



Selectivity versus Reach : Flattening the Curve of Covid 19 for Joint Health and Economic Prosperity

Authors

Jacques Bughin

Solvay Brussels School of Economics and Management, Université Libre de Bruxelles,
iCite – email : : bughinjacquesrenejean@gmail.com

iCite Working Paper 2020-034

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Jacques Bughin¹

We calibrate a SEIR contagion framework, with an extended transmission rate, R_t that accounts for two different classes of infections, and for the effect of both risk perception on citizens behaviors as well as government actions on Belgium data. We demonstrate that curbing the Covid 19 pandemic and enough hospital care capacity structurally with untargetted social distancing is challenging, requiring to sustain at least a recurrent cut of 50% of contagious contacts on top of creating unknown economic risks. With sufficient specificity (>70%) of tests, a model of selective tracing and quarantine is likely to be more effective, and much less costly to the economy. In practice, however, testing the full population is not possible, and would need to be randomized, requiring for safety a mix of testing and tracing, together with some other measures, like protection of more susceptible population either by nature (elderly), or by economic necessity (eg front lines workers).

1. Introduction

By April 14th, the number of recorded infections by the covid 19 is being close to pass the bar of 2 million individuals worldwide. The death toll is just below 120,000 deaths, according to Worldometer's compiled statistics (2020). The current virus, which originated in China, has now spread to more than 200 countries worldwide, leaving virtually no one safe.

It has put hospital systems struggling to cope with the flow of heavy infected cases, as feared by Richard Baldwin in a recent post on Vox (2020). Hospital, human, equipment as well as ICU beds capacity have been under excess demand in a matter of just a few weeks following the outbreak in Wuhan. By end of January, Chinese authorities stimulated the import of many health workers into the city, and managed to build up extra bed capacity in a record time to alleviate the pressure. The health system issue has been apparent as well in Europe, in particular Spain, Northern Italy and France. Major consequences of this disequilibrium are a higher fatality rate and a slower pace of recovery rate of contaminated people than average (Ji et al., 2020). Medical resources are also witnessing a high toll of illness and excessive stress levels (see Day (2020), Fang (2020), or Bughin (2020a)).

Accordingly, many countries have taken a radical « suppression » approach by imposing major economic standstill in the hope of flattening the epidemic curve of covid 19 and spread the demand for care more in line with health system abilities. This suppression strategy went crescendo, first forbidding large crowds meetings (theatres, mega-events, etc). It moved quickly to closing schools and to asking people to work remotely, then went to ask the population to stay home. In general, this suppression strategy is a powerful response to a first wave of an unknown pandemic emerging globally, like the Covid-19. There is emerging evidence, that if containment measures are followed by population, pandemic diffusion may slow (Lin et al, 2020, Gros et al. 2020). This also buys time to know more about the still unknown features of the Covid-19, and to work on means to battle it via the discovery of powerful antiviral and vaccines.

The challenge is that the suppression strategy also implies an economic shutdown, with likely large ripple effects if kept too long, and without massive macroeconomic support (Gourinchas, 2020). Such ripple effects might lead to a negative spiral of demand and supply reduction, bringing a major economic recession worldwide. Otherwise stated, the suppression strategy is not sustainable in the long term, as it creates a growing conflict between health solidarity and economic prosperity. Furthermore, from a behavioral perspective, it is known from past behavioral studies that people have a growing negative feeling, of the burden of prolonged quarantines (e.g., Person et al, 2004).

At the end of the day, the fundamental question is to find a more effective model in-between the two extremes of « no » and « full containment », while still allowing to flatten the curve before the population becomes immune, either naturally, or through medical support. Here, we shed some light on this issue. We look at the question of

¹ Solvay Business School. Fellow Applied Economics KUL, Portulans Institute

lockdown as the typical media business question of reach versus selectivity. Reach is to get all people contained, but may be this is costly and ineffective, versus selectivity, whereby one targets segments or personalizes responses more much effectively. One simple way to assess the best policy model is to merge economics with epidemiology, and look at the interplay between the drivers of the reproduction rate with economic actions. By calibrating a model of contagion to Belgian data, we shows that the reach strategy is usually very sensitive to the ability of sustaining a structural reduction of contacts, on average by more than 50% than current. A selective strategy, whereby infected people and its close ties accept a complete quarantine, is most effective, to the extent that we have at hand a sufficiently powerful test, above 70% specificity.

