

# Coronaviruses are an established cause of severe respiratory, enteric and systemic infections

Four genera:

## 1) *Alphacoronavirus* = infect mammals

[**human coronavirus NL63 (HCoV-NL63)**, **HCoV-229E**, porcine transmissible gastroenteritis coronavirus (TGEV), PEDV, and porcine respiratory coronavirus (PRCV)]

↓  
{ACE2 receptor}

## 2) *Betacoronavirus* = infect mammals

[**SARS-CoV**, **COVID-19**, **MERS-CoV**, bat coronavirus HKU4, **mouse hepatitis coronavirus (MHV)**, bovine coronavirus (BCoV), and **HCoV-OC43**, **HCoV-HKU1**]

↓ ↓ ↓  
{ACE2 receptor} {DPP4 receptor}

↓  
{CEACAM1 receptor}

## 3) *Gammacoronavirus* = infect avian species

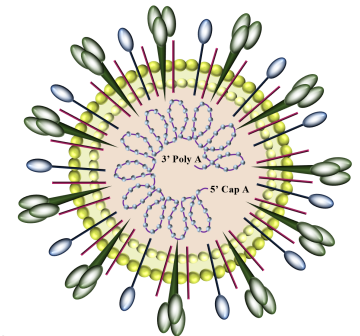
[avian infectious bronchitis coronavirus (IBV)]

## 4) *Deltacoronavirus* = infect both mammalian and avian species

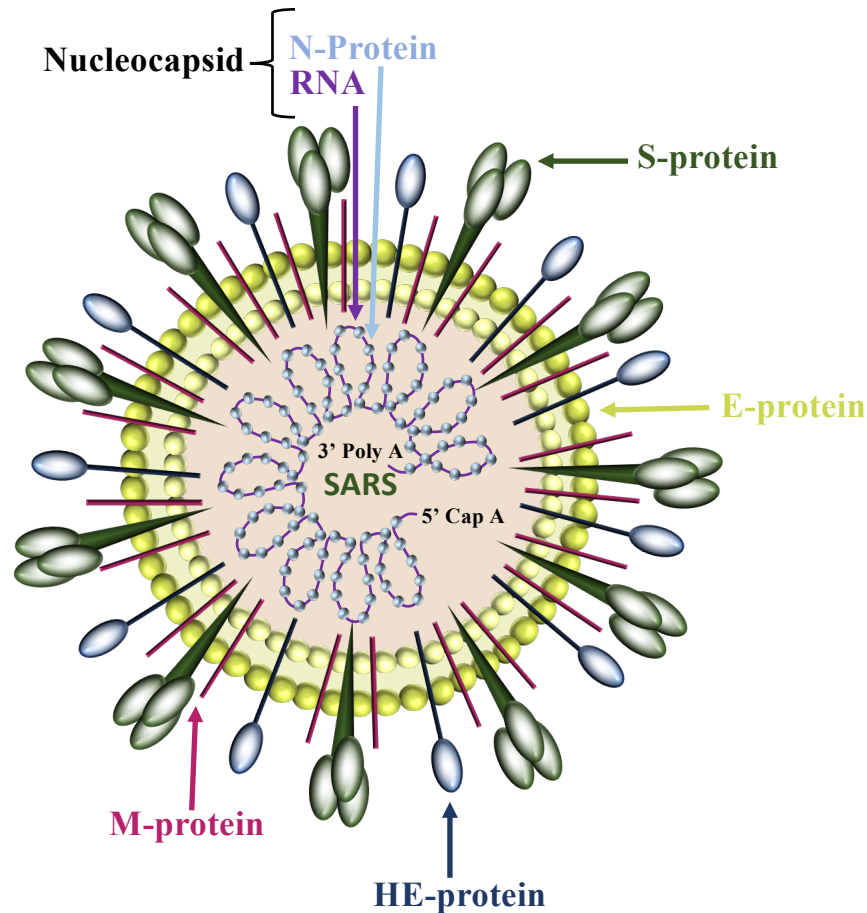
[porcine deltacoronavirus (PdCV)]

**\*The diversity of receptor usage is an outstanding feature of coronaviruses**

{e.g. Angiotensin-converting enzyme 2 gene (ACE2), carcinoembryonic antigen-related cell adhesion molecule 1 (CEACAM1), dipeptidyl peptidase 4 (DPP4), sugar receptors}



PMID: 27578435 [Annu Rev Virol](#). 2016 Sep 29;3(1):237-261.  
PMID: 30646565 [Viruses](#). 2019 Jan 14;11(1).



