

Appendix B

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1 ACATGGGGACGTCACACTCAGCCATAGTTCACGTCACAAATCCAACTCTCATGTTCTTTCTAGTCCTGTTAATCTCTTCATCTCGCTGCTACAGATTACATGCCCAAGCAACGG
121 AGTAAACCACCAGAAAGAACTCCAGATTGAGGGGATATCGAATCGAATCGAAATGATGTCGAAAGTCGATCATGTCCTTTGCGCCTTGTGACAGTATCAGAGCTGAACCTGCCGAC
M I A K S I I V L V A L L E T V F R A E L F D SP
241 CTGGAAATGTCGGGATGACTGCGACTGCTACTTTCGCGTCACTGGGTGACAGATTGTTCCGAGAGAACCTCAGGAGATCCGACTCTGACAGATGGACTCAGCTCAAGGCTCAG
L R R N T
361 ATTTCTCAACATGAAACCCAAATGAGTGTGACTGAGATCGCGCCCTTCCGCGCAGCATCAACTGAGAGCCCTTGCAGCTGGCTGATACAAAGCTGACAAATCACCAGAGATGACTTCGCT
I L N M N A N E L T E I A F P P A D I K L R R L Q L A D N K L L T Q I T K D D P A
481 GGCTCGGGTCTCCGCTGATGACTGACTGAGAAATCAATCAAGACTGTCATCCTGATGCTTTGAGGAGACAGTCCGGGCTGATAACTGGAAATGGAGATGAATCCTCTACAG
G L G F L L E I D L S E N Q I T T V H P D A F R D S P G L I T L E M Q M N P L H
601 CCGGTCGAAGTCCATCTCTGCTGCTCTACTACTTGGATCTCAGCGAATGTAAGCTGGAAAAGCTTACACCGCAATCTTTGAGAACTTGACAACACTGAACAATGGAC
P V E G P F L L S R S L L Y L D L S E C K L E K L T P Q F F Q N L T T L N K L D
721 CTGTCTGAAATCCACTGAAGAGCTGAGTCTCCATCTCGATCCCTTGGTGTAGCTTGGAGGTTTGAACCTGAACAGATGCAACCTCATTCTTCTGATGATGCTTCAATAC
L S G R F L R E L O S S I L D P F V S L E V L R L E R C N L T F I S D D A F F I
841 ACTGAACTTTGAAATACCTGGAGATGCTGAGAACACTTAAGGTACCAACTGATGGATTACTGTGTAGCAAAATGGGAGCTCTCGAGTACTCGATCTTGSCAATCAGCCATA
T H L K Y L E I A E N N L V V E T D N I T V L A K L G R L E Y L D L R O S I
961 TCAAATCCCTCGAAGTCCCTCAACAACAACACTGGCTTAGGAGTCTTATTTAGCTGGAATAATCTCATGCGTGTGAAATATCATCTATCTTGGTGGACACCCCAACACTTC
S N L P E D A F N N N T W L R S L I L A G N N L M R L E I S S I L G D N P K H L LRR-1
1081 GAAATCTAGATCTCAACTGCTCATTTCAGTGGAGCTGCTGACTGCAAAATGCAATCCCAACTGCACAAAACCTGAAAACACTCATATTTGCTGGTAATCTTATCAGCCACTGATCTA
E Y L D L S N C H F S G R L T A N A F T N C T K L K T L I L S G N L L S A T D L
1201 GCTGATGCTTACAGCCGTTATCAAACTCCGAAACTGGGGTGAACACTGAACTTGACAAGACTCCCTGCAAACTTCCCAAACTTGAAAACCTTCAAGAGCTGGACATCTCT
A D A L F P L S K L Q K L G L K N C N L T R L P A N T F H N L E N L Q E L D I S
1321 AGGACCCACTCAGCAAGCATTACAGACTCTTCCACCCTTACTACTTGAACACTTATGATATGAGTACAGTAATCTAGGACAGATATCCGAGCAAGCTTTCAAGATGAC
S H P F L N H A F T E L L S F D T T L E R L D M S H S N L D Q L S G L T F S K R N
1441 GCATAGAACTTAAATCTGTCAGTACATGTTGACAGTCTTGTAGTCAAGCTTCCGAAACTTACACTTTAGAAACCTTAGAGCTCAACAATTTGGATGAAACCTGTTGAGT
A L R T L I L S G N M I T S D S L P F O K K I T H L E T L E L I N N C G L K S I S
1561 TCCTCTGATTTCCCGTCAATTCACCTACCCTGATTTAGAAGAGTGTAGACTTGCCTGATCCTCTAGAGATCTCTAAAGAGGAATGTTGCTTCTGTTCAATAAAACCGCTCAA
S V P F V N T Y F D L E E L R L A G N P L E I S K E G M L L P V Q I K R L K
