

ORF M A Q G R S E L S I D S D R I G L F L Y H P T V I N S F P D  
1 ATGGCCCAAGGAAGAAGTGAACCTCAATTGATTGATCGGATCGGATTATTCCTATACCACCCAACGTGCATCAATTCATTTCCCGAC

ORF D N H H H H L R H H H H H H R P K L K F E P M E M E P T T  
91 GACAACCACCACCACCCTCCGCCACCATCACCACCATCACCACCGTCCCAAGCTCAAGTTTGAACCCATGGAAATGGAACCAACCACC

ORF T R S P I K T I Q F P V N L N C T T T T T S D Q D I P T P S  
181 ACCAGGTACCCGATAAAAACCATCCAATCCAGTCAATCTCACTGACCACCACCACCCTCCGATCAAGACATTCACACGCCGTCC

ORF D H N H R T V I D E M D F F A Q N K K H D D D S K A T A T T  
271 GATCACAATCATCGCAGGTCATCGACGAGATGGACTTCTTCGCCAGAATAAAAAACATGACGATGACTCCAAGGCTACAGCCACCACC

ORF V A A D R P P P K L D F N V N T G L H L L T A N T S S D Q S  
361 GTTCCGCCGATCGACCACCTCCCAAATGGATTCAACGTTAACACTGGACTGCACCTTCTGACTGCGAACACCAGTAGTGATCAATCA

ORF M V D D G I S P N S D D K R A R T E L A V L Q A E I E R M N  
451 ATGGTGGACGATGGAATATCTCCCAACTCCGATGATAAAAGAGCTAGGACTGAGCTAGCTGTTCTTCAAGCGGAGATAGAACGAATGAAC

ORF A E N Q R L R D M L N Q V T T N Y N T L Q M H L M T L M Q Q  
541 GCGGAGAATCAACGCTCGGAGACATGCTCAATCAAGTTACAACCAATTACAACACGCTTCAAATGCATCTGATGACACTGATGCAACAA

ORF Q Q E R E H Q Q D D R N N N N M L D G K V E V D D N K K R G  
631 CAACAAGAACGAGAACATCAACAAGACGATCGGAACAATAACAACATGCTAGATGGGAAAGTGGAAAGTTGATGATAATAAGAAACGTGGT

ORF G L M V P R Q F M D L G L A A E T E E T E L S S S E G R S K  
721 GGCTAATGGTGCCTAGACAGTTCATGGACCTTGATTTAGCTGCGGAGACAGAAGAGACTGAGCTGTCTTCTCCGAGGGGAGGAGCAAA

ORF D R S G S P V N D N G E A A A S K E L C D D Q G R E E S P D  
811 GATCGGTGAGGATCGCCGTTGAATGATAATGGGGAGCGCGGCTTCGAAAGAGTTGTGTGATGATCAGGGAAGAGAGGAGAGTCCAGAT

ORF N G S S Q G W C G G G G G P S K V A R L N N S S K N V D Q A  
901 AATGGATCGAGCAAGGATGGTGTGGGGTGGTGGTGGTCTAGTAAGGTTGCTAGATTGAATAATTCTTCTAAAAATGTTGATCAAGCT

ORF T E A T M R K A R V S V R A R S E A P M I S D G C Q W R K Y  
991 ACCGAGGCCACCATGAGGAAAGCCGAGTCTCCGTACGAGCACGATCCGAGGCACCCATGATCAGTGTGGATGCCAATGGCGGAAGTAT

ORF G Q K M A K G N P C P R A Y Y R C T M A A G C P V R K Q V Q  
1081 GGGCAAAAAATGGCAAAAGGAAACCCGTCGCCCTCGTGCCTATTATCGTTGCACCATGGCTGCCGGCTGCCAGTTCGAAAGCAAGTACAG

ORF R C A E D R T I L I T T Y E G N H N H P L P P A A M A M A S  
1171 AGATGTGAGAAGACAGAACCATCCTCATTACAACCTACGAAGCAATCACAACCACCCATTGCCCTCCGGCTGCCATGGCCATGGCCTCC

ORF T T S S A A R M L L S G S M P S A D G L I N S N F L A R T L  
1261 ACAACCTCTCAGCCGCCGAATGCTCCTCTCAGGCTCAATGCCGAGTGCCGACGGCTAATTAACCTCCAATTTCTAGCCAGGACACTC

ORF L P C S S N M A T I S A S A P F P T V T L D L T Q S P N P L  
1351 CTCCCATGCTCATCAACATGGCCACAATTTAGCCCTCAGCCCATCCCTACTGTCACTCTAGACCTCACCCAAAGCCCAACCCATTA

ORF Q H F Q R P P N P F H V P F P N P Q P P A N L L P Q A F G Q  
1441 CAACACTTCAAAGGCCACCAAAACCCATTTATGTCGCATTCCTTAACCCACAACCACCAGCCAATCTACTGCCCCAAGCATTGGCCAG

ORF A L Y N Q S K F S G L Q M S Q D A Q L G H Q A V P L P P L N  
1531 GCATTGTATAACCAATCAAATTTCTGGGCTTCAAATGTCACAAGATGCCAATGGGCCACCAAGCCGTGCCACTGCCGCCACTGAAT

ORF Q Q N S L A D T V T A L T A D P N F T A A I A A A I S S L I  
1621 CAGCAGAACTCGTTGGCAGACACCGTACTGCCCTCACTGCCACCCCAACTTACCAGCAGCTATAGCGGAGCCATCTTCTCTAATT

ORF G G G N N N G N V A A T T T T S N N N G G V S S S N N N S G  
1711 GGTGGAGGTAACAATAATGGAATGTCCGGCTACCACCACCACCAGCAACAACAATGGCGGCGTTTCGAGCAGTAATAACAATAGCGGT

ORF D N N K V N N S S F M G N \*  
1801 GACAATAATAAAGTCAATAATTCAAGTTTATGGGGAATTAG

**Supplemental Figure 7. Cloned coding sequence and open reading frame (ORF) prediction of *CsWRKY29* gene of tea plants.** The putative ORF of *CsWRKY29* was predicted using NCBI Open Reading Frame Finder ([www.ncbi.nlm.nih.gov/orffinder](http://www.ncbi.nlm.nih.gov/orffinder)).