

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	217460
TACCTTCTTTCTAATACAAGCATATGTTAG M	AGTGGAAATGGGCAACCCTTCTGTTTTTTGC M
ATTAAGTTCTAGGCATACTTTCAAAGCT	GCGCTCTTTGTTCTGATGGAGGTAAGATTT
217486	554363
TGGGCATTTCTAAAATGTTAAACATAAAC W	TCRCATGCCCTRCAYCACTGCCGTGATTCA Y
CATTTCATTCATGGATATTTGTCAACAGA	GAGGTGCAGCGCTTTGGGGACATCGTCCCC
217487	554368
TAAAGAAAACCACAGTTATTAATTAAGAA [AATT]	GGCYACTGCCAGGTGGGCCCSTCTAGGAA M
AATTAATTATGTGTAGTTATAACCAATGA	CCTGGCCACCYAGTCTCAATGCCACCACA
217489	554370
GTATTTTCCAAGTAAAATATTAACATATTA Y	ACCCGCCCTGGCCTGACTCTGCCACTGGCA R
TTCATTGGTCTTCTTTTTTATCTGGTTCTA	CACAGTCAACACAGCAGGTTCACTCACAGC
217468	554371
ATGTGTCAATGGATGCACTGCTTGGGGGAT M	CACCGGGCCCAACGCTGGGCTGCACGCTAC Y
TGAAATCTGGAGAGACATTGATTTT [TCT] GC	CACCAGGCCCCCTGCCACTGCCCGGGCTGG
217473	554353
TCCTCTGCAGTATTTTTGAGCAGTGGCTCC R	ATTTTGCTGTAAAATTAAGCACTTGAATAG R
AAGGCACCGTCTCTTCAAGAAGTTTATCC	TAAAATTTGYATTAGTTTCTAAACTGGAGT
217480	664784
CATTTGCAAAATTGTAACCTAATACAAAGT R	GCCGACCGCCCGCTGYGCCCATCACCCAG R
TAGCCTTCTTCCAACCTCAGGTAGAACACAC	TCCTGGGYTTCGGGCCGCGYTCCAAGGCA
217438	664785
TTGTCGGASCTGCTGGTGAGCGGGASSAAC Y	CCCGCCTGYGCCCATCACCCAGRTCTGGG Y
TGCTGGAGACGGCCGTATCCTCTGTCTGG	TTCGGGCCGCGYTCCAAGGCAAGCRGGG
217439	664793
GACCCTACATCTCCATCTTCTACGCACTG Y	TTGGACACAATGGATTAGGCTGATATGAC M
GCTACCACAGCAYCGTGACCCTGCCGYGGG	AAAGAGTTTGGAAAAGACCAATTAATAA
217441	664802
CTGYGCTACCACAGCAYCGTGACCCTGCCG Y	GATCACTAGCACATCATTGGAGTGAACAT Y
GGGCGSGCRASSCGTTGGCGCCATCTGGG	GACTCTCTCAACAATCCACAAGACCCCTTT
217452	664803
TTCTGCAGAGAGACGGTGTCCATCAGCATC Y	TGTGGACTACTATTTCTTTAATTTATCTT K
GGGCCTCCYTGCAGCAGACCCAGGCTGTCC	CTCTCTTAAAAATAACTGCTTTATTGAGAT
217455	712047
TTCTTTCCAGATCGTGCACAGAACTCTGGC R	TGTCCTTAAACTCTTCTCATTGCCTTAC Y
GCCATGCTGGGTCCCTTGCAGCACTGGCA	TATGATGTATTTTAAACTGGCAAAATA
217456	712043
GTGTGTGTGGCCAGGCATAACGGCTCTCCC R	CCCAATGCCCATGTTCCAGTTCAGAACTGT Y
GGGACGGGTGTGGCCATGATCATCATGTCT	GGGCTATTCAGGCTGTCTTCTTGGTGAAG
217459	712064
CGGGATTCTGCTCGCCAAATGCCTGACAGT K	GTAATGAGCTGTGGTTTCTCTTTACAGC R
TTGGGATTTGTTATCTTTCATGTTTTTCCTC	TAGGATATCTGACGGGATTCTGCTCGCCAA