**Coronavirus virus particles contain four main structural proteins encoded within the 3' end of the viral genome.**

**1) Spike (S)-protein:** trimeric protein that mediates attachment to the host receptor, can be cleaved by a host cell furin-like protease into two separate polypeptides noted S1 (binding domain) and S2 (stalk)

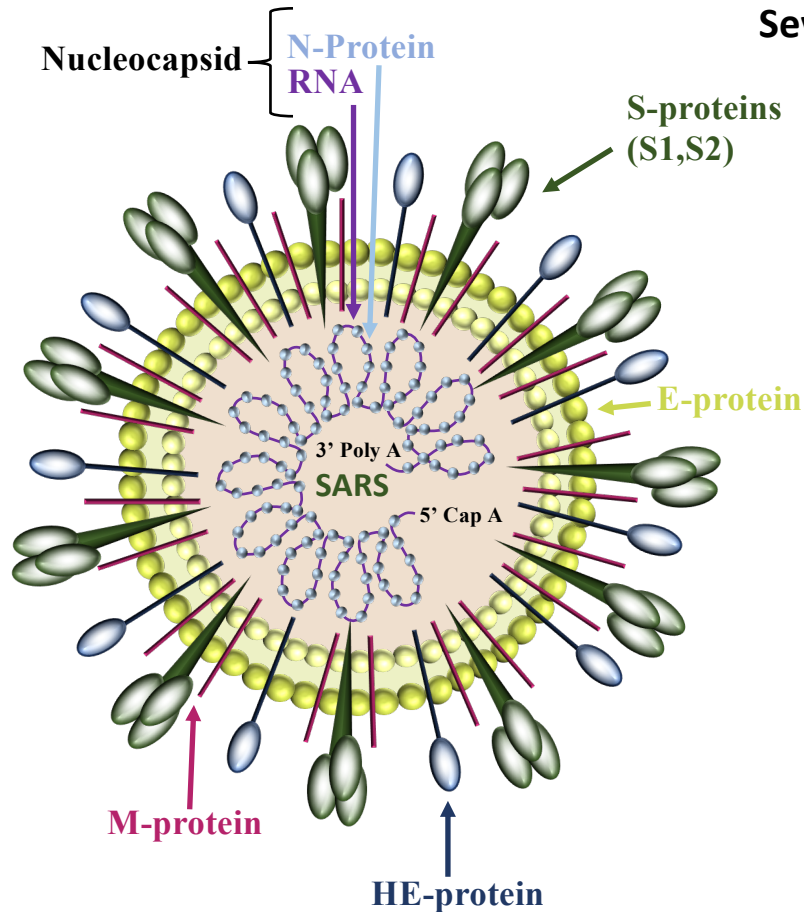
**2) Membrane (M)-protein:** most abundant structural protein in the virion.

**3) Envelope (E)-protein:** facilitates assembly and release of the virus, the ion channel activity in SARS-CoV E protein is required for pathogenesis

**4) N-Protein:** constitutes the only protein present in the nucleocapsid composed of 2 domains that bind RNA

**5) Hemagglutinin-esterase (HE)-protein:** present in a subset of  $\beta$ -coronaviruses, binds sialic acids on surface glycoproteins and contains acetyl-esterase activity

## Severe acute respiratory syndrome coronavirus 2 (SARS-Cov2): COVID-19



➔ Sourced from Huanan Seafood Wholesale Market, which trades in fish and a variety of live animal species including poultry, bats, marmots, and snakes

➔ Positive-strand RNA betacoronavirus

➔ Highly affects older patients with multiple comorbidities

➔ S1 directly binds to the peptidase domain of ACE2, exposing an S2 cleavage site critical for viral infection

➔ S2 is subsequently cleaved by endosomal cysteine proteases cathepsin B and L (CatB/L) and/or serine protease TMPRSS2 prior to fusion where the latter may be preferred.

