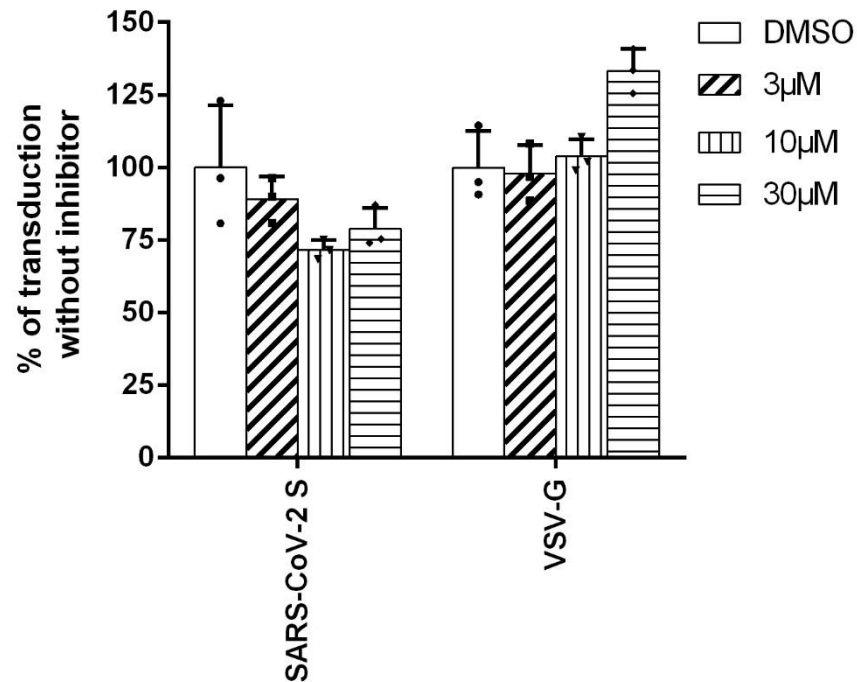


Supplementary information

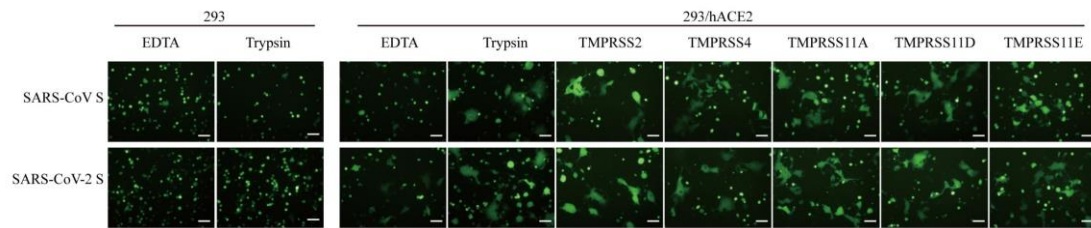
**Characterization of spike glycoprotein of SARS-CoV-2 on virus entry
and its immune cross-reactivity with SARS-CoV**

