

Erratum

Erratum/Correction of Frudakis et al. A Classifier for the SNP-based Inference of Ancestry. J Forensic Sci 2003 July;48(4):771–782.

It has come to the attention of the Journal that there is incorrect 5' and 3' flanking sequence appearing around the IUB variant for each SNP in Appendix 1. Below is the new/correct Appendix 1.

The Journal regrets this error. Note: Any and all future citations of the above-referenced paper should read Frudakis et al. A Classifier for the SNP-based Inference of Ancestry. [published erratum appears in J Forensic Sci 2004 Sept;49(5)] J Forensic Sci 2003 July;48(4):771–782.

217485 TACCTTCTTTCTAATACAAGCATATGTTAG M ATTAAGTTCTAGGCATACTTTCAAAGCT	217460 AGTGGAAATGGGCAACCCTTCTGTTTTTTGC M GCGCTCTTTGTTCTGATGGAGGTAAGATTT
217486 TGGGCATTTCTAAAATGTTAAACATAAAC W CATTTCATTCATGGATATTTGTCAACAGA	554363 TCRCATGCCCTRCAYCACTGCCGTGATTCA Y GAGGTGCAGCGCTTTGGGGACATCGTCCCC
217487 TAAAGAAAACCACAGTTATTAATTAAGAA [AATT] AATTAATTATGTGTAGTTATAACCAATGA	554368 GGCYACTGCCAGGTGGGCCASTCTAGGAA M CCTGGCCACCYAGTCTCAATGCCACCACA
217489 GTATTTTCCAAGTAAAATATTAACATATTA Y TTCATTGGTCTTCTTTTTATCTGGTTCTA	554370 ACCCGCCCTGGCCTGACTCTGCCACTGGCA R CACAGTCAACACAGCAGGTTCACTCACAGC
217468 ATGTGTCAATGGATGCACTGCTTGGGGGAT M TGAAATCTGGAGAGACATTGATTTT [TCT] GC	554371 CACCGGGCCCAACGCTGGGCTGCACGCTAC Y CACCAGGCCCCCTGCCACTGCCCGGGCTGG
217473 TCCTCTGCAGTATTTTTGAGCAGTGGCTCC R AAGGCACCGTCTCTTCAAGAAGTTATCC	554353 ATTTTGCTGTAAAATTAAGCACTTGAATAG R TAAAATTTGYATTAGTTTCTAAACTGGAGT
217480 CATTTGCAAAATTGTAACCTAATACAAAGT R TAGCCTTCTTCCAACCTCAGGTAGAACACAC	664784 GCCGACCGCCCGCTGYGCCCATCACCCAG R TCCTGGGYTTCGGGCCGCGYTCCCAAGGCA
217438 TTGTCGGASCTGCTGGTGAGCGGGASSAAC Y TGCTGGAGACGGCCGTATCCTCTGTCTGG	664785 CCCGCCTGYGCCCATCACCCAGRTCTGGG Y TTCGGGCCGCGYTCCCAAGGCAAGCRGGG
217439 GACCCTACATCTCCATCTTCTACGCACTG Y GCTACCACAGCAYCGTGACCCTGCCGYGGG	664793 TTGGACACAATGGATTAGGCTGATATGAC M AAAGAGTTTGGAAAAGACCAATTAATAA
217441 CTGYGCTACCACAGCAYCGTGACCCTGCCG Y GGGCGSGCRASSCGTTGGGGCCATCTGGG	664802 GATCACTAGCACATCATTGGAGTGAACAT Y GACTCTCTCAACAATCCACAAGACCCCTTT
217452 TTCTGCAGAGAGACGGTGTCCATCAGCATC Y GGGCCTCCYTGACAGACCCAGGCTGTCC	664803 TGTGGACTACTATTTCTTTAATTTATCTT K CTCTCTTAAAAATAACTGCTTTATTGAGAT
217455 TTCTTTCCAGATCGTGCACAGAACTCTGGC R GCCATGCTGGGTCCCTTGCAGCACTGGCA	712047 TGTCCTTAAACTCTTCTCATTGCCTTAC Y TATGATGTATTTTAAACTGGCAAAATA
217456 GTGTGTGTGGCCAGGCATACCGGCTCTCCC R GGGACGGGTGTGGCCATGATCATCATGCT	712043 CCCAATGCCCATGTTCCAGTTCAGAACTGT Y GGGCTATTCAGGCTGTCTTCTTGGTGAAG
217459 CGGGATTCTGCTCGCCAAATGCCTGACAGT K TTGGGATTTGTTATCTTTCATGTTTTCTCT	712064 GTAAATGAGCTGTGGTTTCTCTTACAGC R TAGGATATCTGACGGGATTCTGCTCGCCAA

