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Supplemental Data

Mutability of Y-Chromosomal Microsatellites:

Rates, Characteristics, Molecular Bases,

and Forensic Implications

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Table S1. Bayesian Median Mutation Rates, Mutation Summaries, Repeat Structures, and Genotyping Details of 186 Y-STRs from Analysing DNA-Confirmed Father-Son Pairs (See Footnote for Explanation)

| GBD ID | Repeat Structure (as defined in Methods and Materials) | Bayesian median mutation rate | Bayesian 95% credible interval | Gains/Losses | Total Mutations | Total Meioses | Primer 1 Sequence 5' - 3' | Primer 2 Sequence 5' - 3' | Ta in °C | Multiplex | Ref |
|---------------|---|-------------------------------|--|--------------|-----------------|---------------|-----------------------------|------------------------------|-------------|-----------|------|
| DYF380S1 | (AAT) ₈₋₁₁ | 3.84x10 ⁻⁴ | 1.42x10 ⁻⁵ – 2.06x10 ⁻³ | 0/0 | 0 | 1790 | AGCCATGTGGAT TCACCACT | GACAAACCCATC CTGTCTCC | TD 70-50 | 28 | [29] |
| DYF381S1 | (TTG) ₇₋₈ | 3.95 x10 ⁻⁴ | 1.43 x10 ⁻⁵ – 2.07x10 ⁻³ | 0/0 | 0 | 1774 | TCCATCCATCAA TCCATCAA | CAACCCAAACA CTTGCAGAA | TD 60-50 | 45 | [29] |
| DYF381S2 | (AAC) ₇₋₈ | 3.91 x10 ⁻⁴ | 1.44 x10 ⁻⁵ – 2.11x10 ⁻³ | 0/0 | 0 | 1756 | CAACCCAAACA CTTGCAGAA | TCCATCCATCAA TCCATCAA | TD 60-50 | 20 | [29] |
| DYF382S1 | (GGAT) ₉₋₁₆ (AGAT) ₁ (GGAT) ₃ N ₈ (GGAC) ₃ | 1.05 x10 ⁻³ | 1.52 x10 ⁻⁴ – 3.47x10 ⁻³ | 1/0 | 1 | 1609 | TTGTAAAATGGG CATGTGGA | CCCAAAGTGCTA CCCACCTA | TD 70-50 | 54 | [29] |
| DYF386S1 | (AAT) ₇₋₁₆ | 6.02 x10 ⁻³ | 3.10 x10 ⁻³ – 1.04x10 ⁻² | 3/7 | 10 | 1772 | GACTGCTCAACT GCACTCCA | CCAATGTTACTC ACTATGCTGCTT | TD 70-50 | 29 | [29] |
| DYF387S1 | (AAAG) ₃ (GTAG) ₁ (GAAG) ₄ N ₁₆ (GAAG) ₉ (AAAG) ₁₃ | 1.59 x10 ⁻² | 1.08 x10 ⁻² – 2.24x10 ⁻² | 15/13 | 28 | 1804 | GCCTGGGTGAC AGAGCTAGA | GCCACAGTGTGA GAAGTGTGA | TD 70-50 | 49 | [29] |
| DYF388S1 | (CTTC) ₆ (CTTT) ₅₋₁₉ N ₁₈ (CTTC) ₃ (TTTC) ₁ (CTTC) ₃ N ₈ (CTTC) ₃ N ₃₂ (CTTC) ₃ | 6.85 x10 ⁻³ | 3.64 x10 ⁻³ – 1.15x10 ⁻² | 5/6 | 11 | 1702 | TTCTAGGAAGAT TAGCCACAACA | CCCAGACAACA GAGCAAAAC | TD 65-50 | 52 | [29] |
| DYF390S1 | (TTTA) ₉₋₁₄ | 9.95 x10 ⁻⁴ | 1.42 x10 ⁻⁴ – 3.34x10 ⁻³ | 0/1 | 1 | 1680 | AGCATTCCCTTT CTCATTGC | TGACGAGTTAGT GGGTGCAG | TD 70-50 | 12 | [29] |
| DYF393S1 | (AAG) ₄ (AA) ₁ (AAG) ₁₆₋₃₀ (CAG) ₁₋₂ | 8.57 x10 ⁻³ | 4.91 x10 ⁻³ – 1.37x10 ⁻² | 9/5 | 14 | 1712 | GCAACCAAAAAG GTTTGGAGA | GTGGAGCCTGCT TAAAGGAA | TD 70-50 | 31 | [29] |
| DYF394S1 | (ATT) ₃ (GTT) ₁ (ATT) ₆₋₉ | 3.91 x10 ⁻⁴ | 1.40 x10 ⁻⁵ – 2.09x10 ⁻³ | 0/0 | 0 | 1768 | GCCCTGAACAA AATCTGGAG | GCAGTGAGCTG AGATGGTGA | TD 70-50 | 24 | [29] |
| DYF396S1 | (TCT) ₆₋₉ | 3.86 x10 ⁻⁴ | 1.37 x10 ⁻⁵ – 2.06x10 ⁻³ | 0/0 | 0 | 1785 | TGCACGTCTTCA TACATAGAGC | TTGAATGCCAAG TTATGTAGCA | TD 70-50 | 36 | [29] |
| DYF399S1 | (GAAA) ₃ N ₇₋₈ (GAAA) ₁₀₋₂₃ | 7.73 x10 ⁻² | 6.51 x10 ⁻² – 9.09x10 ⁻² | 55/84 | 139 | 1794 | GGGTTTTACCA GTTTGCAT | CCATGTTTTGGG ACATTCCT | TD 70-50 | 49 | [29] |
