

Journal of
Social and Administrative Sciences

www.kspjournals.org

Volume 9

September 2022

Issue 3

**Determinants in the emergence of viral agents: the
SARS-CoV-2**

By Mario COCCIA [†]

Abstract. The objective of this paper is to clarify, whenever possible, the determinants in the emergence of biological agents to improve aspects connected with public health and biosecurity. Case study of the Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) is investigated to assess the likely emergence from a wildlife spillover and/or scientific research in labs with unexpected accident. Using a meta-analysis, results suggest that a natural spillover of SARS-CoV-2 that has generated more than 5.2 million of deaths, in analogy with natural disaster, seems to have a remote probability, instead a lab accident in the process of scientific research has a probability of occurrence of about 15-30%. These results here are important to support decision making of policymakers for global biosecurity strategies with appropriate responses to prevent the future diffusion of vital agents similar to SARS-CoV-2 in environment and society.

Keywords. Viral agent; Biological agent; SARS-CoV-2; Novel coronavirus; Zoonoses; Natural disaster; Lab accident; Laboratory biosafety; Biosecurity risks.

JEL. C52; L25; M14.

1. Introduction

Coronavirus disease 2019 (COVID-19) is an infectious illness caused by the novel Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2), which appeared in late 2019 (Anand et al., 2021; Bontempi et al., 2021; Bontempi and Coccia, 2021; Coccia, 2021, 2021a). One of the main questions in science and society is if the origins of SARS-CoV-2 is due to natural event of spillover from wildlife or associated with human activity of scientific research (Andersen et al., 2020; Boni et al., 2020; Frutos et al., 2021; 2022; Relman, 2020; Sachs et al., 2020; Segreto et al., 2021; Wolfe et al., 2007). Bloom et al. (2021) argue that initially information provided by Chinese scholars does not clarify if the cause is due to a natural (zoonotic) spillover of bats through an intermediate host or a possible lab incident. The latter hypothesis may be associated with Mojiang mine (China) incident in 2012 when six miners died with an unexplained viral pneumonia (cf., Rahalkar and Bahulikar, 2020). Frutos et al. (2022) discuss about pros and cons factors of the natural origin of the unexplained viral pneumonia in Wuhan (China) over 2019. In this context, Sirotkin and Sirotkin (2020) argue that the etiology of this novel coronavirus is hardly known because the intermediate host for

[†] CNR, National research Council of Italy & Yale University School of Medicine, 310 Cedar Street, Lauder Hall, Suite 118, New Haven, CT 06520, USA.

completing a natural zoonotic jump is not identified, and the application of research techniques of gain-of-function of viral serial passage is one of the possible sources of this novel coronavirus. In fact, the molecular analyses of specimens raise further questions that suggest further investigations of origins of SARS-CoV-2 and factors of risk associated with gain-of-function research (Sirotkin and Sirotkin, 2020). Relman (2020) maintain that to avoid next pandemics like COVID-19, it is important to unravel the origins of SARS-CoV-2. Sirotkin and Sirotkin (2020) also argue that the origin of SARS-CoV-2 plays an important aspect to develop effective drugs and apply appropriate treatments. Overall, then, COVID-19 is still circulating in 2021 with mutations of the novel coronavirus but the origins of SARS-CoV-2 are still unknown.

The present study confronts this problem here by developing a meta-analysis to clarify, whenever possible, the origins of SARS-CoV-2 considering the two principal hypotheses given by a natural spillover and factors associated with human activity in research laboratories. This study is part of a large research project directed to explain the origins of SARS-CoV-2, factors determining transmission dynamics of COVID-19 and best practices to design effective policy responses to cope with and/or to prevent pandemic threats in society (Coccia, 2020, 2021b, 2021c, 2021i, 2021d, 2022).

2. Methodology

The deductive approach of this study is as follows.

Firstly, the social and health phenomenon observed is the COVID-19 that is still circulating with mutations of the novel coronavirus (SARS-CoV-2) and generating continuous infections and deaths in manifold countries (Johns Hopkins Center for System Science and Engineering, 2021).

Secondly, multiple working hypotheses about possible origins of the SARS-CoV-2, based on literature, are proposed (Coccia, 2018):

Hypothesis 1. Natural (zoonotic) spillover from of the novel coronavirus from bats through intermediate host

Hypothesis 2. The application of scientific research for science advances (and consequently lab accident)

Thirdly, the proposed hypotheses are tested by metanalysis and statistical evidence to accept or reject them. In particular:

- The first hypothesis is assessed with estimates of the probability of occurrence used for big natural disasters that generate a lot of fatalities, considering COVID-19 pandemic as a natural disaster (USGS, 2021).
- The second hypothesis is analyzed with the approach of backward chaining: an inference method used in many artificial intelligence applications (Russell and Norvig, 2010). Backward chaining starts with proposed hypothesis and works backwards from consequent facts to antecedent events to assess if any data supports any of these consequents (Figure 1).

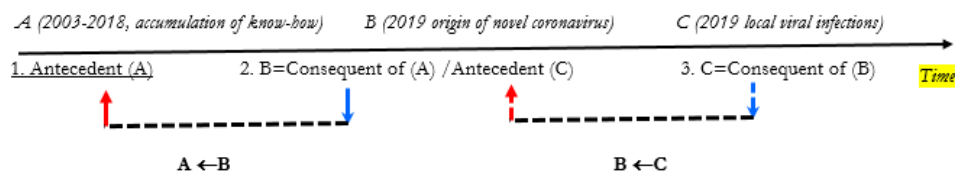


Figure 1. Backwards changing to explain the emergence of the novel viral agent

- Fact (consequent C): The novel virus was first identified in the Chinese city of Wuhan in December 2019
- (B=antecedent of C and consequent of A): the existence of laboratories in Wuhan able to apply scientific research of gain-of-function to support science advances in virology
- (A=antecedent of B). If Chinese lab located in Wuhan has technical know-how to support science advances in coronavirus. In fact, one of the most important approaches to explain scientific development is the theory of the accumulation of knowledge (Science, 1965). The cumulative theory states that scientific development is due to a gradual growth of knowledge based on a sum of facts accumulated by scholars, institutions and other subjects (Haskins, 1965; Seidman, 1987).

In this context of the accumulation of knowledge, basic and applied sciences evolve and converge creating discoveries and path-breaking innovations (Coccia and Wang, 2016; Coccia and Finardi, 2012, 2013; Coccia and Bellitto, 2018; Coccia, 2018b, 2018c; 2020a, 2020b, 2020e, Haskins, 1965). Therefore, discoveries are driven by an activity of accumulation in science and this approach of the evolution of science is irreversible and can never go back (Science, 1965).

Accumulation of knowledge in this specific field of research is measured with total document results (articles, conference papers, conference reviews, book chapters, short surveys, letters, etc.) before the emergence of SARS-CoV-2 in 2019: i.e., from 2005 (first year) to 2018 in Scopus (2021), which is a multidisciplinary database that allows scientometrics analysis to explain characteristics of science and scientific research. Data under study to support the hypothesis are:

- Number of publications using as keywords in search documents of Scopus (2021): bat and SARS-CoV from 2005 to 2018 period
- Journals on which these studies are published
- Affiliations of these publications
- Funding sponsors of publications and scientific research
- Leading countries in these specific studies
- Key papers on these topics and vital subjects

Statistical analyses are performed with the Statistics Software SPSS® version 26.