➔ In vitro, camostat mesylate inhibits TMPRSS2 & ammonium chloride inhibits CatB/L

PMID: 32112977 [Int J Surg](#). 2020 Feb 26;76:71-76.  
PMID: 32142651 [Cell](#). 2020 Mar 4. pii: S0092-8674(20)30229-4  
PMID: 29217279 [Virology](#). 2018 Apr;517:9-15..

## Identified ACE2 immunostaining in HUMANS is most prominent in lung and intestinal epithelia

### Cell types:

**Lung:** type I and type II alveolar epithelial cells

**Nasal:** non-keratinizing squamous epithelium

**GI:** smooth muscle cells and endothelium of vessels from the stomach, small intestine, and colon

**Skin:** basal cell layer of the epidermis extending to the basal cell layer of hair

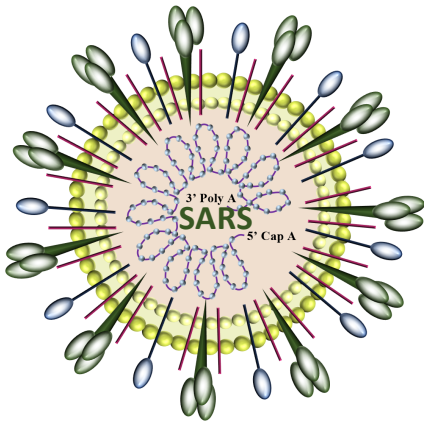
**Kidney:** glomerular and epithelial cells

**All organs:** arterial and venous endothelial cells

Immune cells of the spleen, thymus, lymph nodes, and bone marrow were consistently negative for ACE2

PMID: 15141377 [J Pathol](#). 2004 Jun;203(2):631-7

## Transmembrane protease, serine 2 (TMPRSS2, epitheliasin)



CSCURRAN (2020)

### Human transcript is observed in the following order:

prostate > colon > small intestine > pancreas > kidney > lung > liver

Epitheliasin transcripts in fetal tissues are observed only in kidney and lung.

Epitheliasin is induced by androgens

PMID: 11322890 [Eur J Biochem](#). 2001 May;268(9):2687-99.

Expressed primarily in the apical surfaces of **mouse** renal tubular and airway epithelial cells.

PMID: 10683448 [FEBS Lett](#). 2000 Feb 18;468(1):93-100.

A discernible phenotype in *Tmprss2*<sup>-/-</sup> mice is not identified, suggesting possible redundant functions in the absence of infection

PMID: 16428450 [Mol Cell Biol](#). 2006 Feb;26(3):965-75.



## ACE2 function is primarily identified in the renin/angiotensin system

In an ACE2 knockout model of lung injury, mice exhibited increased vascular permeability, lung edema and neutrophil influx in association with decreased lung function & this phenotype was rescued with a pharmacological inhibitor of the angiotensin II type 1 receptor (**AT<sub>1</sub>R**)

