

A Markov Model to Assess Covid-19 Vaccine Herd Immunization Patterns

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1.0 Introduction

This Markov model *assesses different patterns of vaccination*, which will affect achieving *Herd Immunity*. Such immunization approach assumes the virus will affect a large percentage of the general population, starving it for new *customers* and thus preventing community spread.

Herd Immunization can be acquired by two methods: (a) letting the *virus infect most of the population*. Weaker members (the elderly, those with co-morbidities, etc.) will die, while those surviving will become immunized. Alternatively, *Herd Immunization* can be acquired by (b) *vaccinating a large percent of the general population*. Such is achieved by inoculating a small, controlled virus infection. In either case, a large percent of the population must be involved.

The *urgency of the present situation* is that many people still refuse to take the vaccine. It is imperative to address the need for the population to agree. For this reason this paper will be very short and straight to the point. For *Markov Chain background material* and discussions, readers are sent to *our previous papers: A Markov Chain Model for Covid-19 Survival Analysis*: <https://www.researchgate.net/publication/343021113> [A Markov Chain Model for Covid-19 Survival Analysis](https://www.researchgate.net/publication/343021113) and *A Markov Model to Study Covid-19 Herd Immunization*, found in: <https://www.researchgate.net/publication/343345908> [A Markov Model to Study Covid-19 Herd Immunization?channel=doi&linkId=5f244905458515b729f78487&showFulltext=true](https://www.researchgate.net/publication/343345908)

We will *assume* that, either by *infection or by vaccination*, *individuals* will become *immunized* and will, therefore, *not become sick again nor pass down the infection* to anyone else. We will consider *two levels of vaccination: acceptable* (70% of the *available*¹ population) *and low* (40%). All *other model parameters* including vaccine effectiveness (70%) remain the *same*. We *develop Markov models* for both situations, *obtain and compare* their respective *performance measures*, and discuss the results. We conclude with some general remarks about vaccination policies.