Those results are obviously only illustrative ; but they signal that a selective model with testing and tracing might be a more powerful strategy than any other extreme. This strategic shift is a way to make health and economic prosperity rule converge.

2. Conceptual model

We rely on a classical SIR model, whereby the basic reproduction rate R_0 is given by (see Kermack and McKendrick, 1927) :

$$(1) R_0 = c/v,$$

where c is the contamination rate between infected I and susceptible, S , individuals, and v represents the rate at which infected I people recover.

Using the SIR, it follows that growth of infections through time t , dI/dt is given by the difference between the contamination rate, c times the stock of infected, and the portion v of recovering patients at time t :

$$(2) dI/dt = c(I/N) - vI$$

where N is total population

Our modelling strategy is to move from a constant « c » as a scaling factor of infection growth, to a more general form of the reproduction rate R_t , of the total population through time, consistent with He et al., (2013), or Lin et al. (2020). We, in particular consider that contamination depends on the distribution network of contacts, that people are trying to insulate themselves from infection during contacts, and that we have two segments of infected, one of proportion r , that includes the recorded cases ; and the balance is the non recorded. Typically, as covid 19 has unequal effect on infected, r represents the segment of those tested at time of more severe conditions, especially when people are brought to hospital. Hence, equation (3) posits a formulation of R_t that makes more explicit, a set of crucial, drivers of the dynamics of the virus diffusion :

$$(3) R_t = (c/v) * ((1-ut)d+ut)) * r * (1-sr) * (1-Dt/Nt) \exp(er) + (1-r) * b * (1-s) * (1-Dt/Nt) \exp(e)$$

where $*$ is the multiplier function, and Dt/Nt is the portion of severely infected people, so that e measures the sensitivity of people to adapt their behavior in function of the risk of the disease.

The vector of key parameters, (d, r, s, e, b) is a representation of five important drivers of how the coronavirus disease, covid 19. As we have two segments, some of the parameters, s, e can be segment specific, but we have no empirical basis to do this, so we will now assume, $s=sr$ and $e=e$. Possible differences in those parameters will be absorbed by the relative factor b . Regarding those parameters, we note :

a) R_0 is usually used to define the entire risk of the population infected by the virus, in the limit $R_0 \rightarrow \infty$. This result however relies on an uniform distribution of contagious events, while most outbreaks are not shaped by the “average” individual but possibly by a minority of superspreading events. In particular, the probability of containing an outbreak is significantly lower if there is large heterogeneity in secondary infections, as social spread is weakened across the population (Hebert-Dufresne et al., 2020). To account for the asymetry of

contagion, we hence posit: $0 < u < 1$, where $1-u$ the portion of superspreaders, and $d > 1$ is a multiplicateur effect. For instance in the case of a Pareto distribution of infection, $u=80\%$ and $d=16$, and thus the term $(1-u)d+u=4$, reflecting that the network asymetry reduces the power of spreading to the population significantly, for a given R_0 . We do not know the exact distribution associated with the covid 19, but we have a glimpse from triangulating from other virus distributions of contagion. Measles infection looks like a Pareto distribution; the top 20% typically contributes like 87% for SARS, while the figure is 93% for HIV (Lloyd Smith et al, 2005). For influenza, the distribution looks more like top 20% makes 50% of the secondary infections (Brauer, 2019).