Ou *et al.*



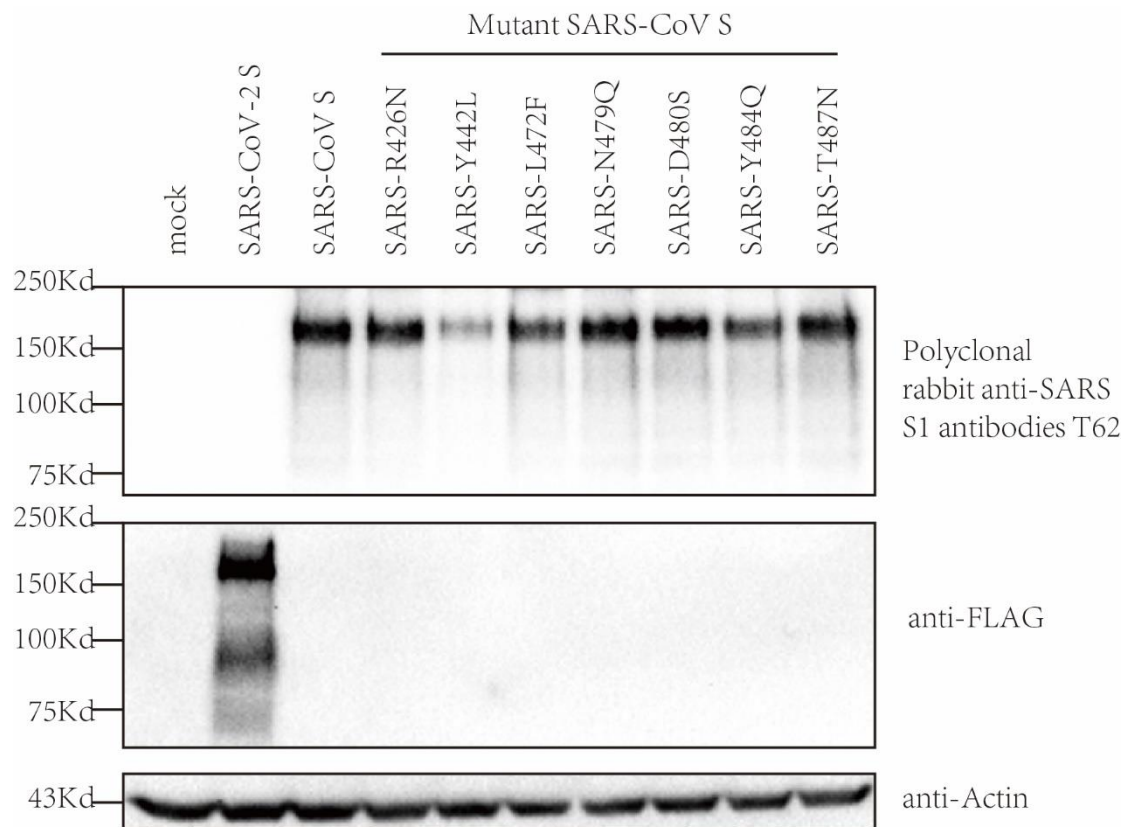
Supplementary Figure 1. Effect of entry of SARS-CoV-2 S pseudovirions on 293/hACE2 by TRPML1 inhibitor.

HEK 293/ACE2 cells were transduced with either SARS-CoV-2 S pseudovirions or VSV-G pseudovirions in the presence of 3, 10, 30μM 130. The luciferase activity were measured at 40 hrs post inoculation. The experiments were done triplicate and repeated three times.



Supplementary Figure 2. Effect of type II membrane serine proteases (TMPRSS) on SARS-CoV-2 S protein mediated cell-cell fusion.

HEK-293T cells transiently expressing CoV S glycoprotein and eGFP were lifted with either trypsin (0.25%) or 1mM EDTA, then overlaid on 293/hACE2 cells transiently expressing type II membrane serine protease (TMPRSS) 2, 4, 11A, 11D, or 11E. After 4 hrs incubation, images of syncytia were captured.



Supplementary Figure 3. Western blot analysis of expression of SARS-CoV mutant S proteins.

Top panel: polyclonal rabbit anti-SARS S1 antibodies T62; middle panel: anti-FLAG
bottom panel: anti-actin.

Supplementary Table 1. Primers used in this study.

