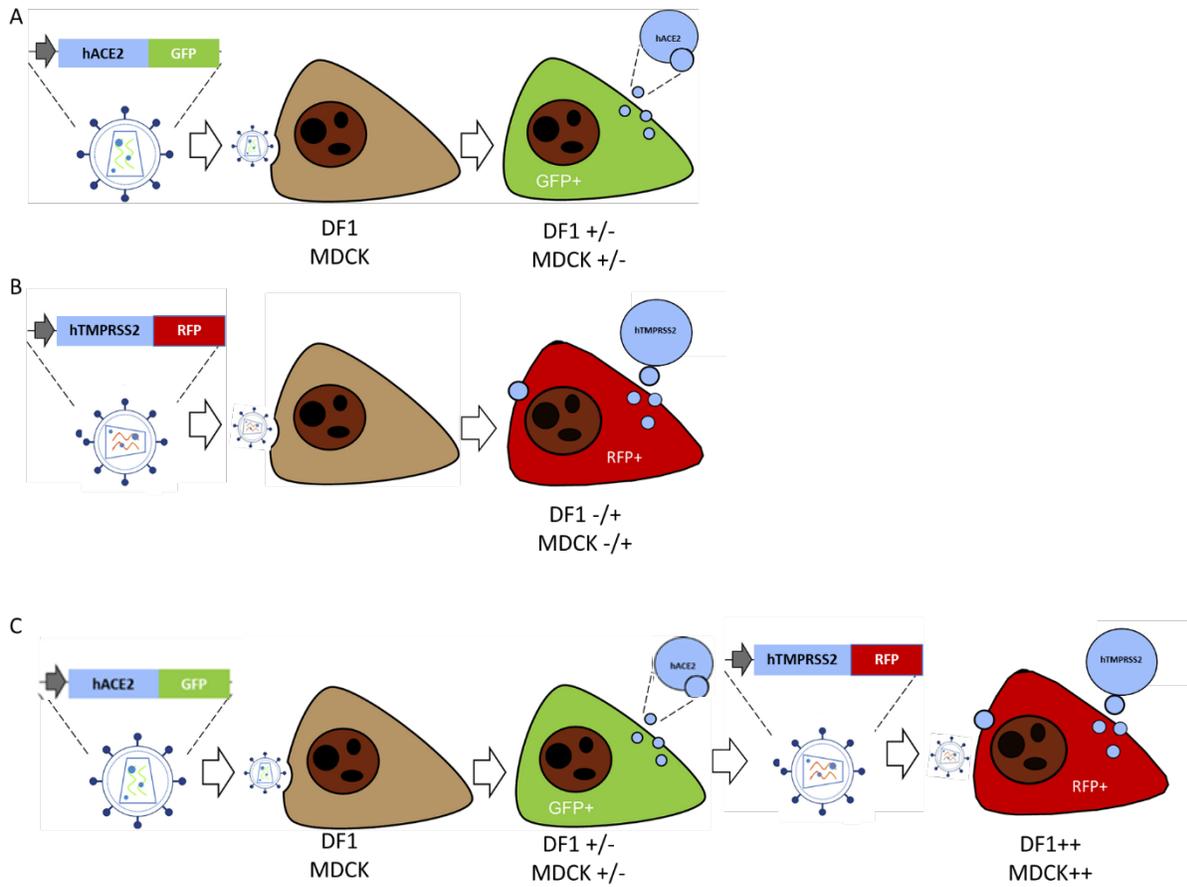
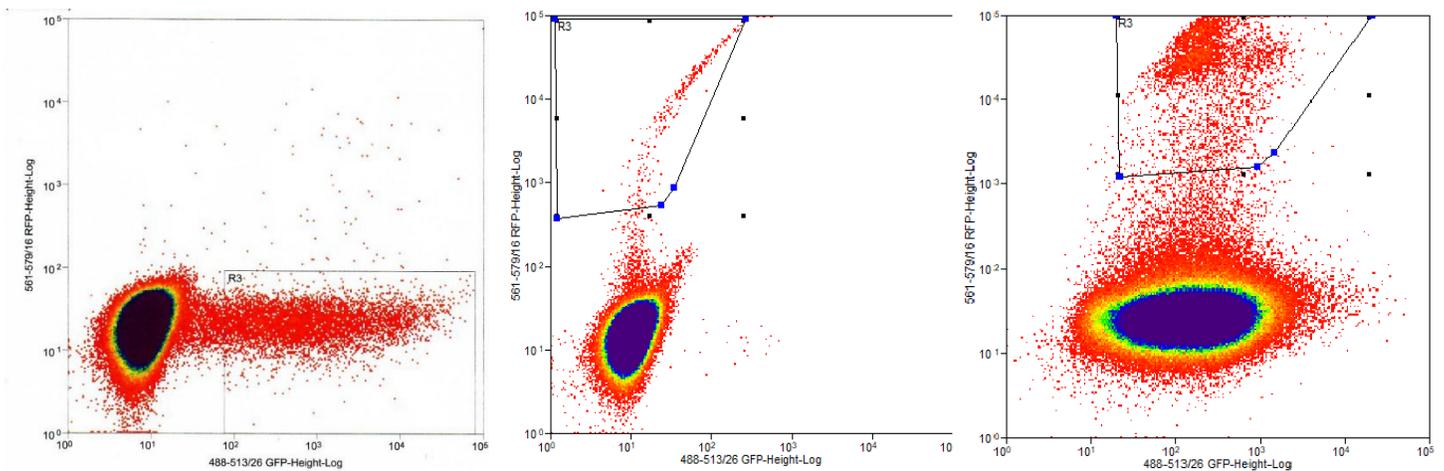


