

Due to an error in production, on pages 82 and 83, the nucleotide sequence in Fig. 2 appeared out of order. The correct sequence appears below.

AGAGGACACACCAATGACGACGAACAACAAAAAAAACCAGGTCTCTCTTCATGATCTCTTCCCTACTTTGGTTTATATATACAAACAAAGGCTTAACGTAGTGCTTTTGCCTTCA -124

M D P S G V K V L E T A E D I Q E R R Q Q V L D R Y H R F K E L S S L R R Q K 39
GAAATGGATCGAAGTGGGTAAAGATTGTTGGAACACGAGAGATATCCAGAAGCAGGCTGCAGCAGGTTTGGACCGTTACCAACAGGTTCAAGAGGCTGTCTCTCGAGGCCCAAAA +126

L E D S Y R Q F Q F F O R D A D E L G K W I Q E K L Q I A S D N E Y K D P S N L Q 79
ACTGAGGATTCCTATGGTTCAGATTCTCCAGCATGAGTAGELGGGAAATGGATCGAAGAGAACCTCCAGATAGCATCTGATGAAATTTACAAAGACCAACCAATTTACA 236

G K L Q K H Q A F E A E V Q A N S G A I V K L D E T G N Q M I N E G B F A S E T 119
GGGAGACTGCAGAGCACCAAGGCTTTGAGCTGAGGTGAGGCCAATTGAGGGCTATCGTTAAGCTGGATGAGCTGGAATCGATGATCAATGAAGGCCATTTCGATCTGAAC 356

I R T R L Q E L H R L W E L L L E K M R E K G V K L L Q A Q K L V Q F L R E C E 159
CATAGAAGATCGTCTACAGAGGCTGCACCGGACTGATGGAGATTACTGCTGGAATAATGAGAGAGAGAGGAGTGAAGCTGTTGCAGACACAGAGGCTGGTGCNAATTCTACGGGAATTGSA 476

D V M D W I N D K E A I A T V T S E E L G Q D L E H V E V L Q K K F E E F Q T D L A 195
AGATGTCTGAGCTGATCAATGACAGGAGCAATAGTGACTTCAGAGAGCTTCGACAGGCTTAGAGCATGTTGAGGTTTTCAGAGAGAAATTTGAAGAGTTTCAAAACAGATCTGCG 596

A H E E R V N E V N Q F A G K L I Q E Q B P E E L I K S K Q D E V N A S W O R 239
AGCTCATGAGGAGAGGTGAATGAAGTGAACNGTTTGGTGGGAACCTTATCCAGGAGCAGACCTCGAGGAGGAATCTATAAGTCCAACAGGATGAAGTAAATGAAGTCAAGCTGGCAGCG 716

L K G L A Q Q R Q G K L F G A A E V Q R P N R D V D E T I S W I R E F G Q L M A 279
TCTTAAGGGCTTGGCCACACAGAGGACAGGAAUACTCTTGGGGCAGCTGAAGTTCGAGGCTTCACAGGGATGTGGATGAATCATCAGCTGGATTGAAGAGAAAGGGCAGTTGATGGC 836

S D D F G R D L A S V Q A L L R K H E G L E R D L A A F H H K V R K A L C A E A D 319
CTCAGATGACTTTGGCAGGACTTGGCCAGCTGCAGGCTTACTACGTGAAGGLGAGGCTGGAAGAGDAGATCTTCAGGATTTCCACATGAAGTTTAAGGCCCTCTGTGCAGAGAGCTGA 956

R L Q Q S H P I N A S O I Q V K R E E L I A N W E Q I R T L A A E R H A R L N D 359
CCGTTTCAGCAATCTCACCTCATATAAGTCTCTCAATTCAGATGAAGAGGAGGAGTGAATTCGCACTGGGAGCAGATCGCAACTCTGGCAGCAGAGAGGCAATGCTGCTCATTAATGA 1076