Constructs used in mutagenesis study	Forward primer sequence (5'-3')	Reverse primer sequence (5'-3')
SARS-CoV-2 S	CCGAAGCTTCACCATGTTC GTTTCCTTGTTTC	CCGGGATCCGCAACATGATCCGCAA GAGCAGC
SARS-CoV S- R426N	GTGCTGGCTTGGAACACTA ACAACATTGATGCTACCTC	GAGGTAGCATCAATGTTGTTAGTGT TCCAAGCCAGCAC
SARS-CoV S- Y442L	CTACAACACTACAAGTATCGT CTGCTGCGTCATGGTAAGC TG	CAGCTTACCATGACGCAGCAGACG ATACTTGTAGTTGTAG
SARS-CoV S- L472F	CTTGCACTCCTCCAGCTTT CAACTGCTACTGGCCTC	GAGGCCAGTAGCAGTTGAAAGCTG GAGGAGTGCAAG
SARS-CoV S- N479Q	CTGCTACTGGCCTCTGCAG GATTACGGTTTCTACAC	GTGTAGAAACCGTAATCCTGCAGA GGCCAGTAGCAG
SARS-CoV S- D480S	CTACTGGCCTCTGAACTCT TACGGTTTCTACACC	GGTGTAGAAACCGTAAGAGTTCAG AGGCCAGTAG
SARS-CoV S- Y484Q	GAACGATTACGGTTTCCAG ACCACTACCGGCATTGG	CCAATGCCGGTAGTGGTCTGGAAA CCGTAATCGTTC
SARS-CoV S- T487N	GTTTCTACACCACTAACGG CATTGGCTACC	GGTAGCCAATGCCGTTAGTGGTGTA GAAAC

Supplementary Table 2. Codon-optimized DNA sequences of spike gene.

