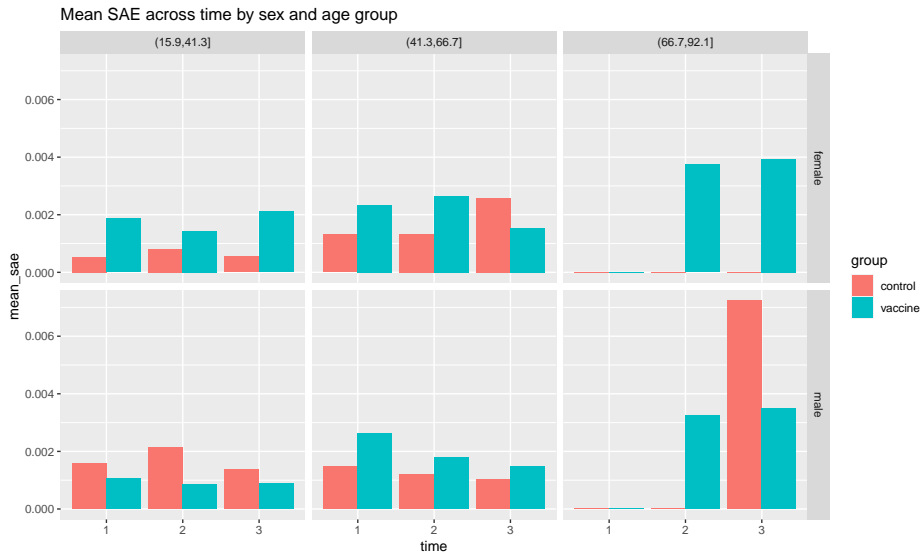
| | 0.01 | 0.02 | 0.03 | 0.04 | 0.05 | 0.06 | 0.07 | 0.08 | 0.09 |
|------|-------|-------|------|------|------|------|------|------|------|
| 0.00 | 18952 | 9400 | 6216 | 4624 | 3668 | 3032 | 2578 | 2236 | 1970 |
| 0.05 | 19950 | 9894 | 6544 | 4868 | 3862 | 3192 | 2712 | 2354 | 2074 |
| 0.10 | 21058 | 10444 | 6906 | 5138 | 4076 | 3368 | 2864 | 2484 | 2190 |
| 0.15 | 22296 | 11058 | 7312 | 5440 | 4316 | 3568 | 3032 | 2630 | 2318 |
| 0.20 | 23690 | 11750 | 7770 | 5780 | 4586 | 3790 | 3222 | 2794 | 2464 |

Table 1: Sample size calculation changing missing rate(row) and control prevalence(column)

Adverse effect analysis for Vaccine v.s. Control

| Characteristic | control, N = 20,625 ¹ | vaccine, N = 20,625 ¹ | p-value ² |
|---|----------------------------------|----------------------------------|----------------------|
| id | 10,313 (5,157, 15,469) | 60,313 (55,157, 65,469) | <0.001 |
| sae | 25 (0.1%) | 37 (0.2%) | 0.069 |
| Unknown | 611 | 2,051 | |
| site | 50 (25, 75) | 50 (25, 75) | >0.9 |
| sex | | | 0.2 |
| female | 10,313 (50%) | 10,190 (49%) | |
| male | 10,312 (50%) | 10,435 (51%) | |
| age | 45 (38, 51) | 45 (36, 53) | 0.3 |
| ¹ Median (IQR); n (%) | | | |
| ² Wilcoxon rank sum test; Pearson's Chi-squared test | | | |

Adverse effect analysis for Vaccine v.s. Control



Adverse effect analysis for Vaccine v.s. Control

Table 2: Missing Data Pattern

| | time1 | time2 | time3 | |
|-------|-------|-------|-------|-------|
| 30342 | 1 | 1 | 1 | 0 |
| 4552 | 1 | 1 | 0 | 1 |
| 3089 | 1 | 0 | 1 | 1 |
| 605 | 1 | 0 | 0 | 2 |
| 1933 | 0 | 1 | 1 | 1 |
| 385 | 0 | 1 | 0 | 2 |
| 288 | 0 | 0 | 1 | 2 |
| 56 | 0 | 0 | 0 | 3 |
| | 2662 | 4038 | 5598 | 12298 |

Adverse effect analysis for Vaccine v.s. Control – missing pattern

GLM: `missing_id ~ sae+sex+age+site+time`

| | Estimate | Std. Error | z value | Pr(> z) |
|-------------|----------|------------|---------|----------|
| (Intercept) | -1.1853 | 0.0372 | -31.83 | 0.0000 |
| sae | 0.0731 | 0.1917 | 0.38 | 0.7030 |
| sexmale | -0.0046 | 0.0155 | -0.30 | 0.7645 |
| age | -0.0023 | 0.0007 | -3.23 | 0.0013 |
| site | -0.0003 | 0.0003 | -1.02 | 0.3093 |
| time2 | -0.1826 | 0.0182 | -10.01 | 0.0000 |
| time3 | -0.4402 | 0.0194 | -22.71 | 0.0000 |

- Missing_id: =0 if no missing follow up, =1 if any missing follow up.
- Missing pattern is not related to the outcome;
- Assuming Missing at random and parameter separability;