Supplemental Figure 1. Schematic representation of approach to generate transgenic cell lines expressing human or animal ACE2 (A) or TMPRSS2 (B), or both (C).



Supplemental Figure 2. FACS purification scheme for transgenic cells expressing ACE2 only (+/-) with GFP gating, TMPRSS2 only (-/+) with RFP gating, or both genes (+/+). Shown are results of purifications of DF1 cells expressing human ACE2 and TMPRSS2 genes. Similar results were obtained with all cell lines constructed.



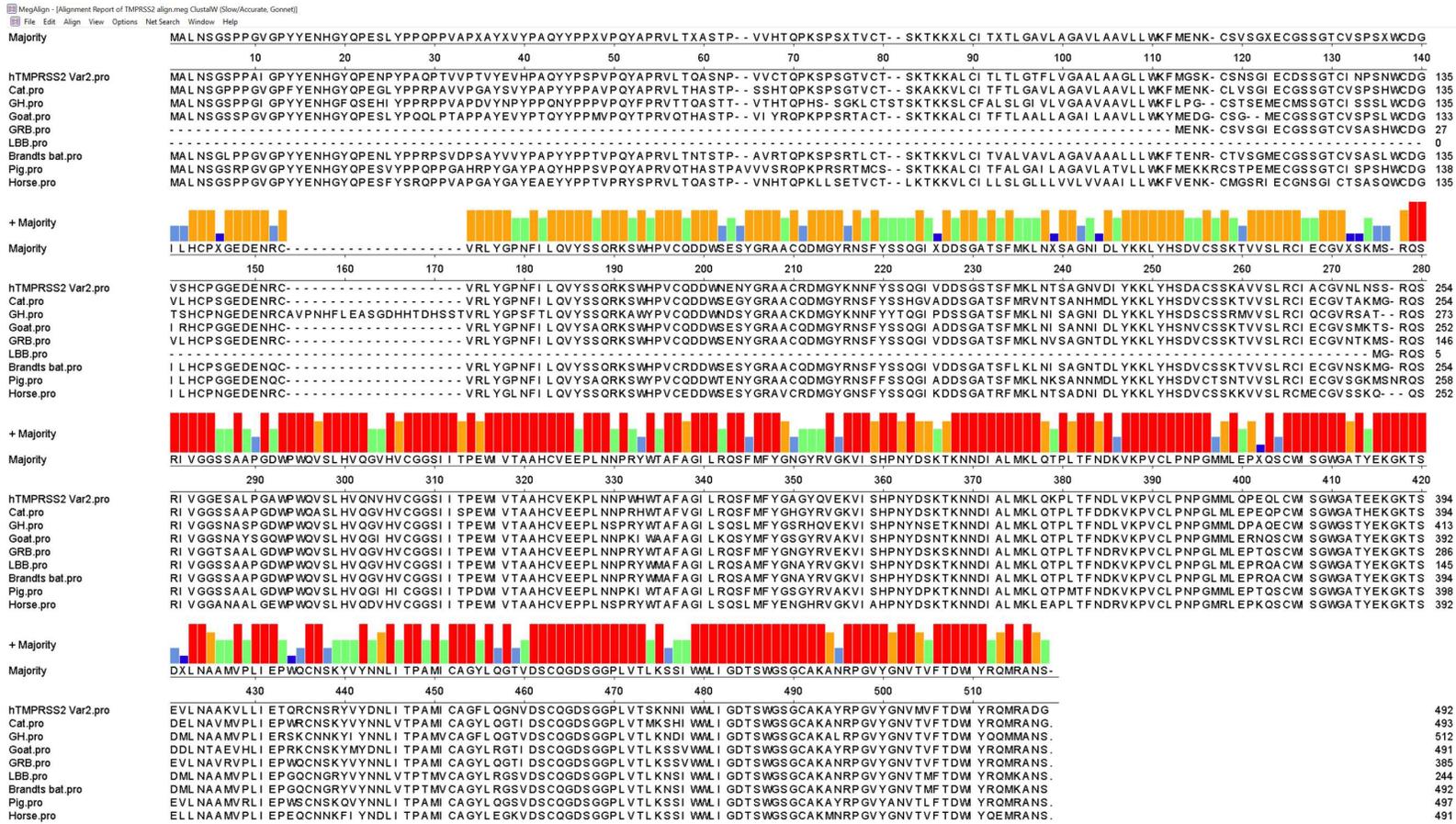
ACE2 + / TMPRSS2 -

ACE2 - / TMPRSS2 +

ACE2 +/- / TMPRSS2 +

Supplemental Figure 3. Amino acid (A) and architectural comparison (B) of TMPRSS2 between human and bat species demonstrating differences at N-terminal section.

A.



B.

Species	Gene	Architecture	aa
<input type="checkbox"/> <i>Homo sapiens</i> human	TMPRSS2 transmembrane serine protease 2		492
<input type="checkbox"/> <i>Hipposideros armiger</i> great roundleaf bat	TMPRSS2 transmembrane serine protease 2		384
<input type="checkbox"/> <i>Myotis lucifugus</i> little brown bat	TMPRSS2 transmembrane serine protease 2		243

