1	Effectiveness of drugs for COVID-19 inpatients in Japanese medical claim data as
2	average treatment effects with inverse probability weighted regression adjustment:
3	Retrospective observational study
4	
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NOTE: This preprint reports new research that has not been certified by peer review and should not be used to guide clinical practice.

19 Abstract

20	Background: Prior stud	es have indicated	l that drugs against	t coronavirus disease 2019
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21 (COVID-19) such as antiviral drugs, anti-inflammatory drugs, steroid and antibody

22 cocktails are expected to prevent severe COVID-19outcomes and death.

- 23 Object: We analyzed medical claim data in Japan to assess the effectiveness of drugs
- againstCOVID-19.

Method: We applied an average treatment effect model with inverse probability 25 26 weighted regression adjustment, to the Medical Information Analysis Databank 27 managed by National Hospital Organization in Japan. The outcome was death during 28 hospitalization. Subjects were all inpatients, inpatients with oxygen therapy, and inpatients with respiratory ventilators, by three age classes: all ages, 65 years old or 29 older, and younger than 65 years old. Data on physical characteristics, underlying 30 diseases, administered drugs, the proportion of mutated strains, and vaccine coverage 31 32 were used as explanatory variables for logistic regression.

33 Result: Estimated results indicated that only an antibody cocktails (sotrovimab,

34 casirivimab and imdevimab) raised the probability of saving life, even though these

- 35 drugs were administered in few cases. On the other hand, other drugs might raise the
- 36 probability of death.

as effective to save life
is effective to save li

- 38 using an average treatment effect model with inverse probability weighted regression
- 39 adjustment. No other drugs such as remdesivir, dexamethasone, baricitinib and
- 40 tocilizumab were found to be effective to save life, even in the pseudo-situation of
- 41 random assignment.

43 Introduction

44	Coronavirus disease 2019 (COVID-19) is an infectious disease that results
45	from severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2). The first case of
46	COVID-19 was detected in December 2019 in Wuhan, China. It subsequently spread
47	worldwide. The drug kinds, which included antiviral drugs, antibody cocktails, and
48	steroid and anti-inflammatory drugs, were developed, leading to decreased incidence of
49	severe COVID-19 and death in Japan [1].
50	Earlier studies and clinical trials have investigated the effectiveness of drugs
51	used against COVID-19 such as an antiviral drug (remdesivir) [2], antibody cocktails
52	(casirivimab/imdevimab, sotrovimab and tixagevimab/cilgavimab) [3-7], a steroid
53	(dexamethasone) [8], RNA-dependent RNA polymerase inhibitors (molnupiravir and
54	favipiravir) [9-12], protease inhibitors (nirmatrelvir/ritonavir) [9, 13], a JAK inhibitor
55	(ruxolitinib) [14], and anti-inflammatory drugs (baricitinib and tocilizumab) [15-17]. All
56	are expected to prevent severe COVID-19 outcomes and death. For this study, we
57	considered the real-world effectiveness of drugs against COVID-19.
58	Medical Information Analysis Databank (MIA) is operated by the National
59	Hospital Organization (NHO) in Japan, which provides a database of medical claims
60	from 60 representative NHO hospitals [18]. NHO is one of the biggest hospital

organization in Japan, it has about 3.4% of all beds in Japan [19]. We used MIA data to
analyze drug effectiveness.

63	In reality, the decision to administer drugs depends on the patient's condition:
64	that is not a random assignment. Generally speaking, patients who are more likely to
65	develop severe illness have higher probability of being administered drugs, which
66	implies a negative association between drugs and death because of selection bias in the
67	observational data. Therefore, non-random choice of drug administration is regarded as
68	an estimate of drug effectiveness.
69	The average treatment effect model with inverse probability weighted
70	regression adjustment might resolve this difficulty statistically but not experimentally
71	[20]. This procedure predicts the likelihood of administration for the drug. Then
72	outcomes were estimated, using logistic regression including dummy variables, whether
73	a drug was administered or not, weighted with the inverse probability of each subject
74	actually belonging to the group estimated by first-step logistic regression for drug
75	administration. It focused more on subjects whose probability of belonging to the group
76	were lower, in other words, less likelihood was regarded as a more nearly random
77	assignment.

