Supplementary 1. Nucleotide sequences and phylogenetic analysis of five CRL promoters.

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          -590      -580      -570      -560      -550      -540      -530      -520      -510      -500
     ..|....|....|....|....|....|....|....|....|....|....|....|....|....|....|....|....|....|....|..
L1P  CGCCACGGGGAAGGCCGTTCCAGGCCCGGTCTGTGCGCCGTGCCCCACTTGTGATGCATCATGACGCACCGCCGAAAAGTCCGTGCTGCACGACGCATGC
L2P  GGCGGCACGGCGCTTATCTGTGCTGTTTTGCGATCACGAACGCCGCCCCGCTGCGGATCCCGAGGTTCCCGAGCCTGTCTCGGCGGCTGGCGAGTGCGGG
L3P  GTATCCACGACCACTTCCGTGCATCCCCTCACTCAACAGGGGGACAATTGCCGCGTAACGTCGAGCGTTTTCGGATATGCATCGACGTGCAGCGACAATC
L4P  ACAGAGGGGTTGACAACCCTCGCATTCTGGAGTGCGACGCGGTTCCATTCGCTGCATCCCCTCGTGGTCATCTACCCAGCAGATCAAGCCGCGGTGACAA
L5P  CAGACACCCACTGCTGTATACGCCCTGTCCGTCGTGCTTCTTTTCAGATCCCCGTCCACACCGTGTGCATTTTTTGCTGGTGTTGGCAACGGATACAAAG

          -490      -480      -470      -460      -450      -440      -430      -420      -410      -400
     ..|....|....|....|....|....|....|....|....|....|....|....|....|....|....|....|....|....|....|..
L1P  CGTTGTTCTGTGTCTTGCAACGGTACACAGATCTCACCTAGCCCTGCGCGGCTACGGGGCTTTGGGTGCTTCCTAACGCAAGGCCCGCATGCTACAGGCA
L2P  GTGCAGGCAACGCCAAGATAATCGCTGGGGCCAGGCGCAGACCATCAGTGACCAAAGCGGGCAGCACACTGACTATGCCGGCGTAAACAATGCCACATGC
L3P  TTTGCATGCACCTAAACACAGCATGTGAACGAGCGCCAATCGCTTCTTGGTCCCAATGCTTGTGAGGCACGGCCCAAGGCGTCCTGATCGTCGGTTATGT
L4P  TAAAAGCTCCTGCAACCAGCTGCAGTCTTGGCCCATCTGACAGGGCACTTCTGGAGCATGTGAAGCCAACCAAGACAAACCTCATTTCTACCCATAGAGC
L5P  CCATGGGGAGTCAACAAACCCCCCAAACACTCCGGGTTCTATCAGGCACACGCAAACGGCATTTGCACCCCCTTTCCGATGATCCGCACATGTGGACCCG

          -390      -380      -370      -360      -350      -340      -330      -320      -310      -300
     ..|....|....|....|....|....|....|....|....|....|....|....|....|....|....|....|....|....|....|..
L1P  CTGGCCAGCTGCCTGGCGCATGCCTCTCCGCAAACCGTCACCCCCATACACCCACAACAGCCACCACCACGCACTTGCCGCACAAGCTTGTCGATCGCTC
L2P  GGCGGTTGGTCTGCCCCGCGACTCTCCCCCGCGACTCTCCCCCGCTTCTCGGCACGCGTAGACCAGCCTTGTCGGGGCACGCGGCGTGGACACAGACCCG
L3P  GGTGTGATGGCCCCACGGTAGGCTCCGGATCGTCGATCAGTCTCTCCGGGCCCTTCCCGCGTGCAGGACGGCGCGTTGTGTGACTGCCGGAGTTTGCAAC
L4P  CATCACAATCGACCGAGGCACTGGTCACAGAGCGGGGCGCACGCAATAGACGCCTTTGCCATTTACGAACCACACGTCGAGGTGTTGCAAAACGGACAAT
L5P  CCACCAAGCCTGCAGGACCATTAACGGCGGAGATCACTGCACCGCAGACGAGCATGTATCCGCCCGTTCCGCAGCGTGCTCTGGATAAAGGTACTCAAAG