SARS-CoV-2 S	ATGTTTCGTTTTTCCTTGTCTGTTGCCTCTCGTTAGTAGCCAATGCGTCAAC CTTACTACTAGAACCCAGCTCCCTCCAGCATATACCAACTCTTTCACCAGG GGCGTATATTACCCGGACAAAGTGTTCCGCTCAAGTGTGCTGCATTCTAC GCAGGACCTTTTCTTGCCCTTTTTTCAGTAATGTTACTTGGTTTCATGCTATC CATGTGTCTGGAACATAACGGAACCAAGCGCTTTGACAACCCCGTCCTCC CTTTCAACGATGGCGTGTACTTCGCTTCCACGGAAAAGTCAAACATAATT CGCGGCTGGATCTTTGGTACAACACTCGACTCAAAGACGCAGAGCCTGC TGATCGTTAATAACGCTACAAATGTTGTGATAAAGGTGTGTGAATTTCAAGT TCTGCAATGATCCCTTCCTGGGTGTGTACTACCATAAGAATAACAAGAGC TGGATGGAATCCGAATTTAGGGTTTACAGTTCCGCTAACAACCTGCACATT CGAATACGTAAGCCAGCCATTTCTTATGGATCTTGAGGGCAAGCAAGGAA ACTTCAAGAACTTGAGGGGAGTTCGTGTTCAAAAATATCGACGGCTATTTT AAGATATATAGCAAGCACACTCCAATAAACTTGGTGCGCGACCTGCCCCA GGGATTCTCTGCTCTGGAGCCCCTGGTGGATCTGCCCATTGGAATAAACA TAACTCGCTTTCAAACACTGCTCGCCCTGCATCGCAGTTACCTCACCCCT GGTGATAGTAGTTCAGGATGGACAGCAGGAGCCGCCGCATACTACGTCG GCTACCTGCAGCCTAGGACCTTCTTGCTGAAGTACAACGAGAACGGTAC AATAACTGACGCTGTGGACTGCGCTCTGGACCCTCTGTCCGAGACGAAG TGCACCCTGAAGAGCTTTACTGTTGAAAAAGGCATTTACCAAACCAGCA ACTTCCGCGTCCAGCCAACCGAGAGCATCGTCAGATTTCCCAACATTACA AATCTGTGTCCCTTCGGCGAGGTGTTCAACGCCACACGCTTCGCTTCAGT GTACGCATGGAACCGCAAGCGCATATCTAACTGCGTCGCGGATTATTCTG TCCTCTACAACCTCCGCCTCTTTCTCCACCTTCAAGTGCTACGGAGTGTC CCGACTAAGCTGAACGATCTCTGCTTTACCAACGTCTACGCGGACTCCTT CGTGATAAGAGGTGATGAAGTGAGACAAATAGCCCCAGGTCAGACTGGT AAGATCGCAGATTACAACCTACAAATTGCCTGATGATTTCACTGGTTGCGT TATCGCGTGGAACCTCTAATAACCTCGATTCTAAGGTCGGTGGTAACCTACA ATTACCTGTACCGCTTGTTTAGGAAGTCAAACCTGAAGCCTTTCGAGAGG GATATTTCAACCGAAATCTATCAAGCGGGTTCAACACCGTGTAACGGTGT GGAAGGATTTAACTGCTACTTCCCCCTGCAGTCTTACGGATTCCAGCCAA CCAATGGCGTGGGTACCAACCTTATCGCGTGGTGGTTCTGAGTTTCGAA CTGTTGCACGCTCCCGCCACGGTATGCGGTCCCAAGAAGAGCACTAACT TGGTGAAGAATAAGTGCGTGAATTTCAATTTCAATGGCCTCACTGGAACCT GGAGTGCTGACCGAATCCAATAAGAAGTTCTTGCCCTTCCAGCAGTTTCGG AAGAGACATTGCTGACACAACCGACGCGGTGCGCGATCCTCAGACTCTG GAGATATTGGACATTACACCATGTTCTTTCGGCGGTGTGTCTGTCATTACT CCGGGCACGAATACTAGCAACCAGGTAGCCGTGCTGTACCAAGACGTGA ATTGCACAGAGGTTCCCGTCGCAATTCACGCTGACCAGCTGACCCCCAC GTGGAGGGTTTACAGCACTGGTAGTAACGTCTTCCAGACGAGAGCCGGT TGCTTGATCGGAGCGGAACATGTGAATAACTCCTACGAGTGCGACATCCC CATCGGAGCCGGTATATGCGCCTCTTATCAGACACAACTAACTCACCCA GGAGAGCCCGCAGTGTGGCTTCTCAAAGCATTATAGCATACACTATGTCT CTTGGTGCCGAAAATTCCGTGGCCTATTCTAACAATTCAATCGCCATCCCA