712037
TGCAGTTTTACCCAATAAGGTGAGTGGATG R
TACATGGAGAAGGAGGGAGGAGTGAACC

712051
CAGCCTAATTACATTCCATAAATGTTGGTA W
CCTTTCCTTTGCTATCTCCGCAAATGTAAG

712052
AGAGAGAATAGACCAGACACCTAGACTTTA R
CAACATTCTCAAAGAACAAGCCATGAAAAG

712058
GTCCCTGCACGTTGCAGGGCCCGCCCTCTG R
CGGGTGTCCCTGACCACCAGCCCGCTCTG

712054
TAATTAGGATATCCATAACCTCAAACATTT R
TTGTTACTTGTGTTGGGAACATTCCAAGTC

712055
ATCTATCATTTCATTTGGTTCTTTTTTCT R
TATCTTTTGTCTTCCCATAGTTTTTCAT

712057
TGTGTGCACCAGTGTGAACTGTGTAGGTT K
TGTGTGGTCCCTGTGCCTGTGTATGTC

756251
GCGACCCCTTACCCGCATCTCCACCCCA R
GACGCCCTTTCCGCCAACGGTCTCTTGG

615921
GCACAGGATGACCTGGGACCCAGCCAGCM M
CCCCGAGACCTGACTGAGGCCTTCCTGGCA

615926
ACCTCGACCACGCTGGCCTGGGGCCTCCTG Y
TCATGATCCTACATCCGGATGTGCAGCGTG

756239
AGTCTCACTATTATCCAGTGCAGTGGTGCA R
TCACAGCTCACTGTAGCCTCTAACTCCCAG

809125
GCAATTCTTAAATCTTGTGCTATGAAGAAA Y
GCTATTAATCCTTCTATTAATGTAACCTG

869787
GAGACTCAAGGCTCTTAATAAAATCTTAAA K
CATTTGAGCTGGAGGAATACCTGAAATCA

869777
CGCGTGGCGGAGCAGAGGCGCTTCTCCRT S
TCCACCTTGGCCTTGGGCTGGGCAAG

869784
GACTGGGGCCTCGGAAGAGCAGGATTTGC R
TAGATGGGTTTGGGAAAGGACATTCYAGG

869785
GCRTAGATGGGTTTGGGAAAGGACATTC Y
AGGAGACCCCACTGTAAGAAGGGCCTGG

869772
TGTCATTAACCTTTTTAAAAATCTACCAA Y
GTGGAACCAGATTCRGCAAGAAGAACAA

869745
AATCACAGAAATTAACCTTGCTGGAAATC Y
GTTCCCAATTCTTCTTCAGCTCCAAGG

869769
TGGCAGCTCCCCAGATAACTCCCACCC M
GCCTTAGCCAGAGTGCCCTCCCTCTT

869794
AATAGTAACTTCGTTTGCTGTTATCTCT S
TCTACTTTCCTAGCTCTCAAAGGTCTAT

869797
GAGGATGGAAAACAGAGACTTACAGAGC Y
CCTCGGGCAGAGCTTGGCCATCCACAT

869798
TGGGATCTCCCTCCTAGTTTCGTTTCTC W
TCCTGTTAGGAATTGTTTTTCAGCAATGG

869802
GTAAGATAATTTCTAACTACTATTATCT S
TTAACAAATACAGTGTGTTTATATCTAAAG

869809
TTGTACAACAAATGTGCCATTTTTCTCC Y
TTTCCATCAGTTTTTACTTGTGTCTTATC

869810
ATGCTGTGGTGCACGAGGTCCAGAGATAC M
TTGACCTTCTCCCACCAGCCTGCCCAT

869813
TGCCTCCTCTGCAGTGGTACAATTACTCT Y
TGTACATGATCAAGAGCACTGTTCTGAAT

886933
CCAGTACCTTATTGTCTGAAGAGAGCTAA Y
AGAAATAGACTGTCAGAGAGTAGACAAA

886937
TAAACACCTAGAATGTTCAAGGTACTCTA K
AAGTTGCTCCAGGGGAAACAGAAAGTGCC

886895
TAAGGTACGCAAAGCACCTCTGCCGTGGG R
GTTGCGGCCAGTTCTGGCAGGAGGGGC

886896
CCATTCTCAGGTGCATGAAAAGGTGGGGG R
GTTGAGCCACAGCTCACTGCATTCCAGTC

886894
TAGTACATTTTATCTAACCTCACTGAGCT Y
TGCAGGGGTACACAGCCGAGTTTAAAGGAC

886892
GCTTTGGTTCATAGGCTTTGTACATTCTG S
ATGGGAAGGTTTTCAGAGCCTGTTCCAGAC

886934
TGTGATCTAGGAAATGTTGCAAGAGCCTTC Y
TTCTCCCTCCTTACTGGAATTTGCAACG

886993
AAGATTATCCTTGTCTTCTTTTCCCC R
TAGATGATCTTAGTAGCCATATTTTCAGA

886994
TTGCTTTTTGGTGAATAAATTTCCATGATT M
CTTCTAAATATTGAATATATACACATTTA

951497
AAATAACGTGCTCATTGGATTTAAATAGA R
GGTGCTATCAAATGTGATTTAAGTTATT

951526
CCCCGCAGACACAAGTCCCCAGCCCTCC R
GGACAGCAATAAGGGTCTTACAAGGCCAG