78

Researchers in natural science including medicine can conduct the experiments

79	and thus delete selection bias in the choice of subjects, even though experiments are
80	expensive and require a longer period, usually. Conversely, researchers in social
81	sciences cannot conduct experiments to individual's choice. Instead of that, they
82	sometimes use the average treatment effect model with propensity score matching,
83	mainly, for the evaluation of programs to control participants choose to join the
84	program.
85	However, even in medicine, random assignment experiments after launch are
86	difficult. Therefore, situations change after a trial: mutated strains emerge, and vaccine
87	coverage and/or developed treatments might affect drug effectiveness, rendering
88	evaluation impossible. Particularly, mortality is often not used as an outcome for
89	evaluation, even though it is the most important endpoint. Therefore, experiments in
90	medicine for changing situations and mortality might be difficult. To resolve this
91	difficulty, the average treatment effect model was used for evaluation in a changing
92	situation through a statistically pseudo-random assignment experiment. Particularly, it
93	was applied for orthopedic surgery and cardiovascular research [21, 22]. However, these
94	studies used propensity scoring matching method: a simple comparison of outcomes
95	among the treatment group and non-treatment group with almost identical probability to
96	receive treatment. In other words, this method did not control for outcomes other than

97	treatment. It might bias the result by the potential confounder. Therefore, we used
98	another average treatment model with inverse probability weighted regression
99	adjustment. Its second step was weighted regression for the outcome weighted with
100	inverse probability of the first step for whether the patient was treated, or not [23,24]. It
101	can control covariates for outcomes other than whether the patient was treated, or not.
102	Conversely, average treatment effect model with propensity score matching cannot
103	control other condition than whether the patient was treated, or not.
104	
105	Materials and Methods
106	Data sources
106 107	Data sources This study used MIA for confirmed inpatients including their age, sex,
107	This study used MIA for confirmed inpatients including their age, sex,
107 108	This study used MIA for confirmed inpatients including their age, sex, underlying diseases, hospitalization week, administered drugs against COVID-19,
107 108 109	This study used MIA for confirmed inpatients including their age, sex, underlying diseases, hospitalization week, administered drugs against COVID-19, outcome and oxygen therapy and/or use of respiratory ventilator. We had accessed to
107 108 109 110	This study used MIA for confirmed inpatients including their age, sex, underlying diseases, hospitalization week, administered drugs against COVID-19, outcome and oxygen therapy and/or use of respiratory ventilator. We had accessed to these data which could identify individual participants.
107 108 109 110 111	This study used MIA for confirmed inpatients including their age, sex, underlying diseases, hospitalization week, administered drugs against COVID-19, outcome and oxygen therapy and/or use of respiratory ventilator. We had accessed to these data which could identify individual participants. We used data for vaccine administration published by the Cabinet Secretariat

The study period was January 2020 through March 2022, using data recorded
as of May 2022, which was the date we conducted this study. The study area was the
entirety of Japan.

118

119 Subjects

	3
120	Subjects were all inpatients who were diagnosed as COVID-19 (U07 in ICD10) from
121	January 2020 through March, 2022. We collected the data on all COVID-19 inpatients
122	in MIA data. Some inpatients who were still hospitalized at the end of study period
123	were excluded. The number of inpatients who were diagnosed as COVID-19 in MIA
124	data moves like 6 waves. These waves were classified from the trough of the prior wave
125	to the trough of the current wave using national data [27]. We defined the 1st wave
126	extended from week 1 of 2020 through week 23 of 2020; the 2nd wave lasted from
127	week 24 of 2020 through week 39 of 2020; the 3rd wave was recorded from week 40 of
128	2020 through week 8 of 2021; the 4th wave occurred from week 9 of 2021 through
129	week 24 of 2021; the 5th wave was from week 25 of 2021 through week 47 of 2021;
130	and the 6 th wave was from week 48 of 2021 through the end of this study.
131	However, apart from purely medical criteria, the criteria for hospitalization of
132	asymptomatic patients or mild patients who do not require oxygen therapy may be

133	strongly influenced by social conditions, such as lack of medical resources or support
134	for staying and recovering at home. Therefore, we considered not only the inclusion of
135	all subjects but also the limitation of subjects on oxygen therapy or respiratory
136	ventilation.
137	Note that we did not distinguish some patients with initiation drug before
138	oxygen therapy or respiratory ventilation and patients with initiation drug after oxygen
139	therapy or respiratory ventilation. Because timing of drug initiation, and oxygen therapy
140	or respiratory ventilation using were not available in this study. In other words, patient
141	condition when initiation drug was not accounted for drug effectiveness.
142	

143 **Definitions of Variables**

144 Physical characteristics: Age and sex.

145 Underlying diseases: We examined cancer (C00-C90 in ICD10), asthma (J45), chronic

146 obstructive pulmonary disease (COPD) (J44), hypertension (I10), heart failure (I50),

147 and DM (E10).

