

Extended Data Fig. 1. Sequence alignment of ACE2 proteins (1-420) from human, macaca, hamster and mouse. Amino acid residues of human ACE2 that are experientially shown to interact with the receptor binding domain (RBD) of SARS-CoV-2 are denoted by * (<https://www.biorxiv.org/content/10.1101/2020.02.19.956235v1>). Amino acid residues that are important for the interaction between human ACE2 and RBD of SARS-CoV are highlighted in red boxes (<https://doi-org.eproxy.lib.hku.hk/10.1038/sj.emboj.7600640>).

