

Degenerate TACATCATTAATTACATCGGGAGTTGGTTGAATATATATCTAGCTCAACCAGATTTTCC-
Wild-type TACATCATTAATTACATCGGGAGTTGGTTGAATATATATCTAGCTCAACCAGATTTTCC-
KACC42780 TACATCACCAATTACATCGGGAGTTGGTTGAATGTATATTCAGCCAGCCTGACATTCTT

Degenerate TTCGAGTCCAATACTACAGTTTTACACCCTCCACGTAGTTAGCTGGTGTGTTTCATTGAGG
Wild-type TTCGAGTC-----
KACC42780 TTTAAGTCCAATACTACAGCTTTGCACCCTCCCATGGCTAGAAGGTGTGTTTCATTGATC
** ****

Degenerate CAGAGCATACACATCAGGCCGGATCAAAGGGCGAATTCTGCAGATATCCATCACACTGG
Wild-type -----
KACC42780 AGGAGGGACGATGTAGATGTG-----

Degenerate CGGCCGCTCGAGCATGCATCTAGAGGGCCCAATTCTTGGCTAGAGCCTTAATACTTCATT
Wild-type -----
KACC42780 -----GCGTTGTATCTTTCTGAGGGACGATTTACTTTGATAGAGCCTTAATACTTCATT

Degenerate ATGATTGTTATCCAGTTTTCTAGTATTTTGGGACCATTTCTTAATGTCAAGATCAATCGG
Wild-type -----
KACC42780 ATGTTTGTATC-----

Degenerate TATATACACCAGCAAATACGCTCTAATGTGTTGCGTTGACTTTAATTAGGCCTTTCGCAG
Wild-type -----
KACC42780 -----

Degenerate TCTCCGAACGTGTCTACAATGGTCTTCAATCGGGATCCTCGTCTGGCCGTAAGTTCCGC
Wild-type -----
KACC42780 -----

Degenerate CGCTATTCATGTCCGCCGTCTAATCCCGCCAGGGAGCGTTTATGAGCCGACCCGTAAGTT
Wild-type -----
KACC42780 -----

Degenerate TTTGGCGGTAGACCCCGGTGCACCCCGCAGCGACGTGGAACATTACGTTACGTCCCGGG
Wild-type -----
KACC42780 -----

Degenerate CAGTATCTGCCGGCCACCTCCATTTGACCGCCCGGATGCTTACCGGATTGTGGCCGTCG
Wild-type -----
KACC42780 -----

Degenerate CACGTTTACCCCGTCGTGGGAAAGTCCTTCCGATGTGGCGCAACAGGTCGTGATCGTGGG
Wild-type -----
KACC42780 -----

Degenerate AAAGGCAGGTGGCGGTGGGTATAAAAGCTGGATCCTCAATAACCAGGGTCTTTTGTGCGT
Wild-type -----
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Wild-type -----
KACC42780 -----

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Wild-type -----
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Wild-type -----
KACC42780 -----

Degenerate TCACTGCTAAAGATCCCCAAGTACATCCGCCGTTTCATATCTCTTGCCCGTTGGCCAG
Wild-type -----
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Degenerate CTTTCAGTCTGCGCCTCGGAACAACAGTGGTCCTATTGGCACGTCTGTTTGCATCACTG
Wild-type -----
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Degenerate ATAACGTGTGGAGATCGTTCGTTCCGCTTTACACTCGTGTGCTAAACGAGTCCCCGTGG
Wild-type -----
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Degenerate ACCTCCCGGACCACTTGTGGTGAAGTATCTCCGGCCCAAAGGCTCCCACCGTATTGACTC
Wild-type -----
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Degenerate TTGCTTTGATTTAGGCACACCGCTCCGAACAACTAAGTGGCAACCGTTACGTGACAC
Wild-type -----
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Wild-type -----
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Degenerate GAATACCGCCGACTCTTCAAAAAGTATCTTGCCTAACGTGAGTAATGCTGCGTTCTTTC
Wild-type -----
KACC42780 -----

Degenerate TTGGACGTGGTCGATATATGATACTCTAAATAGATGCATTGCGACGAAGGACTGGCGACG
Wild-type -----
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Degenerate ACGATACGCTTCATTCACACCAATCTTGAGCTTCTTGACGACGTACAAGTCAACGTCGGT
Wild-type -----
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Wild-type -----
KACC42780 -----