148 Pharmaceutical therapy: We examined the effects of an antiviral drug (remdesivir), an

- 149 antibody cocktails (sotrovimab and casirivimab/imdevimab), a steroid (dexamethasone),
- and anti-inflammatory drugs (baricitinib and tocilizumab). The antibody cocktails were

151	not divided by drug such as sotrovimab and casirivimab/imdevimab because the number
152	of antibody cocktails was not sufficient. The overall effects of the antibody cocktails
153	were examined. We also collected the number of prescriptions such as RNA-dependent
154	RNA polymerase inhibitors (molnupiravir and favipiravir), protease inhibitors
155	(nirmatrelvir/ritonavir). Remdesivir, antibody cocktails (sotrovimab and
156	casirivimab/imdevimab), dexamethasone, baricitinib, and tocilizumab were the top five
157	drugs which earlier studies had shown as effective.
158	Vaccine coverage: The complete rate of second dose of vaccine received two weeks prior
159	by age class, as younger than 65 years old (young patients) or 65 years old or older
160	(elderly patients).
160 161	(elderly patients). Mutated strains: Mutated strains were measured by percentage at one week before
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161 162 163	Mutated strains: Mutated strains were measured by percentage at one week before admission. Omicron included BA.2 or a later sublineage. Alternatively, we used a dummy variable during the 4th–6th wave instead of the proportion of the mutated
161 162 163 164	Mutated strains: Mutated strains were measured by percentage at one week before admission. Omicron included BA.2 or a later sublineage. Alternatively, we used a dummy variable during the 4th–6th wave instead of the proportion of the mutated strains as an explanatory variable, to check robustness. By this specification, the Alpha
161 162 163 164 165	Mutated strains: Mutated strains were measured by percentage at one week before admission. Omicron included BA.2 or a later sublineage. Alternatively, we used a dummy variable during the 4th–6th wave instead of the proportion of the mutated strains as an explanatory variable, to check robustness. By this specification, the Alpha variant strain emerged and then dominated in the 4th wave. Similarly, the Delta variant

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170	Statistical	ana	lysis
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171	We conducted estimates separately by drug type: remdesivir, antibody cocktails
172	(sotrovimab, casirivimab and imdevimab), dexamethasone, baricitinib, and tocilizumab
173	The estimation procedure was the average treatment effect model with inverse
174	probability weighted regression adjustment. It consists of two steps. The first step to
175	assess whether the patient was administered a type of drug or not was performed
176	through logistic regression on their physical condition, underlying diseases, drugs,
177	vaccine coverage, and prevalence in mutated strains as explanatory variables. The
178	second step was logistic regression for fatality weighted with inverse probability
179	estimated by the first step. Explanatory variables in this step were binomial variables:
180	whether the patients were administered the considered drug or not.
181	We adopted 5% level as significance level. All statistical analyses were
182	conducted using software (STATA SE 17.0; Stata Corp.).
183	

Ethical considerations 184

This study was approved by the Ethics Committee of Mie Hospital (Approval No. 185

2020-89). Permission to use MIA data was obtained using the NHO (Registration No. 186

187 1201003).