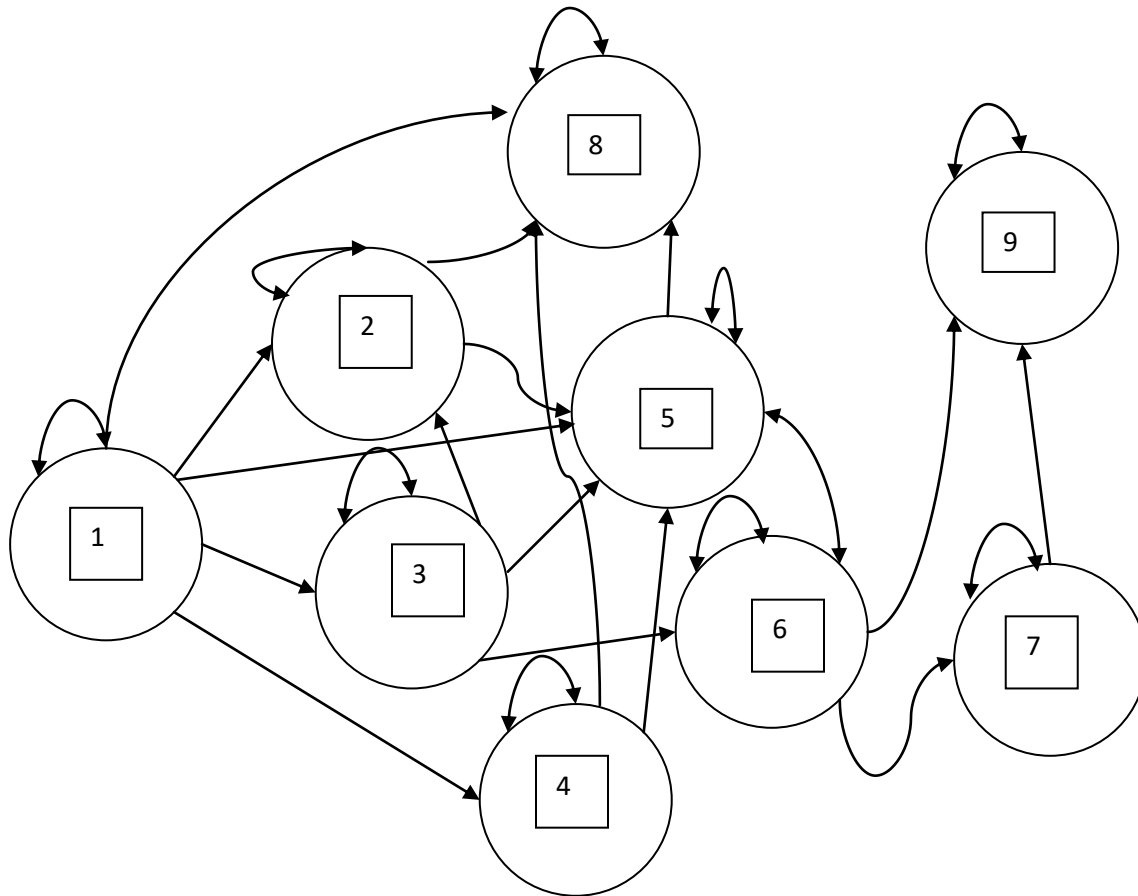
2.0 Previous Covid-19 work

Following our proposal to the retired Academic and Research Communities, to fight Covid-19: <https://www.researchgate.net/publication/341282217> [A Proposal for Fighting Covid-19 and its Economic Fallout](https://www.researchgate.net/publication/341282217) we have written a number of articles on statistical models:

¹ Total population, minus those individuals already infected and thence naturally immunized, etc.

<https://www.researchgate.net/publication/346956247> Logistic Regression in Factor Identification of Covid-19 Vaccine Clinical Trials on identifying important key factors observed in Clinical Trials, <https://www.researchgate.net/publication/346305686> A Digression on Covid-19 Vaccine Clinical Trials and its Consequences and ICUs and hospital staffing using the Negative Binomial distribution: <https://www.researchgate.net/publication/345914205> Covid-19 ICU Staff and Equipment Requirements using the Negative Binomial screening DOEs: <https://www.researchgate.net/publication/344924536> Design of Experiments DOE in Covid-19 Factor Screening and Assessment using statistical methods to establish a new *Vaccine Life*: <https://www.researchgate.net/publication/344495955> Survival Analysis Methods Applied to Establishing Covid-19 Vaccine Life as well as to help *accelerate vaccine testing*: <https://www.researchgate.net/publication/344193195> Some Statistical Methods to Accelerate Covid-19 Vaccine Testing and a *Markov model to study problems of reopening college*: <https://www.researchgate.net/publication/343825461> A Markov Model to Study College Reopening Under Covid-19 and *Markov Model to study the effects of Herd Immunization*: <https://www.researchgate.net/publication/343345908> A Markov Model to Study Covid-19 Herd Immunization?channel=doi&linkId=5f244905458515b729f78487&showFulltext=true as well as of *general survival*: <https://www.researchgate.net/publication/343021113> A Markov Chain Model for Covid-19 Survival Analysis about socio-economic and racial issues affected by Covid-19: <https://www.researchgate.net/publication/343700072> A Digression About Race Ethnicity Class and Covid-19 and developing *A Markov Chain Model for Covid-19 Survival Analysis*: <https://www.researchgate.net/publication/343021113> A Markov Chain Model for Covid-19 Survival Analysis and *An Example of Survival Analysis Applied to analyzing Covid-19 Data*: <https://www.researchgate.net/publication/342583500> An Example of Survival Analysis Data Applied to Covid-19, and *Multivariate Statistics in the Analysis of Covid-19 Data*, and *More on Applying Multivariate Statistics to Covid-19 Data*, both of which can also be found in: <https://www.researchgate.net/publication/341385856> Multivariate Stats PC Discrimination in the Analysis of Covid-19, and the implementation of *multivariate analyses* methods such as: <https://www.researchgate.net/publication/342154667> More on Applying Principal Components Discrimination Analysis to Covid-19 *Design of Experiments* to the Assessment of Covid-19: <https://www.researchgate.net/publication/341532612> Example of a DOE Application to Coronavirus Data Analysis *Offshoring*: <https://www.researchgate.net/publication/341685776> Off-Shoring Taxpayers and the Coronavirus Pandemic and *reliability methods* in ICU assessment: <https://www.researchgate.net/publication/342449617> Example of the Design and Operation of an ICU using Reliability Principles and *Quality Control methods for monitoring Covid-19*: <https://web.cortland.edu/matresearch/ApplicatSPCtoCovid19MFE2020.pdf> *Numerical Example* <https://www.researchgate.net/publication/339936386> A simple numerical example that illustrates the dangers of the Coronavirus epidemic

