

Supplementary Figure S1. *IQCB1* mRNA sequence and predicted protein in normal and affected dogs. **A.** Normal *IQCB1* mRNA (Accession number KF366421). Boxed are the first Methionin and the stop. **B.** *IQCB1* predicted protein in a normal dog, 598 amino acids. **C.** Affected *IQCB1* mRNA. Boxed are the first methionin and the premature stop codon. Underline is the stop codon in a wild-type allele. Bolded and enlarged are the 3 cytosines, which represent the insertional mutation site. **D.** The predicted *IQCB1* protein in the affected dog. Bold is the sequence that is different from the normal. The predicted protein is 330 amino acids long.

A.

GGAGGCAGTAACGGCAGGCCAGGAAGACGACTTCCCCAAGCCTCAGGCCCCAGCTCGTG
CTGTGACAACAGGGTGATGTAAGAAATGAAAGCCAACAGGTACAGACCCAAGGATCTTAT
CTCTAGCTGCTGAAGTTGCAAAAAGTCTGAGCAAAATGTCCCTGTTATACTATTGAAGT
TAAAAGAAATAATAACAACACACCTTTAGGAAGCTCAGAGTTGAAGAAAATCAAACAA
GATATATATTGTTATGACCTCATTCAGTATTGCCTTTTGGTGCTCAGTCAAGATTGTTCT
CGAATCCAGGGAGGTTGGACTACAATATCCCAACTTACACAGATATTAAGCCACTGCTGT
GTGGGCTTGGAGCCAGGAGAAGATGCAGAGGAATTTTACAATGAATTACTCCCATCAGC
TGCAGAAAATTTTCTGGTTTTGGGGAGACGATTGCAAACATGTTTCATCAATTCAGCTA
AGGGTGAAGAAAAGATGAATTACTACACTCGTTCCAAATTGTGACCGATTCTCTCTTCT
GGCTCTTAGGAGGCCATGTTCAACTCATCCAAAATGTACTACAAAGTGATCATTCTTGC
ACTACTGCAAACGACAATGTTCAAATAGGATCTACAGTCATGACTATGCTACAGAAC
ATACTACAGATCAACAGTGGTGATTTACTCAGAATAGAAGGAAAAATCCTACATTCAAT
TTTAGATGAAGTTGTTTTCAAGCTTTTATCAACTCCTAACCAGTCATAAGAAGTACTGC
TACAAAGCTCCTACTGCTGATGACTGAATCCCATCAGGAAATTTTGATTTTACTGAGACT
AAGTGCCTGCTACAAAGGACTCAGAAGTCTATTAACAACATGAGCCTGGGACAGAGT
TTAGTCAAGAACTTGGACAGCTTATTGCCCTTTTAACCCCTAAGGTCTATCAGGAAGTAG
AAGATCAGAACTACATCAAGCAGCTTGCTTGATTCAAGCTTATTGGAAGGGTTTCCAA
ACTAGAAAAGATTAAGAAGCTTCCATCTGCTGTGATTACTTTGCAGAGGAGTTTCAG
ATCTAAACGAACCAAGATATTACTAAAGCTAAATAAGCAGAAAGAAGAAGAGGACCGCA
GATTACAGTTGCAACTTCAAAGACAGAGGCCATGAGATTGTCCCGAGAATTACGGCTG
AGTATGCTCGAAATAGTTCATCCAGGTCAGGTGGAATAATAATCGGGAAATAGAAGA
GAAATCAGCCTTGATTATCCAGAAACACTGGAGAGGGTACAGGGAAAGGAAAAATTTTC
GCCAACAGAGGCCATCTCTCACGGAATATAAAGCAGCTGTCATACTTCAAAGAGCAACTC
TTAAATTCCTAGCAAAGTGCCGTAAGAAAAAGAACTATTTGCTCCTTGGCGAGGACTT
CAAGATCTCACCGATGCACGGAGAGTTGAATTAAGCAACAAGTGGATGACTATCTCAG
AAGACATCCGAGCTCTCAAATGTCAGATATGACTAGCAGAGAGCTCCATTCCCAAGCTCA
AGAACAACCTGCAACACTACCTTATGGGCAGGGCCCTAGAAGAGAGAGCCAGCAGCACAG
GGAGGCTCTGATGGCTCAGATCAGCACCAACATTGAACAGTTAATGAAGGCACCGAGTCT
GAAGGAGGCAGAAGGGAAAGAACCTGAACTCTTCCCTAAGTAGATCCAGGCCTGTGGCAG
CTAAGGCCAAGCAGGCCCATCTTACTGCCCTGAAGCATATACAGGCACCTTGGTGGAAGA
AGCTTGGGGAAGAAGCAGGAGATGAGATTGATGTTCCAAAGGATGAGTTAGTTAGAA
TTAGGAACCTTATTCATTGGTGAACCAACCCCTTAGGAAGCTATCCAGAGAAATGA
CACAAATCCTTTATTTTAGATTATTTTGGTTCTGCCTCTGGCATGCTAGTAGACTAGAGC
TATCTTTTACTCATGGCTTCCAGAGATTCCTCTCCAAATAAGATTTTCAGGCAGTAAGC
AAAGA

B.