| DYF401S1 | (AAGG) ₃ (AAGC) ₁ (AAGG) ₃ N ₃₉ (AAGG) ₃ N ₈ (AAGG) ₃ (AAAG) ₁ (AAGG) ₃ N ₁₃ (AAAG) ₈₋₂₃ G(AAGG) ₆ | 6.50 x10 ⁻³ | 3.08 x10 ⁻³ – 1.18x10 ⁻² | 3/5 | 8 | 1333 | TCGCAACATA GCACTTCAG | TTCTAGGAAGAT TAGCCACAACA | TD 70-50 | 48 | [29] |
| DYF403S1 a | (TTCT) ₁₀₋₁₇ N ₂₋₃ (TTCT) ₃₋₁₇ | 3.10 x10 ⁻² | 2.30 x10 ⁻² – 4.07x10 ⁻² | 29/17 | 46 | 1504 | CAAAATTCATGT GGATAATGAG | ACAGAGCAGGA TTCCATCTA | TD 70-50 | 54 | [29] |
| DYF403S1 b | (TTCT) ₁₂ N ₂ (TTCT) ₈ (TTCC) ₉ (TTCT) ₁₄ N ₂ (TTCT) ₃ | 1.19 x10 ⁻² | 7.05 x10 ⁻³ – 1.86x10 ⁻² | 5/11 | 16 | 1402 | CAAAATTCATGT GGATAATGAG | ACAGAGCAGGA TTCCATCTA | TD 70-50 | 54 | [29] |
| DYF404S1 | (TTTC) ₁₀₋₂₀ N ₄₂ (TTTC) ₃ | 1.25 x10 ⁻² | 7.92 x10 ⁻³ – 1.84x10 ⁻² | 14/7 | 21 | 1739 | GGCTTAAGAAA TTTCAACGCATA | CCATGATGGAAC AATTGCAG | TD 70-50 | 38 | [29] |
| DYF405S1 | (GGAA) ₄₋₁₄ N ₁₁₅ (GGAA) ₃ (GAAA) ₁ (GGAA) ₃ | 1.52 x10 ⁻³ | 3.54 x10 ⁻⁴ – 4.13x10 ⁻³ | 1/1 | 2 | 1756 | CCGTGGTGTCTG AAGCATAG | CACATCAAGTTG CCTGTTTCA | TD 70-50 | 26 | [29] |
| DYF406S1 | (TATC) ₈₋₁₄ | 3.82 x10 ⁻³ | 1.61 x10 ⁻³ – 7.48x10 ⁻³ | 3/3 | 6 | 1744 | CCTGGGTGACAC AGTGAGACT | TCCACCAAAATT CCATGACA | TD 70-50 | 28 | [29] |

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|-------------------|--|-------------------------|--|------|----|------|---------------------------------|-----------------------------|-------------|--------|------|
| DYF410S1 | (AAAT) ₇₋₁₃ | 2.04 x 10 ⁻³ | 4.82 x 10 ⁻⁴ – 5.52x10 ⁻³ | 2/0 | 2 | 1309 | TGACGAGTTAGT GGGTGCAG | GCGGCTAGGGT AGAATCCAT | TD 70-50 | 48 | [29] |
| DYS19/ DYS394 | (TAGA) ₃ (TAGG) ₁ (TAGA) ₆₋₁₆ | 4.37 x 10 ⁻³ | 1.98 x 10 ⁻³ – 8.23x10 ⁻³ | 4/3 | 7 | 1756 | Yfiler | Yfiler | Yfiler | Yfiler | [29] |
| DYS385a | (AAGG) ₄ N ₁₄ (AAAG) ₃ N ₁₂ (AAA G) ₃ N ₂₉ (AAGG) ₆₋₇ (GAAA) ₇₋₂₃ | 2.08 x 10 ⁻³ | 6.24 x 10 ⁻⁴ – 5.06x10 ⁻³ | 2/1 | 3 | 1762 | Yfiler | Yfiler | Yfiler | Yfiler | [29] |
| DYS385b | (AAGG) ₄ N ₁₄ (AAAG) ₃ N ₁₂ (AAAG) ₃ N ₂₉ (AAGG) ₆₋₇ (GAAA) ₇₋₂₃ | 4.14 x 10 ⁻³ | 1.75 x 10 ⁻³ – 8.09x10 ⁻³ | 6/0 | 6 | 1615 | Yfiler | Yfiler | Yfiler | Yfiler | [29] |
| DYS388 | (ATT) ₉₋₁₈ | 4.25 x 10 ⁻⁴ | 1.51 x 10 ⁻⁵ – 2.26x10 ⁻³ | 0/0 | 0 | 1635 | GTGAGTTAGCCG TTTAGCGA | CAGATCGCAACC ACTGCG | TD 60-50 | 50 | [29] |
| DYS389I | (TCTG) ₃ (TCTA) ₆₋₁₄ | 5.51 x 10 ⁻³ | 2.72 x 10 ⁻³ – 9.74x10 ⁻³ | 4/5 | 9 | 1751 | Yfiler | Yfiler | Yfiler | Yfiler | [29] |
| DYS389II | (TCTG) ₄₋₅ (TCTA) ₁₀₋₁₄ N ₂₈ (TCTG) ₃ (TCTA) ₆₋₁₄ | 3.83 x 10 ⁻³ | 1.61 x 10 ⁻³ – 7.49x10 ⁻³ | 2/4 | 6 | 1743 | Yfiler | Yfiler | Yfiler | Yfiler | [29] |
| DYS390 | (TCTG) ₈ (TCTA) ₉₋₄ (TCTG) ₁ (TCTG) ₄ | 1.52 x 10 ⁻³ | 3.52 x 10 ⁻⁴ – 4.09x10 ⁻³ | 0/2 | 2 | 1758 | Yfiler | Yfiler | Yfiler | Yfiler | [29] |
| DYS391 | (TCTG) ₃ (TCTA) ₆₋₁₅ | 3.23 x 10 ⁻³ | 1.26 x 10 ⁻³ – 6.65x10 ⁻³ | 3/2 | 5 | 1759 | Yfiler | Yfiler | Yfiler | Yfiler | [29] |
| DYS392 | (TAT) ₄₋₂₀ | 9.70 x 10 ⁻⁴ | 1.43 x 10 ⁻⁴ – 3.23x10 ⁻³ | 1/0 | 1 | 1728 | Yfiler | Yfiler | Yfiler | Yfiler | [29] |