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MERS-CoV S	ATGATTCACTCCGTGTTCTCCTGATGTTCTGCTCACTCCTACCGAAAGT TATGTGGATGTGGGTCCCGATAGCGTCAAGTCCGCATGCATCGAGGTGGA TATTCAGCAGACATTCTTTGACAAGACCTGGCCTCGGCCAATCGATGTCT CCAAAGCCGACGGCATCATTACCCTCAGGGACGCACCTATAGCAACATC ACTATTACATACCAGGGCCTGTTCCCATATCAGGGCGACACGGAGATATG TACGTGTATTCTGCCGGACATGCTACTGGCACCACTCCTCAGAAACTGTT CGTGGCAAATTACTCCCAGGACGTCAAGCAGTTCGCCAACGGTTTTGTG GTCAGGATCGGCGCCGCTGCAAACCTCTACCGGCACTGTGATCATTAGCCC TTCTACTTCAGCCACAATCAGAAAAATTTACCCAGCTTTTATGCTGGGCTC CAGCGTGGGAAACTTCTCCGATGGCAAGATGGGCCGTTTCTTTAATCACA CTCTGGTCCTGCTCCCTGACGGATGCGGTACACTGCTCCGCGCTTTCTAC TGTATCCTGGAGCCCAGGAGCGGAAATCACTGCCCTGCCGGTAACTCTTA CACCTCATTCGCCACCTATCATACTCCAGCTACAGACTGTTCTGATGGCAA TTATAACAGAAATGCCAGTCTGAACTCCTTCAAGGAATACTTTAATCTCCG TAACTGCACATTCATGTACACCTATAATATCACTGAGGACGAAATTCGGA GTGGTTCGGTATCACACAGACCGCTCAGGGCGTGCACCTGTTTTCTTCAC GCTACGTCGATCTCTATGGCGGAAACATGTTCCAGTTTGCCACTCTGCCC GTGTACGACACAATCAAGTACTATTCCATCATTCCCTCATAGCATCCGGTCT ATTCAGTCAGACCGCAAGGCTTGGGCCGCTTTCTACGTGTATAAACTGCA GCCTCTCACCTTCCTGCTCGATTTTTTCCGTGGACGGATACATCCGCAGGG CCATTGATTGCGGTTTCAACGACCTGAGCCAGCTCCACTGTAGTTATGAA TCCTTCGATGTGGAGAGTGGTGTGTACTCCGTCAGTTCCTTTGAGGCTAA GCCCAGTGGTTCGTCGAGCAGGCAGAAGGCGTGGAGTGCAGCTTC AGCCCTCTGCTCTCTGGCACACCCCCTCAGGTGTATAACTTCAAAAGGCT GGTCTTTACCAACTGTAATTACAACCTGACTAAGCTGCTCAGCCTCTTCTC TGTGAACGACTTTACCTGCTCACAGATCAGTCCCGCAGCCATTGCTTCAA ATTGTTACAGCTCTCTGATCCTCGATTACTTCAGTTATCCTCTGTCCATGAA AAGCGACCTCTCTGTGTCAAGTGCTGGCCCAATCTCCAGTTTAATTACA AGCAGAGCTTCTCTAACCCTACCTGCCTGATTCTCGCCACTGTGCCACAC AACCTGACAACCATCACAAAGCCCCCTCAAGTACAGCTACATCAACAAGT GCTCAAGGCTGCTCAGTGACGATAGAACCGAAGTGCCACAGCTGGTCAA TGCCAACCAGTACTCCCCATGTGTGAGCATCGTCCCCTCTACCGTGTGGG AAGACGGAGATTACTATAGGAAGCAGCTGTCCCCCTGGAGGGTGGAGG ATGGCTGGTGGCATCAGGTAGTACAGTCGCCATGACCGAGCAGCTCCAG ATGGGCTTCGGAATCACTGTGCAGTACGGCACTGATACAAATTCCGTCTG CCCAAAGCTGGAATTTGCTAACGACACAAAAATCGCAAGCCAGCTGGGC AATTGCGTGGAGTACTCCCTCTATGGAGTGAGCGGTCTGTGGCGTCTTCCA GAACTGTACTGCAGTGGGAGTCAGGCAGCAGAGGTTTGTGTACGATGCT TATCAGAACCTGGTCGGCTACTATAGCGACGATGGAAATTACTATTGCCTC CGCGCTTGTGTGTCAGTCCCCGTGAGTGTATCTACGACAAGGAAACCA AAACTCACGCAACCCTGTTTCGGCTCTGTGGCCTGCGAGCATATTTCCAGC ACTATGTCACAGTACTCCAGGAGCACAAGAAGTATGCTGAAGAGACGTG ACTCCACATACGGACCCCTCCAGACCCCTGTGGGTTGCGTCCTGGGCCTC