PMID: 24776703

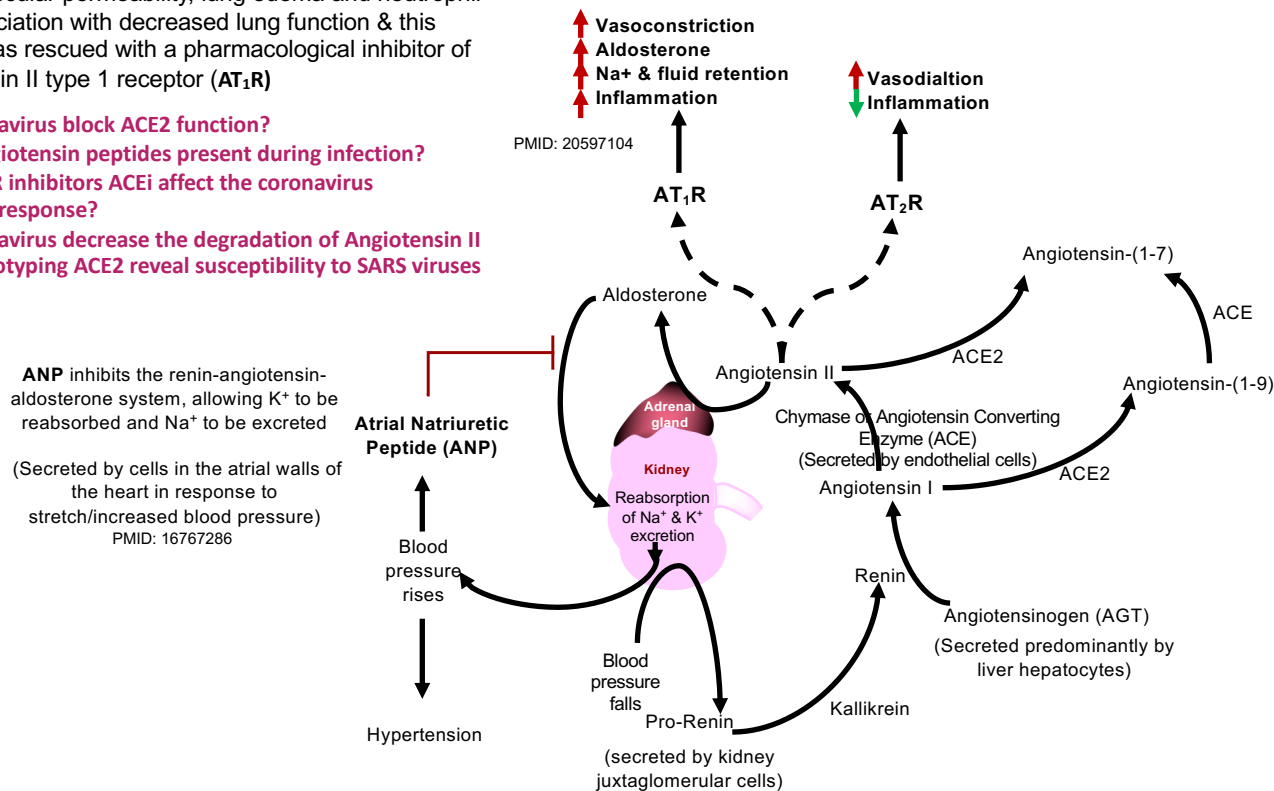
**Q: Does coronavirus block ACE2 function?**

**Q: Are the angiotensin peptides present during infection?**

**Q: Would AT<sub>1</sub>R inhibitors ACEi affect the coronavirus inflammatory response?**

**Q: Does coronavirus decrease the degradation of Angiotensin II**

**Q: Would genotyping ACE2 reveal susceptibility to SARS viruses**



\*Both ACE and ACE2 bind integrins in an RGD-independent manner  
 \*ACE2 increases cellular adhesion and affects integrin signaling  
 \*The RGD motif present in the ectodomain of ACE2 is inaccessible  
 \*sACE2 significantly reduces FAK phosphorylation levels.  
 PMID: 22523556 [PLoS One](#). 2012;7(4):e34747.

**Q: Does sACE2 antagonize COVID-19?**

**Q: Would increasing ACE2 expression (e.g. ACEi) promote viral entry and/OR antagonize inflammation**

**Q: Would a COX2i dampen inflammation?**

**An evolutionary RGD motif in the spike protein of SARS-CoV-2 may serve as a potential high risk factor for virus infection? PREPRINT PAPER:** Notably, we found that S protein of SARS-CoV-2 produced an evolutionary mutation K403R, located at site 403, forming an RGD motif (Arg-Gly-Asp) in the S protein next to the binding site, which has neither been found in RaTG13 nor in SARS-CoV---**Q: Are integrins co-receptors for COVID-19? Is FAK required for viral entry?**

ACE2 purified from failing human heart was found to form a complex with integrin beta1 by immunoprecipitation  
 PMID: 15276642 [Biochim Biophys Acta](#). 2004 Aug 4;1689(3):175-8.

Overexpression of ACE2 in neurons/brain antagonized COX2 and hypertension  
 PMID: 25489058 [Hypertension](#). 2015 Mar; 65(3): 577-586.

PGI<sub>2</sub> upregulates  $\beta$ 1-integrin expression and cell migration in HCC cells  
 PMID: 25289898 [Sci Rep](#). 2014; 4: 6538.