| DYS393/ DYS395 | (AGAT) ₇₋₁₈ | 2.11 x 10 ⁻³ | 6.21 x 10 ⁻⁴ – 5.00x10 ⁻³ | 2/1 | 3 | 1750 | Yfiler | Yfiler | Yfiler | Yfiler | [29] |
| DYS425/ DYF371 | (TGT) ₈₋₁₄ | 1.51 x 10 ⁻³ | 3.48 x 10 ⁻⁴ – 4.08x10 ⁻³ | 1/1 | 2 | 1778 | TGGAGAGAAGA AGAGAGAAAT | AGTAATTCTGGA GGTAAAAATGG | TD 60-50 | 20 | [29] |
| DYS426 | (GTT) ₉₋₁₂ | 3.98 x 10 ⁻⁴ | 1.49 x 10 ⁻⁵ – 2.11x10 ⁻³ | 0/0 | 0 | 1735 | CTCAAAGTATGA AAGCATGACCA | GGTGACAAGAC GAGACTTTGTG | TD 70-50 | 38 | [29] |
| DYS434 | (ATCT) ₉₋₁₂ | 4.04 x 10 ⁻⁴ | 1.47 x 10 ⁻⁵ – 2.14x10 ⁻³ | 0/0 | 0 | 1715 | CACTCCCTGAGT GCTGGATT | GGAGATGAATG AATGGATGGA | TD 60-50 | 40 | [29] |
| DYS435 | (TGGA) ₁₀₋₁₂ | 1.00 x 10 ⁻³ | 1.47 x 10 ⁻⁴ – 3.33x10 ⁻³ | 0/1 | 1 | 1676 | AGCATCTCCACA CAGCACAC | TTCTCTCTCCCC CTCCTCTC | TD 60-50 | 41 | [29] |
| DYS436 | (GTT) ₁₁₋₁₃ | 3.84 x 10 ⁻⁴ | 1.38 x 10 ⁻⁵ – 2.05x10 ⁻³ | 0/0 | 0 | 1798 | CCAGGAGAGCA CACACAAAA | GCAATCCAACCT CAGCCAAT | TD 60-50 | 18 | [29] |
| DYS437 | (TCTA) ₄₋₁₂ (TCTG) ₂ (TCTA) ₄ | 1.53 x 10 ⁻³ | 3.54 x 10 ⁻⁴ – 4.10x10 ⁻³ | 2/0 | 2 | 1760 | Yfiler | Yfiler | Yfiler | Yfiler | [29] |
| DYS438 | (TTTTTC) ₇₋₁₆ | 9.56 x 10 ⁻⁴ | 1.37 x 10 ⁻⁴ – 3.18x10 ⁻³ | 0/1 | 1 | 1751 | Yfiler | Yfiler | Yfiler | Yfiler | [29] |
| DYS439 | (GATA) ₃ N ₃₂ (GATA) ₅₋₁₉ | 3.84 x 10 ⁻³ | 1.63 x 10 ⁻³ – 7.54x10 ⁻³ | 2/4 | 6 | 1736 | Yfiler | Yfiler | Yfiler | Yfiler | [29] |
| DYS441 | (TTCC) _{12-21.2} | 1.18 x 10 ⁻³ | 1.66 x 10 ⁻⁴ – 3.93x10 ⁻³ | 1/0 | 1 | 1419 | ATGTACCTGTAG CCCCAGTGAAC | AAGTTGCAGTGA GCGAAGATTG | TD 70-50 | 27 | [45] |
| DYS442 | (GATA) ₉₋₁₆ (GACA) ₃ | 9.78 x 10 ⁻³ | 5.59 x 10 ⁻³ – 1.57x10 ⁻² | 2/12 | 14 | 1497 | AAACGCCCATC AATCAATGAGT G | CCCCAAGTCCCC AAAGTGTGT | TD 70-50 | 16 | [45] |
| DYS443 | (TTCC) ₁₁₋₁₇ (CTT) ₃ | 2.10 x 10 ⁻³ | 6.24 x 10 ⁻⁴ – 5.01x10 ⁻³ | 2/1 | 3 | 1745 | GAGTTCATGCTG ATGACAAGC | TCATTGGCCACC TGACATTA | TD 70-50 | 29 | [29] |
| DYS444 | (TAGA) ₉₋₁₆ | 5.45 x 10 ⁻³ | 2.68 x 10 ⁻³ – 9.65x10 ⁻³ | 3/6 | 9 | 1775 | TGTGAACCATTT GGCATGTT | TCACGTTGTTCA AGGGTCAA | TD 60-50 | 45 | [29] |

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|--------|---|-------------------------|--|------|----|------|--------------------------------------|------------------------------------|------------------|------------------|------|
| DYS445 | (TTTA) ₆₋₁₄ | 2.16 x 10 ⁻³ | 6.38 x 10 ⁻⁴ – 5.15x10 ⁻³ | 2/1 | 3 | 1704 | GAGCTGAGATT ATGCCACCAAA A | AGTTAAGAGCCC CACCTTCCTG | TD 70-50 | 32 | [45] |
| DYS446 | (TCTCT) ₈₋₂₁ | 2.67 x 10 ⁻³ | 9.38 x 10 ⁻⁴ – 5.87x10 ⁻³ | 2/2 | 4 | 1747 | TATTTTCAGTCT TGTCCCTGTC | AAATGTATGGCC AACATAGCAAA ACC | TD 70-50 | 30 | [45] |
| DYS447 | (TTATA) ₆₋₇ (TTATT) ₁ (TTATA) ₈₋₁₃ (TTATT) ₁ (TTATA) ₅₋₉ | 2.12 x 10 ⁻³ | 6.28 x 10 ⁻⁴ – 5.11x10 ⁻³ | 1/2 | 3 | 1722 | GGGCTTGCTTTG CGTTATCT | GGTCACAGCATG GCTTGGTT | TD 70-50 | 27 | [45] |
| DYS448 | (AGAGAT) ₁₁₋₁₃ N ₄₂ (AGAGAT) _{8,9} | 3.94 x 10 ⁻⁴ | 1.41 x 10 ⁻⁵ – 2.11x10 ⁻³ | 0/0 | 0 | 1747 | Yfiler | Yfiler | Yfiler Yfiler | Yfiler Yfiler | [45] |