At the level of influenza, the term $(1-u)d+u=1,56$. As typically, one estimates R_0^{\wedge} from observed diffusion data that already embeddes the effect from the distribution asymetry of contagion, such estimate R_0^{\wedge} suffers from a downward bias. For an estimate R_0^{\wedge} , say of 1,3, the true R_0 , for influenza, will then be $1,3^*/1,56$, or more like 2. Assume still that one can target the portion $(1-u)$, and get half of them adequately identified and fully isolated (or 10% of population) by time t , then $(1-u)d+u$ declines from 1,56 to 1,28, and R_t by 18%. This effect is twice bigger than isolating 10% of any average individual ; if the asymetry will be like Pareto, the reduction will then be more like 40%. In large asymmetric infections, finding the superspreaders is rather effective, but slightly less in the case of influenza type of viruses.

b) Many countries have been in short supply of testing capacity, while also claiming that tests available in the market to date are suffering from low specificity. Thus, the majority of countries, at the exception of some Asian ones, did limited testing so far. Often, they reallocated the available stock of tests, for those emergency cases showing up at hospitals to sort out true infected cases from others. We take a view then, of two segments of population, one which is tested and recorded, and another one, which is not tested. Accordingly, we may posit : $0 < r < 1$ where r the portion of recorded infected cases to total cases.

The value of r , in practice varies with times. At the outbreak of the virus, it might be close to zero. This is especially true in the case of the covid19, where the contagion may span over 20 days. r is also likely increasing with time, as pre-symptomatic cases become ill, and countries build up more testing capacity and test specificity increases. We rely on other scholar attempts to have an idea of the actual range of value for r . We have estimated r to be in the range of 10% to 20% by March 20th, for most of European countries after 4 weeks into the pandemic implying that a significant amount of cases has been unrecorded (Bughin, 2020b). Li and colleagues (2020), came to the same conclusion for the case of Wuhan, showing that those non recorded cases may be responsible for a rather rapid dissemination of novel coronavirus. A powerful case study is the Italian village of Vo, with 3% by early march of its population infected by the coronavirus, or roughly three weeks after being the host of « patient zero » in Italy, and after all the population got tested and strict containments were put in order. This level of contagion implies a true number of infections about roughly 8 times what was recorded in the 11 towns most affected by the outbreak in Italy. A large part of under-reporting is that covid 19 may high high asymptomatic effect ; 50% of the individuals infected, did not get any symptom, in the case of Vo.

c) The contagion spreads though physical contacts, due to the mechanism of entries of the virus into the respiratory system of individuals. We posit ; $0 < s < 1$ where s is the portion of reduction in contagious social contacts. This reduction is of course a weighted average of actions between different sociodemographic segments—for example, someone in its late thirties, who has a front line sales service job, one teenager kid at high school, and a good social life, might have roughly 5 times more contacts than any retired single person of 60 years old, according to our computation. In general, also, older persons' contacts become only social in a closely tight community setting, making them more at risk, but less contagious given their tied networks. Young adults have all sources of, and all types, of close contacts possible, -friends, work, school kiss and drives, concerts, parties, love partners, etc. Most studies of contact estimation suggest that school and work contacts account for roughly 40% of total contacts, 1/3 extra arises from community circles, and the balance is from own family (see Kelso et al. 2013).

Getting s larger than 0,5 is however challenging. This is because some work from front lines remain necessary in critical sectors, such as healthcare, equipment manufactruring and food and pharmaceutical services for instance . Those sectors typically amount to 25% to 30% of a developed economy. Citizens are usually not fully compliant with social distiancing. In fact, most academic studies suggest that contact rates for non infected individuals, tend to decrease by 30 to 40% during periods of large influenza, but this is rarely higher, if not strictly imposed (see Caley, et al 2008).

d) The term, $(1 - D_t/N_t) \exp(e)$, where D_t/N_t is the ratio of (death and patients in critical conditions) to population N_t , at time t . We posit that $e > 0$ where e is a proxy for risk aversion towards the virus. It plays out as a multiplier of extra caution taken by the population as far as the severe contagion spreads into the economy. During the 2019 H1N1 outbreak, 25% of Americans were avoided crowded area, as part of the risk perception to catch the disease (Steelfisher, et al, 2010). For a flu-type, a resulting reduction of the attack between 20 to 40% is not implausible as people start to wear mask, and other protective equipments (Tyson, et al 2020). There might also be evidence that people distort information, and overstate the risk of the disease (Brahmbhatt and Dutta, 2008). This adds to the idea of e being large. For example, during the 2002 SARS, more than 25% of Asian citizens thought they could be contaminated, even if the ex post rate happened to be less than 0.1%.

