201216 Brussels



# COVID-19 through the eyes of an immunologist

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# - The virus: SARS-CoV-2

very high infectivityrare severe/letal cases

- The Spike protein
- Receptor and host factors
- The disease: COVID-19
- Innate and adaptive immune responses
- Vaccines





## SARS-CoV-2



Origin of SARS-CoV-2: bats (closest relative: Bat-CoV-RATG13 96.2% genome identity)

 $\hat{\mathbb{U}}$ 

man

Tang D, et al. PLoS Pathog 2020;16:e1008536.

#### **SARS-CoV-2** structural proteins





## SARS-CoV-2 genome





## SARS-CoV-2 genome



- structural proteins
- proteins required for viral replication and production (in ORF1a and 1b)
- 'accessory factors', functions not completely known, participation in pathogenesis?



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#### SARS-CoV-2 spike protein

- ±40 per virion
- 20 nm
- trimer
- heavily glycosylated
- flexible





#### SARS-CoV-2 spike protein

- ±40 per virion
- 20 nm





#### Monomer of SARS-CoV-2 spike protein





#### Trimer of SARS-CoV-2 spike protein



- 20 nm
- trimer
- heavily glycosylated
- flexible







Bangaru S, et al. Science 2020;370:1089-94.

#### SARS-CoV-2 spike protein: polysaccharides in color





(T323: unique site of O-glycosylation)

Multiple sites of N-glycosylation

- $\rightarrow$  many proteic epitopes are masked
  - = virus partial protection against neutralizing antibodies
- $\rightarrow$  protective effect of blood group O

#### Model of membrane fusion by ß-coronavirus spike protein



This process is THE main target for antibody-mediated virus neutralization

Khao J. and McGill G. Digizyme Inc.

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#### The receptor: angiotensin converting enzyme 2 (ACE2)



Adapted from Groß S et al. J Mol Cell Cardiol 2020;144:47-53.

#### **ACE2 expression profile is key to COVID-19 symptoms**





## SARS-CoV-2 spike protein: processing required

- ±40 per virion
- 20 nm



- heavily glycosylated
- flexible





#### S1 domain: attachment S2 domain: fusion



Li F, et al. Annu Rev Virol 2016;3:237-61.

## SARS-CoV-2 spike protein: processing required for virus entry

- ß-coronavirus S protein requires 2 proteolytic cleavages:

- between domains S1 and S2

to favor changes of S1 conformations: closed conformation: RBD buried open conformation: RBD accessible to ACE2

to allow for the detachment of S1, unmasking S2

- upstream of the fusion peptide, to allow its insertion in the plasma membrane
- efficacy and extent of these activation steps regulate cellular tropism and viral pathogenesis





#### Host factor: TMPRSS2

- transmembrane serine protease 2
- cleaves at S1/S2 and at S2'
- surface co-expression of ACE2 and TMPRSS2 on the same cells (in cis) appears optimal for SARS-CoV-2 infection of human cell lines *in vitro*

*in vivo*, coexpression by nasal epithelial cells, bronchial transient secretory cells (Sungnak et al. 2020 Nat Med 26:681) (Lukassen et al. 2020 EMBO J e105114)

- ACE2 and TMPRSS2 genes are induced in various proinflammatory conditions (obesity, diabetes, ...)



#### A unique feature of SARS-CoV-2, absent in other members of its clade



- insertion of 12 nucleotides upstream of S1/S2 in SARS-CoV-2

- absent in SARS-CoV-1, MERS-CoV and Bat-CoV-RATG13
- creates a **furin**-like cleavage site

(furin: endoprotease in the secretory pathway compartments)

- furin is highly expressed in lungs
- participates in SARS-CoV-2 pathogenesis ? (very high contagiousness and rapid spread)



#### Host factor: neuropilin 1



- neuropilins (NRP1 and NRP2):

- surface receptor for semaphorins and VEGF
- highly expressed in almost all pulmonary and olfactory cells
- highest levels of expression in endothelial cells

- neuropilin 1 binds furin-cleaved sustrates (the polybasic RRAR carboxyterminal sequence on S1)

- blocking this interaction reduces SARS-CoV-2 infectivity in vitro





Daly J, et al. Science 2020;370:861-65; Canti-Castelvetri L, et al. Science 2020;370:856-60.