| DYS449 | (TTCT) ₁₃₋₉ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₃₋₁₉ | 1.22 x 10 ⁻² | 7.54 x 10 ⁻³ – 1.85x10 ⁻² | 14/5 | 19 | 1617 | TGGAGTCTCTCA AGCCTGTTC | CCATTGCACTCT AGGTTGGAC | TD 60-50 | 46 | [29] |
| DYS450 | (TTTTA) ₇₋₁₁ N ₁₂ (TTTTA) ₃ | 1.04 x 10 ⁻³ | 1.54 x 10 ⁻⁴ – 3.50x10 ⁻³ | 0/1 | 1 | 1598 | GCCTTTCCAATT TCAATTTCTGA | TGGAATATGATG CAGCTGTTTGT | TD 70-50 | 39 | [45] |
| DYS452 | (TATAC) ₅₋₄ [(CATAC) ₁ (TATAC) ₁] ₂₋₄ N ₂₀ (TATAC) ₃ (CATAC) ₁ (TATAC) ₃ | 4.02 x 10 ⁻³ | 1.56 x 10 ⁻³ – 8.28x10 ⁻³ | 2/3 | 5 | 1411 | TTTATTATACTC AGCTAATTAATT GGTT | GTGGTGTCTGA TGAGGATAAT | TD 70-50 | 16 | [45] |
| DYS453 | (AAAT) ₉₋₁₅ | 3.89 x 10 ⁻⁴ | 1.43 x 10 ⁻⁵ – 2.08x10 ⁻³ | 0/0 | 0 | 1782 | GGGTAACAGAA CAAGACAGT | CTAAAAGTATGG ATATTCTTCG | TD 60-50 | 45 | [45] |
| DYS454 | (AAAT) ₇₋₁₃ | 4.75 x 10 ⁻⁴ | 1.71 x 10 ⁻⁵ – 2.55x10 ⁻³ | 0/0 | 0 | 1458 | TCACAATGACCC TTTTGTGC | GTTCTTTGGCCC TGCATTTA | TD 60-50 | 46 | [29] |
| DYS455 | (ATTT) _{6,2-11} | 4.26 x 10 ⁻⁴ | 1.59 x 10 ⁻⁵ – 2.28x10 ⁻³ | 0/0 | 0 | 1618 | ATCTGAGCCGA GAGAATGATA | GGGGTGGAAAC GAGTGTT | TD 70-50 | 39 | [45] |
| DYS456 | (AGAT) ₁₁₋₂₃ | 4.94 x 10 ⁻³ | 2.35 x 10 ⁻³ – 8.97x10 ⁻³ | 6/2 | 8 | 1757 | Yfiler | Yfiler | Yfiler Yfiler | Yfiler Yfiler | [45] |
| DYS458 | (GAAA) ₁₁₋₂₄ | 8.36 x 10 ⁻³ | 4.80 x 10 ⁻³ – 1.34x10 ⁻² | 7/7 | 14 | 1756 | Yfiler | Yfiler | Yfiler Yfiler | Yfiler Yfiler | [45] |
| DYS459 | (ATTT) ₆₋₁₁ | 2.67 x 10 ⁻³ | 9.36 x 10 ⁻⁴ – 5.86x10 ⁻³ | 2/2 | 4 | 1741 | CAGGTGAACTG GGGTAAATAAT | TTGAGCAACAG AGCAAGACTTA | TD 70-50 | 27 | [45] |
| DYS460 | (TAGA) ₈₋₁₃ | 6.22 x 10 ⁻³ | 3.19 x 10 ⁻³ – 1.07x10 ⁻² | 2/8 | 10 | 1717 | GCCAAACTCTTT CCAAGAAG | TCTATCCTCTGC CTATCATTTATT A | TD 60-50 | 20 | [29] |
| DYS461 | (TAGA) ₈₋₁₃ | 9.89 x 10 ⁻⁴ | 1.40 x 10 ⁻⁴ – 3.29x10 ⁻³ | 0/1 | 1 | 1695 | AGGCAGAGGAT AGATGATATGG AT | TTCAGGTAAATC TGTCCAGTAGTG A | TD 60-50 | 19 | [29] |
| DYS462 | (CATA) ₉₋₁₄ | 2.65 x 10 ⁻³ | 9.20 x 10 ⁻⁴ – 5.80x10 ⁻³ | 1/3 | 4 | 1771 | TGTGCTGTACCA GTTGCCTA | CCAGCCTGAGCA AGAGAGTA | TD 70-50 | 30 | [45] |
| DYS463 | (AAAGG) ₆₋₇ (AAGGG) ₉₋₁₉ | 1.51 x 10 ⁻³ | 3.49 x 10 ⁻⁴ – 4.07x10 ⁻³ | 2/0 | 2 | 1776 | AATTCTAGGTTT GAGCAAAGACA | ATGAGGTTGTGT GACTTGACTG | TD 70-50 | 33 | [45] |
| DYS464 | (CCTT) ₉₋₂₀ N ₄₆ (CCTT) ₃ N ₈ (CCTT) ₄ | 7.27 x 10 ⁻³ | 3.96 x 10 ⁻³ – 1.20x10 ⁻² | 5/7 | 12 | 1745 | TTACGAGCTTTG GGCTATG | CCTGGGTAACAG AGAGACTCTT | TD 70-50 | 31 | [45] |
| DYS468 | (CTG) ₄ N ₄₄ (CCT) ₃ N ₄₀ (CTT) ₃ N ₃₅ (CCT) ₄ N ₈ (CTC) ₄ (CTT) ₇₋₉ (ATTCAT) ₈₋₁₀ | 1.74 x 10 ⁻³ | 4.03 x 10 ⁻⁴ – 4.69x10 ⁻³ | 1/1 | 2 | 1535 | GGGAGTTCCAA ACTTTTTTCA | GGGGGAAGATG ACAATGATG | TD 70-50 | 37 | [29] |
| DYS469 | (CTT) ₃ N ₃₉ (CTT) ₄ (GTT) ₁ (CTT) ₁₀ | 2.99 x 10 ⁻³ | 1.04 x 10 ⁻³ – | 3/1 | 4 | 1555 | TTTGGGGACTGA | CCCCAGCTGGTA | TD | 41 | [29] |

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|--------|--|-----------------------|--|-----|---|------|----------------------------|-----------------------------|-------------|----|------|
