

Highly efficient inhibition of SARS-CoV-2 entry by a biologically unique ACE2-IgG4-Fc fusion protein with a stabilized hinge region

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Introduction

Severe acute respiratory syndrome (SARS)-like coronavirus 2 (SARS-CoV-2) and other human pathogenic coronaviruses use the angiotensin-converting enzyme 2 (ACE2) to enter human host cells.¹ ACE2-Fc fusion proteins composed of the human immunoglobulin G (IgG) fragment crystallizable (Fc) part fused to the extracellular domain of ACE2 have been suggested as a highpriority treatment option for COVID-19², however bear the risk of unwanted Fc-receptor activation and antibody-dependent disease enhancement. We have designed an innovative ACE2-IgG4-Fc fusion protein as a potent SARS-CoV-2 entry blocker.³ First, the ACE2-IgG4-Fc fusion protein acts as a highly efficient virus trap by targeting the entire receptor-binding domain (RBD) of SARS-CoV-2. Second, ACE2-IgG4-Fc cleaves angiotensin II to angiotensin 1-7, potentially providing additional protection to the lung and cardiovascular system. Third, the risk of antibody-dependent disease enhancement is minimized by using the IgG4 Fc region.



Materials and methods

ACE2-Fc construct design

Constr.	ACE2 sequence (active site mutation)	IgG isotype (hinge region modification)
1	Q18-G732 (no)	IgG4 (S228P)
2	Q18-S740 (no)	IgG4 (S228P)
3	Q18-G732 (H374N, H378N)	IgG4 (S228P)
4	Q18-S740 (H374N, H378N)	IgG4 (S228P)
5	Q18-G732 (no)	IgG1 (truncated)
6	Q18-S740 (no)	IgG1 (truncated)
7	Q18-G732 (H374N, H378N)	IgG1 (truncated)
8	Q18-S740 (H374N, H378N)	IgG1 (truncated)

Protein expression and purification – Transient expression in HEK293 cells. Protein A followed by size-exclusion chromatography.

Circular dichroism (CD) - J-1500 spectropolarimeter (Jasco). Size-exclusion chromatography with multi-angle light scattering (SEC-MALS) – Shimadzu HPLC, Superdex 200 Increase 10/300 GL column and HELEOS II detector (Wyatt Technology).

Enzymatic activity - Cleavage of a peptidyl-4-methylcoumaryl-7-amide (MCA) and fluorimetric detection of the free MCA. Assay kit from Abcam (Cat.No. ab273297).

Surface plasmon resonance (SPR) - Biacore X100 (Cytiva), Biotin CAPture kit (Cytiva), captured SARS-CoV-2 RBD with an AviTag™. ELISA - ACE2:SARS-CoV-2 spike S1 inhibitor screening assay kit (BPS Bioscience; Cat.No. 79945).

Virus neutralization assays - SARS-CoV-2-GFP⁴ is based on the original Wuhan SARS-CoV-2 isolate (GenBank accession MT108784). SARS-CoV-2-Jan (EPI_ISL_582134), SARS-CoV-2-April (EPI_ISL_466888), and SARS-CoV (AY291315.1) were isolated from patient material in Germany. Vero E6 cells were used in the assays.



b. SEC-MALS shows that the ACE2-Fc molecules exist as homodimers.







SPR experiments.

b. ELISA shows that ACE2-Fc proteins bind with high affinity to the spike protein of SARS-CoV-2.

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- ACE2-Fc proteins neutralize SARS-CoV and different SARS-CoV-2 strains.

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replication in Vero E6 cells (MOI = 0.3).

ACE2-Fc proteins neutralize:

- a. SARS-CoV with IC50 between 90 and 250 nM (MOI = 0.03).
- b. SARS-CoV-2-Jan with IC50 between 7 and 11 nM (MOI = 0.03).
- c. SARS-CoV-2-April with IC50 between 0.3 and 0.8 nM (MOI = 0.03).

Conclusions

- ACE2-IgG4-Fc proteins have favorable biophysical and pharmaceutical characteristics. ACE2-Fc molecules with WT ACE2 preserve enzymatic activity.
- All ACE2-Fc proteins bind to SARS-CoV-2 spike proteins with high affinity.
- ACE2-IgG4-Fc efficiently inhibits SARS-CoV-2-GFP cell entry.

References