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khr mutant 1 (1 strains) khr mutant 4 (1 strains) khr mutant 3 (1 strains) khr mutant 5 (2 strains) khr mutant 2 (3 strains) khr mutant 9 (19 strains) khr mutant 10 (1 strains) khr mutant 3 (5 strains) khr mutant 8 (2 strains) khr mutant 7 (6 strains) khr mutant 11 (1 strains) khr mutant 6 (1 strains) khr mutant 2 (1 strains) khr mutant 13 (1 strains) khr mutant 1 (1 strains) khr mutant 12 (1 strains) consensus

ATGGGCCACTTAGCGATCCTTTTCAGTATTATCGCTGTATGGAATATAGCTACAGCTGTTGCATCGAGCGACAGCATTTACCTTAAGGGACACAGAGTAG ATGGG<mark>T</mark>CACTTAGCGATCCTTTTCAGTATTATCGCTGTATGGAATATAGCTACAGCTGTTGCATCGAGCGACAGCATTTACCTTAAGGGACACAGAGTAG ATGGGCCACTTAGCGATCCTTTTCAGTATTATCGCTGTATGGAATATAGCTACAGCTGTTGCATCGAGCGACAGCATTTACCTTAAGGGACACAGAGTAG ATGGGCCACTTAGCGATCCTTTTCAGTATTATCGCTGTATGGAATATAGCTACAGCTGTTGCATCGAGCGACAGC<u>ATTTACCTTAAGGGACACAGAGTAG</u> ATGGGCCACTTAGCGATCCTTTTCAGTATTATCGCTGTATGGAATATAGCTACAGCTGTTGCATCGAGCGACAGCATTTACCTTAAGGGACACAGAGTAG ATGGGCCACTTAGCGATCCTTTTCAGTATTATCGCTGTATGGAATATAGCTACAGCTGTTGCATCGAGCGACAGCATTTACCTTAAGGGACACAGAGTAG ATGGGCCACTTAGCGATCCTTTTCAGTATTATCGCTGTATGGAATATAGCTACAGCTGTTGCATCGAGCGACAGCATTTACCTTAAGGGACACAGAGTAG

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GACAGGATATCGACAGTCTATACAGAGTGTACGATAATGGTACTATGTACCCTGTCACTTTCAATGAGTGGTTAAATGATCTAACTGGGATG<mark>G</mark>ATGACT GACAGGATATCGACAGTCTATACAGAGTGTACGATAATGGTACTATGTACCCTGTCACTTTCAATGAGTGGTTAAATGATCTAACTGGGATG<mark>G</mark>ATGACTT GACAGGATATCGACAGTCTATACAGAGTGTACGATAATGGTACTATGTACCCTGTCACTTTCAATGAGTGGTTAAATGATCTAACTGGGATG<mark>G</mark>ATGACTT GACAGGATATCGACAGTCTATACAGAGTGTACGATAATGGTACTATGTACCCTGTCACTTTCAATGAGTGGTTAAATGATCTAACTGGGATG<mark>G</mark>ATGACTT GACAGGATATCGACAGTCTATACAGAGTGTACGATAATGGTACTATGTACCCTGTCACTTTCAATGAGTGGTTAAATGATCTAACTGGGATG<mark>G</mark>ATGACTT

khr mutant 1 (1 strains) khr mutant 4 (1 strains) khr mutant 3 (1 strains) khr mutant 5 (2 strains) khr mutant 2 (3 strains) khr mutant 9 (19 strains) khr mutant 10 (1 strains) khr mutant 3 (5 strains) khr mutant 8 (2 strains) khr mutant 7 (6 strains) khr mutant 11 (1 strains) khr mutant 6 (1 strains) khr mutant 2 (1 strains) khr mutant 13 (1 strains) khr mutant 1 (1 strains) khr mutant 12 (1 strains) consensus