| | ${}_{30}\text{T}(\text{CTT})_3\text{N}_{17}(\text{CTT})_5\text{N}_{37}(\text{CTT})_3$ $\text{N}_{12}(\text{CTT})_4\text{N}_{12}(\text{CTT})_3\text{N}_{12}(\text{CTT})_5$ $(\text{CCT})_4\text{N}_9(\text{CTT})_3(\text{CCT})_3$ | | 6.54×10^{-3} | | | | ATTCAAAA | AAATGAGT | 60-50 | | |
| DYS470 | $(\text{GTT})_{8-12}\text{N}_{33}(\text{GTT})_3$ | 4.20×10^{-4} | 1.51×10^{-5} – 2.23×10^{-3} | 0/0 | 0 | 1651 | GGTCCTTCAGGA ACCAGTTG | TGGCTGTAAAAC AAATATCAGCA | TD 60-50 | 44 | [29] |
| DYS472 | $(\text{AAT})_{7-9}$ | 4.46×10^{-4} | 1.62×10^{-5} – 2.37×10^{-3} | 0/0 | 0 | 1549 | AGATTGTCCAC CTGCACTC | GAGGCACTGTGT TCAGCAAA | TD 70-50 | 1 | [29] |
| DYS473 | $(\text{AAT})_8\text{N}_{12}(\text{AAT})_{9-13}$ | 4.13×10^{-4} | 1.47×10^{-5} – 2.21×10^{-3} | 0/0 | 0 | 1676 | CAGCCTGGATA GCAGAGTGA | CCTCTTTTCTTTG CTGGTTCCTT | TD 60-50 | 44 | [29] |
| DYS474 | $(\text{AAC})_{9-10}$ | 3.92×10^{-4} | 1.41×10^{-5} – 2.08×10^{-3} | 0/0 | 0 | 1766 | CCCCTGAACTTA AAAGGTGGA | GGCATCTAGGTT TACTGTGAGGA | TD 60-50 | 22 | [29] |
| DYS475 | $(\text{TAA})_{7-9}(\text{CAA})_1(\text{TAA})_3$ | 4.14×10^{-4} | 1.54×10^{-5} – 2.21×10^{-3} | 0/0 | 0 | 1681 | CCCACCAAGGG TTTTCAGA | CCCACAGAAAG ATGTTGAGG | TD 60-50 | 23 | [29] |
| DYS476 | $(\text{TGA})_{7-13}$ | 9.40×10^{-4} | 1.35×10^{-4} – 3.12×10^{-3} | 1/0 | 1 | 1779 | CGACTATGATTT GGGCTGTG | AGCTGGGAAGT ACTCAATGCTC | TD 70-50 | 7 | [29] |
| DYS477 | $(\text{TTG})_{8-9}$ | 3.91×10^{-4} | 1.37×10^{-5} – 2.07×10^{-3} | 0/0 | 0 | 1765 | TAACTTACAGAA AAGCTCAGGG | AAGTGAATCGA GTGCCTAGC | TD 60-50 | 42 | [29] |
| DYS478 | $(\text{CAG})_4(\text{CAA})_1(\text{CAG})_8$ | 4.04×10^{-4} | 1.46×10^{-5} – 2.17×10^{-3} | 0/0 | 0 | 1718 | ACAGGCAACAA ATTGGGTA | TCAGGATAAGCT AGCAGTCTATG | TD 60-50 | 20 | [29] |
| DYS480 | $(\text{TTA})_{6-10}$ | 3.91×10^{-4} | 1.44×10^{-5} – 2.09×10^{-3} | 0/0 | 0 | 1783 | CCAGCACCTAG GTTGAGGTA | CAGCACTCCAAA ATGACAGA | TD 70-50 | 9 | [29] |
| DYS481 | $(\text{CTT})_{22-32}$ | 4.97×10^{-3} | 2.36×10^{-3} – 9.03×10^{-3} | 3/5 | 8 | 1744 | AGGAATGTGGC TAACGTGTG | ACAGCTCACCAG AAGTTGC | TD 70-50 | 12 | [29] |
| DYS484 | $(\text{AAT})_{10-16}\text{N}_{12}(\text{AAT})_3(\text{TAT})_3$ | 2.61×10^{-3} | 9.09×10^{-4} – 5.73×10^{-3} | 2/2 | 4 | 1792 | CCTATCATCCGC ATGGACTT | CCTGGTTGACAA AGCCAGAT | TD 60-50 | 21 | [29] |
| DYS485 | $(\text{TTT})_{0-1}(\text{TAA})_{11-21}$ | 4.04×10^{-4} | 1.53×10^{-5} – 2.13×10^{-3} | 0/0 | 0 | 1730 | AAAGCAGACTT CGCCACTACA | AAAAATTAGCTG GGCCTGGT | TD 70-50 | 9 | [29] |
| DYS487 | $(\text{AAT})_{10-16}$ | 1.77×10^{-3} | 4.08×10^{-4} – 4.78×10^{-3} | 1/1 | 2 | 1511 | TGTGGGAGGCC TAAGAAAA | CCTGGGCAACA GAGAAAGAC | TD 70-50 | 1 | [29] |
| DYS488 | $(\text{ATA})_{10-16}$ | 4.40×10^{-4} | 1.60×10^{-5} – 2.32×10^{-3} | 0/0 | 0 | 1576 | GGGGAGGGATA GCATTAGGA | TACCCTGGTCCA CTTCAACC | TD 70-50 | 3 | [29] |
| DYS489 | $(\text{TAA})_{10-15}$ | 4.48×10^{-4} | 1.66×10^{-5} – 2.38×10^{-3} | 0/0 | 0 | 1552 | ACCCAAAGATTT GTCGGCTA | AAAATTAGCCG AGCATGGTG | TD 70-50 | 37 | [29] |
| DYS490 | $(\text{TAA})_{8-16}$ | 3.95×10^{-4} | 1.48×10^{-5} – 2.10×10^{-3} | 0/0 | 0 | 1759 | CCTGGCAGGAA TTATCCAGA | GCAGAGCTTGCA CTGAGCT | TD 70-50 | 10 | [29] |