3. Results

- Hypothesis 1. Origin of SARS-CoV-2 with natural (zoonotic) spillover from bats through an intermediate host

The novel Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) has generated from December 2019 to November 2021 more than 5,210,000 deaths worldwide ([Johns Hopkins University, 2021](#)). This novel coronavirus and COVID-19 pandemic are assumed to be a natural disaster. U.S. Geological Survey (USGS) assesses natural disasters forecasting life loses. USGS (2021) calculates probability estimates for the occurrence of earthquake, hurricane, flood, and tornado disasters with 1,000 fatalities per event in the United States for 1 year exposure times (Table 1).

Table 1. *Forecasting Life Losses with Natural Disasters*

Disaster	1,000 fatalities per event	
	Exposure time	
	1 year Probability of occurrence %	2 years ≈Probability of occurrence %
Earthquakes	1.0	2.0
Hurricanes	6.0	12.0
Floods	0.4	0.8
Tornadoes	0.6	1.2
Arithmetic mean of all disasters	2.0	4.0

The average probability of occurrence of a big natural disaster that generates in 2 years 1,000 fatalities is roughly 4.0%; *mutatis mutandis* a natural disaster that generates over 2 years almost 5,000,000 fatalities is infinitely small (i.e., probability of occurrence is almost 0%) or impossible event. This basic analysis leads to reject the hypothesis of natural spillover of the novel coronavirus from bats to worldwide society.

- Alternative Hypothesis 2. Application of scientific research to coronaviruses for science advances

The second hypothesis is analyzed with the approach of backward chaining:

- November-December 2019. Fact (consequent C): the novel Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) was first identified in the Chinese city of Wuhan in December 2019. At the end of 2019, medical professionals in Wuhan (China) were treating cases of pneumonia that had an unknown source ([Backer et al., 2020](#); [Li et al., 2020](#); [Public Health England, 2020](#); [Riou and Alhaus, 2020](#)). Days later, researchers confirmed that the illnesses (called COVID-19) were caused by a new coronavirus (SARS-CoV-2).

- 2018 year (fact B as antecedent of C): in Wuhan (China) there is a principal laboratory able to support science advances in virology: Wuhan Institute of Virology Chinese Academy of Sciences (CAS), that was founded in 1956. It is the only institute in the country focused exclusively on carrying out fundamental research in general virology. Its research has expanded

from general virology to encompass clinical related virology and research on emerging disease (WIV, 2021). Fact B as antecedent of C is also a main consequent of A.

- 2005-2018 (A is antecedent B). This period, before the emergence of the novel coronavirus, has a lot of scientific research concerning the relationship between bats and SARS-CoV as detected with an in-depth search in Scopus (2021). At global level from 2005 (first year available) to 2018, there are 133 document results in this specific topic. The leading laboratories to perform scientific research on these specific topics are Chinese Academy of Sciences, Wuhan Institute of Virology-Chinese Academy of Sciences and The University of Hong Kong (Figure 2). These studies have been published in international journals, such as Journal of virology, Mbio, Archives of biology and Journal of general Virology (Figure 3). Instead, the most important funding sponsors of these studies are in figure 4, such as National Institute of Allergy and Infectious Diseases, National Institutes of Health, National Natural Science Foundation of China, U.S. Department of Health and Human Services, Chinese Academy of Sciences and Ministry of Science and

Technology of the People's Republic of China. Finally, the proactive countries in the studies on these topics are in figure 5.

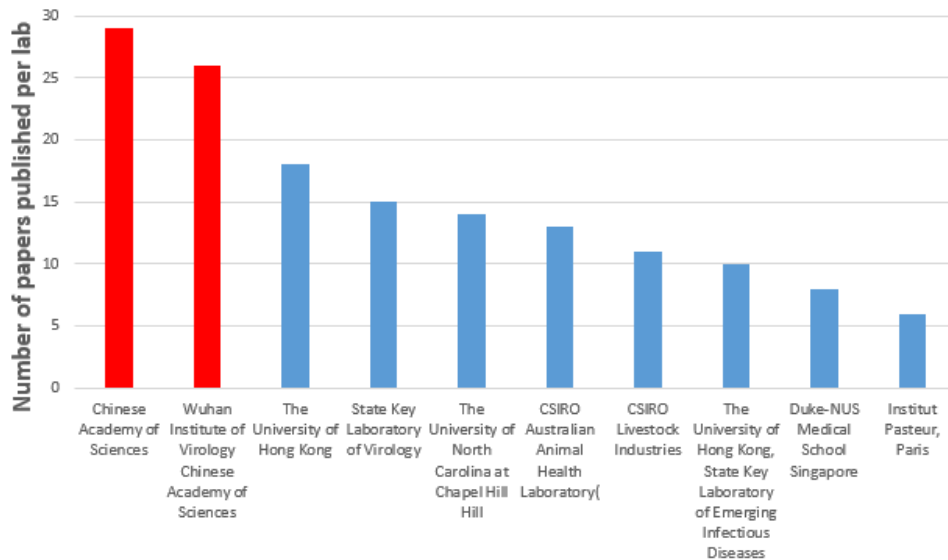


Figure 2. Leading top 10 laboratories in performing scientific research on SARS-CoV from 2005 to 2018.

Journal of Social and Administrative Sciences

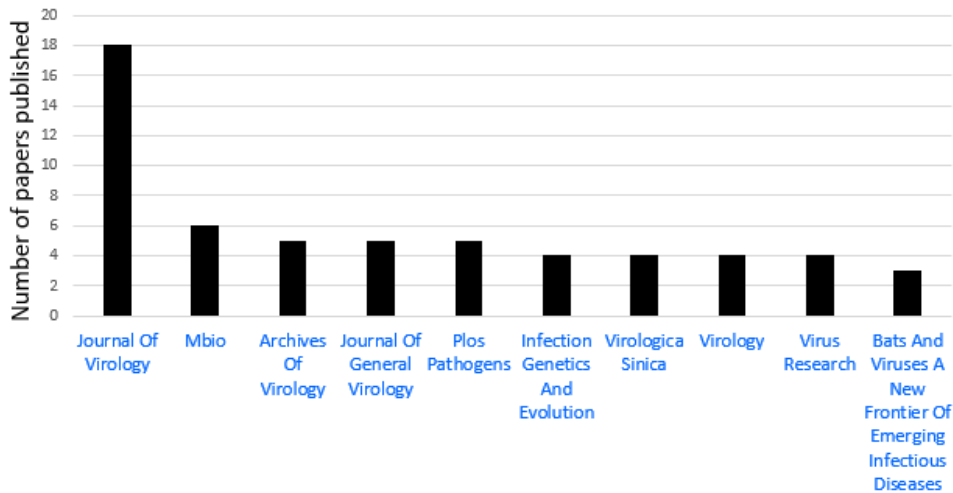


Figure 3. Top 10 journals publishing scientific research on SARS-CoV from 2005 to 2018.