ACEi and ibuprofen purportedly increase ACE2

<https://www.thelancet.com/action/showPdf?pii=S2213-2600%2820%2930116-8>

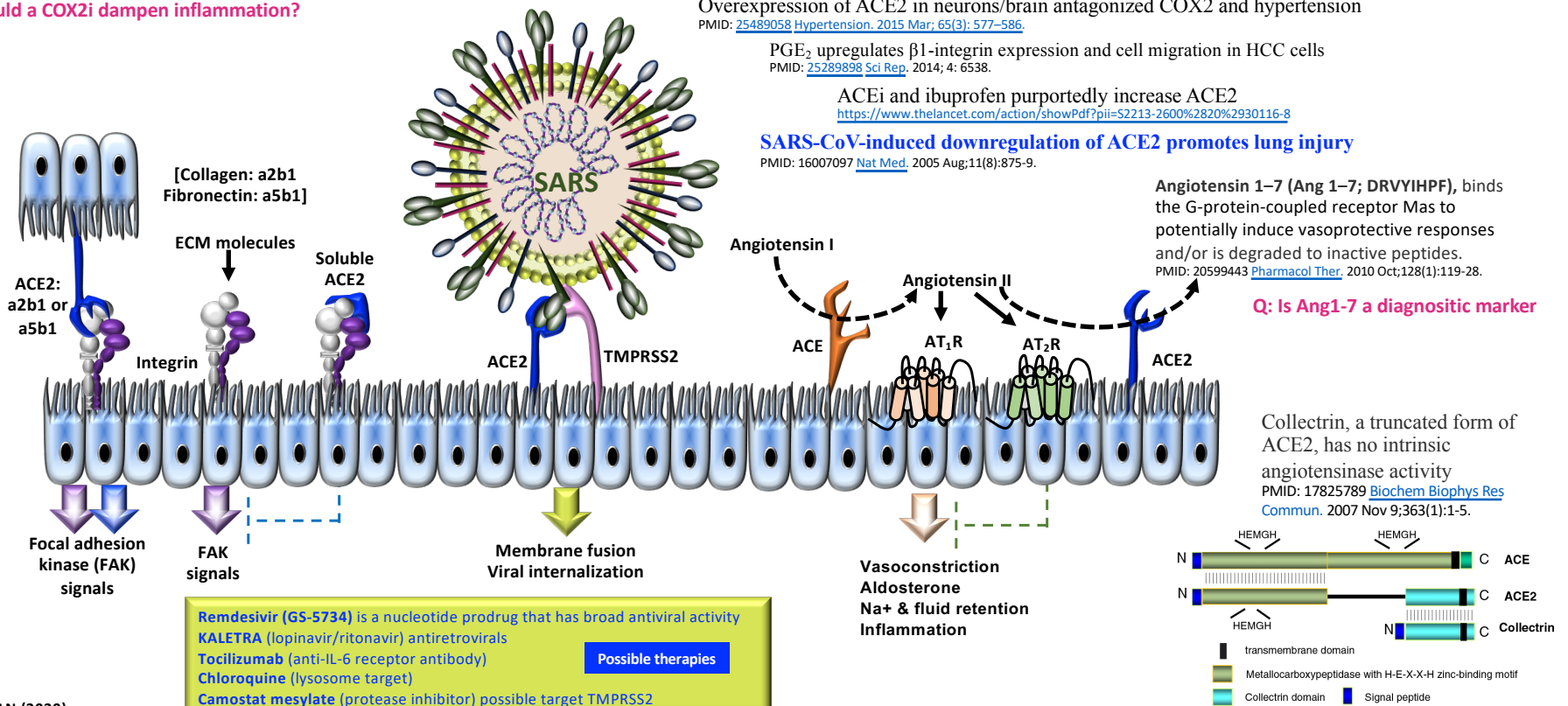
**SARS-CoV-induced downregulation of ACE2 promotes lung injury**

PMID: 16007097 [Nat Med](#). 2005 Aug;11(8):875-9.

Angiotensin 1-7 (Ang 1-7; DRVYIHPF), binds the G-protein-coupled receptor Mas to potentially induce vasoprotective responses and/or is degraded to inactive peptides.  
 PMID: 20599443 [Pharmacol Ther](#). 2010 Oct;128(1):119-28.

