<u>Declaration form on Nothing to Declare or Nothing New to Declare for use in the information exchange</u>

Measure	Nothing to declare	Nothing new to declare	Year of last declaration if nothing new to declare
A, part 1		X	2014
A, part 2 (i)		X	2014
A, part 2 (ii)		X	2014
A, part 2 (iii)	X		
В			
C			
E		X	2018
F		X	1992
G		X	2016

(Please mark the appropriate box(es) for each measure with a tick, and fill in the year of last declaration in the last column where applicable.)

Date: April, 2021

State Party to the Convention: Czech Republic

Date of ratification/accession to the Convention: 30 April 1973 (former Czechoslovakia)

24 March 1993 (Czech Republic)

National point of contact: State Office for Nuclear Safety

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Director

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B. Information on outbreaks of infectious diseases and similar occurrences, that seem to deviate from the normal pattern

In 2020, the following infectious diseases have been reported in the Czech Republic:

Infectious disease	Number of cases	
Highly Pathogenic Avian Influenza (H5N8)	2 outbreaks: o 15 (back-yard flock) o 137 500 (commercial farm)	
COVID-19	718 380	

More available on:

 $\underline{https://wahis.oie.int/\#/dashboards/country-or-disease-dashboard}$

http://www.szu.cz/publikace/data/2020

C. Encouragement of publication of results and promotion of use of knowledge

List of the most important publications, which appeared during the year 2020:

Konkolova E., Klima M., Nencka R., Boura E. (2020) Structural analysis of the putative SARS-CoV-2 primase complex. *Journal of Structural Biology*. Volume 211, Issue 2, 107548 https://doi.org/10.1016/j.jsb.2020.107548Get rights and content

Krafcikova P., Silha J., Nencka R. *et al.* (2020) Structural analysis of the SARS-CoV-2 methyltransferase complex involved in RNA cap creation bound to sinefungin. *Nature Communication*. 11, 3717. https://doi.org/10.1038/s41467-020-17495-9

Kriegova E., Fillerova R., Kvapil P. (2020): Direct-RT-qPCR Detection of SARS-CoV-2 without RNA Extraction as Part of a COVID-19 Testing Strategy: From Sample to Result in One Hour. *Diagnostics*. 10(8), 605; https://doi.org/10.3390/diagnostics10080605

Matyasek R., Kovarik A. (2020): Mutation Patterns of Human SARS-CoV-2 and Bat RaTG13 Coronavirus Genomes Are Strongly Biased Towards C>U Transitions, Indicating Rapid Evolution in Their Hosts. *Genes.* 11(7), 761; https://doi.org/10.3390/genes11070761

Bartas M., Brazda V., Bohalova N., Cantara A., Volna A., Stachurova T., Malachova K., Jagelska E., Porubiakova O., Cerven J., Pecinka P. (2020) In-Depth Bioinformatic Analyses of Nidovirales Including Human SARS-CoV-2, SARS-CoV, MERS-CoV Viruses Suggest Important Roles of Non-canonical Nucleic Acid Structures in Their Lifecycles. *Frontiers in Microbiology*. 11, 1583. https://doi.org/10.3389/fmicb.2020.01583

Pohanka M. (2020) *Bacillus anthracis* as a biological warfare agent: infection, diagnosis and countermeasures. *Bratislava Medical Journal*. 121 (3)175 – 181. https://doi.org/10.4149/BLL_2020_026

Pohanka M. (2020) Chapter Eleven - Biosensors commercial off the shelf in biological warfare attack. *Commercial Biosensors and Their Applications*. Pages 287-300. https://doi.org/10.1016/B978-0-12-818592-6.00011-6

Spidlova P., Stojkova P., Sjöstedt A., Stulik, J. (2020) Control of *Francisella tularensis* Virulence at Gene Level: Network of Transcription Factors. *Microorganisms*. 8(10), 1622 https://doi.org/10.3390/microorganisms8101622

Ostry V., Dofkova M., Blahova J., Malir F., Kavrik R., Rehurkova I., Ruprich J. (2020) Dietary exposure assessment of sum deoxynivalenol forms, sum T-2/HT-2 toxins and zearalenone from cereal-based foods and beer. *Food and Chemical Toxicology*. 139; 111280. https://doi.org/10.1016/j.fct.2020.111280