Figure 1: The State Space Diagram for the Markov Chain



RATIONALIZATION

The Markov Chain *unit time* is a week. Transitions refer to the State changes that occur from *one Monday to the following Monday*. State (1): the *General population*. Asymptomatically infected individuals, unaware of their condition, pass the virus on to others and then become immunized. State (2): *Mild Covid-19*; detected persons are quarantined at home. State (3): *Non Vaccinated* persons (refused vaccination). Some may spread the virus in the community. Some may develop symptoms, are tested, and are quarantined at home. If they become very ill, they are sent to the hospital. State (4): *Vaccinated*; some individuals may suffer mild Covid-19, stay at home, and eventually 70% become immunized. Some others may get sick and require medical attention. State (5): *infected and requiring medical attention at home*. State (6): persons with *serious illness* which require *hospitalization and treatment*. Some recover and are sent home; others get worse, and are placed in ICUs. State (7): *patients in ICU*. Some improve and are sent back to the ward, and others are placed on Ventilators or die. State (8): *herd immunization*; and State (9): *death*. These two latter states are *Absorbing*: *once you enter them, you cannot leave*.

We obtain, from the TPM, Sub-Matrix Q, corresponding to the seven Transient states:

Matrix Q of the Seven Transient States:

<u>Pop</u>	<u>MildInf</u>	<u>NonVac</u>	<u>Vaccine</u>	<u>SevereCov</u>	<u>Hospital</u>	<u>ICU</u>
0.05	0.05	0.23	0.54	0.03	0.00	0.00
0.00	0.70	0.00	0.00	0.05	0.00	0.00
0.00	0.10	0.70	0.00	0.10	0.10	0.00
0.00	0.20	0.00	0.07	0.03	0.00	0.00
0.00	0.00	0.00	0.00	0.30	0.20	0.00
0.00	0.00	0.00	0.00	0.20	0.57	0.20
0.00	0.00	0.00	0.00	0.00	0.20	0.65

We then subtract matrix Q from the Identity Matrix, yielding (I-Q),

0.95	-0.05	-0.23	-0.54	-0.03	0.00	0.00
0.00	0.30	0.00	0.00	-0.05	0.00	0.00
0.00	-0.10	0.30	0.00	-0.10	-0.10	0.00
0.00	-0.20	0.00	0.93	-0.03	0.00	0.00
0.00	0.00	0.00	0.00	0.70	-0.20	0.00
0.00	0.00	0.00	0.00	-0.20	0.43	-0.20
0.00	0.00	0.00	0.00	0.00	-0.20	0.35