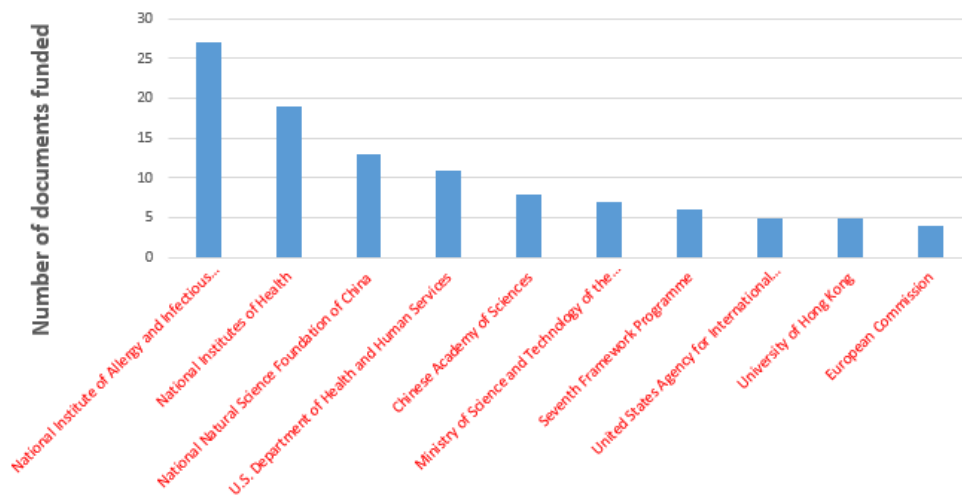


Figure 4. Top 10 funding sponsor of scientific research on SARS-CoV from 2005 to 2018.

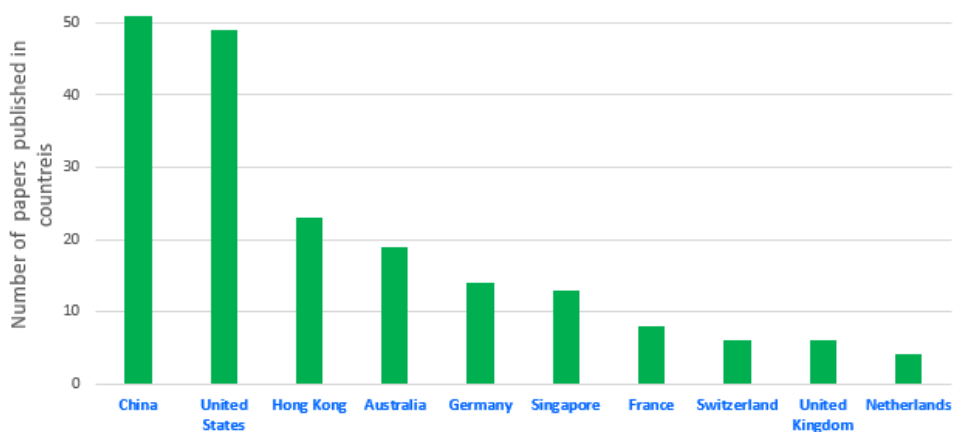


Figure 5. Top 10 countries performing scientific research on SARS-CoV from 2005 to 2018.

A focus on scientific research concerning bat and SARS-CoV over 2005-2018 period in Wuhan Institute of Virology-Chinese Academy of Sciences (Scopus, 2021), which is in the epicenter city of COVID-19 shows that has 26

M. Coccia, JSAS, 9(3), 2022, p.174-190.

Journal of Social and Administrative Sciences

papers published on international journals (e.g., Journal of General Virology, Journal of Virology, Virologica Sinica, Archives of Virology, BMC Evolutionary Biology, Bats and Viruses-A New Frontier of Emerging Infectious Diseases, etc.). The 26 paper represents 20% of total publications over 2005-2018. If we consider also Chinese Academy of Science, having 29 papers, The University of Hong Kong with 28 papers (with State Key Laboratory of Emerging Infectious Diseases) and State Key Laboratory of Virology (Center for Emerging Infectious Diseases, Wuhan Institute of Virology-WIV) with 15, in China there is more than 74% of scientific research in these topics over 2005-2018. These studies at WIV are done mainly in collaboration with Chinese Academy of Sciences, CSIRO Australian Animal Health Laboratory, CSIRO Livestock Industries, Duke-NUS Medical School Singapore, East China Normal University, University of Chinese Academy of Sciences, etc.

Main funding sponsors of these studies are: Ministry of Science and Technology of the People's Republic of China, National Institutes of Health, Chinese Academy of Sciences, National Institute of Allergy and Infectious Diseases, National Natural Science Foundation of China, U.S. Department of Health and Human Services, Fogarty International Center, Commonwealth Scientific and Industrial Research Organisation, European Commission and National Research Foundation Singapore (cf., Appendix A).

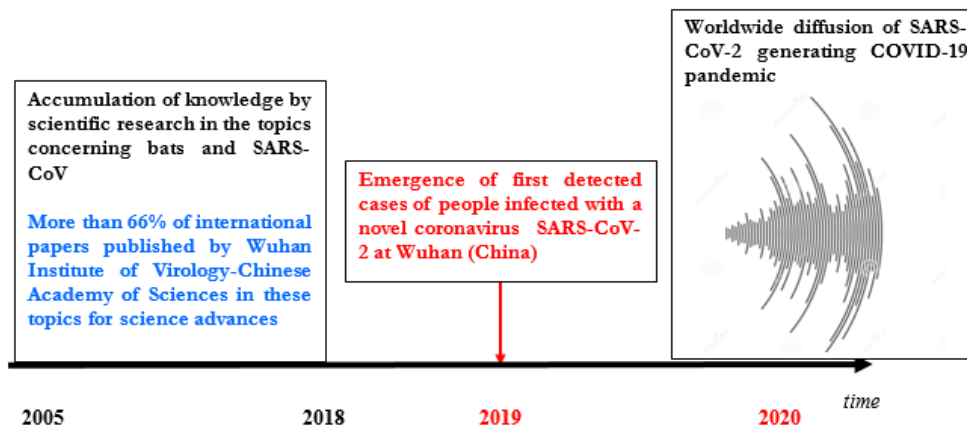


Figure 6. Chronology of events associated with COVID-19 pandemic shock

Overall, the leading role worldwide of Wuhan Institute of Virology Chinese Academy of Sciences in the research on bat and SARS-CoV to produce science advances suggests that this laboratory has accumulated technical knowledge and know-how over 2005-2018 period to support a gradual growth of knowledge in fundamental research in this field of research (Figure 6). Hence, this backward reasoning seems to support the hypothesis that the novel coronavirus may be originated with a process of accumulation of knowledge (2005-2018=13 years) in a specific place over time and space with research directed to science advances.

4. Discussions

Relman (2020) argues that: "A deliberative process for investigating the origins of this pandemic must be representative of all relevant disciplines, expertise, and stakeholders; must achieve political neutrality, scientific balance, and access to all relevant information and samples; and must operate with transparency and independent oversight. Without these features, it will not be credible, trustworthy, or effective". In this context, the findings of the study here suggest that natural spillover of the novel coronavirus is a rare or impossible event, whereas the creation of a novel coronavirus with a research activity to produce science advances is a reasonable hypothesis because of the accumulation of knowledge in this specific research filed in the principal laboratories over 2005-2018 period. In particular, the hypothesis of a (natural) zoonotic spillover from bats, through an intermediate host to humans, is a rare event because some scholars wrongly compare nature to an engineer. Jacobs (1977) argues that this is a misleading comparison because unlike natural evolution, the engineers work with a conceived plan to achieve goals (products) with their endeavors. Moreover, engineers work to produce a new product using specific materials and equipment designed to achieve the task. Relman (2020) argue that the explanation of the origin of SARS-CoV plays a vital role in forecasting future pandemics. If the hypothesis of natural spillover from bats is true with strong evidence of the casual event of SARS CoV- 2 passing directly from bat to human, or by an intermediate host, then efforts of prevention must be directed to improve the management of the interactions between bats (and in general wildlife) and human (cf., [Latinne et al., 2020](#)). Daszak et al. (2020) argue that to prevent the next epidemic and pandemic like COVID-19, research and investment of nations should focus on:

- 1) surveillance among wildlife to identify the high-risk pathogens they carry
- 2) surveillance among people who have contact with wildlife to identify early spillover events
- 3) improvement of market biosecurity regarding the wildlife trade.