e) We finally posit $0 < b < 1$, where b is the ratio of the level of contagion of non recorded cases to total cases. We note that b is linked to milder cases or asymptomatic cases of the Covid 19. There are only a few studies trying to estimate b , and they seem to conclude that b might be in the range of 0.4- 0.5 for Covid 19. If the portion of asymptomatic cases may be as large as 50%, b is likely to be in the range of 0,4 to 0,6. As said earlier, this rate of asymptomatic to total cases has been noticed in fully tested populations like the village of Vo in Northern Italy, or in the cases of random sampling in South Korea..

3. Experiments

We have played with the five parameters above, after calibrating the SIR model to Belgium as of April 1, at the time Belgium had recorded just below 14,000 covid-19 infections, and suffered 828 hospital fatalities. Belgium took a decision relatively early for a comprehensive lockdown, already the day after the 10th recorded deaths in the country. It is however notably suffering from supply chain scarcity, as well as from test availability (just above 6/1000 of population has been tested by April 1) so that the country early chose to only test people at hospital venues. Belgium does not trace people, even if it may be looking at possible means, at current stage.

We focus on hospitals only, not home care, as we wish to look at the adequation between hospital capacity and hospitalizations. Home care is a rather large sector in Belgium, and is likely the largest pool of infections, given the demographics of home care residents. It is estimated to represent to date up to 50% of the fatalities in the country. Most home care are put in quarantine, and will start to be extensively tested by now.

Sensitivity analysis

If we look at the daily change of recorded cases and fatalities, on that day, Belgium recorded cases might imply that the pandemic could have been spreading, like the flu ($R_0=1,3$) by April 1, as crude estimate based on daily recorded cases for 20 days before, implies that R^t looks like 1,5. We however know that this R^t suffers from mismeasurement errors and possibly asymmetry in contagion. For our simulation, we thus have computed R_0 for three assumptions of infected cases, eg with $r=0,7$; $r=0,2$, and $r=0,1$ by April 1.

R_0 is computed up to March 18th before the containment measures, and before risk aversion may kick in. Assuming that $d > 1$ is like the influenza, we compute that the adjusted $R_0=3,8$ for $r=0,7$; $R_0=2,7$ for $r=0,2$, and $R_0=2,55$ for $r=0,1$. Not surprisingly, our estimate of R_0 decreases with r , as r implies a large volume of unnoticed infections at the start. At those levels of R_0 , Covid 19 might look to be twice more infectious than the typical flu, at $R=1,3$. This range of R_0 is also consistent with multiple studies on the covid 19 and also hints at the fact that we must induce large risk aversion and/or reduce a large portion of social contacts to hope to curb the disease, with $R_t < 1$.

Table 1 presents the simulation results, based on *one final assumption* that unrecorded cases have 3,5 times lower probability to end up as a fatality in hospital than recorded cases. This assumption is consistent with the portion of large asymptomatic cases. Our simulation is illustrated the following discrete values: $s=0,4$ and $s=0,7$ (As said, $s=0,7$ is difficult to achieve without very strict, and coercitive, rules, especially if the effort must be sustained more than 4 to 6 weeks); $b=0,4$ and $b=0,6$, as well as $e=220,1100$. The value of the risk aversion e (1100) has been computed from different studies (He et al., 2013). Regarding r , we present the two extremes case of $r=0,1$ (most likely) and $r=0,7$ (current understated cases). We posit as well that $u=80\%$, and u is increased to 90%, as a result of stopping mega events, vacation gathering, etc. We also compute the implied hospital ICU

capacity, as the last column, based on a capacity of 2500 ICU hospital beds. The results in **Table 1** demonstrate that :

a) The main driver of the outcome of the covid 19 among the 5 variables discussed is **the level of change in social contacts, s**. Given the high R_0 for Covid 19, it is usually good to have social contacts on average reduced by more than 50%, in order to reach a peak in infection. If this is maintained, then pandemic is stopped, but a too fast relaxation of measures that makes $s < 40\%$ will lead to a new exploding wave of contagion.