We invert the above: **Matrix inverse S = (I-Q)⁻¹**

<u>Populat</u>	<u>MildInf</u>	<u>NonVac</u>	<u>Vaccine</u>	<u>SevCov</u>	<u>Hospital</u>	<u>ICU.</u>
1.05263	0.85191	0.80702	0.61121	0.39130	0.50350	0.28772
0.00000	3.33333	0.00000	0.00000	0.29071	0.18416	0.10524
0.00000	1.11111	3.33333	0.00000	1.04665	1.71885	0.98220
0.00000	0.71685	0.00000	1.07527	0.11879	0.07525	0.04300
0.00000	0.00000	0.00000	0.00000	1.74428	1.10497	0.63141
0.00000	0.00000	0.00000	0.00000	1.10497	3.86740	2.20994
0.00000	0.00000	0.00000	0.00000	0.63141	2.20994	4.11997

The Potential Matrix, of which S (above) is a sub-matrix, contains the Long-Run Sojourns, or the average number of visits to each of the States (matrix columns), when starting from any other of the States (represented by the matrix rows). Sojourns that occur outside matrix S are either Zero (if the states cannot be reached) or Infinite (if the transition is into an Absorbing state).

We then obtain Matrix $G = SB$ where S is as above, and B is the 7x2 matrix formed with the last two columns of the initial TPM, corresponding to the two Absorbing states. Such matrix G then provides the probabilities of ever reaching said two absorbing states (immunization or death), when starting from any of the seven transient states (i=1, ... 7) below, denoted by:

Populat(1); MildInf(2) ;NonVaccin(3) ;Vaccin(4) ;SevereCovid(5) ;Hospitalized(6) ;ICU(7)

St Immunizat Deaths.

1.	0.941738	0.058262
2.	0.978690	0.021310
3.	0.801105	0.198895
4.	0.991293	0.008707
5.	0.872139	0.127861
6.	0.552486	0.447514
7.	0.315706	0.684294

We will interpret this table of results through a numerical example. Assume there is a population of one million adults, aged 18 and above (which is the age for taking the Covid vaccine). From Line 1 (i.e. Population) we have that 94.2% of these individuals (or 942,000) will be immunized, while 5.8% of them (or 58,200) will die. If we look at Line 2 (i.e. Mildly Infected and naturally immunized) we see that, from those individuals, 97.8% will survive and 2.1% will die. If we look at Line 3 (i.e. Non Vaccinated² sub-population, because they rejected the vaccine), 80.1% will be immunized and 19.8% will die. Regarding Line 4 (i.e. Vaccinated³ sub-population), 99% will survive immunized, and 0.87% (less than 1%) will die of Covid-19. For Line 7 (i.e. those in an ICU), 31.5% will survive and become immunized, while the remaining 68.4% will die.

Beware that, except Line 1 (which is the only one that concerns the total population), each of the percents of other Lines apply only to the sub-totals of such sub-population. For example, for the Non-Vaccinated sub-population of $0.23 \times 1,000,000 = 231,000$, 19.8% will die ($231000 \times 0.198 = 45,738$ persons) and $231,000 - 45,738 = 185,262$ will survive and become naturally immunized.

Follow exactly the same calculations above, for the pessimistic framework of 40% vaccinated:

The Transition Probability Matrix P for the Second (40% vaccination) Markov Chain is:

Pop;MildInf;NonVac;Vaccine;SevereCov;Hospital;ICU;Immune;Dead.

0.05	0.05	0.46	0.31	0.03	0.00	0.00	0.10	0.00
0.00	0.70	0.00	0.00	0.05	0.00	0.00	0.25	0.00
0.00	0.10	0.70	0.00	0.10	0.10	0.00	0.00	0.00
0.00	0.20	0.00	0.07	0.03	0.00	0.00	0.70	0.00
0.00	0.00	0.00	0.00	0.30	0.20	0.00	0.50	0.00
0.00	0.00	0.00	0.00	0.20	0.57	0.20	0.00	0.03
0.00	0.00	0.00	0.00	0.00	0.20	0.65	0.00	0.15
0.00	0.00	0.00	0.00	0.00	0.00	0.00	1.00	0.00
0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	1.00

After implementing the same algorithmic process developed above, we obtain:

² Non-Vaccinated transition rate is: $0.3 \times (1 - 0.1 - 0.05 - 0.05 - 0.03) = 0.3 \times 0.77 = 0.231$