However, if the novel coronavirus is created by scholars by research for science advances and then "SARSCoV-2 escaped from a lab" ([Relman, 2020](#)) causing a pandemic crisis, critical aspects of prevention are the improvement of biosecurity in laboratory testing of hazardous pathogens. A further comparative meta-analysis of two hypotheses shows that natural spillover of SARSCoV-2 (that is generating high numbers of COVID-19 related infections and deaths in two years) is almost an impossible event (using the analogy with the probability of occurrence of other natural disasters; cf., [USGS, 2021](#)), whereas manifold studies shows that there have been several high-profile accidents in research laboratories worldwide ([Ménard and Trant, 2020](#)). In fact, Hellman et al. (1986) examining almost 600 accidents between 1966 and 1984, found that 13% of accidents occurred in research labs and 2% in fabrication rooms. Van Noorden (2013), with a survey from Nature and UCLA of about 2,500 scientists, reveals that 30% of interviewed

M. Coccia, JSAS, 9(3), 2022, p.174-190.

reported having witnessed a severe lab injury. Another study in Canadian chemistry and biology labs reports that 15% of scholars surveyed had at least one injury (Ayi and Hon, 2018). Simmons et al. (2018) found that lab accidents represented 18.4% of the total incidents reported at Iowa State university. In fact, the support of the hypothesis of human factors in the origins of SARS-CoV-2, and likely accident lab and consequential diffusion in society, leads to basic aspects of improving the technical guidelines at all levels for biosafety of laboratories conducting testing of hazardous pathogens similar to SARS-CoV-2 that generate pandemic crisis (Figure 7).

Hypothesis 1	Hypothesis 2
<i>Natural spillover from bats through an intermediate host of the novel coronavirus</i>	<i>Process of research for science advances and lab accident with consequential spatial-temporal diffusion</i>
Probability of occurrence of a natural event generation with more than 5,200,000 deaths	Probability of a lab accident
≈ 0 %	≈ 13-30 %

Figure 7. Comparative probability of proposed hypotheses based on a meta-analysis.

Ménard and Trant (2020, p. 18) maintain that factors determining lab accidents can be due to: "risks associated with the materials or equipment being used, risks related to the skills, knowledge and choices of the research personnel doing the study, characteristics or qualities of the PI and the research lab in which the research is occurring and risk factors arising from the departmental or institutional level".

Hence, a priori the epidemic/pandemic, in responding to a constant pandemic threat of novel viral agents in future, the international communities must reinforce surveillance and proper biosafety procedures in public and private institutes of virology that study viruses and new viruses to avoid that may be accidentally spread in surrounding environments with damages for population and vegetation. In this context, international collaboration among scientists is a basic aspect to address these risks and support decisions of policymakers to prevent lab accidents and threats for future pandemics that create huge socioeconomic issues worldwide (National Health Commission of The People's Republic of China , 2020). Yuan et al. (2020) argue that in China, information of lab safety should be internally linked to the national intelligent syndromic surveillance system, which could help different levels of organizations to better coordinate and allocate resources for targeted investigations and interventions to improve the biosafety of labs at the greatest need and facilitate more comprehensive surveillance of risk for disease outbreak (Jia and Yang, 2020).

A posteriori in the initial phase of the epidemic/pandemic, prevention and preparedness of pandemic threats have to directed to design and implement strategic actions given by improvements of the early warning systems in the international community using existing infrastructure to ensure rapid detection of suspected cases in humans based on reliable international laboratories that receive all the data and clinical specimens needed for an

accurate evaluation of an emergence of pandemic risk for applying timely containment operations at local and global level (Coccia, 2021d, 2022).

5. Conclusions

The origins of novel viral agents associated with future epidemics/pandemics pose, more and more, serious questions and policy responses to security, biosecurity and public health of nations (Relman, 2020). A pandemic like COVID-19 can occur at any time with little warning; any delay in detecting and sharing novel virus samples; and in developing, producing, distributing, or administering a therapeutic or vaccine could result in significant additional morbidity and mortality, and deterioration of socioeconomic systems (Coccia, 2020c, 2021a, 2021e, 2021i; Huang et al., 2021). The findings of the study here suggest that natural spillover of the novel coronavirus is a rare or impossible event, whereas the creation of a viral agent with a research activity to produce science advances is a reasonable hypothesis because of the accumulation of knowledge in specific research fields of principal laboratories worldwide. Although this study has provided interesting results about the origin of SARS-CoV-2, that are of course tentative, it has several limitations. First, a limitation of the study is the lack of data about specific scientific activity of laboratory testing for hazardous pathogens also because of information that are classified for national security. Second, not all possible confounding factors that affect the origins of this novel coronavirus are taken into consideration and in future these factors deserve to be analyzed for supporting results here. Third, the lack of integration of data to find parents additional genome sequences of coronaviruses and measurements of SARS-CoV-2 evolution under a variety of defined conditions. Future research should consider these aspects, new data, when available, and when possible, to examine also other factors associated with the origins of this novel coronavirus. Despite these limitations, the results presented here suggests the critical aspect of the accumulation of scientific knowledge that is a main factor to support science advances in the field of virology for the creation of novel viral agents (Wu et al., 2016; Zhang and Holmes, 2020). However, there is need for much more detailed research in these topics and this study encourages further investigations that should be collaborative between scholars of different disciplines and nations to have access to relevant information and to design appropriate policy responses to prevent similar pandemic both if the novel coronavirus has a natural spillover from wildlife and if it is due to scientific research for science advance and consequential laboratory accident.

Overall, then, different factors of the origins of SARS-CoV-2 are not only related to medicine but also to other social, political and economic aspects, as well as leadership in international system, play a critical role to clarify the truth and to improve the preparedness of countries to prevent similar pandemic or to control negative impact of pandemic crisis on public health, economy and society (cf., Coccia, 2019, 2020a; Coccia, 2021f; 2021h). To conclude, Relman (2020) argues that: "A deliberative process for

M. Coccia, JSAS, 9(3), 2022, p.174-190.

Journal of Social and Administrative Sciences

investigating the origins of this pandemic must be representative of all relevant disciplines, expertise, and stakeholders; must achieve political neutrality, scientific balance, and access to all relevant information and samples; and must operate with transparency and independent oversight... A more complete understanding of the origins of COVID-19 clearly serves the interests of every person in every country on this planet. ... it will lead to more effective responses to this pandemic, as well as efforts to anticipate and prevent the next one. It will also advance our discussions about risky science".