**Q: Is Ang1-7 a diagnostic marker**

Collectrin, a truncated form of ACE2, has no intrinsic angiotensinase activity  
 PMID: 17825789 [Biochem Biophys Res Commun](#). 2007 Nov 9;363(1):1-5.



## Middle East respiratory syndrome coronavirus (MERS-CoV)

\*MERS-CoV uses the N-terminal part of its spike (S)—the so called S1 protein to bind to two host cell surface molecules, dipeptidyl peptidase-4 (DPP4) and  $\alpha$ 2,3-sialic acids.

\*Post attachment, MERS-CoV uses the C-terminal part of its S protein—known as S2—to interact with host proteases, such as furin, TMPRSS2, and cathepsins

\*TMPRSS2 and DPP4 are held in one complex at the cell surface by a scaffolding protein, the tetraspanin CD9, leading to a rapid and efficient entry of MERS-CoV into the susceptible cells

PMID: 30893947 [Viruses](#). 2019 Mar 19;11(3). pii: E280.

\*DPP-4 is a cell surface aminopeptidase that was originally characterized as a T cell differentiation antigen (CD26)

\*DPP-4 is expressed ubiquitously and found in many cell types, including the endothelial cells in multiple vascular beds, rendering the enzyme highly accessible to the peptide substrates circulating through the gut, liver, lung, and kidney----- the kidney expresses the highest levels of DPP-4 per organ weight

\*The loss of DPP-4 cell surface expression has shown to be associated with decreased phosphorylation of integrin  $\beta$ 1 at the S785 residue

PMID: 26877767 [Fibrogenesis Tissue Repair](#). 2016 Feb 13;9:1.

Within COVID-19, many different and unique residues (R408, Q409, T445, V417, L461, D467, S469, L491, N492, D493, Y 494, T497, T150, Y504) are predicted to interact with CD26

[Emerg Microbes Infect.](#) 2020 Dec;9(1):601-604. PMID: 32178593

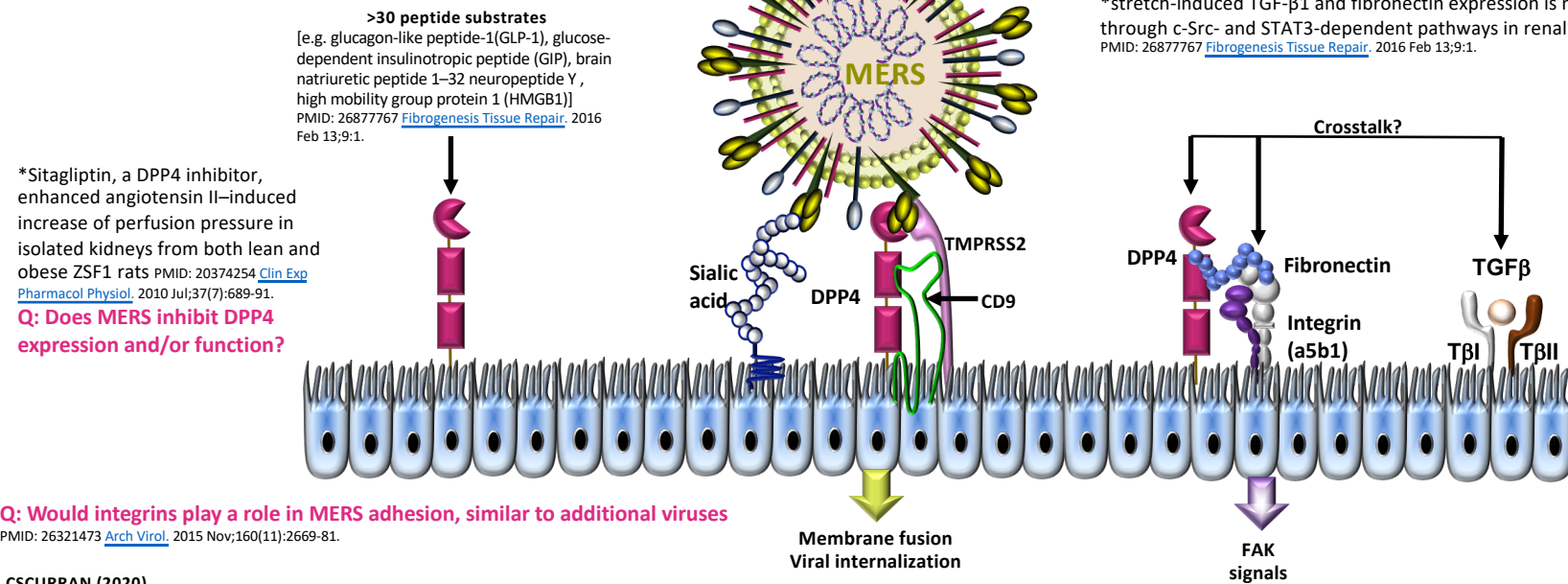
\*CD26/DPP4 has shown the ability to act as a non-integrin receptor, being able to bind fibronectin & collagen & is implicated in autoimmune, neurological, & metabolic diseases