b) Given Belgium high ICU bed capacity, - above average of EU-27 countries-, and the fact that Belgium went quickly into containment actions, ICU capacity is still ok, even at $s=0,4$; but there are also a few cases when ICU capacity becomes tight, **especially when risk aversion is low, and does not induce enough of self-protection of people in face of poor social distancing by third parties (e.g, when $e=220$)**

c) Superspreading plays a role, **but especially when risk aversion as well as contact reductions are limited- alas, it also happens when time ICU capacity becomes tight.**

b) Adequate measure, r , of true infections is a must, as it drives the dynamics of healthcare resources, in particular in the long-term ; this is because a better understanding of total infections makes R_0 more accurate and typically not overstated, but it also implies a longer tail of cases. In our simulations, should r be truly 0,7, we know that R_0 should then be higher (R_0 at $r=0,7$ reaches 3,8, or roughly 60% higher than R_0 for $r=0,1$ to fit the data dynamics of infections). This means, that for the same proportional reduction of contacts, s and same level of self protection, e , R_t will be higher in absolute value, at same t , for $r=0,7$ than for $r=0,1$. This means that the run rate, after May1, will be higher in that case.

Further, when total infections are much higher than recorded ones ($r=0,1$), a large part of infections may be due to lighter cases than cases requiring long hospitalisations ; but in reverse, when those cases are requiring hospitalizations, they tend to be more serious cases in absolute terms, and with higher mortality². As mortality further shortens time spent in ICU and peak is happening slightly faster, capacity usage after May is lower in case of $r=0,1$. However, this does not say that this is better managed, as this may be arising at the expense of a tighter capacity in late April, and larger mortality.

Table 1.- Belgium SIR model for covid 19, rollout from April 1, 2020 to May 1, estimates

<div style="border: 1px solid black; padding: 2px; display: inline-block;"> $r=0,1,$ $R_0=2,4$ </div>		e	s	B	u	total infection	Total Deaths	Infection peak by (10 days period)	run rate death (10 days ca	ICU
										May
		1100	0,7	0,6	0,9	103000	2660	April 11 to 21	290	
		1100	0,7	0,6	0,8	107250	2740	April 11 to 21	300	
		1100	0,7	0,4	0,9	97800	2565	April 11 to 21	280	
		1100	0,7	0,4	0,8	100250	2600	April 11 to 21	290	
		1100	0,4	0,6	0,9	125000	3060	No peak	370	
		1100	0,4	0,6	0,8	137000	3260	No peak	420	
		1100	0,4	0,4	0,9	110500	2810	No peak	330	
		1100	0,4	0,4	0,8	118000	2930	No peak	360	
		220	0,7	0,6	0,9	116000	2890	April 11 to 21	340	
		220	0,7	0,6	0,8	125000	3045	April 11 to 21	385	
		220	0,4	0,6	0,9	173000	3780	No peak	685	
		220	0,4	0,6	0,8	211000	4300	No peak	970	

² Hospitalization rates are 3,5 lower for non recorded cases, but there are 7 times more infections than in the case of $r=0,7$. Thus, the system gets more people into the hospital gate

r=0,7 RO=3,8					total	Total	peak by	run rate	ICU
E	S	B	U	infection	Deaths	(10 days period)	death (10 days ca	May	
1100		0,7	0,6	0,9	29000	1340	April 11	490	
1100		0,7	0,6	0,8	37000	1435	April 11	600	
1100		0,7	0,4	0,9	26500	1300	April 11	470	
1100		0,7	0,4	0,8	31500	1380	April 11	570	
1100		0,4	0,6	0,9	71000	1815	No peak	1210	
1100		0,4	0,6	0,8	99000	2060	No peak	1680	
1100		0,4	0,4	0,9	60000	1680	No peak	1095	
1100		0,4	0,4	0,8	82500	1890	No peak	360	
220		0,7	0,6	0,9	32000	1375	April 11	540	
220		0,7	0,6	0,8	40400	1480	April 11	685	
220		0,4	0,6	0,9	91500	1920	No peak	1520	
220		0,4	0,6	0,8	140000	2210	No peak	2220	