Appendix

Publications concerning bats and SARS-CoV from 2015 to 2018 period

- Ge, X. Y., Wang, N., Zhang, W., Hu, B., Li, B., Zhang, Y. Z., Zhou, J. H., Luo, C. M., Yang, X. L., Wu, L. J., Wang, B., Zhang, Y., Li, Z. X., & Shi, Z. L. (2016). Coexistence of multiple coronaviruses in several bat colonies in an abandoned mineshaft. *Virologica Sinica*, 31(1), 31–40. <https://doi.org/10.1007/s12250-016-3713-9>
- Ge, X. Y., Yang, W. H., Zhou, J. H., Li, B., Zhang, W., Shi, Z. L., & Zhang, Y. Z. (2017). Detection of alpha- and betacoronaviruses in rodents from Yunnan, China. *Virology journal*, 14(1), 98. <https://doi.org/10.1186/s12985-017-0766-9>
- Hu, B., Zeng, L. P., Yang, X. L., Ge, X. Y., Zhang, W., Li, B., Xie, J. Z., Shen, X. R., Zhang, Y. Z., Wang, N., Luo, D. S., Zheng, X. S., Wang, M. N., Daszak, P., Wang, L. F., Cui, J., & Shi, Z. L. (2017). Discovery of a rich gene pool of bat SARS-related coronaviruses provides new insights into the origin of SARS coronavirus. *PLoS pathogens*, 13(11), e1006698. <https://doi.org/10.1371/journal.ppat.1006698>
- Lau, S. K., & Chan, J. F. (2015). Coronaviruses: emerging and re-emerging pathogens in humans and animals. *Virology journal*, 12, 209. <https://doi.org/10.1186/s12985-015-0432-z>
- Liang, J., Yang, X. L., Li, B., Liu, Q., Zhang, Q., Liu, H., Kan, H. P., Wong, K. C., Chek, S. N., He, X., Peng, X., Shi, Z. L., Wu, Y., & Zhang, L. (2017). Detection of diverse viruses in alimentary specimens of bats in Macau. *Virologica Sinica*, 32(3), 226–234. <https://doi.org/10.1007/s12250-017-3976-9>
- Luo, C. M., Wang, N., Yang, X. L., Liu, H. Z., Zhang, W., Li, B., Hu, B., Peng, C., Geng, Q. B., Zhu, G. J., Li, F., & Shi, Z. L. (2018). Discovery of Novel Bat Coronaviruses in South China That Use the Same Receptor as Middle East Respiratory Syndrome Coronavirus. *Journal of virology*, 92(13), e00116-18. <https://doi.org/10.1128/JVI.00116-18>
- Luo, Y., Li, B., Jiang, R. D., Hu, B. J., Luo, D. S., Zhu, G. J., Hu, B., Liu, H. Z., Zhang, Y. Z., Yang, X. L., & Shi, Z. L. (2018). Longitudinal Surveillance of Betacoronaviruses in Fruit Bats in Yunnan Province, China During 2009-2016. *Virologica Sinica*, 33(1), 87–95. <https://doi.org/10.1007/s12250-018-0017-2>
- Menachery, V. D., Yount, B. L., Jr, Debbink, K., Agnihothram, S., Gralinski, L. E., Plante, J. A., Graham, R. L., Scobey, T., Ge, X. Y., Donaldson, E. F., Randell, S. H., Lanzavecchia, A., Marasco, W. A., Shi, Z. L., & Baric, R. S. (2015). A SARS-like cluster of circulating bat coronaviruses shows potential for human emergence. *Nature medicine*, 21(12), 1508–1513. <https://doi.org/10.1038/nm.3985>
- Wang, M. N., Zhang, W., Gao, Y. T., Hu, B., Ge, X. Y., Yang, X. L., Zhang, Y. Z., & Shi, Z. L. (2016). Longitudinal surveillance of SARS-like coronaviruses in bats by quantitative real-time PCR. *Virologica Sinica*, 31(1), 78–80. <https://doi.org/10.1007/s12250-015-3703-3>
- Wang, N., Li, S. Y., Yang, X. L., Huang, H. M., Zhang, Y. J., Guo, H., Luo, C. M., Miller, M., Zhu, G., Chmura, A. A., Hagan, E., Zhou, J. H., Zhang, Y. Z., Wang, L. F., Daszak, P., & Shi, Z. L. (2018). Serological Evidence of Bat SARS-Related Coronavirus Infection in Humans, China. *Virologica Sinica*, 33(1), 104–107. <https://doi.org/10.1007/s12250-018-0012-7>
- Waruhiu, C., Ommeh, S., Obanda, V., Agwanda, B., Gakuya, F., Ge, X. Y., Yang, X. L., Wu, L. J., Zohaib, A., Hu, B., & Shi, Z. L. (2017). Molecular detection of viruses in Kenyan bats and discovery of novel astroviruses, caliciviruses and rotaviruses. *Virologica Sinica*, 32(2), 101–114. <https://doi.org/10.1007/s12250-016-3930-2>
- Yang, X. L., Hu, B., Wang, B., Wang, M. N., Zhang, Q., Zhang, W., Wu, L. J., Ge, X. Y., Zhang, Y. Z., Daszak, P., Wang, L. F., & Shi, Z. L. (2015). Isolation and Characterization of a Novel Bat Coronavirus Closely Related to the Direct Progenitor of Severe Acute Respiratory Syndrome Coronavirus. *Journal of virology*, 90(6), 3253–3256. <https://doi.org/10.1128/JVI.02582-15>
- Zeng, L. P., Gao, Y. T., Ge, X. Y., Zhang, Q., Peng, C., Yang, X. L., Tan, B., Chen, J., Chmura, A. A., Daszak, P., & Shi, Z. L. (2016). Bat Severe Acute Respiratory Syndrome-Like Coronavirus WIV1 Encodes an Extra Accessory Protein, ORFX, Involved in Modulation of the Host Immune Response. *Journal of virology*, 90(14), 6573–6582. <https://doi.org/10.1128/JVI.03079-15>
- Zeng, L. P., Ge, X. Y., Peng, C., Tai, W., Jiang, S., Du, L., & Shi, Z. L. (2017). Cross-neutralization of SARS coronavirus-specific antibodies against bat SARS-like coronaviruses. *Science China. Life sciences*, 60(12), 1399–1402. <https://doi.org/10.1007/s11427-017-9189-3>
- Zhang, H., Peng, C., Liu, B., (...), Yuan, Z., Shi, Z. (2018) Evaluation of MICRO-CHEM PLUS as a Disinfectant for Biosafety Level 4 Laboratory in China, *Applied Biosafety* 23(1), pp. 32-38
- Zhou, P., Fan, H., Lan, T., Yang, X. L., Shi, W. F., Zhang, W., Zhu, Y., Zhang, Y. W., Xie, Q. M., Mani, S., Zheng, X. S., Li, B., Li, J. M., Guo, H., Pei, G. Q., An, X. P., Chen, J. W., Zhou, L., Mai, K. J., Wu, Z. X., ... Ma, J. Y. (2018). Fatal swine acute diarrhoea syndrome caused by an HKU2-related coronavirus of bat origin. *Nature*, 556(7700), 255–258. <https://doi.org/10.1038/s41586-018-0010-9>