<u>GGCAACAAATAACGCGACAATATTA</u>AAACGTGACAGTAGCGATGTTTCTTGCGTTAATGAAACATGCCAATACGTGGATTACCACGTGGATGACGAAGGA GGCAACAAATAACGCGACAATATTAAAACGTGACAGTAGCGATGTTTCTTGCGTTAATGAAACATGCCAATACGTGGATTACCACGTGGATGACGAAGGA GGCAACAAATAACGCGACAATATTAAAACGTGACAGTAGCGATGTTTCTTGCGTTAATGAAACATGCCAATA<mark>T</mark>GTGGATTACCACGTGGATGACGAAGGA GGCAACAAATAACGCGACAATATTAAAACGTGACAGTAGCGATGTTTCTTGCGTTAATGAAACATGCCAATA<mark>T</mark>GTGGATTACCACGTGGATGACGAAGGA GGCAACAAATAACGCGACAATATTAAAACGTGACAGTAGCGATGTTTCTTGCGTTAATGAAACATGCCAATACGTGGATTACCACGTGGATGACGAAGGA

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khr mutant 1 (1 strains) khr mutant 4 (1 strains) khr mutant 3 (1 strains) khr mutant 5 (2 strains) khr mutant 2 (3 strains) khr mutant 9 (19 strains) khr mutant 10 (1 strains) khr mutant 3 (5 strains) khr mutant 8 (2 strains) khr mutant 7 (6 strains) khr mutant 11 (1 strains) khr mutant 6 (1 strains) khr mutant 2 (1 strains) khr mutant 13 (1 strains) khr mutant 1 (1 strains) khr mutant 12 (1 strains) consensus

GTTATAA<mark>A</mark>TATAGACATATCTACGTATCGTATCCCTGTCGAATGGGATAATGGTTCTGCAGGCAACGCATCATATGGAGTCTCAAAGCGTGATACAAAAT ${ t GTTATAACTATAGACATATCTACGTATCGTATCCCTGTCGAATGGGATAATGGTTCTGCAGGCAACGCATCATATGGAGTCTCAAAGCGTGATACAAAAT$ GTTATAACTATAGACATATCTACGTATCGTATCCCTGTCGAATGGGATAATGGTTCTGCAGGCAACGCATCATATGGAGTCTCAAAGCGTGATACAAAA GTTATAACTATAGACATATCTACGTATCGTATCCCTGTCGAATGGGATAATGGTTCTGCAGGCAACGCATCATATGGAGTCTCAAAGCGTGATACAAAA GTTATAACTATAGACATATCTACGTATCGTATCCCTGTCGAATGGGATAATGGTTCTGCAGGCAACGCATCATATGGAGTCTCAAAGCGTGATACAAAAT GTTATAACTATAGACATATCTACGTATCGTATCCCTGTCGAATGGGATAATGGTTCTGCAGGCAACGCATCATATGGAGTCTCAAAGCGTGATACAAAA GTTATAACTATAGACATATCTACGTATCGTATCCCTGTCGAATGGGATAATGGTTCTGCAGGCAACGCATCATATGGAGTCTCAAAGCGTGATACAAAA GTTATAACTATAGACATATCTACGTATCGTATCCCTGTCGAATGGGATAATGGTTCTGCAGGCAACGCATCATATGGAGTCTCAAAGCGTGATACAAAAT GTTATAACTATAGACATATCTACGTATCGTATCCCTGTCGAATGGGATAATGGTTCTGCAGGCAACGCATCATATGGAGTCTCAAAGCGTGATACAAAA GTTATAACTATAGACATATCTACGTATCGTATCCCTGTCGAATGGGATAATGGTTCTGCAGGCAACGCATCATATGGAGTCTCAAAGCGTGATACAAAA GTTATAACTATAGACATATCTACGTATCGTATCCCTGTCGAATGGGATAATGGTTCTGCAGGCAACGCATCATATGGAGTCTCAAAGCGTGATACAAAA <u>GTTATAACTATAGACATATCTACGT</u>ATCGTATCCCTGTCGAATGGGATAATGGTTCTGCAG<mark>A</mark>CAACGCATCATATGGAGTCTCAAAGCGTGATACAAAAT GTTATAACTATAGACATATCTACGTATCGTATCCCTGTCGAATGGGATAATGGTTCTGCAGGCAACGCATCATATGGAGTCTCAAAGCGTGATACAAAAT GTTATAACTATAGACATATCTACGTATCGTATCCCTGTCGAATGGGATAATGGTTCTGCAGGCAACGCATCATATGGAGTCTCAAAGCGTGATACAAAA GTTATAACTATAGACATATCTACGTATCGTATCCCTGTCGAATGGGATAATGGTTCTGCAGGCAACGCATCATATGGAGTCTCAAAGCGTGATACAAAAT GTTATAACTATAGACATATCTACGTATCGTATCCCTGTCGAATGGGATAATGGTTCTGCAGGCAACGCATCATATGGAGTCTCAAAGCGTGATACAAAAT