          -290      -280      -270      -260      -250      -240      -230      -220      -210      -200
     ..|....|....|....|....|....|....|....|....|....|....|....|....|....|....|....|....|....|....|..
L1P  GGGGACCGAATGCAACGCGCGGGGTGAATGACCCCATGCTTGTTTACTGCAGCGGGCGGCTCGGCTCAGCCTGATCCCGGCGAATATGCACAGACAAGGC
L2P  GTGCCAGGCCCAACCGCACACCGCTATTCCCAGCGCACAGTGGGGTGCTGCGCGCGTTCCCCGTCCAAGCATCGTGCGCCTCGCGAACGCCGCGGCCTCG
L3P  TGTTTTTGTCCGTTTGATGTTCCAATACAGCGTTGCATTTCTCCACTGCCCCCACCCACCACCACGGCCCCCGGTCACGCTCCCCCTCCCCAACTCCGTG
L4P  GATCACGCGCCGCGGCTGTTGCCTTAAGCGCCGTAAAAGCCCGGGCTAAAGGAATGCTGACATGAACGTAGGGCCACCAGGCAAGGCGTTATGTTGCATT
L5P  TCGACAGAATCTCAGCCAGGTGGCGTACAGAGAATAGCCATACCCACCAAGCAATGCCCACCAAGCCATGTCCCACCAAGTCACATTTGGACACAAACGC

          -190      -180      -170      -160      -150      -140      -130      -120      -110      -100
     ..|....|....|....|....|....|....|....|....|....|....|....|....|....|....|....|....|....|....|..
L1P  AACGCTGTGTGCGTTAGTGAGGACAGATGCACACAGACCAACCGCGTCAATATACACTGGTCACTTGTTGTGACTGAGGCAGGAAGGCAGCGCAGGAGCC
L2P  CCATAACCACCGGCTAGTCTCCGGCGCGTGCGGGGAGGCTTGTCCGCACGTCACTGCTGCTTGGCTTTGGCCAGCATGTGGCCGCCACGCCCCGTCTGCA
L3P  TGCGACATGTGAACTCCCCTCATGCGCTGCGGGCGGGGCCGGATCAGTGCGTCTGCCGCCTGGGCAATCCGTGCGCCCCGCCATTGCCCGGACGATCCCC
L4P  GCCAGTGATGGGGCCTTACAATCAAGGTAATCCCAACTTTCCGGATGCGTACTCGTCAATGCAGCCCCCCTTGCCGCATGTTGCACTGTCCAATCCAAGT
L5P  AGTGCCAGGGTTTCACAACTGGCTCCTACCTGACGCATCTCGGCAAATGCCACTCTGTAGTTGGTCCAGACATGGCAGGCATAAACAAGCACCAGGCGGT

          -90       -80       -70       -60       -50       -40       -30       -20       -10       +1
     ..|....|....|....|....|....|....|....|....|....|....|....|....|....|....|....|....|....|....|..
L1P  GTTAATCCCGGGCATGACGATCACGGTTGGAGGTTGACCCGCTAATCCCGTTTCCATTGACATAAATAGGGAGTGGTTTCCTCCCCACTCGCTCCCCATG
L2P  CGTTGTGTGGTCTTTACTGCATTGGAGCCAGGACCGTAGTGCAACCCCGCGCCGGTCCGGGGTATAAAAGCGGAAGCACTCTCACTCCAGTCTCTCCATG
L3P  TCCGTGTGCTAACGGTTGACAACAATGCGGCCTGATCCGGACTACCGCACACCGCTCCGAGTATAAAAGCAGAAGCATTCTCACCTGCTCGCTCCCCATG
L4P  CAACAACAACTAGCTGATCGGCGGCGGGGGCTATGCGCCGCGCCTAAAATGTTCCGCTTATAACCGCGGACTCTCCACTCGCTCCAGCTCACTCCCCATG
L5P  CAGACACAAGAGCGTGGGGAGGTGACAATGCCGATCGCGGACCAGTTTACACCGCTCCGAGTATAAAAGCAGAAGCATTCTCACCTGCTCGCTCCCCATG
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Supplementary 1: Nucleotide sequences and phylogenetic analysis of five CRL promoters. Only 600-bp sequences of each promoter are shown. ORE elements, TATAA and CAAT like boxes are underlined. The nucleotides are numbered starting from the start codon as +1 (position A).