712037
TGCAGTTTTACCCAATAAGGTGAGTGGATG R
TACATGGAGAAGGAGGGAGGAGTGAACC

712051
CAGCCTAATTACATTCCATAAATGTTGGTA W
CCTTTCCTTTGCTATCTCCGCAAATGTAAG

712052
AGAGAGAATAGACCAGACACCTAGACTTTA R
CAACATTCTCAAAGAACAAGCCATGGAAAAG

712058
GTCCCTGCACGTTGCAGGGCCCGCCCTCTG R
CGGGTGTCCCTGACCACCAGCCCGCTCTCG

712054
TAATTAGGATATCCATAACCTCAAACATTT R
TTGTTACTTGTGTTGGGAACATTCCAAGTC

712055
ATCTATCATTTCATTTGGTTCTTTTTTCT R
TATCTTTTGTTCCTCCCATAGTTTTTCAT

712057
TGTGTGCACCAGTGTGAACTGTGTAGGTT K
TGTGTGGTCCCTGTGCCTGTGTATGTC

756251
GCGACCCCTTACCCGCATCTCCACCCCA R
GACGCCCTTTCCGCCCAACGGTCTCTTGG

615921
GCACAGGATGACCTGGGACCCAGCCAGCM M
CCCCGAGACCTGACTGAGGCCTTCCTGGCA

615926
ACCTCGACCACGCTGGCCTGGGGCCTCCTG Y
TCATGATCCTACATCCGGATGTGCACGGTG

756239
AGTCTCACTATTATCCAGTGCAGTGGTGCA R
TCACAGCTCACTGTAGCCTCTAACTCCCAG

809125
GCAATTCTTAAATCTTGTGCTATGAAGAAA Y
GCTATTAATCCTTCTATTAATGTAACCTG

869787
GAGACTCAAGGCTCTTAATAAAATCTTAAA K
CATTGAGCTGGAGGAATACCTGAAATCA

869777
CGCGTGGCGGAGCAGAGGCGCTTCTCCRT S
TCCACCTTGGCCTTGGGCTGGGCAAG

869784
GACTGGGGCCTCGGAAGAGCAGGATTTGC R
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869785
GCRTAGATGGGTTTGGGAAAGGACATTC Y
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869772
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GTGGAACCAGATTCRGCAAGAAGAACAA

869745
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GTTCCCAATTCTTCTTCAGCTCCAAGG

869769
TGGCAGCTCCCCAGATAACTCCCACCC M
GCCTTAGCCAGAGTGCCCTCCCTCTT

869794
AATAGTAACTTCGTTTGCTGTTATCTCT S
TCTACTTTCCTAGCTCTCAAAGGTCTAT

869797
GAGGATGGAAAACAGAGACTTACAGAGC Y
CCTCGGGCAGAGCTTGGCCCATCCACAT

869798
TGGGATCTCCCTCCTAGTTTCGTTTCTC W
TCCTGTTAGGAATTGTTTTTCAGCAATGG

869802
GTAAGATAATTTCTAACTACTATTATCT S
TTAACAAATACAGTGTGTTTATATCTAAAAG

869809
TTGTACAACAAATGTGCCATTTTTCTCC Y
TTTCCATCAGTTTTTACTTGTGTCTTATC

869810
ATGCTGTGGTGCACGAGGTCCAGAGATAC M
TTGACCTTCTCCCACCAGCCTGCCCAT

869813
TGCCTCCTCTGCAGTGGTACAATTACTCT Y
TGTACATGATCAAGAGCACTGTTCTGAAT

886933
CCAGTACCTTATTGTCTGAAGAGAGCTAA Y
AGAAATAGACTGTCAGAGAGTAGACAAA

886937
TAAACACCTAGAATGTTCAAGGTACTCTA K
AAGTTGCTCCAGGGGAAACAGAAAAGTGCC

886895
TAAGGTACGCAAAGCACCTCTGCCGTGGG R
GTTGCGGCCAGGTTCTGGCAGGAGGGGC

886896
CCATTCTCAGGTGCATGAAAAGGTGGGGGC R
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886894
TAGTACATTTTATCTAACCTCACTGAGCT Y
TGCAGGGGTACACAGCCGAGTTAAGGAC

886892
GCTTTGGTTCATAGGCTTTGTACATTCTG S
ATGGGAAGGTTTTCAGAGCCTGTTCCAGAC

886934
TGTGATCTAGGAAATGTTGCAAGGCCTTC Y
TTCTCCCTCCTTACTGGAATTTGCAACG

886993
AAGATTATCCTTGTCTTCTTTTCCCC R
TAGATGATCTTAGTAGCCATATTTTCAGA

886994
TTGCTTTTTGGTGAATAAATTTCCATGATT M
CTTCTAAATATTGAATATATACACATTTA

951497
AAATAACGTGCTCATTGGATTTAAATAGA R
GGTGCTATCAAATGTGATTTAAGTTATT

951526
CCCCGCAGACACAAGTCCCCAGCCCTCC R
GGACAGCAATAAGGGTCTTACAAGGCCAG