References

- Anand, U., Cabrerros, C., Mal, J., Ballesteros, F., Jr, Sillanpää, M., Tripathi, V., Bontempi, E. 2021. Novel coronavirus disease 2019 (COVID-19) pandemic: From transmission to control with an interdisciplinary vision. *Environmental research*, 197, 111126. [10.1016/j.envres.2021.111126](https://doi.org/10.1016/j.envres.2021.111126)
- Andersen K. G., A. Rambaut, W. I. Lipkin, E. C. Holmes, R. F. Garry, 2020. The proximal origin of SARS-CoV-2. *Nat. Med.* 26, 450–452 (2020).
- Ardito L., Coccia M., Messeni Petruzzelli A. 2021. Technological exaptation and crisis management: Evidence from COVID-19 outbreaks. *R&D Management*, vol. 51, n. 4, pp. 381-392. Special Issue: Providing solutions in emergencies: R&D and innovation management during Covid-19 Part-2, September 2021, [10.1111/radm.12455](https://doi.org/10.1111/radm.12455)
- Ayi, H.-R., Hon, C.-Y. 2018. Safety culture and safety compliance in academic laboratories: A Canadian perspective. *J. Chem. Health Saf.* 25, 6–12.
- Backer J.A., Klinkenberg D., Wallinga J. 2020. Incubation period of 2019 novel coronavirus (2019-nCoV) infections among travellers from Wuhan, China, 20–28 January 2020. *Euro Surveill* 2020;25: 2000062.
- Bloom, J. D., Chan, Y. A., Baric, R. S., Bjorkman, P. J., Cobey, S., Deverman, B. E., Fisman, D. N., Gupta, R., Iwasaki, A., Lipsitch, M., Medzhitov, R., Neher, R. A., Nielsen, R., Patterson, N., Stearns, T., van Nimwegen, E., Worobey, M., & Relman, D. A. (2021). Investigate the origins of COVID-19. *Science (New York, N.Y.)*, 372(6543), 694. [10.1126/science.abj0016](https://doi.org/10.1126/science.abj0016)
- Boni, M.F., Lemey, P., Jiang, X. et al. 2020. Evolutionary origins of the SARS-CoV-2 sarbecovirus lineage responsible for the COVID-19 pandemic. *Nat Microbiol* 5, 1408–1417 (2020). [10.1038/s41564-020-0771-4](https://doi.org/10.1038/s41564-020-0771-4)
- Bontempi E., Coccia M., 2021. International trade as critical parameter of COVID-19 spread that outclasses demographic, economic, environmental, and pollution factors, *Environmental Research*, vol. 201, Article number 111514, [10.1016/j.envres.2021.111514](https://doi.org/10.1016/j.envres.2021.111514)
- Bontempi E., Coccia M., Vergalli S., Zanoletti A. 2021. Can commercial trade represent the main indicator of the COVID-19 diffusion due to human-to-human interactions? A comparative analysis between Italy, France, and Spain, *Environmental Research*, vol. 201, Article number 111529, [10.1016/j.envres.2021.111529](https://doi.org/10.1016/j.envres.2021.111529)
- Calabrese G., Coccia M., Rolfo S. 2005. Strategy and market management of new product development: evidence from Italian SMEs. *International Journal of Product Development*, vol. 2, n. 1-2, pp. 170-189. [10.1504/IJPD.2005.006675](https://doi.org/10.1504/IJPD.2005.006675)
- Coccia, M. (2003). Metrics of R&D performance and management of public research institute, *Proceedings of IEEE- IEMC 03, Piscataway*, pp. 231-236.
- Coccia, M. (2005). A taxonomy of public research bodies: a systemic approach, *Prometheus*, 23(1), 63-82. doi. [10.1080/0810902042000331322](https://doi.org/10.1080/0810902042000331322)
- Coccia, M. (2005a). Countrymetrics: valutazione della performance economica e tecnologica dei paesi e posizionamento dell'Italia, *Rivista Internazionale di Scienze Sociali*, 113(3), 377-412.
- Coccia, M. (2008). Measuring scientific performance of public research units for strategic change. *Journal of Informetrics*, 2(3), 183-194. doi. [10.1016/j.joi.2008.04.001](https://doi.org/10.1016/j.joi.2008.04.001)
- Coccia, M. (2013). Population and technological innovation: the optimal interaction across modern countries, *Working Paper Ceris del Consiglio Nazionale delle Ricerche*, vol.15, n.7.
- Coccia, M. (2014). Steel market and global trends of leading geo-economic players. *International Journal of Trade and Global Markets*, 7(1), 36-52. doi. [10.1504/IJTGM.2014.058714](https://doi.org/10.1504/IJTGM.2014.058714)
- Coccia, M. (2015). Spatial relation between geo-climate zones and technological outputs to explain the evolution of technology. *Int. J. Transitions and Innovation Systems*, 4(1), 5-21. doi. [10.1504/IJTIS.2015.074642](https://doi.org/10.1504/IJTIS.2015.074642)
- Coccia, M. (2016). Problem-driven innovations in drug discovery: co-evolution of the patterns of radical innovation with the evolution of problems, *Health Policy and Technology*, 5(2), 143-155. doi. [10.1016/j.hlpt.2016.02.003](https://doi.org/10.1016/j.hlpt.2016.02.003)
- Coccia, M. (2017). Varieties of capitalism's theory of innovation and a conceptual integration with leadership-oriented executives: the relation between typologies of executive,

Journal of Social and Administrative Sciences

- technological and socioeconomic performances. *Int. J. Public Sector Performance Management*, 3(2), 148–168. doi. [10.1504/IJPSPM.2017.084672](https://doi.org/10.1504/IJPSPM.2017.084672)
- Coccia, M. (2017a). Disruptive firms and industrial change, *Journal of Economic and Social Thought*, 4(4), 437-450. doi. [10.1453/jest.v4i4.1511](https://doi.org/10.1453/jest.v4i4.1511)
- Coccia, M. (2017b). New directions in measurement of economic growth, development and under development, *Journal of Economics and Political Economy*, 4(4), 382-395. doi. [10.1453/jepe.v4i4.1533](https://doi.org/10.1453/jepe.v4i4.1533)
- Coccia, M. (2017c). Sources of disruptive technologies for industrial change. *L'industria – Rivista di Economia e Politica Industriale*, 38(1), 97-120. doi. [10.1430/87140](https://doi.org/10.1430/87140)
- Coccia, M. (2017d). Sources of technological innovation: Radical and incremental innovation problem-driven to support competitive advantage of firms. *Technology Analysis & Strategic Management*, 29(9), 1048-1061. doi. [10.1080/09537325.2016.1268682](https://doi.org/10.1080/09537325.2016.1268682)
- Coccia, M. (2018). An introduction to the methods of inquiry in social sciences, *Journal of Social and Administrative Sciences*, 5(2), 116-126. doi. [10.1453/jsas.v5i2.1651](https://doi.org/10.1453/jsas.v5i2.1651)
- Coccia, M. (2018a). An introduction to the theories of institutional change, *Journal of Economics Library*, 5(4), 337-344. doi. [10.1453/jel.v5i4.1788](https://doi.org/10.1453/jel.v5i4.1788)
- Coccia, M. (2018b). General properties of the evolution of research fields: a scientometric study of human microbiome, evolutionary robotics and astrobiology, *Scientometrics*, 117(2), 1265-1283. doi. [10.1007/s11192-018-2902-8](https://doi.org/10.1007/s11192-018-2902-8)
- Coccia, M. (2018c). The origins of the economics of Innovation, *Journal of Economic and Social Thought*, 5(1), 9-28. doi. [10.1453/jest.v5i1.1574](https://doi.org/10.1453/jest.v5i1.1574)
- Coccia, M. (2018d). The relation between terrorism and high population growth, *Journal of Economics and Political Economy*, 5(1), 84-104. doi. [10.1453/jepe.v5i1.1575](https://doi.org/10.1453/jepe.v5i1.1575)
- Coccia, M. (2018e). Classification of innovation considering technological interaction, *Journal of Economics Bibliography*, 5(2), 76-93. doi. [10.1453/jeb.v5i2.1650](https://doi.org/10.1453/jeb.v5i2.1650)
- Coccia, M. (2018f). An introduction to the theories of national and regional economic development, *Turkish Economic Review*, 5(4), 350-358. doi. [10.1453/ter.v5i4.1794](https://doi.org/10.1453/ter.v5i4.1794)
- Coccia, M. (2019). Metabolism of public organizations: A case study, *Journal of Social and Administrative Sciences*, 6(1), 1-9. doi. [10.1453/jsas.v6i1.1793](https://doi.org/10.1453/jsas.v6i1.1793)
- Coccia, M. (2019a). The theory of technological parasitism for the measurement of the evolution of technology and technological forecasting, *Technological Forecasting and Social Change*, 141, 289-304. doi. [10.1016/j.techfore.2018.12.012](https://doi.org/10.1016/j.techfore.2018.12.012)
- Coccia, M. (2019b). A Theory of classification and evolution of technologies within a Generalized Darwinism, *Technology Analysis & Strategic Management*, 31(5), 517-531. doi. [10.1080/09537325.2018.1523385](https://doi.org/10.1080/09537325.2018.1523385)
- Coccia, M. (2019). Theories and the reasons for war: a survey. *Journal of Economic and Social Thought*, 6(2), 115-124. doi. [10.1453/jest.v6i2.1890](https://doi.org/10.1453/jest.v6i2.1890)
- Coccia, M. (2020a). Factors determining the diffusion of COVID-19 and suggested strategy to prevent future accelerated viral infectivity similar to COVID. *Science of The Total Environment*, 729, n.138474. doi. [10.1016/j.scitotenv.2020.138474](https://doi.org/10.1016/j.scitotenv.2020.138474)
- Coccia, M. (2020b). How (Un)sustainable Environments are Related to the Diffusion of COVID-19: The Relation between Coronavirus Disease 2019, Air Pollution, *Wind Resource and Energy. Sustainability*, 12, 9709. doi. [10.3390/su12229709](https://doi.org/10.3390/su12229709)
- Coccia, M. (2020c). How do environmental, demographic, and geographical factors influence the spread of COVID-19. *Journal of Social and Administrative Sciences*, 7(3), 169-209. doi. [10.1453/jsas.v7i3.2018](https://doi.org/10.1453/jsas.v7i3.2018)
- Coccia, M. (2020d). Destructive Technologies for Industrial and Corporate Change. In: Farazmand A. (eds), *Global Encyclopedia of Public Administration, Public Policy, and Governance*. Springer, Cham. doi. [10.1007/978-3-319-31816-5_3972-1](https://doi.org/10.1007/978-3-319-31816-5_3972-1)
- Coccia, M. (2020e). Deep learning technology for improving cancer care in society: New directions in cancer imaging driven by artificial intelligence. *Technology in Society*, 60, 1-11, art. no.101198. doi. [10.1016/j.techsoc.2019.101198](https://doi.org/10.1016/j.techsoc.2019.101198)
- Coccia, M. (2020f). How does science advance? Theories of the evolution of science. *Journal of Economic and Social Thought*, 7(3), 153-180. doi. [10.1453/jest.v7i3.2111](https://doi.org/10.1453/jest.v7i3.2111)