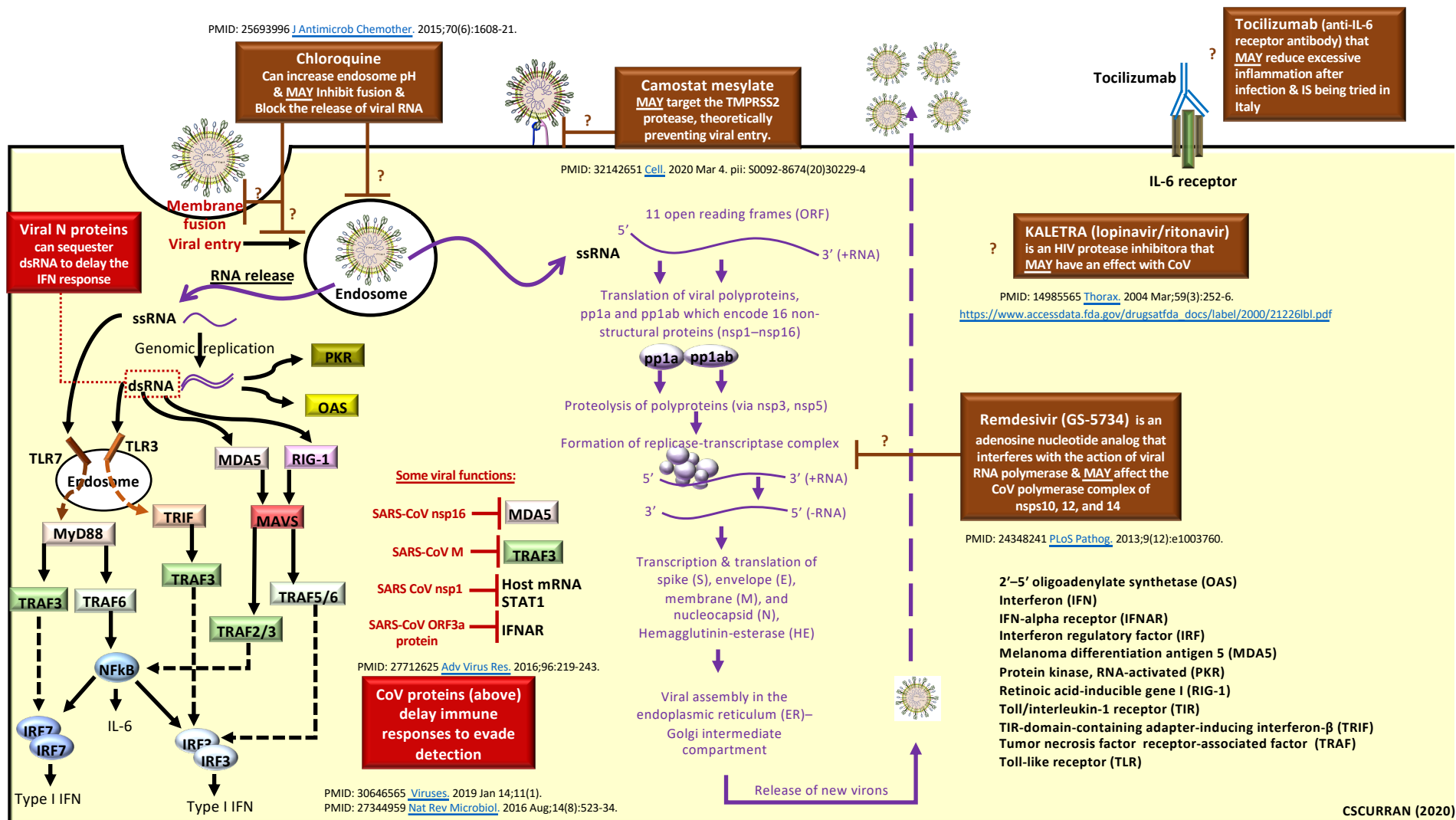
PMID: 26919392 [Clin Exp Immunol](#). 2016 Jul; 185(1): 1–21.