Adverse effect analysis for Vaccine v.s. Control

$$\begin{aligned}\log\left(\frac{\pi_{ijk}}{1 - \pi_{ijk}}\right) = & \beta_0 \\ & + \beta_1 I(\text{time} == 2)_{ijk} + \beta_1 I(\text{time} == 3)_{ijk} \\ & + \beta_3 I(\text{time} == 1)_{ijk} \times I(\text{group} == \text{Vaccine})_{ij} \\ & + \beta_4 I(\text{time} == 2)_{ijk} \times I(\text{group} == \text{Vaccine})_{ij} \\ & + \beta_5 I(\text{time} == 3)_{ijk} \times I(\text{group} == \text{Vaccine})_{ij} \\ & + \beta_6 I(\text{sex} == \text{male})_{ij} \\ & + \beta_7 \text{age}_{ij} \\ & + \alpha_{0i} + \alpha_{1ij} + \epsilon_{ijk}\end{aligned}$$

- i for site, j for subject, k for time measure
- α_{0i} – site-level random intercept
- α_{1ij} – nested random intercept, $\alpha_{1ij} = \alpha_{1ik}$ if $I(\text{group} == \text{Vaccine})_{ij} = I(\text{group} == \text{Vaccine})_{ik}$

Adverse effect analysis for Vaccine v.s. Control

| | Estimate | Std. Error | z value | Pr(> z) |
|--------------------|----------|------------|---------|----------|
| (Intercept) | -7.38 | 0.39 | -19.08 | 0.00 |
| time2 | 0.06 | 0.28 | 0.21 | 0.83 |
| time3 | 0.19 | 0.27 | 0.70 | 0.48 |
| sexmale | -0.10 | 0.15 | -0.64 | 0.52 |
| age | 0.01 | 0.01 | 2.09 | 0.04 |
| time1:groupvaccine | 0.47 | 0.27 | 1.76 | 0.08 |
| time2:groupvaccine | 0.32 | 0.27 | 1.17 | 0.24 |
| time3:groupvaccine | 0.04 | 0.28 | 0.15 | 0.88 |

Table 3: Fixed effect estimation for adverse effect model

Adverse effect analysis – ANOVA for time effect

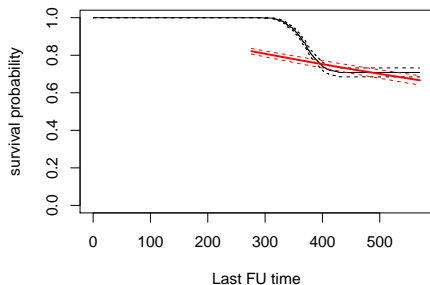
| | Chisq | Df | Pr(>Chisq) |
|------------|-------|----|------------|
| time | 0.12 | 2 | 0.9399 |
| sex | 0.41 | 1 | 0.5206 |
| age | 4.36 | 1 | 0.0368 |
| time:group | 4.28 | 3 | 0.2329 |

Table 4: Deviance test for adverse effect model

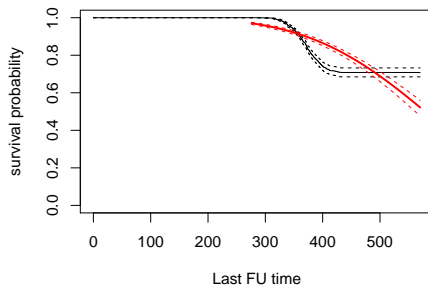
- From the Deviance test using Wald chisquare test statistics, we have $T_{interaction} = 4.28 > \chi^2_{0.05,3}$, P value > 0.05
- There is no significant difference between odds of having SAE in the vaccine group and control group at any of the three assessment time points.

Survival analysis COVID contraction after vaccine shot

KM and exponential estimates of survival curve



KM and Weibull estimates of survival curve



| | time | n.risk | n.event | surv | std.err | lower | upper |
|---|--------|---------|---------|------|---------|-------|-------|
| 1 | 360.00 | 1399.00 | 177.00 | 0.90 | 0.01 | 0.89 | 0.92 |

Table 5: Survival Rate at 12 Month

Survival analysis COVID contraction after vaccine shot

| | par_fitting | est | lcl | ucl |
|---|-------------|--------|--------|---------|
| 1 | Exponential | 974.17 | 892.32 | 1075.15 |
| 2 | Weibull | 974.17 | 892.71 | 1068.79 |
| 3 | K-M | | | |

Table 6: Estimated Median Survival Time

- The survival rate didn't drop to 50% at the end of the study.
- Flat tail of the survival curve, both exponential and Weibull distribution can't fit the trend well.
- Not interpretable parametric model fitting for both median and mean survival time.

Conclusion and discussion

- A pharmaceutical company therefore would like to conduct a phase III randomized (**1-to-1 ratio**), **stratified**, observer-blinded, placebo-controlled trial is designed with sample size calculation changing missing rate and control prevalence.
- No significant difference between odds of having SAE in the vaccine group and control group at any of the three assessment time points.
- The survival rate didn't drop to 50% at the end of the study. No interpretable estimation of mean and median survival time.

Effect of 2 Inactivated SARS-CoV-2 Vaccines on Symptomatic COVID-19 Infection in Adults: A Randomized Clinical Trial | Global Health | JAMA | JAMA Network. (n.d.). Retrieved May 2, 2022, from <https://jamanetwork.com/journals/jama/fullarticle/2780562>

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