Journal of Social and Administrative Sciences

- Coccia, M. (2020g). The evolution of scientific disciplines in applied sciences: dynamics and empirical properties of experimental physics, *Scientometrics*, 124, 451-487. doi. [10.1007/s11192-020-03464-y](https://doi.org/10.1007/s11192-020-03464-y)
- Coccia, M. (2020h). Multiple working hypotheses for technology analysis, *Journal of Economics Bibliography*, 7(2), 111-126. doi. [10.1453/jeb.v7i2.2050](https://doi.org/10.1453/jeb.v7i2.2050)
- Coccia, M. (2020i). Asymmetry of the technological cycle of disruptive innovations. *Technology Analysis & Strategic Management*, 32(12), 1462-1477. doi. [10.1080/09537325.2020.1785415](https://doi.org/10.1080/09537325.2020.1785415)
- Coccia, M., Bellitto, M. (2018). Human progress and its socioeconomic effects in society, *Journal of Economic and Social Thought*, 5(2), 160-178. doi. [10.1453/jest.v5i2.1649](https://doi.org/10.1453/jest.v5i2.1649)
- Coccia, M., Benati, I. (2018). Rewards in public administration: A proposed classification, *Journal of Social and Administrative Sciences*, 5(2), 68-80. doi. [10.1453/jsas.v5i2.1648](https://doi.org/10.1453/jsas.v5i2.1648)
- Coccia, M., Benati, I. (2018a). Comparative Models of Inquiry, A. Farazmand (ed.), *Global Encyclopedia of Public Administration, Public Policy, and Governance*, Springer International Publishing AG, part of Springer Nature. doi. [10.1007/978-3-319-31816-5_1199-1](https://doi.org/10.1007/978-3-319-31816-5_1199-1)
- Coccia, M., Cadario, E. (2014). Organisational (un)learning of public research labs in turbulent context. *International Journal of Innovation and Learning*, 15(2), 115-129. doi. [10.1504/IJIL.2014.059756](https://doi.org/10.1504/IJIL.2014.059756)
- Coccia, M., Finardi, U. (2012). Emerging nanotechnological research for future pathway of biomedicine. *International Journal of Biomedical nanoscience and nanotechnology*, 2(3-4), 299-317. doi. [10.1504/IJBNN.2012.051223](https://doi.org/10.1504/IJBNN.2012.051223)
- Coccia, M., Finardi, U. (2013). New technological trajectories of non-thermal plasma technology in medicine. *Int. J. Biomedical Engineering and Technology*, 11(4), 337-356. doi. [10.1504/IJBET.2013.055665](https://doi.org/10.1504/IJBET.2013.055665)
- Coccia, M., Rolfo, S. (2000). Ricerca pubblica e trasferimento tecnologico: il caso della regione Piemonte in Rolfo S. (eds) *Innovazione e piccole imprese in Piemonte*, Franco Angeli Editore, Milano (Italy).
- Coccia, M., Rolfo, S. (2008). Strategic change of public research units in their scientific activity, *Technovation*, 28(8), 485-494. doi. [10.1016/j.technovation.2008.02.005](https://doi.org/10.1016/j.technovation.2008.02.005)
- Coccia, M., Wang, L. (2015). Path-breaking directions of nanotechnology-based chemotherapy and molecular cancer therapy, *Technological Forecasting & Social Change*, 94(1), 155-169. doi. [10.1016/j.techfore.2014.09.007](https://doi.org/10.1016/j.techfore.2014.09.007)
- Coccia, M., Wang, L. (2016). Evolution and convergence of the patterns of international scientific collaboration, *Proceedings of the National Academy of Sciences of the United States of America*, 113(8), 2057-2061. doi. [10.1073/pnas.1510820113](https://doi.org/10.1073/pnas.1510820113)
- Coccia, M., Watts, J. (2020). A theory of the evolution of technology: technological parasitism and the implications for innovation management, *Journal of Engineering and Technology Management*, 55(2020), 101552. doi. [10.1016/j.jengtecman.2019.11.003](https://doi.org/10.1016/j.jengtecman.2019.11.003)
- Daszak P., Olival K. J., Li H. 2020. A strategy to prevent future epidemics similar to the 2019-nCoV outbreak, *Biosafety and Health*, [10.1016/j.bsheal.2020.01.003](https://doi.org/10.1016/j.bsheal.2020.01.003)
- ECDC 2021. SARS-CoV-2 variants of concern as of 3 June 2021. European Centre for Disease Prevention and Control, <https://www.ecdc.europa.eu/en/covid-19/variants-concern> (accessed on 10 June 2021).
- Frutos, R., Gavotte, L., Devaux, C. A. (2021). Understanding the origin of COVID-19 requires to change the paradigm on zoonotic emergence from the spillover to the circulation model. *Infection, genetics and evolution : journal of molecular epidemiology and evolutionary genetics in infectious diseases*, 95, 104812. [10.1016/j.meegid.2021.104812](https://doi.org/10.1016/j.meegid.2021.104812)
- Frutos, R., Javelle, E., Barberot, C., Gavotte, L., Tissot-Dupont, H., Devaux, C. A. (2022). Origin of COVID-19: Dismissing the Mojiang mine theory and the laboratory accident narrative. *Environmental research*, 204, 112141. Advance online publication. [10.1016/j.envres.2021.112141](https://doi.org/10.1016/j.envres.2021.112141)
- Haskins C. P. 1965. Report of the President by Carnegie Institution of Washington Yearbook 63, 1963-64, Washington, D.C. (USA)
- Hellman, M. A., Savage, E. P. & Keefe, T. J. 1986. Epidemiology of accidents in academic chemistry laboratories. Part 1. Accident data survey. *J. Chem. Educ.* 63, A267.