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Wild-type -----
KACC42780 -----

Degenerate **ATCGCCAAGAGGAAAGGGGTACCAATCAGGGGGCCGTCGTATACCTCGACTGGACGCCC**
Wild-type -----
KACC42780 -----

Degenerate **GACGCTTGGGAGTATGGCTGCCAGGTGGCGAGTGATCTCGTACAGCTGGGGTTGCGCCAG**
Wild-type -----
KACC42780 -----

Degenerate **GACATGACACAATACCGTTTCGACAATACGATGCCGGGCGGGAAGCAGTTCTTGCATAAT**
Wild-type -----
KACC42780 -----

Degenerate **ATGAAGTCTACGGTACAAACATGCCTTCGACTCCGCCGTACGTTATTTCTTTTTTCT**
Wild-type -----
KACC42780 -----

Degenerate **TCTTCCTCTTTAAGAATGTATTTGCCTAGATCATCCGCAGTCTGCGAGCACGCAAGTG**
Wild-type -----
KACC42780 -----

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Wild-type -----
KACC42780 -----

Degenerate **CCCTGCCGCCCGTCCTCGCCGGCGACCGGAAAGAAAGAAAGAAGCGGGGGGAGA**
Wild-type -----
KACC42780 -----

Degenerate **AGGGGAACAGATGATCACGCCATGCCCGGCTCCGCCATGTGGAATGCCGCCAGCAACGG**
Wild-type -----
KACC42780 -----

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Wild-type -----
KACC42780 -----

Degenerate **CAGGACGTTCTCACCCTCTTCTCGATCGTTATCGCTCCAACCTCCCTTCGGAACAAAGG**
Wild-type -----
KACC42780 -----

Degenerate **GCATTGTGAAAAGAAGCTCGCAGCGGCCAGAGCAAACGTCATTGCCCTTCTCGGAGCGGA**
Wild-type -----
KACC42780 -----

Degenerate **CTTCCTCACATCCGGCAAGGTTCCCGCCGACTTTACGCCACAACAAGCGGAAAGGCCCT**
Wild-type -----
KACC42780 -----

Degenerate TGGTGAAGTACCAAGTATTTTGTATGTTATTTGCTTCTTTCTTTGTGATCCCTATCTT
Wild-type -----
KACC42780 -----

Degenerate ACTCCAGTGTACACAGAATTGTCTAAAGTGGAACTTCCTGTCAAAGGCTGAACGTGCGGC
Wild-type -----
KACC42780 -----

Degenerate TATGGCACTTCCCGTCGACGTAGACTCCCCGGTGCTTTGACCGTCTCGT**CAATCGTTG**
Wild-type -----
KACC42780 -----

Degenerate **GAACGCAATGTCGAGCGTGCCCAAGCGCATGAGTGGTGTGCAACATCAATCGGCTGGAT**
Wild-type -----
KACC42780 -----

Degenerate **GGTTGAGATTTTGTGCGGGCGGTCCCGATCCCCGGTCAGAGA**GCCCGACATATATCCCGA
Wild-type -----
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Wild-type -----
KACC42780 -----

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Wild-type -----
KACC42780 -----

Degenerate GCCGAGACGCAGGAGCAAGACGTGCATGCACTTCCAAC**GACATATGGAAGACTTC**GCCATG
Wild-type -----
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Wild-type -----
KACC42780 -----

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Wild-type -----
KACC42780 -----

Degenerate GAGTTTGCCACTAAAGATGAGCAGGAGTTGCGGGAGTACTTGACAATCCGGGAC**GGATGC**
Wild-type -----
KACC42780 -----

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Wild-type -----
KACC42780 -----

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Wild-type -----
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Degenerate GCTATATTCAAACGCCTGACTTTTTACCGACCAGGCCGATGTCCACGACACGCTCCGGCAC
Wild-type -----
KACC42780 -----

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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
KACC42780 -----



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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
KACC42780 -----



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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Degenerate GTGGCGGACTGGAACACTTACTATAAACAACCTCAGATTTTCCAGACTC**CCGACGCAT**
Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
KACC42780 -----

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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
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Wild-type -----
KACC42780 -----

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Wild-type -----
KACC42780 -----

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Wild-type -----
KACC42780 -----

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Wild-type -----
KACC42780 -----