khr mutant 1 (1 strains) khr mutant 4 (1 strains) khr mutant 3 (1 strains) khr mutant 5 (2 strains) khr mutant 2 (3 strains) khr mutant 9 (19 strains) khr mutant 10 (1 strains) khr mutant 3 (5 strains) khr mutant 8 (2 strains) khr mutant 7 (6 strains) khr mutant 11 (1 strains) khr mutant 6 (1 strains) khr mutant 2 (1 strains) khr mutant 13 (1 strains) khr mutant 1 (1 strains) khr mutant 12 (1 strains) consensus

 ${f A}{f T}{f G}{f A}{f G}{f A}{f C}{f A}{f T}{f C}{f C}{f G}{f T}{f T}{f C}{f C}{f G}{f T}{f T}{f C}{f T}{f G}{f C}{f G}{f T}{f A}{f C}{f G}{f C}{f G}{f T}{f T}{f C}{f G}{f C}{f G}{f T}{f T}{f C}{f G}{f C}{f A}{f T}{f G}$ ATGAGACATTCTGTAAGAAGAAATATGCGGTATAAACGTTTCCGGTTTCTGTAACGCGTATGACTTCGCCGTTCCTGCTTTCGACTTCGGTGGCAATG1 ATGAGACATTCTGTAAGAAGAAATATGCGGTATAAACGTTTCCGGTTTCTGTAACGCGTATGACTTCGCCGTTCCTGCTTTCGACTTCGGTGGCAATG1 ATGAGACATTCTGTAAGAAGAAATATGCGGTATAAACGTTTCCGGTTTCTGTAACGCGTATGACTTCGCCGTTCCTGCTTTCGACTTCGGTGGCAATG1 ATGAGACATTCTGTAAGAAGAAATATGCGGTATAAACGTTTCCGGTTTCTGTAACGCGTATGACTTCGCCGTTCCTGCTTTCGACTTCGGTGGCAATG1 ATGAGACATTCTGTAAGAAGAAAATATGCGGTATAAACGTTTCCGGTTTCTGTAACGCGTATGACTTCGCCGTTCCTGCTTTCGACTTCGGTGGCAATG1 ATGAGACATTCTGTAAGAAGAAAATATGCGGTATAAACGTTTCCGGTTTCTGTAACGCGTATGACTTCGCCGTTCCTGCTTTCGACTTCGGTGGCAATG1 ATGAGACATTCTGTAAGAAGAAATATGCGGTATAAACGTTTCCGGTTTCTGTAACGCGTATGACTTCGCCGTTCCTGCTTTCGACTTCGGTGGCAATG1 ATGAGACATTCTGTAAGAAGAAATATGCGGTATAAACGTTTCCGGTTTCTGTAACGCGTATGACTTCGCCGTTCCTGCTTTCGACTTCGGTGGCAATG ATGAGACATTCTGTAAGAAGAAATATGCGGTATAAAC<mark>A</mark>TTTCCGGTTTCTGTAACGCGTATGACTTCGCCGTTCCTGCTTTCGACTTCGGTGGCAATG1 ATGAGACATTCTGTAAGAAGAAAATATGCGGTATAAACGTTTCCGGTTTCTGTAACGCGTATGACTTCGCCGTTCCTGCTTTCGACTTCGGTGGCAATG1 ATGAGACATTCTGTAAGAAGAAATATGCGGTATAAACGTTTCCGGTTTCTGTAACGCGTATGACTTCGCCGTTCCTGCTTTCGACTTCGGTGGCAATG1 ATGAGACATTCTGTAAGAAGAAATATGCGGTATAAACGTTTCCGGTTTCTGTAACGCGTATGACTTCGCCGTTCCTGCTTTCGACTTCGGTGGCAATG1 ATGAGACATTCTGTAAGAAGAAAATATG<mark>T</mark>GGTATAAACGTTTCCGGTTTCTGTAACGCGTATGACTTCGCCGTTCCTGCTTTCGACTTCGGTGGCAATG1 ATGAGACATTCTGTAAGAAGAAATATGCGGTATAAACGTTTCCGGTTTCTGTAACGCGTATGACTTCGCCGTTCCTGCTTTCGACTTCGGTGGCAATG1 ATGAGACATTCTGTAAGAAGAAAATATGCGGTATAAACGTTTCCGGTTTCTGTAACGCGTATGACTTCGCCGTTCCTGCTTTCGACTTCGGTGGCAATG1