	GTGAACTCTTCACTGTTTCGTGCGAAGATTGCAAGCTGCCTCTCGGACAGTC CCTGTGTGCCCTCCCAGACACCCCCAGCACACTGACCCACGTTCAAGTG CGGAGTGTCCCCGGCGAGATGCGTCTGGCATCTATCGCCTTCAATCACCC AATTCAGGTGGATCAGCTGAACAGTTCCTACTTTAAGCTCTCTATCCCCAC AAACTTCTCATTTGGCGTGACCCAGGAGTACATCCAGACTACAATTCAGA AGGTGACCGTGGACTGCAAACAGTACGTGTGTAACGGATTCCAGAAATG CGAACAGCTGCTCCGCGAGTACGGCCAGTTTTGTAGCAAGATCAATCAG GCACTGCATGGCGCCAACCTCCGTCAGGACGATAGCGTGCGGAACCTGT TCGCTTCTGTCAAGAGCTCTCAGTCAAGTCCTATCATTCCAGGTTTCGGT GGCGACTTTAACCTGACCCTGCTCGAACCAGTGTCTATCTCAACTGGCAG TCGCTCCGCTAGGAGCGCAATCGAGGATCTGCTCTTCGACAAGGTGACC ATTGCCGACCCCGGATACATGCAGGGTTATGACGATTGCATGCAGCAGGG ACCTGCCTCTGCTAGGGATCTGATCTGTGCTCAGTACGTGGCAGGTTATA AGGTCTTGCCACCCCTCATGGACGTGAACATGGAAGCTGCATACACCTCC AGCCTGCTCGGTTCCATCGCCGGAGTCGGTTGGACTGCTGGCCTGTCTTC ATTCGCCGCTATCCCCTTTGCTCAGTCTATTTTCTACAGACTGAACGGCGT GGGAATCACCCAGCAGGTCTGTGTCAGAGAATCAGAAGCTCATTGCCAAT AAATTCAACCAGGCCCTGGGAGCTATGCAGACTGGTTTTACCACTACAAA CGAGGCCTTCCAGAAAGTGCAGGACGCGAGTCAACAATAACGCACAGGC CCTGAGCAAGCTCGCTTCCGAGCTGAGCAACACCTTCGGAGCAATCAGC GCCTCTATTGGTGATATCATTAGAGGCTGGACGTGCTCGAACAGGATGC CCAGATCGACAGACTGATTAATGGCCGTCTGACCACTCTCAACGCTTTCG TGGCACAGCAGCTGGTCCGGTCAGAAAGTGCAGCCCTGTCCGCCAGCT CGCTAAGGACAAAGTGAACGAGTGCCTCAAGGCTCAGTCTAAAAGATCA GGATTTTGTGGTCAGGGCACCCACATCGTGAGCTTCGTGGTCAATGCCCC AAACGGCCTGTACTTTATGCACGTCGGATACTATCCCTCAAACCATATCGA GGTCGTGAGTGCTTACGGCCTGTGCGATGCTGCAAATCCTACAAACTGTA TTGCCCCAGTGAACGGATACTTCATCAAGACCAACAACACTAGAATTGTG GACGAATGGTCTTACACCGGCAGTTCCTTCTATGCACCCGAGCCTATCAC TTCCTGAAACACAAAGTACGTGGCCCCCAGGTCACTTATCAGAATATCA GTACAAACCTGCCTCCACCACTGCTCGGCAACTCCACCGGAATTGACTTC CAGGATGAACTGGACGAGTTCTTTAAGAATGTGTCCACAAGCATCCCTAA CTTTGGAAGCCTGACCCAGATTAACACAACCTGCTCGATCTCACTTACG AGATGCTGAGCCTCCAGCAGGTGGTCAAGGCTCTGAATGAATCTTACATC GACCTGAAAGAGCTCGGTAATTATACATACTATAACAAGTGGCCCTGGTA CATCTGGCTGGGTTTCATTGCAGGACTCGTGGCTCTGGCACTCTGCGTCT TCTTTATCCTGTGCTGTACCGGATGCGGTACTAATTGTATGGGCAAGCTGA AATGTAACCGCTGTTGCGACCGGTACGAGGAGTACGACCTGGAGCCCCA CAAGGTGCACGTGCAC
MHV S	ATGCTGTTTCGTGTTTCATCCTGTTTCCTGCCCAGCTGCCTGGGCTACATCGGC GACTTCCGGTGCATCCAGCTGGTGAACAGCAACGGAGCAAACGTGAGC GCACCAAGCATCAGCACCGAGACCGTGGAGGTGAGCCAGGGACTGGGC ACCTACTACGTGCTGGACAGAGTGACCTGAACGCCACCCTGCTGCTGA CCGGCTACTACCCCGTGGACGGCAGCAAGTTCAGGAACCTGGCCCTGAC