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189 Resul	ts
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190	The number of	of COVID-19	inpatients in	this study was	21727. Fig 1	l shows the

191 number of COVID-19 inpatients, COVID-19 inpatients with oxygen therapy,

- 192 COVID-19 inpatients with respiratory ventilation and fatalities due to COVID-19 by the
- 193 hospitalized week in MIA data. Fig 2 presents the drugs against COVID-19 by the week
- 194 in which the COVID-19 inpatient administered the drug was admitted to the hospital in
- 195 MIA data. Fig 1 shows similar trends in COVID-19 inpatients, COVID-19 inpatients
- 196 with oxygen therapy, COVID-19 inpatients with respiratory ventilation and fatalities
- due to COVID-19. Fig 2 also shows that all drugs for treating COVID-19 have a similar
- 198 trend. The most prescribed drugs were dexamethasone, followed by remdesivir,
- 199 baricitinib, tocilizumab and antibody cocktails (sotrovimab, casirivimab/imdevimab).
- 200
- Fig 1. COVID-19 inpatients, COVID-19 inpatients with oxygen therapy,
- 202 COVID-19 inpatients using respiratory ventilation and fatalities attributed to

203 COVID-19 by the hospitalized week in MIA data

204 Notes: Blue line and orange line show the numbers of COVID-19 inpatients and

205	inpatients	with oxy	gen therapy	respectively	in MIA d	lata (right	scale). G	ray line an	d
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- 206 yellow line show the numbers of COVID-19 inpatients with respiratory ventilator and
- 207 fatalities due to COVID-19 respectively in MIA data (left scale).

208

209 Fig 2. Drugs for treating COVID-19 in MIA data

- 210 Notes: Blue line and orange line show the numbers of remdesivir and dexamethasone
- 211 respectively in MIA data (left scale). Gray line, yellow line and light blue line show the

212 numbers of tocilizumab, baricitinib and antibody cocktails (sotrovimab,

213 casirivimab/imdevimab) respectively in MIA data (right scale).

- 214
- 215

216	Table 1 shows the characteristics 21727 inpatients. The mean of their age was
217	54.3. The proportion of sex of male and female was almost same. The most common
218	underlying disease was diabetes mellitus followed by hypertension, cancer, heart failure,
219	asthma, and COPD. Dexamethasone was the most prescribed drug, followed by
220	remdesivir, baricitinib, tocilizumab, and antibody cocktails. Fig 3 presents proportions
221	of mutated strains (Alpha, Delta and Omicron variants) in Tokyo and vaccination
222	coverage with two and three doses in Japan. The predominant variant changed from

Alpha to Delta in July 2021 and from Delta to Omicron in December 2021. Vaccination

coverage with two doses has increased since May 2021 and reached 80% by the end of

225 2021.

226

	All inpatients		
Characteristics	n	%	
Age			
Mean (SD)	54.3	(25.6)	
Sex			
Female	10083	46.4	
Underlying diseases			
Cancer	1036	4.8	
Hypertension	2820	13	
Diabetes mellitus	3241	14.9	
Heart failure	825	3.8	
Asthma	575	2.6	
COPD	217	1.0	
Drugs against COVID-19			
Remdesivir	1910	8.8	
Dexamethasone	6041	27.8	
Tocilizumab	562	2.6	
Baricitinib	1087	5.0	
Antibody cocktails	525	2.4	

227 Table 1. Characteristics of subjects

228 Note: Range of age was 0-108.

229 Abbreviation: SD: standard deviation, COPD: chronic obstructive pulmonary disease

231 Fig3. Proportions of Alpha, Delta and Omicron variants in Tokyo, in addition to

232 vaccination coverage with two and three doses in Japan

- 233 Notes: Blue, orange, and gray lines respectively represent the proportions of the Alpha,
- Delta and Omicron variants. Data of the proportions of the Alpha and Delta variants 234
- were published by the Cabinet Secretariat. Yellow and light blue lines respectively 235
- 236 represent vaccination coverage with two and three doses in the entire Japanese
- 237 population. Data of the vaccination coverage were published by the monitoring meeting
- in Tokyo. 238
- 239

240	Table 2 presents the estimation results. Overall, four drugs except for antibody
241	cocktails (sotrovimab, casirivimab/imdevimab), remdesivir, dexamethasone, baricitinib,
242	and tocilizumab had positive treatment effects among those found to be significant. That
243	finding might indicate that these drugs reduce the probability of saving life. Conversely,
244	antibody cocktails had negative treatment effects which indicated a contribution to
245	saving life. Only four pairs were identified to estimate parameters; other combinations
246	were not.
247	
248	Table 2. Estimation Results of Average Treatment using Inverse Probability
249	Weighted Regression Adjustment
	Age class All 65 years old or older Younger than 65 years old