S Y R L Q R F L A D F R D L T S W V T E M K A L I N A D E L A N D V A G A E A L 399
CTCTACAGGTTGCAAGGCTTCTTCGAGATTTCGGAGCTCTAGTCTGGTAACTGAGATGAAGGCTGCTATAAATGCTGATGAGCTTCCCAATGATGTGGCTGGAGCAGAGGCCCT 1196

T D R H Q E A K K E I D A H E D S F R S A D E S G Q A L L A A G H Y A S D E V K 439
LCTGATGATCATCAGAGCAATGAAGGGAATATGATGCTCAGAGATAGCTTCAGATCTGCTGATGAGTGTGGCAGGCTTTGCTGCGAGCTGAGGACATGCTTCTGATGAAGTTAA 1416

E K L T I L S D E R S A L L E L W E L R R Q Q Y E Q C M D L Q L F Y R D T E Q V 479
AGAAAGCTGACTATCTCTCAGATGAAGATCTGCTTCTGCGAATCATGGAGACTTCGACGCCAACGATGAGCAGCTGCATGGATTCGAGCTTTTTACAGAGATCTGAGCAAGT 1436

D N W M S E Q E A F L L N E A D L G D S L D S V E A L L K K H E D F E K S L S Q 519
TGACAACTGAGTGACCAAGAGAAATTTCTGCTGAACGAAGGACTTGGTGAATCTCTCGATAGGTGGAGGCTCTCTCAAGAGAGATGAAGATTTGAGAAATCCCTAGATGCTCA 1626

E E K I T A L D E F A T K L I Q N N H Y A M D D V A T R R D A L L S R R N A L H 559
AGAGAGAAATTCACAGCATGATGAGTTTGTACTAAGCTGATCAAAATTAACATTTATGCCATGATGAGTGTGCTACGCCAGAGATGCTCTCTGAGTGGCAGAAATGCTCTTCA 14676

E R A M K R R A Q L A D S F H L Q Q F R S D S D E L K S W N E K M K T A T D E 599
TGAAGAGCCCATGAAGGCCGCTGCCAATCTGCGAGACTCTTTCATCTCCAGCAGTTTTTCAGGGACTCTGATGAATGAAGAGTGGGTTAATGAAGAAATGAAGACTGCTACCGATGA 1796

A Y K D P S N L Q G K V Q K H Q A F E A E A L S A N Q S R I D A L E K A G O K L I 639
GGCTTACAGGATCCATCAACTTGCAGGTGAAGTTCAGAGKCATCAGGCTTTGAAGCAGACTTCCTGCTAATCAGAGCTGATATGATGCACTGGAAGAGGCTGGCCAGAGTTGAT 1916

D V N H Y A S D E V A A R M N E V I S L W K K L L E A T E L K G I K L R E A N Q 679
TGATGTCAATCACTATGTCATCTGATGAGGTGGCAGCTGCAATGAATGAATCATCAGCTTTGGAGAAACTCTTGAAGACCCAGGAGCTCAAGAGTATAAAGCTGGTGAAGGCCAATCA 2036

Q Q Q F N R N V E D I E L W L Y E V E G R L A S D D Y G K D L T S V Q N L O K K 719
ACAGCAGCAATTTAATGCAATGTGAAGACATTTGATGTGGCTGTATGAAGTGGAGGGTCACTTGGCTTCTGATGATTTAGAAAGATTTACAGAGGCTTCAGAAATCTCTCAGAGAA 2156

H A L L E A D V A A C H A D G P I D G I T T I Q A R Q F Q D A G H F D A D N I K K K Q 759
ACATGGCTCTGAGGGCAGATGTGCTGCCATACCGATTCGATAGTATGGGATTACCATTCAGGACCGGATTCAGAGATGCTGGGCACTTTGATCTGACATATCAAGAGAAACA 2276

E A L V A R Y E A L K Q D P M V A R K Q K L A D S L R L Q Q L F R D I E D E E T W 799
AGAAGCTTGTATGAGCTGTGATGAGCTTGAAGGATTCATGTGGCTGCGACAGGAAACTTCGAGATTTCTCTGCTCGAGCAGGTTTTCGCTGACATGAGGATGAAGAGGCTG 2396