| DYS491 | $(\text{ATA})_{8-14}$ | 4.09×10^{-4} | 1.45×10^{-5} – 2.17×10^{-3} | 0/0 | 0 | 1706 | GGAATGGGGAG GGATAACAT | GGAGAAAATTC AATGCAGATACC | TD 70-50 | 15 | [29] |
| DYS492 | $(\text{ATA})_{9-15}$ | 3.92×10^{-4} | 1.45×10^{-5} – 2.09×10^{-3} | 0/0 | 0 | 1770 | AGATGAGCCAG GCTTCAGAC | AGTAGGGGTCA GGCACAATG | TD 70-50 | 7 | [29] |
| DYS493 | $(\text{AAC})_{8-11}$ | 3.86×10^{-4} | 1.45×10^{-5} – 2.06×10^{-3} | 0/0 | 0 | 1800 | ACTCCAGTCTGG GTGGACAG | CCCTGGGATTAT AGGCATGA | TD 70-50 | 36 | [29] |
| DYS494 | $(\text{TA})_{4-6}(\text{TAA})_{7-11}$ | 3.89×10^{-4} | 1.43×10^{-5} – 2.07×10^{-3} | 0/0 | 0 | 1783 | TTGCAACACTGT TCATTTGGA | AACAAACCTGC ATGTTCTTCAA | TD 70-50 | 11 | [29] |
| DYS495 | $(\text{AAT})_{12-19}$ | 2.09×10^{-3} | 6.19×10^{-4} – 4.97×10^{-3} | 2/1 | 3 | 1755 | CCCAGCTATTCA GGAGGTTG | GCCAGAAAGTG TGAGTCATCC | TD 70-50 | 10 | [29] |
| DYS497 | $(\text{TAA})_{9-16}$ | 1.49×10^{-3} | 3.46×10^{-4} – 4.05×10^{-3} | 1/1 | 2 | 1786 | AACATGTGCGTT TTCAACCA | GCATGTTGTGCA CATGTAACC | TD 70-50 | 13 | [29] |

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|---------|--|-------------------------|--|------|----|------|----------------------------|----------------------------|-------------|----|------|
| DYS499 | (TTG) ₈ | 3.93 x 10 ⁻⁴ | 1.40 x 10 ⁻⁵ – 2.09x10 ⁻³ | 0/0 | 0 | 1771 | TGGGTCAGAGA AAGGATTGC | GGAGGAAGAGG TTGCAATGA | TD 70-50 | 47 | [29] |
| DYS502 | (AAT) ₄ (TGC) ₁ (CAT) ₆₋₉ | 3.85 x 10 ⁻⁴ | 1.43 x 10 ⁻⁵ – 2.05x10 ⁻³ | 0/0 | 0 | 1792 | CAGCAAGCCAC CATACCATA | TGTGCTTTTGGGA GTTTGGAG | TD 70-50 | 34 | [29] |
| DYS504 | (CCTT) ₁₀₋₂₀ N ₇ (CCCT) ₃ | 3.24 x 10 ⁻³ | 1.26 x 10 ⁻³ – 6.62x10 ⁻³ | 1/4 | 5 | 1746 | TCTACACCACTG TGCCAAGC | GGCAACAGAGC AACCTCT | TD 70-50 | 8 | [29] |
| DYS505 | (TCCT) ₉₋₁₅ | 1.51 x 10 ⁻³ | 3.50 x 10 ⁻⁴ – 4.07x10 ⁻³ | 1/1 | 2 | 1760 | TCTGGCGAAGTA ACCCAAAC | TCGAGTCAGTTC ACCAGAAGG | TD 70-50 | 6 | [29] |
| DYS508 | (TATC) ₈₋₁₅ | 3.03 x 10 ⁻³ | 1.05 x 10 ⁻³ – 6.63x10 ⁻³ | 2/2 | 4 | 1544 | ACAATGGCAAT CCCAAATTC | GAACAAATAAG GTGGGATGGAT | TD 70-50 | 1 | [29] |
| DYS509 | (AAAT) ₇₋₁₁ (AATAA) ₁ (AAAT) ₃ | 1.06 x 10 ⁻³ | 1.55 x 10 ⁻⁴ – 3.53x10 ⁻³ | 0/1 | 1 | 1590 | AACATGGTGAA TCCCTGTCTCT | TGTCCCCAGGGC TTTTTAAT | TD 70-50 | 37 | [29] |
| DYS510 | (GATA) ₃ N ₁₂ (GATA) ₉₋₁₅ N ₁₃ (GGAT) ₄ N ₉ (GATA) ₃ | 5.99 x 10 ⁻³ | 3.09 x 10 ⁻³ – 1.03x10 ⁻² | 4/6 | 10 | 1779 | TTTTTCCTCCCTT ACCACAGA | TCTGGAGAAGA CAGAACTGTCA | TD 70-50 | 34 | [29] |
| DYS511 | (GATA) ₉₋₁₄ | 1.52 x 10 ⁻³ | 3.51 x 10 ⁻⁴ – 4.11x10 ⁻³ | 1/1 | 2 | 1760 | GATAGGATGGG GTGGATGTG | TGTGAATCCCC TTCTACATCTC | TD 70-50 | 13 | [29] |
| DYS512 | (AGAT) ₇₋₁₃ | 3.96 x 10 ⁻⁴ | 1.44 x 10 ⁻⁵ – 2.11x10 ⁻³ | 0/0 | 0 | 1738 | CACGCCAGCTA ATTTTTGT | GGGAGGAATAA AGGAAGGTTG | TD 70-50 | 26 | [29] |
| DYS513 | (TCTA) ₄ (TCCA) ₁ (TATC) ₃ (CGTA) ₁ (TCTA) ₉₋₁₅ | 6.09 x 10 ⁻³ | 3.14 x 10 ⁻³ – 1.05x10 ⁻² | 6/4 | 10 | 1751 | ATTGATCCATCC GTCTGTCC | GTTGGATGAAG GGAGAGCAG | TD 70-50 | 32 | [29] |
