

AGCAGT T T G T G T A T T T A C A G A A A C A G T A A C C A T T T T G A T G G A A C T A G G A G A G A G C T C T G A T T A C T T T T G C<sup>72</sup>

M E L G R A L S Y F A

TCTAGCAACAATTATTTCAATATTCCTGCAGTCAGACACTGATTGAGTCGGAGTCAGTGATCATCAAACCTGA<sup>144</sup>

L A T I I Q Y S C S Q T L I E S E S V I I K P D

TCAGTCTCATAAACTGACCTGTACAGCCTCTGGATTTAACTTTGGTAACCTATTACATGGCCTGGATTAGACA<sup>216</sup>

Q S H K L T C T A S G F N F G N Y Y M A W I R Q

GGCACCTGAAAAGGGCTGGAATTTGTTGCAACTATCTCAGAAGGCAGTAGTAGCAAGTATTACTCCAGTGC<sup>288</sup>

A P G K G L E F V A T I S E G S S S K Y Y S S A

AGTTAATGGCCGCTTCACCATGTCCAGAGACAACAGTAAGATGCAGGTGTATCTGTACATGACCAGTGTGAG<sup>360</sup>

V N G R F T M S R D N S K M Q V Y L Y M T S V R

GACAGAAGACACTGCAGTGTATTACTGCACTAGACGGACTGGGGTGTACTTTGACTACTGGGGAAAAAGGAAC<sup>432</sup>

T E D T A V Y Y C T R R T G V Y F D Y W G K G T

TTCACTGACCGTGACTTCAGCTGTGCAAAGCGCCCCGAAATCCCTGTTTTCCCCTGTGGCAGTGGCGCTCGGC<sup>504</sup>

S V T V T S A V Q S A P K S L F P V W Q C G S A

CTCGGACGGTTTtagtcaactctttggctgCGTCACGCGGATTTGGCCTCCGCCGACGGACTGAGCTTCATATG<sup>576</sup>

S D G L V T L G C V T R D L A S A D G L S F I W

GAAGGATGCGAGCGGGAGCGCGCTGACTGACGTCGTGCAATACCCGGCGGTGCAAGCGACCGGAGGGTACAC<sup>648</sup>

K D A S G S A L T D V V Q Y P A V Q A T G G Y T

CTCGGTGAGCCATGTGCGCGTCAAGGCTTCTGACTGGAACGGGAACAAGAAGTTCACGTGCGAAGTCAAAAA<sup>720</sup>

S V S H V R V K A S D W N G N K K F T C E V K N

TGGCCTAGGATCTAAAGACGCGTCCCTTGCAAAAAGCCAGTTGAGAGAGAGCTCCATGCATCTCTGCTTCTAAC<sup>792</sup>

G L G S K D A S L Q K P V E R E L H A S L L L T

AACTCCAACCCAAACAGAAATAGACAATGGAACAGCTACCTTCGTCTGCTTAGCTACACCATTTTcaccta<sup>864</sup>

T P T Q T E I D N G T A T F V C L A T P F S P K

ATCACACACATTTAAGTGGACTCTTGAAAAAGACAGACATCAGTAATAAGGTCAAAGAGAACATAGTAAGCCA<sup>936</sup>

S H T F K W T L E K T D I S N K V K E N I V S Q

GAATAAAGGTAACCTTCACTGCCATAAGTGTTTTGGAACCTCAGCGCCAGTGAATGGACAAGCTCAACTTCTCC<sup>1008</sup>

N K G N F T A I S V L E L S A S E W T S S T S P

AGTTAAGTGCGAATTCCAGCAGAAGAACCATAATGTGTTCAAAGAAGCGAGTTATGCACCAGGTGACACAAA<sup>1080</sup>

V K C E F Q Q K N H N V F K E A S Y A P G D T K

ACAGCCACAGGTGAAAATAACTGGACCTTCCACCGAGGACATTCTGATCAAAAAGAGCCGGCCAGCTCGAGTG<sup>1152</sup>

Q P Q V K I T G P S T E D I L I K R A G Q L E C

CAGGGCCGAGGGAGACACGGGTTTCAAGAGCATTAAATGGCTTATTGGAATAGAGAGATCTCTTCTCTATC<sup>1224</sup>

R A E G D T G F K S I K W L I G N R E I S S L S

AAATCTATCTTCGAAAACGACGGTTTCACTCCAAACCCACATCGGTTTCGAAGAGTGGATCAATGGCACCGA<sup>1296</sup>

N L S S K T T V S L Q T H I G F E E W I N G T E

ATTATCTGTGAGGTGGAACATGAAGCATTCACTCAACAGTATGAAAAAGTAACCTTCAAAAAGAGAAAAATGG<sup>1368</sup>

F I C E V E H E A F T Q Q Y E K V T F K R E N G

CAATCCGGAATTCCCCAAGGTTTACTTGCTCGCTCCACCAGAGAGCTCTGGTGAATCAGTGACCCTGACTTG<sup>1440</sup>

N P T E A F P K V Y L L A P P E S S G E S V T L T C

CTATGT TAAAGACTTCTACCCTAAGGAGGTGGCTGTGTCTTGGCTTGTTAACGATAAAACAAGTGAAGAAGT<sup>1512</sup>

Y V K D F Y P K E V A V S W L V N D K Q V E E V

GGTCGGCTATGAGCAGAACCACCTGCAGTTATCGACAGAAACAACCTCTTTTCAGTGACAGCCAGCTGAT<sup>1584</sup>

V G Y E Q N T T A V I D R N N L F S V Y S Q L I

TATCAAAACTGCAGACTGGAACAGTGGCAGTGTGTTcagctgcctggTTTTATCATGAGTCCATCAAGGACTG<sup>1656</sup>

I K T A D W N S G S V F S C L V Y H E S I K D C

TGTGCGCCCCATATCCAGATCCATCGCTAAAGATTCAAAAACGCCACCTTAGTGAATCTCACCTTGACCAA<sup>1728</sup>

V R P I S R S I A K D S K T P T L V N L T L T N

TCCCCAATCTTGCTCTTGCTCTACGTATTAGAATTTGTGTTGTCTGCTATTAATGCTTGATTTATTTTAATT<sup>1800</sup>

P Q S C S C S T Y \*

TTTGCCTCATCTTTTATGTCTTTTTTTTTTTTTTATTAATGCAATGTcattgtCTTGATGTCTGTCTGATGCTTG<sup>1872</sup>

TATGTCTTGGTTTTTTGGAACATGTCTAATGCAATAATACACTGAAACAATGACAATAAAAAAAAAATCCTTCAG<sup>1944</sup>

CATGTACAAA<sup>1980</sup>

Figure S2. The full-length cDNA sequence and translated amino acid of catfish IgM heavy chain (variant 1A32). 5' and 3'-untranslated sequences were shown in *italic*. The stop codon was marked as asterisk. The poly-A signal was underlined.