Core assumption 1: non recorded cases (including asymptomatic cases) leads to 4 times lower hospitalisations;

Core assumption 2: Distribution of contagion follows influenza (top 20% makes 50% of contagion)

Core assumption 3: hospitalisation conversion to ICU and to fatalities calibrated on April 1 and then constant

Core assumption 4: recovered cases are immune in short-term

Core assumption 5: contagion is 15 days for very mild cases, not going to hospitals, 8.5 when going to hospital

Based on SEIR model, and calibrated data on period Mars 21 to April 1; only includes hospitalisation cases so as to me

Maximum likelihood point estimates

On top of the sensitivity analysis, we provide point estimates of where Belgium may lie. We do this by finding the vector that minimizes the residuals prediction for April 11, where data are available for Belgium. To ensure convergence, we first hypothesize the most probable cases. Based on the above, the current range of values should be in the range of $r=0,1$ (multiple press reports had suggested that 100,000 cases could be the true toll of the pandemic by March 23, in Belgium³); $s=0,5^4$; $b=0,4$ (as probably a large part of cases are asymptomatic cases); $e < 1100$ as we also know that e may be « capacity constrained », (very limited mask protection is available for example in Belgium). We also assume that $(1-u).d+u = 1.6$, as for H1N1 influenza type. We can posit as well that u is 95%, as Belgium cancelled most events with more than 100 people, and closed restaurants, etc.

Using maximum likelihood function optimisation, the vector that minimizes the gap versus observed data by April 11, 2020 suggests that $e = 940$; $s = 0,55$; $b = 0,43$; $u = 0,96$ for $r = 0,1$. This is rather well specified as any 10% deviation from this vector on average increase residuals by more than 150%. The fit deteriorates fast if we increase the value of s and b , and decrease e . Social distancing to date has resulted in a decline just above 50%; lower than what has been computed for China (> 80%; see Lin et al. 2020) as enforcement in Wuhan and China were very strict, for example. At this level, given R_0 at 2,6, this is not enough per se to stop the pandemic, as the

³ This extrapolation is based on the number of cases reported with flu symptom, while the pandemics of flu was official phasing out by early March, see <https://www.dhnet.be/actu/belgique/100-000-cas-actifs-de-coronavirus-a-redouter-actuellement-en-belgique-5e7c88c8d8ad5816316d2145>.

⁴ It is said that for instance, about 70% of Belgian people are working from home. Nevertheless, travel outside home has been reduced by about 30% in a radius of 5 kms, versus traditional periods without containment, see <https://www.vrt.be/vrtnws/fr/2020/03/26/enquete-de-l-universite-danvers-les-seniors-suivent-mieux-que>, and <https://plus.lesoir.be/290734/article/2020-03-28/selon-une-analyse-des-donnees-telecoms-les-belges-adaptent-leur-comportement>

new $R_0 = s \cdot 2,6 = 1,4 > 1$). We find that unrecorded cases are less contagious than others, in line with the fact that possibly half of infected people are asymptomatic, and in line with other estimates (Lin, et al, 2020). Risk aversion is present, but lower than what observed for the Spanish Flu with a 20% lower effect on limiting contagion (He, et al., 2013). The value of u means that about 40% of superspreading cases have been spotted.

If we optimise for $r=0,7$, the local optimum implies a doubling in e ($e = 2200$), s increases to 63%, and b declines to 0,25% ; $u=0,94$. We should expect the optimisation to lead to such a result, as it implies that the limited number of cases outside hospitals is much less contagious ($b=0,25$ for $r=0,7$ versus 0,43 for $r=0,7$) than the ones arriving to hospitals, reinforcing the rationale not to focus on those cases when one lacks test ability. The results for $r=0,7$ are however not that credible. They exhibit a significant deterioration in predictions, with a twenty-fold increase in residuals versus the case of $r=0,1$ ⁵.