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#### **COVID-19**







Cevik et al. 2020 BMJ 371:m3862

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- detection by cells of pathogen-associated molecular patterns (PAMP)
  - main sensor for SARS-CoV-2: TLR3
  - detected PAMP: virus-specific aberrant RNA structures (dsRNA, ...)
- transcriptional response: two antiviral programs
  - type-I (IFN- $\alpha$ , IFN- $\beta$ ) and type-III interferons (IFN- $\lambda$ )  $\rightarrow$  cellular antiviral defenses too little, too slow
  - chemokines  $\rightarrow$  recruitment and coordination of subsets of leukocytes too much



**COVID-19** 

#### Viral resistance to innate immunity

- all pathogens, viruses and others, have acquired mechanisms to partially resist immunity (innate and adaptive)





#### **Inflammatory cells**

- local (lungs) production of inflammatory cytokines (IL-1ß, IL-6, TNF, ...) and chemokines
  - levels of gene and protein expression are correlated with disease severity
  - especially in severe cases: associated with impaired type I IFN responses (comparisons with RSV of influenza A)
  - 'cytokine storm' and viral production remain present in the inflamed lungs of deceased patients
- followed by the accumulation of inflammatory cells, including neutrophils, in the lungs
  - immature myeloid cells found in blood (reactive myelopoiesis)
- the higher level of basal inflammation associated with age ('inflammaging') or obesity (inflammasome activation) might be risk factors



Altman DM, et al. Sci Immunol 2020;5:eabd6160; Hadjadj J, et al. Science 2020;369:718.

#### Adaptive immunity to viruses





- T lymphodepletion at late sages of disease (observed also in sepsis from other causes)



## Adaptive immunity to SARS-CoV-2



- activation of CD8 T cells - cytolysis of SARS-CoV-2-infected cells



#### Is there something 'immunologically' remarkable with SARS-CoV-2?

- For the immunologist, SARS-CoV-2 is a perfectly **normal virus** 

It induces innate and adaptive immunity with classical effector mechanisms to which it partially resists.

- Today, the peculiarity of SARS-CoV-2 appears to be a proportionally **low IFN-I response**.

This low IFN response is responsible for more viruses during a longer time, and these viruses continue to trigger the inflammatory response (chemokines and inflammatory cytokines).

Together, these mediators continue to recruit innate immune cells (of myeloid origin) at sites of virus production, notably the lungs.

N.B. Inborn errors of IFN-I induction or signalling were found in adult patients (n=23/659) with life-threatening COVID-19 pneumonia, but these genotypes were silent until infection with SARS-CoV-2. These pathways are however known to be associated with influenza respiratory distress. Thus we appear to depend more on IFN-I to control SARS-CoV-2 than other viruses.



Zhang Q, et al. Science 2020;370:eabd4570.

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(therapeutic antibodies are not discussed here)



#### **SARS-CoV-2** vaccines

#### **Objective:**

Stimulate the production of neutralizing anti-SARS-CoV-2 antibodies (the other option is to transfer such antibodies) All currently proposed vaccines aim at inducing anti-Spike protein antibodies

#### Method:

Inoculate the antigen(s) in such a way that they are immunogenic (stimulate an in vivo adaptive immune response)

Vaccination modality is a key factor for immunogenicity:

- either co-inject antigen and immunological adjuvants (increase antigen half-life and induce local inflammation)
- or inject DNA or RNA encoding the antigen, which is then produced by the vaccinee's own cells

DNA or RNA has to be 'packaged' (lipid nanoparticles, recombinant viruses) Roles of packaging: protect against DNAse/RNAse and promote entry into cells



#### SARS-CoV-2 vaccines for Belgium (dec 2020)



Krammer F. Nature 2020;586:516-27.

## **Expected immune responses with (SARS-CoV-2) vaccines**

	<b>B</b> cells	CD4 T cells	CD8 T cells	
mRNA ?	+	+	+	Pfizer-BioNTech Moderna Curevac
replication-incompetent adenovirus	+	+	+	AstraZeneca Janssen Gamaleya (Spoutnik V)
protein + adj	+	+	-	GSK-Sanofi Novavax
inactivated virion + adj	+	+	-	Sinovac Biotech
live attenuated virus	+	+	+	Pasteur-Merck Measles-spike prot Pasteur-TheraVectys Lentivirus-spike prot

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not for immunocompromised individuals

Krammer F. Nature 2020;586:516-27.