I R E K E P I A A S T N R G K D L I G V Q N L L K F H Q A L Q A E I A G H E P R 839
GATCGAGGAAGAGNCATTTGCGAGCTCACAAACGAGGCGAAGGACTTAATTTGGTTCGAGATCTGCTAAGAAGSACCAAGGCTTTCAGGAGCAHAATTCAGGCCATGAGGCTCG 2516

I K A V T Q K G N A A G T E E G E H F A E D V G A G V R K I K L D A C S G D S L K A K A 879
CATTAAGACTCACACAAAGGAANTGCAATGGTGAAGAGGACATTTTGTCTAGAGATGTGAAGTCAAAATTAATGAATGAATTAACCAAAATGGGACTCTCTGAAGAGAGGCG 2636

S Q R R Q D L E D S L Q A Q Q Y F A D A N E A Q S W M R E K E P I V G S T D Y G 919
ATCTCAAGGGCAGACGATCTAGAGGATTTCTCTCAAGCTCAGCAGTATTTTGTCTGATGCTAATGAGGCAATCATGATGATGAGGGAAGAGAGGCCAATTTGATGAGCAGCAGGATATGG 2756

K D E D S T G E A A L L K K H E A L M S I A Y G S S I G A L R E Q A Q S C R Q Q 959
AAAGACTGAAGACTTCTGAGGCTCTCTCAAGAGCATGAAGCTTTGATGTGATCTTTCTGCTTACGGCAGTATGACATGAGGATTAAGGGAACAGGCGGAGCTCTCGAGGCAACA 2876

V A P T D D E T G K E L V L A L Y D Y O E K S P R E V T M K K G D I L T L L N S 999
AGTGTCTCCCATCTGATGATGAACGTGAAAGAGGCTTGTCTAGCATCTTATGATTACCAAGAGAGAGTCTCTGGGAGGTGACTATGAAGAGGAGATATTCTAACCTCTCTCAACAG 2996

T N K D W N K V E N D R Q G F V P A A Y V K K L D P A O S A S R E N L L E E Q 1039
CAGCAACAGGACTGSGAAGGTGAAGTTAAGCATGTCAGGGCTTTGTACAGGCTGCTGATGTGAATAATAGATCTGCGCCAGTCTGCTATCCGAGAGAAATCTCTGGAAGAGCA 3116

G S I A L R Q E Q I D N Q T L I T K E V G S V S L G M K Q V E E L Y H S L L E L 1079
AGGCAGATAGCATTCGACAGGCAAAATGACAACCAAGCTCTCAATTAGGAAGTGGCAGTGTATCTCTGCTGATGAAGACCTCGAAGAACTGTATCAITCCCTCTTGAAGT 3236

G E K R K G M L E K S K K F M L F R E A N E L Q O W I N E K E A A L T N E E V 1119
GGGAGAAATCTAAGGACTGCTAGAAATAAGCTGCAAGAGATTATGCTTTTCGTGAGGCTAATGAGCTTCAACATGATCAATGAGAAGAGGCTGCACATCAACAGAGAGAT 3356

G A D L E Q V E V L O K K E D D F Q K D L X A N E S R L K D I N K A N D L E S 1159
GGGTGCTGATTGGGAGCTGGGGTCTACAGAGAGGTTGATGATTTTCAGAGGATTTAAAGGCTATGATGTCAGATCAAGAGGATTAACAGGATTTGCAATGATTTGGAGTCT 3476

12.99	12.99
3596	3716
1239	1279
3956	1319
3836	1359
	4076
1399	1439
4196	4239
1479	1479
4409	1519
	1559
1599	1599
4696	1639
	1679
1719	1719
1759	1759
5276	5276
1799	1839
5396	5516
	1879
1939	1939
5596	5676
1999	1999
5996	6116
2039	2079
6236	6236
	2119
6356	2159
	2199
	6476
	6596
2239	2279
6716	6836
	2319
	2359
	7076
2399	2399
	7196
2439	2479
7316	7436
	7556
	7650