LOCUS pJFRC7-PAT3-Halo(V2)-\_C 9839 bp ds-DNA circular 03-JUN-2021

DEFINITION .

FEATURES Location/Qualifiers

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CDS 6232..6324

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CDS 7213..7923

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polyA\_signal complement(8433..8624)

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121 TTGTTTGCCG GATCAAGAGC TACCAACTCT TTTTCCGAAG GTAACTGGCT TCAGCAGAGC

181 GCAGATACCA AATACTGTTC TTCTAGTGTA GCCGTAGTTA GGCCACCACT TCAAGAACTC

241 TGTAGCACCG CCTACATACC TCGCTCTGCT AATCCTGTTA CCAGTGGCTG CTGCCAGTGG

301 CGATAAGTCG TGTCTTACCG GGTTGGACTC AAGACGATAG TTACCGGATA AGGCGCAGCG

361 GTCGGGCTGA ACGGGGGGTT CGTGCACACA GCCCAGCTTG GAGCGAACGA CCTACACCGA

421 ACTGAGATAC CTACAGCGTG AGCTATGAGA AAGCGCCACG CTTCCCGAAG GGAGAAAGGC

481 GGACAGGTAT CCGGTAAGCG GCAGGGTCGG AACAGGAGAG CGCACGAGGG AGCTTCCAGG

541 GGGAAACGCC TGGTATCTTT ATAGTCCTGT CGGGTTTCGC CACCTCTGAC TTGAGCGTCG

601 ATTTTTGTGA TGCTCGTCAG GGGGGCGGAG CCTATGGAAA AACGCCAGCA ACGCGGCCTT

661 TTTACGGTTC CTGGCCTTTT GCTGGCCTTT TGCTCACATG TTACCGTCGA CGATGTAGGT

721 CACGGTCTCG AAGCCGCGGT GCGGGTGCCA GGGCGTGCCC TTGGGCTCCC CGGGCGCGTA

781 CTCCACCTCA CCCATCTGGT CCATCATGAT GAACGGGTCG AGGTGGCGGT AGTTGATCCC

841 GGCGAACGCG CGGCGCACCG GGAAGCCCTC GCCCTCGAAA CCGCTGGGCG CGGTGGTCAC

901 GGTGAGCACG GGACGTGCGA CGGCGTCGGC GGGTGCGGAT ACGCGGGGCA GCGTCAGCGG

961 GTTCTCGACG GTCACGGCGG GCATGTCGAC AAGCCGAACA TATGGGCGCG CCTAGTATGT

1021 ATGTAAGTTA ATAAAACCCA TTTTTGCGGA AAGTAGATAA AAAAAACATT TTTTTTTTTT

1081 ACTGCACTGG ATATCATTGA ACTTATCTGA TCAGTTTTAA ATTTACTTCG ATCCAAGGGT

1141 ATTTGATGTA CCAGGTTCTT TCGATTACCT CTCACTCAAA ATGACATTCC ACTCAAAGTC

1201 AGCGCTGTTT GCCTCCTTCT CTGTCCACAG AAATATCGCC GTCTCTTTCG CCGCTGCGTC

1261 CGCTATCTCT TTCGCCACCG TTTGTAGCGT TACGTAGCGT CAATGTCCGC CTTCAGTTGC

1321 ATTTTGTCAG CGGTTTCGTG ACGAAGCTCC AAGCGGTTTA CGCCATCAAT TAAACACAAA

1381 GTGCTGTGCC AAAACTCCTC TCGCTTCTTA TTTTTGTTTG TTTTTTGAGT GATTGGGGTG

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1981 TTCGCAGAGC TGCATTAACC AGGGCTTCGG GCAGGCCAAA AACTACGGCA CGCTCCGGCC

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2221 ACATTTGCTC AAGAACGGTG AGTTTCTATT CGCAGTCGGC TGATCTGTGT GAAATCTTAA

2281 TAAAGGGTCC AATTACCAAT TTGAAACTCA GTTTGCGGCG TGGCCTATCC GGGCGAACTT

2341 TTGGCCGTGA TGGGCAGTTC CGGTGCCGGA AAGACGACCC TGCTGAATGC CCTTGCCTTT

2401 CGATCGCCGC AGGGCATCCA AGTATCGCCA TCCGGGATGC GACTGCTCAA TGGCCAACCT

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2701 AGGAAGCGTC TGGCATTCGC CTCCGAGGCT CTAACCGATC CGCCGCTTCT GATCTGCGAT

2761 GAGCCCACCT CCGGACTGGA CTCCTTTACC GCCCACAGCG TCGTCCAGGT GCTGAAGAAG

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3121 TCGGATCGCC AAGATATGCG ACAATTTTGC CATTAGCAAA GTAGCCCGGG ATATGGAGCA

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