khr mutant 1 (1 strains) khr mutant 4 (1 strains) khr mutant 3 (1 strains) khr mutant 5 (2 strains) khr mutant 2 (3 strains) khr mutant 9 (19 strains) khr mutant 10 (1 strains) khr mutant 3 (5 strains) khr mutant 8 (2 strains) khr mutant 7 (6 strains) khr mutant 11 (1 strains) khr mutant 6 (1 strains) khr mutant 2 (1 strains) khr mutant 13 (1 strains) khr mutant 1 (1 strains) khr mutant 12 (1 strains) consensus

CTACAACCTTGTTAGTGGTATCACTGATAGGATTAAAGAAGCTACGAAGAGGGATAAGACTGAATGTCTAGGTTACGAACTAGATCATGTGAGAATAGAT CTACAACCTTGTTAGTGGTATCACTGATAGGATTAAAGAAGCTACGAAGAGGGGATAAGACTGAATGTCTAGGTTACGAACTAGATCATGTGAGAATAGAT CTACAACCTTGTTAGTGGTATCACTGATAGGATTAAAGAAGCTACGAAGAGGGGATAAGACTGAATGTCTAGGTTACGAACTAGATCATGTGAGAATAGAT CTACAACCTTGTTAGTGGTATCACTGATAGGATTAAAGAAG<mark>T</mark>TACGAAGAGGGGATAAGACTGAATGTCTAGGTTACGAACTAGATCATGTGAGAATAGAT <u>CTACAACCTTGTT</u>AGTGGTATCACTGATAGGATTAAAGAAGCTACGAAGAGGGGATAAGACTGAATGTCTAGGTTACGAACTAGATCATGTGAGAATAGAT CTACAACCTTGTTAGTGGTATCACTGATAGGATTAAAGAAGCTACGAAGAGGGGATAAGACTGAATGTCTAGGTTACGAACTAGATCATGTGAGAATAGAT CTACAA<mark>T</mark>CTTGTTAGTGGTATCACTGATAGGATTAAAGAAGCTACGAAGAGGGGATAAGACTGAATGTCTAGGTTACGAACTAGATCATGTGAGAATAGAT CTACAACCTTGTTAGTGGTATCACTGATAGGATTAAAGAAGCTACGAAGAGGGATAAGACTGAATGTCTAGGTTACGAACTAGATCATGTGAGAATAGAT CTACAACCTTGTTAGTGGTATCACTGATAGGATTAAAGAAGCTACGAAGAGGGGATAAGACTGAATGTCTAGGTTACGAACTAGATCATGTGAGAATAGAT CTACAACCTTGTTAGTGGTATCACTGATAGGATTAAAGAAGCTACGAAGAGGGGATAAGACTGAATGTCTAGGTTACGAACTAGATCATGTGAGAATAGAT CTACA<mark>G</mark>CCTTG<mark>C</mark>TAGTG<mark>A</mark>TATCA<mark>T</mark>TGATAGGATTAAAGAAGCTACGAAGAGGGGATAAGACTGAATGTCTAGGTTACGAACTAGATCATGTGAGAATAGAT CTACA<mark>G</mark>CCTTG<mark>C</mark>TAGTG<mark>A</mark>TATCA<mark>T</mark>TGATAGGATTAAAG<mark>C</mark>GCTACGAAGAGGGGATAAGAC<mark>C</mark>GAATGTCTA<mark>C</mark>GTTACGAACTAGATCATGTGAGAATAGAT CTACA<mark>G</mark>CCTTG<mark>C</mark>TAGTG<mark>A</mark>TATCA<mark>T</mark>TGATAGGATTAAAG<mark>G</mark>C<mark>A</mark>CTACGAAGAGGGATAAGAC<mark>C</mark>GAATGTCTA<mark>C</mark>GTTACGAACTAGATCATGTGAGAATAGAT CTACA<mark>G</mark>CCTTG<mark>C</mark>TAGTG<mark>A</mark>TATCA<mark>T</mark>TGATAGGATTAAAG<mark>GC</mark>GCTACGAAGAGGGGATAAGAC<mark>C</mark>GAATGTCTA<mark>C</mark>GTTACGAACTAGATCATGTGAGAATAGAT CTACA<mark>G</mark>CCTTG<mark>C</mark>TAGTG<mark>A</mark>TATCA<mark>T</mark>TGATAGGATTAAAG<mark>G</mark>CGCTACGAAGAGGGGATAAGAC<mark>C</mark>GAATGTCTA<mark>C</mark>GTTACGAACTAGATCATGTGAGAATAGAT CTACA<mark>G</mark>CCTTG<mark>C</mark>TAGTG<mark>A</mark>TATCA<mark>T</mark>TGATAGGATTAAAG<mark>G</mark>CGCTACGAAGAGGGATAAGACTGAATGTCTAGGTTACGAACTAGATCATGTGAGAATAGAT