| DYS516 | (TTCT) ₄ N ₃₀ (TTCT) ₉₋₁₈ | 6.66 x 10 ⁻³ | 3.55 x 10 ⁻³ – 1.12x10 ⁻² | 7/4 | 11 | 1753 | TTTCCAATGACC AAGACGTG | CGAACCTGCAA ATTGTTTCC | TD 60-50 | 22 | [29] |
| DYS517 | (AAAG) ₁₀₋₁₈ N ₈ (AAAG) ₃ | 3.21 x 10 ⁻³ | 1.25 x 10 ⁻³ – 6.62x10 ⁻³ | 3/2 | 5 | 1766 | TAATCGTCCCAT TTTGAGCA | TGCAATCCAAA CTCAGAAA | TD 60-50 | 22 | [29] |
| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₄₋₂₂ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₁₋₁₉ N ₂₇ (AAGG) ₄ | 1.84 x 10 ⁻² | 1.25 x 10 ⁻² – 2.60x10 ⁻² | 8/20 | 28 | 1556 | GGCAACACAAG TGAAACTGC | TCAGCTCTTACC ATGGGTGAT | TD 70-50 | 54 | [29] |
| DYS520 | (GATA) ₁₀₋₁₃ (CATA) ₁₀₋₁₁ | 2.66 x 10 ⁻³ | 9.22 x 10 ⁻⁴ – 5.80x10 ⁻³ | 2/2 | 4 | 1760 | AACAGCCTGCC AACATAGT | ACCATCATGCC TGCAATA | TD 70-50 | 24 | [29] |
| DYS521 | (CTTT) ₅ (TCTT) ₃ (TTTT) ₁ (CTTT) ₅ T (CTTT) ₄₋₁₄ | 9.54 x 10 ⁻⁴ | 1.37 x 10 ⁻⁴ – 3.18x10 ⁻³ | 0/1 | 1 | 1751 | GCCACGACACCT GTTCAGTA | GCTGGGAGTGA GACCTGTA | TD 70-50 | 24 | [29] |
| DYS522 | (ATAG) ₈₋₁₅ | 1.04 x 10 ⁻³ | 1.53 x 10 ⁻⁴ – 3.44x10 ⁻³ | 1/0 | 1 | 1620 | CCTTTGAAATCA TTCATAATGC | TCATAAACAGA GGGTTCTGG | TD 70-50 | 4 | [29] |
| DYS525 | (AGAT) ₈₋₁₃ | 9.78 x 10 ⁻⁴ | 1.42 x 10 ⁻⁴ – 3.26x10 ⁻³ | 0/1 | 1 | 1712 | ATTCACACCATT GCACTCCA | CCATCTGTTTAT CTTCCCATCA | TD 70-50 | 2 | [29] |
| DYS526a | (CCTT) ₁₀₋₁₇ | 2.72 x 10 ⁻³ | 9.52 x 10 ⁻⁴ – 5.97x10 ⁻³ | 2/2 | 4 | 1716 | TCTGGTGAAGT ATCCAAACC | GGGTTACTTCGC CAGAAGGT | TD 65-50 | 51 | [29] |
| DYS526b | (CCCT) ₃ N ₂₀ (CTTT) ₁₁₋₁₇ (CCTT) ₆₋₁₀ N ₁₃ (CCTT) ₁₀₋₁₇ | 1.25 x 10 ⁻² | 7.88 x 10 ⁻³ – 1.87x10 ⁻² | 9/11 | 20 | 1651 | TCTGGTGAAGT ATCCAAACC | GGGTTACTTCGC CAGAAGGT | TD 65-50 | 51 | [29] |
| DYS530 | (AAAC) ₈₋₁₁ | 3.94 x 10 ⁻⁴ | 1.45 x 10 ⁻⁵ – 2.10x10 ⁻³ | 0/0 | 0 | 1760 | CAGGGTCAAAA TACCCCTCC | CTGCGGGACAAT GAAACAC | TD 70-50 | 15 | [29] |
| DYS531 | (AAAT) ₉₋₁₃ | 1.00 x 10 ⁻³ | 1.45 x 10 ⁻⁴ – 3.50x10 ⁻³ | 0/1 | 1 | 1682 | GACCCACTGGC ATTCAAATC | TGCTCCCTTCT TTGTAGACG | TD 70-50 | 3 | [29] |
| DYS532 | (TCCC) ₃ N ₅ (TTCC) ₅ N ₉ (TTCT) ₃ (TTCC) ₁ (TTCT) ₆₋₁₇ N ₁₇ (TTCT) ₃ N ₁₃ (TTCC) ₄ N ₇₀ (TTCT) ₃ N ₆ (TTCT) ₃ | 3.24 x 10 ⁻³ | 1.13 x 10 ⁻³ – 7.10x10 ⁻³ | 3/1 | 4 | 1441 | TTGGTTTTATGC CTTTCCT | TAGGTGACAGA GCAGGATTC | TD 70-50 | 39 | [29] |

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|--------|--|------------------------|---|-------|----|------|--------------------------------|-----------------------------------|-------------|----|------|
| DYS533 | (TATC) _{9,14} | 5.01 x10 ⁻³ | 2.39 x10 ⁻³ – 9.11x10 ⁻³ | 4/4 | 8 | 1730 | CATCTAACATCT TTGTCATCTACC | TGATCAGTTCTT AACTCAACCA | TD 70-50 | 5 | [29] |
| DYS534 | (CTTT) ₃ N ₈ (CTTT) ₉₋₂₀ N ₉ (CTTT) ₃ | 6.51 x10 ⁻³ | 3.44 x10 ⁻³ – 1.10x10 ⁻² | 9/2 | 11 | 1794 | CATCTACCCAAC ATCCATCTA | GACAAAGATGTT AGATGAATAGA CA | TD 60-50 | 18 | [29] |
| DYS536 | (TCCT) ₇₋₁₉ N ₈ (TTCT) ₄ | 1.15 x10 ⁻³ | 1.66 x10 ⁻⁴ – 3.83x10 ⁻³ | 1/0 | 1 | 1453 | TTGCTTTTCTGC TTCCCTTC | ATCGCATTCCCC TCTCCTAC | 52 | 53 | [29] |