³ Vaccinated transition rate is: $0.7 \times (1 - 0.1 - 0.05 - 0.05 - 0.03) = 0.3 \times 0.77 = 0.539$

Again the Matrix $G = SB$, where $S = (I - Q)^{-1}$, and B is the 7×2 matrix formed by the last two columns of the initial TPM corresponding to the two Absorbing states. Matrix G provides the probabilities of ever reaching said two absorbing states (immunization or death), when we start from any of the seven transient states ($i=1, \dots, 7$) below, denoted by:

Populat(1); MildInf(2) ;NonVaccin(3) ;Vaccin(4) ;SevereCovid(5) ;Hospitalized(6) ;ICU(7)

<u>St</u>	<u>Inmunizat</u>	<u>Deaths.</u>
1.	0.900908	0.099092
2.	0.978690	0.021310
3.	0.801105	0.198895
4.	0.991293	0.008707
5.	0.872139	0.127861
6.	0.552486	0.447514
7.	0.315706	0.684294

Again we interpret this table of results through a *numerical example*. Assume a population of one million adults, aged 18 and above (which is the age for taking Covid19-19 vaccine). From Line 1 (i.e. *Population*) we have that 90.2% of these individuals (or 902,000) will be *immunized*, while 9.9% of them (or 99,000) *will die*. The results from the rest of the Lines are the same as in the previous example. How, then, do we explain a larger number of deaths in the general population?

The answer is straight forward. *Now, 46% of the total population of one million (i.e. 460,000 individuals) are Non-Vaccinated. Thence, 19.9% of them (i.e. 91,080) will die*, instead of only 45,738 persons, in the previous example, and *mostly from those who refused to take the vaccine*.

4.0 Discussion

We have *implemented a Markov Chain twice, changing only the parameters* that represented the *vaccination pattern*. The model *results are unquestionably different*. We have used this technique before, applying a *Linear Programming (LP) model* to the US manufacturing off-shoring process during the last quarter of a century. We considered, in our LP model, activities such as providing health care, retraining, unemployment benefits, etc. to off-shored-displaced workers, which were not included (or paid for) in off-shoring companies' LPs. They were paid by the US government (tax payers). *Both LP model results* https://www.researchgate.net/publication/341685776_Off-Shoring_Taxpayers_and_the_Coronavirus_Pandemic) were *also unquestionably different*.

In either case, the *reader may challenge* our choice of *parameter values*. What is indisputable is how, in both cases, there is *an obvious difference in model results*, independently of the value of model parameters used. Such *difference stems from the different patterns of behavior modeled*.

Then, vaccination activity carries two aspects: one individual and the other, social. The vaccine *protects the individual*. In addition, if enough individuals in the general population take it, the

vaccine *has an effect over the Pandemic*. With few new customers to infect, *the virus starves* and ends up disappearing! If not enough people become vaccinated this latter, *greater benefit* is lost.

Markov models, due to their State Spaces (E_j) and transition probabilities (X_{ij} for $i,j=0, \dots, k$), are particularly *useful to model Immunization patterns*. For example, State (1) is *General Population* and State (8), *Immunized Population*. Then, $X_{18} = 0.1$ corresponds to the 10% of asymptomatic persons that become immunized; $X_{12} = 0.05$ corresponds to the 5% persons with mild infections that are detected, quarantined, do not require medical attention, and finally become immunized. Also, $X_{15} = 0.03$ represents the 3% of the infections that require treatment at home. If infections are positively resolved, patients become immunized; if not, patients are hospitalized.

There are 70% (or 40%) *persons vaccinated* (resulting in $X_{14} = 0.54$ or 0.31 once persons already naturally immunized, sick, younger than 17 years, etc. are subtracted). And there are 23% (or 46%) persons *Non-Vaccinated* (resulting in $X_{13} = 0.46$ or 0.23 , just as above). These *rates can be changed*, and model results can then be assessed and compared, *to find efficient strategies*.