MKPTGTDPRILSLAAEVAKSPEQNVPVILLKLKEIINNTPLGSSELKKIKQDIYC
YDLIQYCLLVLSQDCSRIQGGWTTISQLTQILSHCCVGLPEGEDAEFFYNELLPS
AAENFLVLGRRLQTCFINSAGEEKDELLHSFQIVTDSLFWLLGGHVQLIQNVLQ
SDHFLHLLQTDNVQIGSTVMTMLQNILQINSGDLLRIEGKILHSILDEVVFKLLS
TPNPVIRSTATKLLLLMTESHQEILILLRLSACYKGLRSLLNKHEPGTEFSQELG
QLIALLLTPKVYQEVEDQKLHQAACLIQAYWKGFQTRKRLKKLPSAVITLQRSFRS
KRTKILLKLNKQKEEEDRRLQLQLQRQRAMRLSRELRLSMLEIVHPGQVEKYNRE
IEKSALI IQKHWRGYRERKNFRQQRPSLTEYKAAVILQRATLKF LAKCRKKKKL
FAPWRGLQDLTDARRVELKQOVDDYLRRHPSSQMSDMTSRELHSQAQEQLQHYLM
GRALEERAQQHREALMAQISTNIEQLMKAPSLKEAEGKEPELFLSRSRPVAAKAK
QAHLTALKHIQAPWWKKLGEEAGDEIDVPKDEFSLELGLTFIGGTPP

C.

AGACGACTTCCCCAAGCCTCAGGCCCCAGCTCGTGCTGTGACAACAGGGTGATGTAAGAA
ATGAAGCCAACAGGTACAGACCCAAGGATCTTATCTCTAGCTGCTGAAGTTGCAAAAAG
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CAGTATTGCCTTTTGGTGCTCAGTCAAGATTGTTCTCGAATCCAGGGAGGTTGGACTACA
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GCAGAGGAATTTTACAATGAATTACTCCCATCAGCTGCAGAAAATTTTCTGGTTTTGGG
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TTACTCAGAATAGAAGGAAAAATCCTACATTCAATTTTAGATGAAGTTGTTTTCAAGCT
TTTATCAACTCCTAACCAGTCATAAGAAGTACTGCTACAAAGCTCCTACTGCTGATGAC
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TTGCCCTTTTAAACCCTAAGGTCTATCAGGAAGTAGAAGATCAGAAACTACATCAAGCAG
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CCCATCTGCTGTGATTACTTTGCAGAGGAGTTT**CAGATCTTAA**ACGAACCAAGATATTAC
TAAAGCTAAATAAGCAGAAAGAAGAAGAGACCGCAGATTACAGTTGCAACTTCAAAGA
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TGGGCAGGGCCCTAGAAGAGAGAGCCAGCAGCACAGGGAGGCTCTGATGGCTCAGATCA
GCACCAACATTGAACAGTTAATGAAGGCACCGAGTCTGAAGGAGGCAGAAGGGAAAGAA
CCTGAACTCTTCTAAGTAGATCCAGGCCTGTGGCAGCTAAGGCCAAGCAGGCCCATCTT
ACTGCCCTGAAGCATATACAGGCACCTTGGTGGAAGAAGCTTGGGGAAGAAGCAGGAGA

TGAGATTGATGTTCCAAAGGATGAGTTTAGTTTAGAATTAGGAACTTTATTCATTGGTG
GAACCAACCCCTTAGGAAGCTATCCAGAGAAATGACACAAATCCTTTATTTTAGATTA
TTTTGGTTCTGCCTCTGGCATGCTAGTAGACTAGAGCTATCTTTTACTCATGGCTTTCCA
GAGATTCCTCTCCAAATAAGATTTTCAGGCAGTAAGCAAAGAGTTGTCTTCCCACCTTTT
GCAGTTC

D.

MKPTGTDPRILSLAAEVAKSPEQNVPVILLKLKEIINNTPLGSSELKKIKQDIYC
YDLIQYCLLVLSQDCSRIQGGWTTISQLTQILSHCCVGLPEGEDAEFYNELLPS
AAENFLVLGRRLQTCFINSAKGEEKDELLHSFQIVTDSLFWLLGGHVQLIQNVLQ
SDHFLHLLQTDNVQIGSTVMTMLQNILQINSGDLLRIEGKILHSILDEVVFKLLS
TPNPVIRSTATKLLLLMTESHQEILILLRLSACYKGLRSLLNKHEPGTEFSQELG
QLIALLLTPKVYQEVEDQKLHQAACLIQAYWKGFO~~TRKRLK~~LP**ICCDYFAEEFQI**