Continuing thus on our favorite case on $r=0,1$, and keeping the values of the vector constant for April, Belgium might reach 105,000 total (recorded and not recorded) infections by May 1st, with a cumulative toll of 2700 *hospital* deaths, and an active ICU utilisation, of 1400 beds. The peak in daily infections happens before April 17 to April 21, while the number of new cases being hospitalized peaks slightly later, as do number of deaths, just before May 1st, reducing the pressure on ICU.

Still, a deterioration in s and e would lead to a plateau and a rebuild of demand for ICU, especially when $s < 0,4$ and $e < 550$. We thus confirm that reduction of contagious interaction, s , is the main first order target, and increased self protection, e . In the absence of specific targetting, and only testing at time of hospital venue, we should aim for a reduction of **50% of interaction** both to flatten the curve, and be compatible with health constraints ; Risk aversion leads to self protective behavior is a good thing, leading people to more systematically wash hands, wear masks etc. In this respect, we believe that the controversy raised in Belgium regarding the wearing or not mask is unfortunate ; the behavior is effective if everyone does it addition to other distancing and self protective behaviors.

4. From reach to selective testing, tracing and enforced quarantine

One can directly derive from above, that a reduction of contact by 50% is difficult to hold, especially, *without* some imposed shut down of education and work in the context of the Covid 19. In such a case, we estimate that $s = 28\%$, and infections, ICU and death toll will increase versus current case by 20%, 18%, and 15% by May first, 2020⁶. Peak will not arise in this time period, or if it has, outbreak will restart for a wave 2.

However, keeping the economy and education close, has large perverse consequence on the economy. Hence, it must imply a few additional amended strategies, for economies to be relaunched, and keeping the disease under control. This includes :

a) **Telecommuting should be promoted as a standard** of practice going forward. Nevertheless, this is only possible for certain sectors, and for some functions, possibly, for about 1/3 of the working population to date⁷ (see also Dingel and Neiman, 2020). Even if done, the task distribution might imply that this will be possible for only a partial time of the work, implying major reorganization of the way one work. Further, the shift will

⁵ For sake of completeness, we also increased the asymmetry of contagion distribution, d from the current case of top 20% = 50%, to top 20% = 65%, which is in line with some influenza metrics. By assuming a higher asymmetry, the prediction deteriorates too, but only two the three fold the original simulation. With this new case, we implicitly state that the actual R_0 is bigger than thought, but contagion loses power across network of individuals. We find that b decreases accordingly, from $b = 0,43$ to $b = 0,36$; s increases from 0,56 to 0,62, and $e = 940$ moves up to 1160. The value of u remains unchanged. Again, this result simply reflects that one may have to push harder to flatten an pandemic with higher R_0

⁶ Multipel studies suggests that about 40% of close contacts is linked to school and work. Given that about 70% of those contacts seem to have been eliminated through containment, about half (29%) of s (57%) is due to this channel ; or otherwise, said, without it, s will decrease to $s = 28\%$, creating a rebuild of the curve

⁷ See reference in <https://bfi.uchicago.edu/working-paper/how-many-jobs-can-be-done-at-home/>

especially play on the lower quartile of the working population, as already telecommuting is easier and more pronounced for higher quartile of workers, (managers, etc).

b) **Public protection must become the new normal**, eg in Asia, most restaurants require individuals to wash hands, most malls have hydroalcohol gel at their entrance to be used by the public etc. Close contact prohibitions make this a difficult strategy for major sportive, cultural, or entertainment events, or major business fares for example. The same issue may apply for most transportations, not only within the mode of transportation, but because excessive mobility may seed new waves of the coronavirus, as currently witnessed by China⁸.

c) **Cyclic shutdown may be a rule for a while**, by which part of economy, part of working days are shutdown in cycle for the 2 weeks of contagion.

c) Finally, **one important policy seems to be** the priority of systematic testing, as acknowledged among others by Dewatripont et al. (2020). Testing can be used for two objectives : ensuring people are not infected, so they can be safe to be interacted with, or be put back to work. One clever idea is group testing to maximize costs and roll out, see Gollier and Gossner, (2020). Or, in our case here, ensuring that one can spot the infected and their close social ties upfront, so that they can be put in strict quarantine, instead of putting everyone in containment.