One important parameter is $X_{48} = 0.7$ (*transition from Vaccinated to Immunized*). We assumed that the *vaccine was 70% efficient* (accounting for all different efficiency rates for diverse ages, co-morbidities, etc.). *Other values can be used* and model results can be assessed and compared. The same can be said about transition rates related to hospitalization, ICU etc. that depend on the load such hospitals and health care practitioners have, which in turn depend on infection rates.

The *time unit* utilized for transitions was *a week* (other units could have been used: hours, days, months, etc.). For example, the *weekly rate of asymptomatic infection* ($X_{18} = 0.1$), for members of the general population who contract Covid-19 and then become immunized, means it is 10%.

Probabilities of persons dying or immunized, combined with population estimates, helps assess the number of cases, deaths, immunized persons, etc. that result from the different scenarios.

Results from the present Markov model, complement results obtained in our previous papers, as different models address the different aspects of Covid-19 problems. For example, the article https://www.researchgate.net/publication/342583500_An_Example_of_Survival_Analysis_Data_Applied_to_Covid-19 can be used in *Triage situations*, to evaluate the probability of a patient dying, given its co-morbidities, and allocate scarce ICU beds or doctors to those patients with a higher chance of making the best possible use of these resources.

One advantage of *achieving a Vaccine Herd Immunization* is the possibility of *avoiding* the bad *triage situation* mentioned above. If the healthcare systems overheat, due to the large number of Covid-19 cases, Triage becomes a reality.

5.0 Conclusions

Again, *the urgency of this paper* stems from recent polls results that suggest that *a large number of people are not yet willing to take the vaccine*. And this is *a very serious problem*. Some argue that *there are still many unanswered questions* regarding the vaccine: How long does immunity last? Does it prevent passing on the virus? What are its secondary effects? *However, there is a key question*: which is the greater risk, taking the vaccine and becoming protected, with all the mentioned caveats, or contracting Covid-19, passing it on to those we love, and perhaps dying.

Vaccination activity carries two aspects: one individual and the other social. First, the vaccine protects the individual. In addition, if enough individuals in the general population become vaccinated, it has an effect over the Pandemic. With few new customers to infect, the Covid-19 virus starves, and ends by disappearing, which is the only real solution to the Covid-19 problem. *If not enough people become vaccinated, this latter, larger benefit is lost*.

Our Markov model, as well as its transition rates and its states, were all based on our readings about Covid-19 and our 45+ years of statistical experience in modeling and data analysis. This author is not a public health specialist, but a statistician. *Our intent is to place these models in the hands of health specialists*, so they can implement them with better information. Models may provide more neutral rules to establish medical procedures, if such is ever necessary.

Finally, *our statistical models* do not intend to compete with, but to complement and enrich, the models developed by Epidemiologists and other Public Health professionals. One example⁴ says:

The total population is partitioned into eight stages of disease: S, susceptible (uninfected); I, infected (asymptomatic or pauci-symptomatic infected, undetected); D, diagnosed (asymptomatic infected, detected); A, ailing (symptomatic infected, undetected); R, recognized (symptomatic infected, detected); T, threatened (infected with life-threatening symptoms, detected); H, healed (recovered); E, extinct (dead). The interactions among these stages are shown in Fig. 1. We omit the probability rate of becoming susceptible again after having recovered from the infection. Although anecdotal cases are found in the literature, re-infection rate value appears negligible.

The reader will recognize how their eight stages of the disease are somewhat similar to our Nine State Space Elements. Also, how said article's Figure 1 is also somewhat similar to our own State Space Diagram, presented in Section 3.0. Modeling tools used are different.

Both models are fine and pursue a similar objective: to help study and overcome Covid-19. Researchers are allies, not competitors.

As statistician George Box once said: *all models are wrong; some models are useful*.

⁴ See: <https://www.nature.com/articles/s41591-020-0883-7#Sec2>

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