	CGGCACCAACAGCGTGAGCCTGAGCTGGTTCCAGCCCCCTACCTGAGC CAGTTCAACGACGGCATCTTCGCCAAGGTGCAGAACCTGAAGACCAGCA CCCCAGCGGGCGCCACCGCCTACTTCCCCACCATCGTGATCGGCAGCCTG TTCGGCTACACCAGCTACACCGTGGTCATCGAGCCCTACAACGGCGTGAT CATGGCCAGCGTGTGCCAGTACACCATCTGCCAGCTGCCCTACACCGACT GCAAGCCCCAACCAACGGCAACAAGCTGATCGGCTTCTGGCACACCGA CGTGAAGCCCCCATCTGCGTGCTGAAGCGCAACTTCACCCTGAACGTG AACGCCGACGCCTTCTACTTCCACTTCTACCAGCACGGCGGCACCTTCTA CGCCTACTACGCCGACAAGCCCAGCGCCACCACCTTCCTGTTACGCGTGT ACATCGGCGACATCCTGACCCAGTACTACGTGCTGCCCTTCATCTGCAAC CCAACCGCAGGCAGCACCTTCGCACCAAGGTACTGGGTGACCCCACTGG TGAAGAGACAGTACCTGTTCAACTTCAACCAGAAGGGCGTGATCACCAG CGCCGTGGACTGCGCCAGCAGCTACACCAGCGAGATCAAGTGCAAGACC CAGAGCATGTGCCCAGCACCGGAGTGTACGAGCTGAGCGGATACACCG TGCAGCCAGTGGGAGTGGTGTACAGGAGGGTGGCAAACCTGCCAGCATG CAACATCGAGGAGTGGCTGACCGCACGGAGCGTGCCAGCCCCCTGAAC TGGGAGAGAAAGACCTTCAGAACTGCAACTTCAACCTGAGCAGCCTGC TGAGATACGTGCAGGCCGAGAGCCTGTTCTGCAACAACATCGACGCCAG CAAGGTGTACGGCCGCTGCTTCGGCAGCATCAGCGTGGACAAAGTTCGCC GTGCCCCGGAGCAGACAGGTGGACCTGCAGCTGGGCAAACAGCGGCTTC CTGCAGACCGCCAACTACAAGATCGACACCGCCGCCACCAGCTGCCAGC TGCACTACACCCTGCCCAAGAACAACGTGACCATCAACAATCACAACCC CAGCAGCTGGAACAGGCGCTACGGCTTCAACGACGCCGGCGTGTTCCGGC AAGAACCAGCACGACGTGGTGTACGCCCAGCAGTGCTTCACCGTGCGGA GCAGCTACTGCCCATGCGCACAGCCAGACATCGTGAGCCCATGCACCAC CCAGACCAAGCCAAAGAGCGCCTTCGTGAACGTGGGCGACCACTGCGA GGGACTGGGAGTGCTGGAGGACAACCTGCGGAAACGCCGACCCCCACAA GGGATGCATCTGCGCCAACAACAGCTTCATCGGCTGGAGCCACGACACC TGCCTGGTGAACGACAGATGCCAAATCTTCGCCAACATCCTGCTGAACG GCATCAACAGCGGCACCACCTGCAGCACCGACCTGCAGCTGCCCAACAC CGAGGTGGTGACCGGCATCTGCGTGAAGTACGACCTGTACGGCATCACC GGCCAGGGCGTGTTCAAGGAGGTGAAGGCCGACTACTACAACAGCTGG CAGACCCTGCTGTACGACGTGAACGGCAACCTGAACGGCTTCAGGGACC TGACCACCAACAAGACCTACACCATCCGCAGCTGCTACAGCGGCCGGGT GAGCGCAGCATTCACAAAGGACGCACCAGAGCCAGCCCTGCTGTACAGA AACATCAACTGCAGCTACGTGTTTCAGCAACAACATCAGCAGGGAGGAGA ACCCCCTGAACTACTTTCGACAGCTACCTGGGATGCGTGTTGAACGCAGA CAACAGGACCGACGAGGCCCTGCCAACTGCGACCTGAGGATGGGAGC AGGACTGTGCGTGGACTACAGCAAGAGCAGAAGGGCCCCACCGCAGCGT GAGCACCGGCTACAGACTGACCACCTTCGAGCCCTACACCCCCATGCTG GTGAACGACAGCGTGCAGAGCGTGGACGGCCTGTACGAGATGCAGATCC CCACCAACTTCACCATCGGCCACCACGAGGAGTTCATCCAGACCCGGAG CCCCAAGGTGACCATCGACTGCGCCGCCTTCGTGTGCGGCGACAACACC GCCTGCAGACAGCAGCTGGTGGAGTACGGCAGCTTCTGCGTGAACGTGA