Supplemental Table 1. List of genes and accession numbers used in this study.

<u>Species:</u>	<u>ACE2</u>	<u>TMPRSS2</u>
Human	NM_021804.1	NM_005656.4
Cat	NM_001039456.1	XM_023238709.1
Goat	NM_001290107.1	XM_005675629.3
Golden Hamster	XM_005074209.3	XM_013116227.3
Great Roundleaf Bat	XM_019667391.1	XM_019626040.1
Horse	XM_001490191.5	XM_005606160.3
Little Brown Bat	XM_023753670.1	XM_006104440.3
Pig	NM_001123070.1	XM_021071009.1

867 Supplemental Table 2. List of primers used in this study

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<u>Primers used in this study:</u>	<u>Forward</u>	<u>Reverse</u>
ACE2 Universal	CTCTTTCTGGCTCCTTCTCAGCTT	TCCAAGAGCTGATTTTAGGCTTATCC
Human ACE2	CTAGCTGTCAAGCTCTTCCTGGCT	GGATCCTAAAAGGAGGTCTGAACATCATCA
Chicken ACE2	ACGCTAGCCGCTTCTCACTAGC	AGCCAATGGATCTGCCAGAA
TMPRSS2 Universal	ATGGCTTTGAACTCAGGGTC	CTGTTTGCCCTCATTTGTCGATA
Human TMPRSS2	GGAAAACCCCTATCCCGCAC	GAATCGACGTTCCCCTGCAG
Chicken TMPRSS2	TGTTACCAGAGGACCTCCGC	TCTGCCAGGCCACAAGTAGG
Bat TMPRSS2	CAGGGATTTTGAGACAATCTTTCAT	CAAAGTGACCAGAGGACCG

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