Age	class
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	difference	p value	difference	p value	difference	<i>p</i> value
Remdesivir						
Proportion						
All	0.042	0.000	0.099	0.000	0.007	0.331
Oxygen therapy	0.043	0.000	0.070	0.000	0.002	0.746
Respiratory ventilator	0.034	0.370	0.071	0.160	0.001	0.987
Period						
All	0.040	0.000	0.094	0.000	0.006	0.335
Oxygen therapy	0.041	0.000	0.067	0.000	0.002	0.765
Respiratory ventilator	0.031	0.419	0.063	0.194	-0.029	0.415
Dexamethasone						
Proportion						
All	0.039	0.000	0.090	0.000	0.004	0.017
Oxygen therapy	0.032	0.000	0.056	0.000	0.002	0.684
Respiratory ventilator	-0.006	0.833	-0.017	0.670	0.005	0.889
Period						
All	0.039	0.000	0.088	0.000	0.004	0.022
Oxygen therapy	0.031	0.000	0.053	0.000	0.002	0.661
Respiratory ventilator	-0.005	0.858	-0.013	0.735	0.005	0.891
Tocilizumab						
Proportion						
All	0.114	0.000	0.255	0.000	0.025	0.010
Oxygen therapy	0.127	0.000	0.219	0.000	N/A	N/A
Respiratory ventilator	0.098	0.027	N/A	N/A	-0.011	0.776
Period						
All	0.117	0.000	0.259	0.000	0.025	0.010
Oxygen therapy	0.129	0.000	0.219	0.000	N/A	N/A
Respiratory ventilator	0.105	0.016	N/A	N/A	-0.030	0.441
Baricitinib						
Proportion						
All	0.072	0.002	0.371	0.000	-0.001	0.766
Oxygen therapy	0.107	0.006	0.316	0.000	-0.004	0.519
Respiratory ventilator	0.281	0.000	0.376	0.000	N/A	N/A
Period						
All	N/A	N/A	N/A	N/A	N/A	N/A
Oxygen therapy	N/A	N/A	N/A	N/A	N/A	N/A

Respiratory ventilator	N/A	N/A	N/A	N/A	N/A	N/A
Sotrovimab or						
casirivimab/imdevimab						
Proportion						
All	-0.042	0.000	-0.097	0.000	-0.001	0.878
Oxygen therapy	-0.074	0.000	-0.125	0.000	N/A	N/A
Respiratory ventilator	N/A	N/A	N/A	N/A	N/A	N/A
Period						
All	N/A	N/A	N/A	N/A	N/A	N/A
Oxygen therapy	N/A	N/A	N/A	N/A	N/A	N/A
Respiratory ventilator	N/A	N/A	N/A	N/A	N/A	N/A

250 **Notes**: Yellow denotes significance. "N/A" denotes not available.

251

252	Overall, all age and older patients had more significant estimators than younger
253	patients had. Moreover, few significant estimates were found when the subjects were
254	limited to patients who had used a respiratory ventilator. Only for tocilizumab in all
255	ages were these found to be significantly positive. Only four parameters were identified
256	for estimation for antibody cocktails, even though these were all negative and
257	significant.
258	Estimation results for mutated strains were defined as the proportion, which
259	was similar with that defined as the period, except for antibody cocktails. Among the
260	results obtained for antibody cocktails, there is no pair to estimate parameters.
261	

262 **Discussion**

263	Results showed that antibody cocktails might contribute to saving life
264	consistently. Our earlier studies, simple logistic regression and estimation using
265	propensity score matching, yielded similar results [28,29]. However, estimation results
266	of propensity score matching method for tocilizumab and baricitinib were, respectively,
267	positive and negative. In this study, we obtained consistent estimated results for
268	tocilizumab and baricitinib. These results indicated that tocilizumab and baricitinib
269	might not contribute to saving life. Even for antibody cocktails, the effects for younger
270	or more severe patients who used respiratory ventilators were not confirmed. Prior study
271	showed that casirivimab/imdevimab, one of antibody cocktails, reduced viral load [3].
272	Antibody cocktails were usually administered for mild COVID-19 patients, that is, they
273	were administered before viral load increased significantly. Therefore, they might be
274	more effective to COVID-19 compared to drugs used for severe COVID-19 patients.
275	The small sample for the younger patients or severe patients might be also another
276	reason.
277	These counterintuitive findings, except for those for antibody cocktails, which
278	are inconsistent with results obtained from earlier studies of the effectiveness of
279	remdesivir and dexamethasone [2, 8], might result from worse matching at the first step.