Journal of Social and Administrative Sciences

- Huang J., Xiaoyue Liu, Li Zhang, Yingjie Zhao, Danfeng Wang, Jinfeng Gao, Xinbo Lian, Chuwei Liu. 2021. The oscillation-outbreaks characteristic of the COVID-19 pandemic, *National Science Review*, vol. 8, n. 8, , nwab100, [10.1093/nsr/nwab100](https://doi.org/10.1093/nsr/nwab100)
- Jacob F. (1977). Evolution and tinkering. *Science (New York, N.Y.)*, 196(4295), 1161–1166. [10.1126/science.860134](https://doi.org/10.1126/science.860134)
- Jia, Peng, Yang, Shujuan, 2020. China needs a national intelligent syndromic surveillance system. *Nature Med.* 26, 990. [10.1038/s41591-020-0921-5](https://doi.org/10.1038/s41591-020-0921-5)
- Johns Hopkins Center for System Science and Engineering, 2021. Coronavirus COVID-19 Global Cases, [Retrieved from] (accessed on 4 October 2021).
- Johns Hopkins University 2021. COVID-19 Dashboard by the Center for Systems Science and Engineering (CSSE) at Johns Hopkins University (JHU)". Retrieved 28 October 2021.
- Latinne A. et al., 2020. Origin and cross-species transmission of bat coronaviruses in China. *Nat. Commun.* 11, 4235.
- Li Q., Guan X., Wu P., et al. 2020. Early transmission dynamics in Wuhan, China, of novel coronavirus-infected pneumonia. *N Engl J Med* 2020; published online Jan 29. Doi. [10.1056/NEJMoa2001316](https://doi.org/10.1056/NEJMoa2001316)
- Ménard, A. D., & Trant, J. F. (2020). A review and critique of academic lab safety research. *Nature chemistry*, 12(1), 17–25. [10.1038/s41557-019-0375-x](https://doi.org/10.1038/s41557-019-0375-x)
- National Health Commission of The People's Republic of China, & Bureau of Disease Prevention and Control, National Health Commission of People's Republic of China chinabiosafetyandhealth@ivdc.chinacdc.cn (2020). Technical guidance for laboratory testing of 2019-nCoV infection (Third Edition). *Biosafety and health*, 2(1), 3–5. [10.1016/j.bsheal.2020.02.001](https://doi.org/10.1016/j.bsheal.2020.02.001)
- Pagliaro M., Coccia M. 2021. How self-determination of scholars outclasses shrinking public research lab budgets, supporting scientific production: a case study and R&D management implications. *Heliyon*. vol. 7, n. 1, e05998. [10.1016/j.heliyon.2021.e05998](https://doi.org/10.1016/j.heliyon.2021.e05998)
- Public Health England, 2020. Novel coronavirus (2019-nCoV) – what you need to know. 2020. [Retrieved from] (accessed Jan 31, 2020).
- Rahalkar, M. C., & Bahulikar, R. A. (2020). Lethal Pneumonia Cases in Mojiang Miners (2012) and the Mineshaft Could Provide Important Clues to the Origin of SARS-CoV-2. *Frontiers in public health*, 8, 581569. [10.3389/fpubh.2020.581569](https://doi.org/10.3389/fpubh.2020.581569)
- Relman D. A. (2020). Opinion: To stop the next pandemic, we need to unravel the origins of COVID-19. *Proceedings of the National Academy of Sciences of the United States of America*, 117(47), 29246–29248. [10.1073/pnas.2021133117](https://doi.org/10.1073/pnas.2021133117)
- Riou J., Althaus C.L. 2020. Pattern of early human-to-human transmission of Wuhan 2019 novel coronavirus (2019-nCoV), December 2019 to January 2020. *Euro Surveill* 2020; 25: 2000058.
- Russell Stuart J., Norvig Peter 2010. *Artificial Intelligence a Modern Approach Third Edition*, Prentice Hall
- Sachs J. D et al. 2020. The Lancet COVID-19 Commission. *Lancet* 396, 454–455 (2020). *Science* 1965. The Evolution of Science, *Science-New Series*, vol. 148, no. 3671, p. 737, Stable
- Scopus 2021. Documents. <https://www.scopus.com/> (accessed 15 October 2021)
- Segreto, R., Deigin, Y., McCairn, K., Sousa, A., Sirotkin, D., Sirotkin, K., Couey, J. J., Jones, A., & Zhang, D. (2021). Should we discount the laboratory origin of COVID-19? *Environmental chemistry letters*, 1–15. Advance online publication. [10.1007/s10311-021-01211-0](https://doi.org/10.1007/s10311-021-01211-0)
- Seidman S. S. 1987. Models of scientific development in sociology. *Humboldt Journal of Social Relations* vol. 15, n.1, pp.119- 139
- Simmons, H. E., Matos, B. & Simpson, S. A. 2017. Analysis of injury data to improve safety and training. *J. Chem. Health Saf.* 24, 21–28.
- Sirotkin, K., & Sirotkin, D. (2020). Might SARS-CoV-2 Have Arisen via Serial Passage through an Animal Host or Cell Culture? A potential explanation for much of the novel coronavirus' distinctive genome. *BioEssays : news and reviews in molecular, cellular and developmental biology*, 42(10), e2000091. [10.1002/bies.202000091](https://doi.org/10.1002/bies.202000091)
- USGS 2021. Natural Disasters—Forecasting Economic and Life Losses. [Retrieved from]. (Accessed 28 October 2021)

Journal of Social and Administrative Sciences

- Van Noorden, R. 2013. Safety survey reveals lab risks. *Nature* 493, 9–10 (2013).
- WIV 2021. History Wuhan Institute of Virology. [[Retrieved from](#)] (Accessed 28 October 2021)
- Wolfe, N., Dunavan, C. & Diamond, J. 2007. Origins of major human infectious diseases. *Nature* 447, 279–283. [10.1038/nature05775](https://doi.org/10.1038/nature05775)
- Wu et al., Z. 2016. Deciphering the bat virome catalog to better understand the ecological diversity of bat viruses and the bat origin of emerging infectious diseases. *ISME J.* 10, 609–620 (2016).
- Yuan, D., Gao, W., Liang, S., Yang, S., & Jia, P. (2020). Biosafety threats of the rapidly established labs for SARS-CoV-2 tests in China. *Environment international*, 143, 105964. [10.1016/j.envint.2020.105964](https://doi.org/10.1016/j.envint.2020.105964)
- Zhang Y.-Z., E. C. Holmes, 2020. A genomic perspective on the origin and emergence of SARS-CoV-2. *Cell* 181, 223–227.



Copyrights

Copyright for this article is retained by the author(s), with first publication rights granted to the journal. This is an open-access article distributed under the terms and conditions of the Creative Commons Attribution license (<http://creativecommons.org/licenses/by-nc/4.0>).