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CCTGCTGTTGATTGGTCCATATTTATTTCTACTTGGAAACAAGGGTCGGCCAATTGTGACACACAGGCATCTGCTGACAGCTTGAAATGTGCTGCCCAAA CCTGCTGTTGATTGGTC<mark>T</mark>ATATTTATTTCTACTTGGAA<mark>G</mark>CAAGGG<mark>C</mark>CGGCCAATT<mark>A</mark>TGACACACAGGCATCTGCTGACAGCTTGAAATGTGCTGCCCAAA CCTGCTGTTGATTGGTC<mark>T</mark>ATATTTATTTCTACTTGGAA<mark>G</mark>CAAGGG<mark>C</mark>CGGCCAATT<mark>A</mark>TGACACACAGGCATCTGCTGACAGCTTGAAATGTGCTGCCCAAA CCTGCTGTTGATTGGTC<mark>T</mark>ATATTTATTTCTACTTGGAA<mark>G</mark>CAAGGG<mark>C</mark>CGGCCAATT<mark>A</mark>TGACACACAGGCATCTGCTGACA<mark>A</mark>CTTGAAATGTGCTGCCCAAA CCTGCTGTTGATTGGTC<mark>T</mark>ATATTTATTTCTACTTGGAA<mark>G</mark>CAAGGG<mark>C</mark>CGGCCAATT<mark>A</mark>TGACACACAGGCATCTGCTGACAGCTTGAAATGTGCTGCCCAAA CCTGCTGTTGATTGGTC<mark>T</mark>ATATTTATTTCTAC<mark>C</mark>TGGAA<mark>G</mark>CAAGGG<mark>C</mark>CGGCCAATT<mark>A</mark>TGACACACAGGCATCTGCTGACAGCTTGAAATGTGCTGCCCAAA

khr mutant 1 (1 strains) khr mutant 4 (1 strains) khr mutant 3 (1 strains) khr mutant 5 (2 strains) khr mutant 2 (3 strains) khr mutant 9 (19 strains) khr mutant 10 (1 strains) khr mutant 3 (5 strains) khr mutant 8 (2 strains) khr mutant 7 (6 strains) khr mutant 11 (1 strains) khr mutant 6 (1 strains) khr mutant 2 (1 strains) khr mutant 13 (1 strains) khr mutant 1 (1 strains) khr mutant 12 (1 strains) consensus