\*DPP-4 or integrin  $\beta$ 1 deficiency resulted in the inhibition of TGF- $\beta$ 2-stimulated hetero-dimerization of TGF- $\beta$ Rs

\*stretch-induced TGF- $\beta$ 1 and fibronectin expression is mediated by  $\beta$ 1-integrin through c-Src- and STAT3-dependent pathways in renal epithelial cells

PMID: 26877767 [Fibrogenesis Tissue Repair](#). 2016 Feb 13;9:1.





## The human immune response to COVID-19

\*Most patients also developed lymphopenia and pneumonia with characteristic pulmonary ground glass opacity changes on chest CT.

**\*Severe case cytokines (41 patients):** IL-2, IL-7, IL-10, G-CSF, IP-10, MCP-1, MIP-1A, and TNF $\alpha$

**\*99 patients:** increased total neutrophils (38%), reduced total lymphocytes (35%), increased serum IL-6 (52%) and increased c-reactive protein (84%) were observed.

\*SARS-Co-V directly infects macrophages and T cells, a key feature in SARS-CoV-mediated pathogenesis, in COVID-19?. Only minimal percentages of monocytes/macrophages in the lung expressed ACE2.

\*SARS-CoV induces delayed type I IFN and loss of viral control in an early phase of infection, in COVID-19?

\*Type I IFN for treatment in mouse models of either SARS-CoV or MERSCoV infection, the timing of administration is key to yield protective response.

\*SARS-CoV infection induces seroconversion as early as day 4 after onset of disease and was found in most patients by 14 days.

\*Long lasting specific IgG and neutralizing antibody are reported as long as 2 years after infection

\*Sera from 5 patients of confirmed COVID-19 show some cross-reactivity with SARS-CoV, but not other coronavirus.

\*The virus specific T cells from the severe group tended to be a central memory phenotype with a significantly higher frequency of polyfunctional CD4+ T cells (IFN $\gamma$ , TNF $\alpha$ , and IL-2) and CD8+ T cells (IFN $\gamma$ , TNF $\alpha$  and degranulated state), as compared with the mild-moderate group.

\*Strong T cell responses correlated significantly with higher neutralizing antibody while more serum Th2 cytokines (IL-4, IL-5, IL-10) were detected in the fatal group

PMID: 32105090 [Asian Pac J Allergy Immunol](https://doi.org/10.1186/s12931-020-0819-2). 2020 Feb 27.

### Breadth of concomitant immune responses prior to patient recovery: a case report of non-severe COVID-19

<https://www.nature.com/articles/s41591-020-0819-2>

\*Increased antibody-secreting cells (ASCs-CD3-CD19<sup>+</sup>CD27<sup>hi</sup>CD38<sup>hi</sup>), follicular helper T cells (T<sub>fh</sub> cells-CD4<sup>+</sup>CXCR5<sup>+</sup>ICOS<sup>+</sup>PD-1<sup>+</sup>), activated CD38<sup>+</sup>HLA-DR<sup>+</sup> CD4<sup>+</sup> T cells and CD38<sup>+</sup>HLA-DR<sup>+</sup>CD8<sup>+</sup> T cells and immunoglobulin M (IgM) and IgG antibodies that bound the COVID-19 were detected in blood before symptomatic recovery.

\*These immunological changes persisted for at least 7 d following full resolution of symptoms.

[detection of SARS-CoV-2 in sputum, nasopharyngeal aspirates and feces but not urine, rectal swab or whole blood]