280	In addition, endogeneity or selection bias might not be controlled well in the decisions
281	to use drugs. Drugs to be administered against COVID-19 depend on the severity of
282	illness in patients. For instance, an earlier report described dexamethasone as effective
283	for severe COVID-19, but as ineffective against mild or moderate COVID-19 [8].
284	Actually, more than antibody cocktails, four other drugs were reportedly more effective
285	for severely ill patients [2-4, 8]. To resolve this apparent contradiction, more
286	information indicating severity might be necessary to make more precise the first step
287	estimation. Moreover, we did not use test results such as those for blood pressure, BMI,
288	or oxygen saturation. Such information must be included in the first step. Inclusion of
289	such data remains as a challenge for future research. The results of antibody cocktails
290	included many "not available", although drug effectiveness was confirmed in matched
291	cases. This result might be attributed to the small number of antibody cocktails used in
292	patients. In Japan, remdesivir, dexamethasone, baricitinib, casirivimab/imdevimab,
293	sotrovimab, and tocilizumab were approved in May 2020, July 2020, April 2021, July
294	2021, September 2021, and January 2022 respectively [30]. More recently approved
295	drugs are likely to be smaller samples. Accumulating data may resolve this shortcoming.
296	We also have to pay attention to apply the results of this study to all COVID-19 patients
297	because the targets of this study were all inpatients.

298

299 Limitations

- 300 First, we estimated drug effectiveness separately. However, the choice of drug
- 301 was not actually independent. Therefore, intercorrelation among drugs should be
- 302 incorporated into the estimation model.
- 303 Second, because MIA is a database of medical claims, data from the most recent
- 304 weeks may change during next few months. For this study, data were collected and
- analyzed from January, 2020 through the end of March, 2022, as of May, 2022. If the
- 306 study period is extended, the data and its estimation may differ over time.
- 307 Third, the drugs examined in this study were not necessarily administered to
- 308 the severest patients. For example, antibody cocktails were often administered to mild
- 309 patients. However, to simplify the analysis, we did not consider the combination of
- 310 drugs or change from one considered drug to another drug. This choice might introduce
- 311 some bias in the results. This process or pattern of drug administration may be
- 312 important in evaluating drug effectiveness.
- Fourth, due to data availability, it was not possible to take into account the timing of drug initiation, and oxygen therapy or respiratory ventilation to estimate drug effectiveness. If this information were available, it would be possible to distinguish

316	between patients who were administered drugs before oxygen therapy or respiratory
317	ventilation and patients who were administered drugs after oxygen therapy or
318	respiratory ventilation. It might be useful for more careful estimation to drug
319	effectiveness.

320

321 Conclusion

322 The obtained results demonstrated that antibody cocktails for all or older patients might

323 contribute to the saving of life. However, more information, including test results, is

necessary for better matching and achievement of definitive conclusions.

325

326 Data availability

The data in this study are available from the National Hospital Organization, but the availability of these data is restricted because of privacy reasons. These data were used under license for this study and are therefore not available to the public. Data are available from the authors with permission of the Ethics Committee and National Hospital Organization.