A A GCA CTTGA A A GTGA A CA CA TCA CCA A A A A A CA GCTTTCTGTA TTCA CCTA GA TA A TGGTGGA TCA TTTA A CTTA GA CA TTA GGCTA A TA TCTGA GCT AAGCACTTGAAAGTGAACACAATCACCAAAAAACAGCTTTCTGTATTCACCTAGATAATGGTGGATCATTTAACTTAGACATTAGGCTAATATCTGAGCT AAGCACTTGAAAGTGAACACAATCACCAAAAAAACAGCTTTCTGTATTCACCTAGATAATGGTGGATCATTTAACTTAGACATTAGGCTAATATCTGAGCT AAGCACTTGAAAGTGAACACAATCACCAAAAAAACAGCTTTCTGTATTCACCTAGATAATGGTGGATCATTTAACTTAGACATTAGGCTAATATCTGAGCT AAGCACTTGAAAGTGAACACAATCACCAAAAAACAGCTTTCTGTATTCACCTAGATAATGGTGGATCATTTAACTTAGACATTAGGCTAATATCTGAGCT AAGCACTTGAAAGTGAACACAATCACCAAAAAAACAGCTTTCTGTATTCACCTAGATAATGGTGGATCATTTAACTTAGACATTAGGCTAATATCTGAGCT AAGCACTTGAAAGTGAACACAATCACCAAAAAACAGCTTTCTGTATTCACCTAGATAATGGTGGATCATTTAACTTAGACATTAGGCTAATATCTGAGCT AAGCACTTGAAAGTGAACACAATCACCAAAAAACAGCTTTCTGTATTCACCTAGATAATGGTGGATCATTTAACTTAGACATTAGGCTAATATCTGAGCT AAGCACTTGAAAGTGAACACAATCACCAAAAAACAGCTTTCTGTATTCACCTAGATAATGGTGGATCATTTAACTTAGACATTAGGCTAATATCTGAGCT AAGCACTTGAAAGTGAACACAATCACCAAAAAACAGCTTTCTGTATTCACCTAGATAATGGTGGATCATTTAACTTAGACATTAGGCTAATATCTGAGCT AAGCACTTGAAAGTGAACACAATCACCAAAAAACAGCTTTCTGTATTCACCTAGATAATGGTGGATCATTTAACTTAGACATTAGGCTAATATCTGAGCT A A GC<mark>C</mark>CTTGA A A GTGA A CA A TCA CCA A A A A A CA GCTTTCTGTATTCA CCTA GA TA A TGGTGGA TC<mark>C</mark>TTTA A CTTA GA CATTA GGCTA A TA TCTGA GCT A A GC<mark>C</mark>CTTG A A A GTG A A CA CA TCA CCA A A A A A CAGCTTTCTGTATTCA CCTAGATA A TGGTGGATC<mark>G</mark>TTTA A CTTAGA CATTAGGCTA A TA TCTGA GCT A A GC<mark>C</mark>CTTGA A A GTGA A CA A TCA CCA A A A A A CA GCTTTCTGTATTCA CCTA GA TA A TGGTGGA TC<mark>C</mark>TTTA A CTTA GA CATTA GGCTA A TA TCTGA GCT A A GC<mark>C</mark>CTTG A A A GTG A A C A C A A T C A C C A A A A A A C A GCTTTCTGTATTCA CCTA G A T A T GGTGG A T C<mark>C</mark>TTTA A CTT A G A CATT A G GCTA A T A T CTG A G C T A A GC<mark>C</mark>CTTGA A A GTGA A CA A TCA CCA A A A A A CA GCTTTCTGTATTCA CCTA GA TA A TGGTGGA TC<mark>C</mark>TTTA A CTTA GA CATTA GGCTA A TA TCTGA GCT