1681 ACTCTAGACTTTCCGACTCAATCTAATACATATCTCCAGAAAACACTGAAATCTTTCATAAACAATCCCGCCCTCAATTAGCAGGCAACAATGAATAGTCAAATAAAAATG
T L D L S D C N L T Y I P P E T L K S F I N I T R L Q L A G N K L N S S N K N S
1801 TTGGAACTCTGAGAAATGTCGCCAGTGGAGTCTCCGATTTGAGTAGAAACGAACTCACTGATACCTCCAGTGTTCGATAATAACACATTTGTGAGTGTGGAACATGTA
L E F L R K F P Q L E L P L D L S R N E L T D I T P S V F D N N T L L S A V K L V
1921 AGTAATCCTTGGAAATGTTGTCATATTTGGACATGTTGGAGTGGCAATACAGTTAAAGGTGATCTGGAGTCTTATTGGTTCAACCACTGATCCAGAAAGCGATATCCACTG
S H P F K C S C H I A D H W E W A I T V K S D L S V L I S T T D E A I S T G
2041 GGAAGGAAAGAGAGAGAGGACTGTTGCTGCTTTGACTCGAAAACCTGCTGATCAGTAGATATTCAGTTGAGAAAGCCGCTCGCAAGGATTTCTGGATATTTAAACAGGAG
G R K K K K K G L L C R P D S E T S P I S K D I Q P R K K P A R E D F V D N V N R T
2161 TGCGTAGTATGAGGGAGGCGAATGGAATCGAACATCGCTGAGACCGCTAGATTTTAAAGTCAAGCGATACAGTTTAAATCAGAGGAAAGTAATAATTAGCAGSTA
W A R Y V R E A D C E S S N R L R P A R L L K S Q P I T V *
2281 GTGATTTATTCGGATGTAACAGCACTACGATTGTTGCATATTATAGACCAAGTTTGGAGACAAGGAAATTTGCTCATGGGTGAAAAAAGTGGATTACAGCTTGCATTTACT
ATGAACTTGAATGAGAATCAAGATTTGTAATGGAATGCCAACAGACATTTGTCAAATTTGGGSCCTTCCACTGTCATTTGGTTATAGACAGAAAGTTGTAATAGCAGATTTGGGTGA
2521 GAAATGAGAGATGATTTCTTCTAGAAAGTAGGACCGCTAACCTAAGAGTAGAGAGTGTGCAAGGCTGACTGAGAAACCTAGAGACCTGTTTATCGTTGAGTTGCATAATGAAATTAATAGACT
TCTTACTTTTAGAAGTGTTTACACTCACATTCAGCTATTCTATTATCATCATATATGCTCAATTTCTCTATTTTCTGTTTATTCTAATCTCATGACGAATTTAGAAATTTACT
2761 ATTGAAATGCTATGTTTATTTTAAAGATTCAAACCTCAAAACATTTTGGATAGAGTTTATATTAATGCTGATTTCCAGATATATTTTGTATGAGAAATGAAATA
2881 ATATGTGATCAGTTAGCTTACTGGTACAGTACTGATGACAAATATTGACAAATATTGACAGAGATTTGATTCAGTCTGGTCACTACTTTTTCCAAATGTTTCTGCTGAAATGCA
3001 ATTTGTCACAAATATTTTCATGGGACATTTATCTATTACAACTTATATGTTTATCATGAAATATATCATGCAACTTGTGACCTCAGGCTCAACATTTATGCTAATCTCA
3121 AATTGATCTCGAAAATATGATGACCAATCACTATTATAACTAATATGCAACCTCACTTTGTTTATGCCAAAATGATGAAATGGCTTTCTACATAAGACTTTAATACCAAAAT
3241 TTTACGTTTGTCTCAATTTGTAATTTACCTTAAATTTGTAATTCATCATGACCGTAAGATGTAAATAAATCATGCTGAAAAGTAAACAAAAAATAA
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Appendix B The nucleotide, deduced amino acid sequences and structure characterization of *LsToll-13* from *L. striatellus* (GenBank ID: KU866525). The deduced amino acid sequence of *LsToll-13* is shown in one letter symbols below the nucleotide sequence. The numeric positions of the nucleotide sequence are shown on the left. The start codon and stop codon are shown in a box. The asterisk (*) represents the stop codon. The letters in red indicate primers to validate the full length cDNA of *LsToll-13*. The signal peptide (sp) domain and the LRRNT/LRR-1/LRR motifs are marked with the single underline. The LRR-TYP (typical) was not labeled.