332

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336

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340

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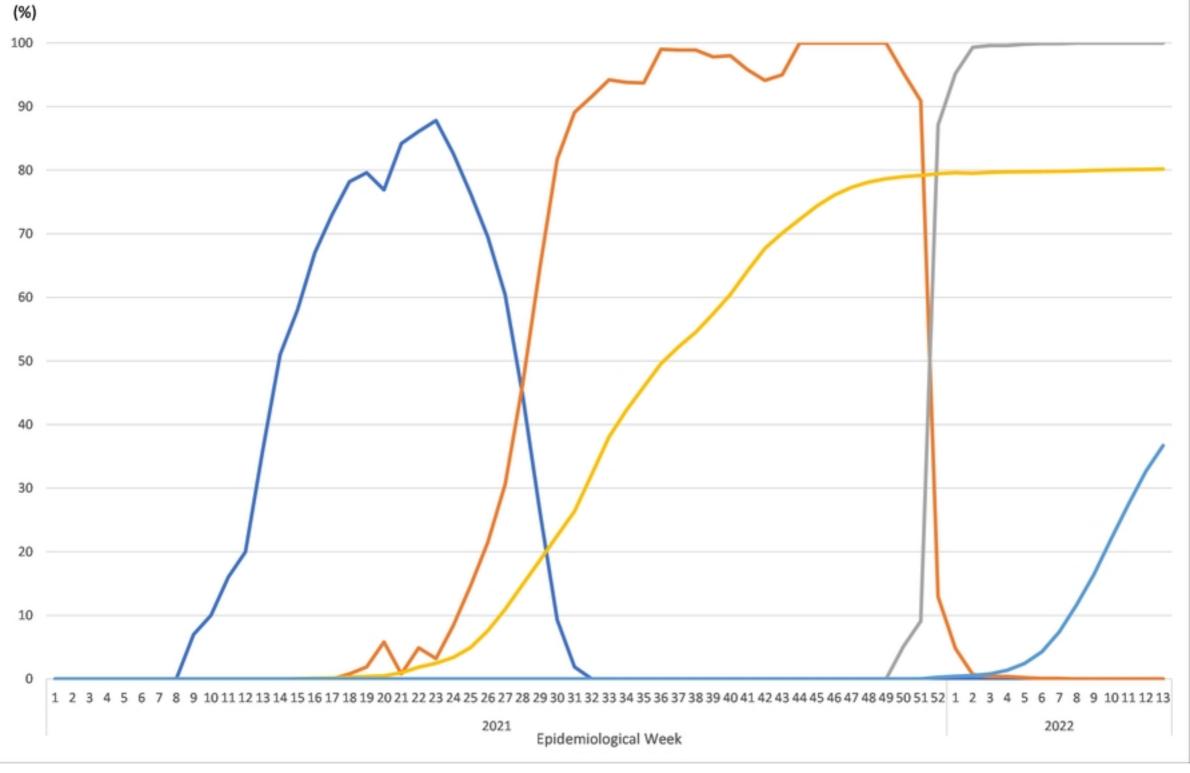