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Wild-type -----
KACC42780 -----

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Wild-type -----
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Degenerate TGGTGACGAGGATCTCGATCGTGACCCCGACTTCAACTACAATCCTGACATTGAAGATGT
Wild-type -----
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Wild-type -----
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Degenerate GCTCGATCCCGCGCGGATTGGGCCGGCATGGACAAGGACGCCGTTTCAGAAGTACATTGC
Wild-type -----
KACC42780 -----

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Wild-type -----
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Degenerate **CAACGCCTACCAGGATGTTTACGACCACCCGACGCCAAAGTCATGAAAGACATTGATTC**
Wild-type -----
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Wild-type -----
KACC42780 -----

Degenerate **CGATGCCAAGTTCACTCTCGGCGGTCATCTTCATTGCAAGGTTCCGTTACAGGTAAGTGA**
Wild-type -----
KACC42780 -----

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Wild-type -----
KACC42780 -----

Degenerate **AGAATGTTTACCGTGATCCCGCGGATGATACAGCTGTCTTCTGGGGCAAATGTGGAATTG**
Wild-type -----
KACC42780 -----

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Wild-type -----
KACC42780 -----

Degenerate **TTCACAGCGGATCTATGCTGCGAATCTCACCGCCGCCAAGGAAATCCTTCCGGACGTGG**
Wild-type -----
KACC42780 -----

Degenerate **TCAAGCAGTGGTCTCCGGATTATCAAACGGAACCTAAGCCGGACTGGGTACTGGCGCCCTA**
Wild-type -----
KACC42780 -----

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Wild-type -----
KACC42780 -----

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Wild-type -----
KACC42780 -----

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Wild-type -----
KACC42780 -----

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Wild-type -----
KACC42780 -----

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Wild-type -----
KACC42780 -----

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Wild-type -----
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Wild-type -----
KACC42780 -----

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Wild-type -----
KACC42780 -----

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Wild-type -----
KACC42780 -----

Degenerate CCAAGTCCATGAAGCTGTATCCACAAACCGGTTCAAATGAAAACAGCGGACGACATTC
Wild-type -----
KACC42780 -----

Degenerate **CGGTGTCGTA**TCCTCTTGCTATGACGCGCTCCTCACCAACCTGGGAGCAAAGACGCTG
Wild-type -----
KACC42780 -----

Degenerate GAGCTGTCCGT**GTCGAAGTACGTGTGCCGCGCAGTCAAGCCAATGCTGT**CTTTGCTCTTG
Wild-type -----
KACC42780 -----

Degenerate CCATGGAACCGCGCGACTTTCAAGGGTACACGCATATAGTGAAGAACGAGATGTTATGGT
Wild-type -----
KACC42780 -----

Degenerate ATGTCATGTGCC**AGAATCTTTCTGGCATATATGCTTACATGGATCGCAAACAGGGCATGG**
Wild-type -----
KACC42780 -----

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Wild-type -----
KACC42780 -----

Degenerate **GACACCCGCGCCATAAGGCC**ACCCCTTCTACTTGTTGCCGAACTACATTTGTCTTCAAC
Wild-type -----
KACC42780 -----

Degenerate TCCATGCATTCACGGCCGATGGAGATGGGAT**GGGACCGGCAGGTGGGAGCT**CTCATTCTC
Wild-type -----
KACC42780 -----

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Degenerate CCTATCGGCACCGTCGCACAGAGCTCGGAGAACAAAGGATCGCATGATTTGGGTTAGGGGG
Wild-type  -----
KACC42780  -----

Degenerate GGTTAGGGTTAGGGGGTTAGGGTTAGGGG
Wild-type  -----
KACC42780  -----

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Supplemental Fig. S2. Alignment of the sequences corresponding degenerate specific region from degenerate (D4), wild type (W1) and KACC42780. Gene was predicted by FGENESH, and function and domain were estimated by UniProt (<http://www.uniprot.org/>) and Protein BLAST in NCBI. Exons of the putative helicase gene are shown in bold and start and stop codons are underlined. * indicates the identical residues in three sequences. ▼ and arrows indicate the sequence amplified by the SCAR marker discriminating degenerates. ▽ and arrows indicate the genomic sequence corresponding to HELICc and HepA domains. Telomeric repeating units (TTAGGG) are in shade box.