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	ACGCCATCCTGAACGAGGTGAACAACCTGCTGGACAACATGCAGCTGCA GGTGGCCAGCGCACTGATGCAGGGAGTGACCATCAGCAGCAGGCTGCCA GACGGAATCAGCGGCCCCATCGACGACATCAACTTCAGCCCCCTGCTGG GATGCATCGGCAGCACCTGCGCAGAGGACGGAAACGGACCAAGCGCAA TCCGCGGCCGGAGCGCCATCGAGGACCTGCTGTTTCGACAAGGTGAAGCT GAGCGACGTGGGCTTCGTGGAGGCCTACAACAACCTGCACCGGCGGCCA GGAGGTGAGAGACCTGCTGTGCGTGCAGAGCTTCAACGGCATCAAGGTG CTGCCACCAGTGCTGAGCGAGAGCCAGATCAGCGGATACACCACCGGAG CAACCGCTGCCGCAATGTTCCCACCCTGGAGCGCTGCCGCAGGCGTGCC CTTCAGCCTGAGCGTGCAGTACAGGATCAACGGCCTGGGCGTGACCATG AACGTGCTGAGCGAGAACCAGAAGATGATCGCCAGCGCCTTCAACAACG CCCTGGGCGCCATCCAGGACGGCTTCGACGCCACCAACAGCGCCCTGGG CAAGATCCAGAGCGTGGTGAACGCCAACGCCGAGGCCCTGAACAACCT GCTGAACCAGCTGAGCAACAGGTTCGGAGCAATCAGCGCCAGCCTGCA GGAGATCCTGACCCGGCTGGAGGCAGTGGAGGCAAAGGCACAGATCGA CAGACTGATCAACGGCAGGCTGACCGCCCTGAACGCCTACATCAGCAAG CAGCTGAGCGACAGCACCCCTGATCAAGGTGAGCGCCGCCAGGCCATCG AGAAGGTGAACGAGTGCGTGAAGAGCCAGACCACCCGCATCAACTTCT GCGGCAACGGCAACCACATCCTGAGCCTGGTGCAGAACGCCCCCTACGG CCTGTACTTCATCCAATTACGCTACGTGCCAATCAGCTTCACCACCGCAA ACGTGAGCCCAGGACTGTGCATCAGCGGCGACAGGGGCCTGGCCCCAA AGGCAGGCTACTTCGTGCAGGACGACGGCGAGTGGAAGTTCACCGGCA GCAGCTACTACTACCCCGAGCCCATCACCGACAAGAACAGCGTGATCATG AGCAGCTGCGCCGTGAACTACACCAAGGCCCCCGAGGTGTTCTGAACA CCAGCATCCCCAACCTCCCGACTTCAAGGAGGAGCTGGACAAGTGGTT CAAGAACCAGACCAGCATCGCCCCGACCTGAGCCTGGACTTCGAGAAG CTGAACGTGACCCTGCTGGACCTGACCTACGAGATGAACAGAATCCAGG ACGCCATCAAGAAGCTGAACGAGAGCTACATCAACCTGAAGGAAGTGG GCACCTACGAGATGTACGTGAAGTGGCCCTGGTACGTGTGGCTGCTGATC GGCCTGGCCGGAGTGGCAGTGTGCGTGCTGCTGTTCTTCATCTGCTGCTG CACCGGCTGCGGCAGCTGCTGCTTCAAGAAGTGCGGCAACTGCTGCGAC GAGTACGGCGGCCACCAGGACAGCATCGTGATCCACAACATCAGCAGCC ACGAGGAC
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