Regarding the last point, we again resort to simulation to show the logic, based on Belgian figures for the Covid-19 pandemic. At the estimated R0 of 2,6, u=96%, and s= 28% as result of no imposed economic actions, we find that the new amended R0 becomes 1,7, and about 69% of population may be affected in the long term by the covid 19 with z= 69% is the result of solving for the equation (4) :

$$(4) \quad R_0 + (1/z) * \log(1-z) = 0$$

where R0= 1,7.

In the case of s=57% (full shutdown), same math leads to 17% of the Belgian population to be infected. We thus infer that the economic lockdown prevents 52% of population infected cases, or a reduction of 52/69= 75% of the infections. Yet, a) it also hurts at least 31% of people who will not have been infected ; b) on top of this, if all infected cases will lead to 3 weeks of work lost, this has to be compared to 70%*(1-31%)=49% of work lost for 8 weeks, or more than two times the true cost of shutdown, which may or may not be justified by the productivity loss of the fatalities.⁹

Now, consider we can target effectively the contaminated individuals, at 70% or 90%, and their social ties obey like today, in Belgium at containment, at 50% rate, or it improves in proportion of the one at risk, at 50%/69%= 72%. At those levels, worse case (70% specificity and 50% rate of compliance) still imply 59% of cases contamination, while the base case (90% and 72% compliance) , leads to less than 5% contamination. Middle case (70%specificity, 72% rate of compliance) would lead to 30% attack rate. We derive from this simulation that a 75% specificity success, with 75% compliance of quarantines for infected and their social ties, may lead to same effect of economic shutdown, but with much lower economic and fairness issues.

⁸ <https://news.yahoo.com/china-reports-record-coronavirus-cases-013153635.html>

⁹ 70% is 100% shutdown with 30% of economy remaining in function as necessary. Note that, if all infected cases will lead to 3 weeks of work lost, this has to be compared to 70% *(1-31%)=49% of work lost for 8 weeks, or more than two times the true cost of shutdown. On the other hand, and assuming at 2% death rate from our Table 1, and statistics that suggests that 40% of total deaths will be for people with still 20 years of work, this is equivalent to 52%*2%*0,4*20=0,08% of a year, or 4 weeks of productivity loss equivalent= thus, at this level, the economic shutdown becomes a valuable return. The labor productivity is only one factor, = if this creates a demand pressure and a spiraling depression, then the shutdown is indeed not returning a positive ROI for society.

The new social norm of being traced

We are not stating from above that we should give up (all measures of) the economic shutdown. We understand that test specificity must be high enough, and jury is still out if it can, especially given high asymptomatic cases of COVID-19. Further, tests must be large, and tracing continuous to control exit. We might have to do a mix of both approaches in those circumstances, even if today, asymptomatic cases may have low contagious power; and evidence shows that medical and non-medical testing combined may make the target of 70% possible (Alibaba claims that its AI-based applications have success at 95%).

We also understand that tracing individuals can be done through mobile and Bluetooth or Beacon-like technology, but this must be done with extreme caution (Christoph and Gunther, 2020). However, given the large externality of contagion, tracing is likely to be raised as an important social norm to comply with, in order to accept a reconciliation of health and economic prosperity.

In practice, this path of combined testing/tracing and selective economic shutdown has been shown today by countries such as Singapore or South Korea, to be the best to control the COVID-19 pandemic to date (Anderson et al, 2020). In those cases, data were anonymized and a civil society governance model for data crunching was made possible to preserve privacy and GDPR compliance. As such, digital technologies may become a great complement to our society, both as prevention to find cures and vaccines to COVID-19, as well as to manage the social contagion process (see Pissarides and Bughin, 2019).

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