khr mutant 1 (1 strains)	${ t TTCATTTTCGAAATATAACCCATGGGCTCTTCCATGTCCGAAGGACAAAGGCTCCAATTCTTGGCAAGTTGTGAGCGACTGTTTTCAATAA$	891
khr mutant 4 (1 strains)	${ t TTCATTTTCGAAATATAACCCATGGGCTCTTCCATGTCCGAAGGACAAAGGCTCCAATTCTTGGCAAGTTGTGAGCGACTGTTTTCAATAA$	891
khr mutant 3 (1 strains)	${ t TTCATTTTCGAAATATAACCCATGGGCTCTTCCATGTCCGAAGGACAAAGGCTCCAATTCTTGGCAAGTTGTGAGCGACTGTTTTCAATAA$	891
khr mutant 5 (2 strains)	${ t TTCATTTTCGAAATATAACCCATGGGCTCTTCCATGTCCGAAGGACAAAGGCTCCAATTCTTGGCAAGTTGTGAGCGACTGTTTTCAATAA$	891
khr mutant 2 (3 strains)	${ t TTCATTTTCGAAATATAACCCATGGGCTCTTCCATGTCCGAAGGACAAAGGCTCCAATTCTTGGCAAGTTGTGAGCGACTGTTTTCAATAA$	891
khr mutant 9 (19 strains)	${ t TTCATTTTCGAAATATAACCCATGGGCTCTTCCATGTCCGAAGGACAAAGGCTCCAATTCTTGGCAAGTTGTGAGCGACTGTTTTCAATAA$	891
khr mutant 10 (1 strains)	${ t TTCATTTTCGAAATATAACCCATGGGCTCTTCCATGTCCGAAGGACAAAGGCTCCAATTCTTGGCAAGTTGTGAGCGACTGTTTTCAATAA$	891
khr mutant 3 (5 strains)	${ t TTCATTTTCGAAATATAACCCATGGGCTCTTCCATGTCCGAAGGACAAAGGCTCCAATTCTTGGCAAGTTGTGAGCGACTGTTTTCAATAA$	891
khr mutant 8 (2 strains)	${ t TTCATTTTCGAAATATAACCCATGGGCTCTTCCATGTCCGAAGGACAAAGGCTCCAATTCTTGGCAAGTTGTGAGCGACTGTTTTCAATAA$	891
khr mutant 7 (6 strains)	${ t TTCATTTTCGAAATATAACCCATGGGCTCTTCCATGTCCGAAGGACAAAGGCTCCAATTCTTGGCAAGTTGTGAGCGACTGTTTTCAATAA$	891
khr mutant 11 (1 strains)	${ t TTCATTTTCGAAATATAACCCATGGGCTCTTCCATGTCCGAAGGACAAAGGCTCCAATTCTTGGCAAGTTGTGAGCGACTGTTTTCAATAA$	891
khr mutant 6 (1 strains)	TTCATTTTCGAAATATAACCCATGGGCTCTTCCATGTCCGAAGGACAAAG <mark>A</mark> CTC <mark>A</mark> AATTCTTGGCA <mark>G</mark> GTTGTGAGCGACTGTTTTCAATAA	891
khr mutant 2 (1 strains)	TTCATTTTCGAAATATAACCCATGGGCTCTTCCATGTCCGAAGGACAAAG <mark>A</mark> CTC <mark>A</mark> AATTCTTGGCA <mark>G</mark> GTTGTGAGCGACTGTTTTCAATAA	891
khr mutant 13 (1 strains)	TTCATTTTCGAAATATAACCCATGGGCTCTTCCATGTCCGAAGGACAAAG <mark>A</mark> CTC <mark>A</mark> AATTCTTGGCA <mark>G</mark> GTTGTGAGCGACTGTTTTCAATAA	891
khr mutant 1 (1 strains)	TTCATTTTCGAAATATAACCCATGGGCTCTTCCATGTCCGAAGGACAAAG <mark>A</mark> CTC <mark>A</mark> AATTCTTGGCA <mark>G</mark> GTTGTGAGCGACTGTTTTCAATAA	891
khr mutant 12 (1 strains)	TTCATTTTCGAAATATAACCCATGGGCTCTTCCATGTCCGAAGGACAAAG <mark>A</mark> CTC <mark>A</mark> AATTCTTGGCA <mark>G</mark> GTTGTGAGCGACTGTTTTCAATAA	891
consensus	\[\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\	

X non-conserved

X similar

 \times \geq 50% conserved