Figure3

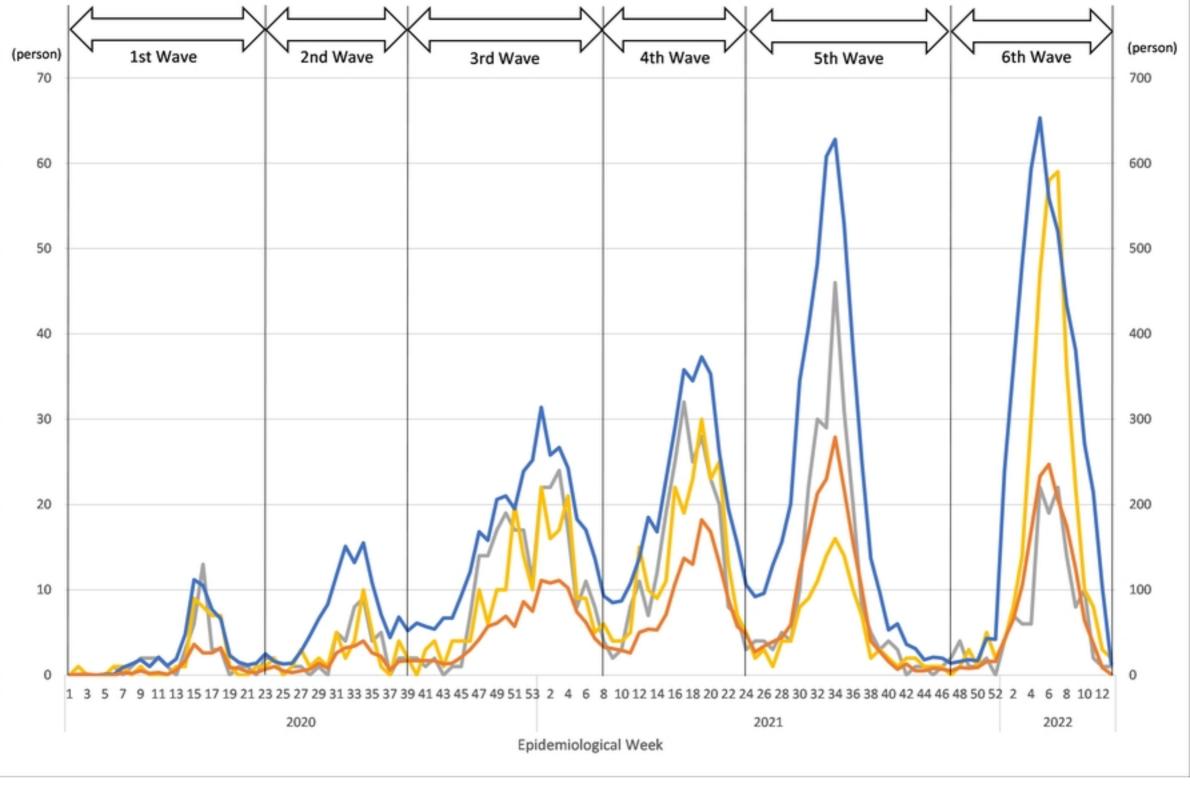


Figure1

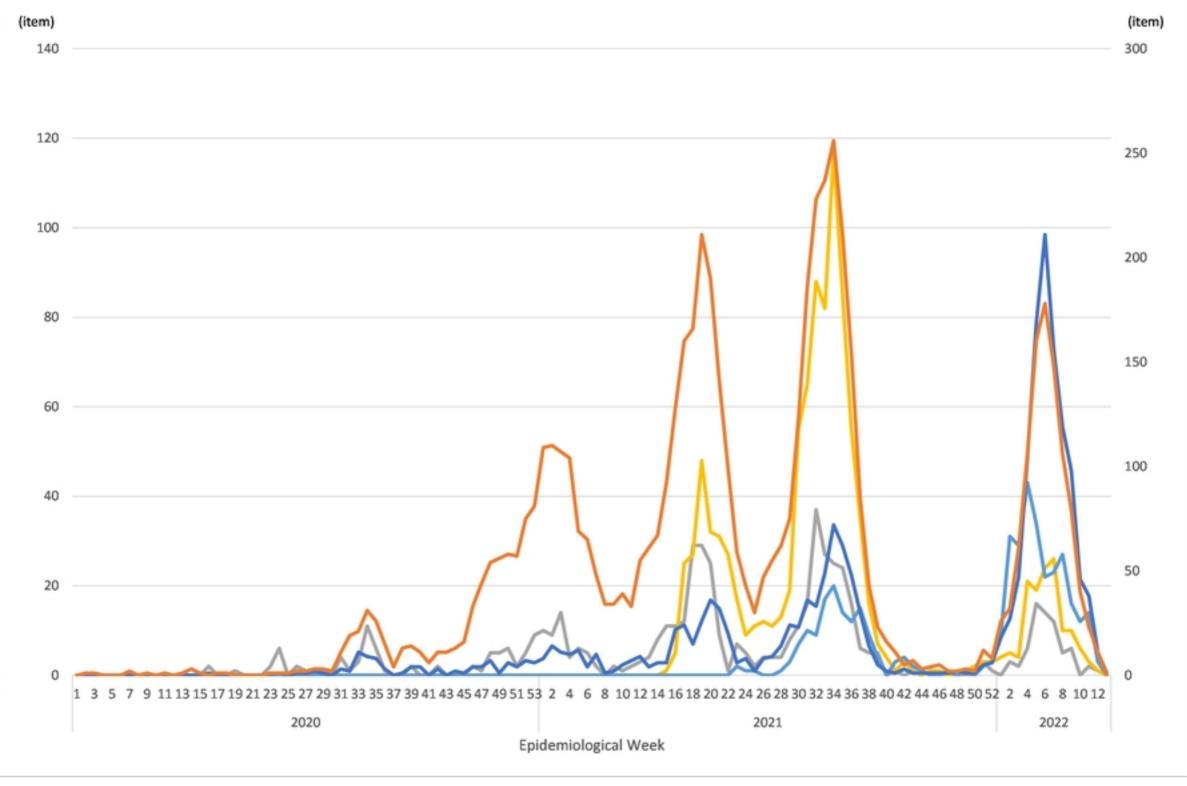


Figure2