| DYS537 | (TCTA) ₈₋₁₃ | 2.38 x10 ⁻³ | 7.12 x10 ⁻⁴ – 5.70x10 ⁻³ | 2/1 | 3 | 1539 | GGTCTCCAATTC CATCCAGA | TGGAACATGCC ATTAATCA | TD 70-50 | 3 | [29] |
| DYS538 | (GATA) ₉₋₁₃ | 3.94 x10 ⁻⁴ | 1.47 x10 ⁻⁵ – 2.10x10 ⁻³ | 0/0 | 0 | 1765 | CCCCTGAATCAC CAGAGTTC | AACCAGCCCAA ATACCCATC | TD 70-50 | 31 | [29] |
| DYS539 | (TAGA) ₈₋₁₄ | 1.00 x10 ⁻³ | 1.46 x10 ⁻⁴ – 3.32x10 ⁻³ | 0/1 | 1 | 1676 | GTTGAAGCCCTC AATCTGGT | GGTGCAGATCTC CCAAATTC | TD 60-50 | 19 | [29] |
| DYS540 | (TTAT) _{9,14} | 3.30 x10 ⁻³ | 1.28 x10 ⁻³ – 6.79x10 ⁻³ | 2/3 | 5 | 1718 | GACCGTGACTC TGGCCAAT | CAGGAGGCTAG CTCAGGAGA | TD 70-50 | 7 | [29] |
| DYS541 | (TATC) ₁₀₋₁₅ (TTC) ₁ (TATC) ₃ | 3.92 x10 ⁻³ | 1.65 x10 ⁻³ – 7.68x10 ⁻³ | 2/4 | 6 | 1700 | TTCTATCTGTTC ATCCATCTAGG | ACCTTTAAGAAG CCTTCACC | TD 60-50 | 20 | [29] |
| DYS543 | (AGAT) ₃ (GATA) _{7,2-16} N ₄₂ (ATGT) ₃₋₄ (ATGG) _{2,3} N ₃₅ (GAAA) ₃ | 7.10 x10 ⁻³ | 3.77 x10 ⁻³ – 3.53x10 ⁻³ | 4/7 | 11 | 1645 | CAAGGGCCAAT TATGTATGT | TGATCTTCCTGG TCACCTTT | TD 60-50 | 23 | [29] |
| DYS544 | (TAGA) ₃ N ₁₅ (TAGA) ₃ (TGGA) ₁ (TAGA) ₆₋₁₂ | 3.96 x10 ⁻⁴ | 1.44 x10 ⁻⁵ – 1.20x10 ⁻² | 0/0 | 0 | 1748 | CTGGGCAACAG AGCAAGATT | AATGCTGGCCAA AACAAAGT | TD 70-50 | 32 | [29] |
| DYS545 | (TGTT) ₈₋₁₁ | 3.90 x10 ⁻⁴ | 1.39 x10 ⁻⁵ – 2.09x10 ⁻³ | 0/0 | 0 | 1779 | GAGGGGAGTGT AGAAAAGATGC | GATCCAAGATG GTGCCATTG | TD 70-50 | 30 | [29] |
| DYS546 | (TTCC) ₃ N ₂₃ (TTCT) ₃ N ₃₃ (TTCC) ₃ N ₁₆ (TTCT) _{9,19} | 4.35 x10 ⁻³ | 1.85 x10 ⁻³ – 8.56x10 ⁻³ | 2/4 | 6 | 1531 | CCTGAGCTATTT TCCCTTTGC | TGCAGTACATCC TGGGGAAT | TD 70-50 | 35 | [29] |
| DYS547 | (CCTT) ₉₋₁₃ T(CTTC) ₄₋₅ N ₅₆ (TTTC) ₁₀₋₂₂ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₉₋₁₆ N ₁₄ (TTTC) ₃ | 2.36 x10 ⁻² | 1.70 x10 ⁻² – 3.18x10 ⁻² | 22/17 | 39 | 1679 | TCCATGTTACTG CAAAATACAC | TGACAGAGCAT AAACGTGTC | TD 60-50 | 17 | [29] |
| DYS549 | (GATA) ₉₋₁₅ | 4.55 x10 ⁻³ | 2.05 x10 ⁻³ – 8.58x10 ⁻³ | 1/6 | 7 | 1684 | AACCAAATTC GGGATGTACTG A | GTCCCCTTTTCC ATTTGTGA | TD 70-50 | 15 | [29] |
| DYS550 | (AAGG) ₄ N ₁₆ (AAGG) ₄ (AAAG) ₁ (AAGG) ₆₋₁₁ | 3.87 x10 ⁻⁴ | 1.41 x10 ⁻⁵ – 2.06x10 ⁻³ | 0/0 | 0 | 1794 | GCCTGGGTAAC AGGAGTGAA | AGCTGAAAAC GTGCTGCTG | TD 70-50 | 34 | [29] |
| DYS551 | (AGAT) ₁₀₋₁₆ N ₈ (AGAC) ₃ (AGGT) ₁ (AGAT) ₄ | 3.26 x10 ⁻³ | 1.26 x10 ⁻³ – 6.72x10 ⁻³ | 1/4 | 5 | 1737 | CCAGCCTGGGTG ACAAAGTA | AAAGTTCCTCCC AGTTGCAC | TD 70-50 | 38 | [29] |
| DYS552 | (TCTA) ₃ (TCTG) ₁ (TCTA) ₇₋₁₂ N ₄₀ (TCTA) ₁₁₋₁₆ | 2.69 x10 ⁻³ | 9.21 x10 ⁻⁴ – 5.87x10 ⁻³ | 3/1 | 4 | 1742 | CCATAGTGCCGA GGTCAAGT | AACACCTGATGC CTGGTTG | TD 70-50 | 38 | [29] |
| DYS554 | (TAAA) ₈₋₁₁ | 9.41 x10 ⁻⁴ | 1.36 x10 ⁻⁴ – 3.15x10 ⁻³ | 1/0 | 1 | 1777 | CTGGGCCACAG AGTGAGAC | GGGCCAGTCTTT GCAATATC | TD 70-50 | 13 | [29] |
| DYS556 | (AAAT) ₈₋₁₂ | 1.59 x10 ⁻³ | 3.70 x10 ⁻⁴ – 4.30x10 ⁻³ | 1/1 | 2 | 1683 | TGCTGTACATC ACCAATGA | TTTGGTTGCTGA AGCATTGA | TD 70-50 | 14 | [29] |
| DYS557 | (TTTC) ₄ (TTCTC) ₁ (TTTC) ₄ (TTC) ₁ (TTTC) ₁₂₋₂₁ | 3.80 x10 ⁻³ | 1.60 x10 ⁻³ – 7.45x10 ⁻³ | 3/3 | 6 | 1758 | TTTTCTGTGCCA AGCCTACA | TCTAATGCACCT TGAGGGATG | TD 60-50 | 21 | [29] |
| DYS558 | (TTTG) ₃ (TTTA) ₅₋₁₀ | 3.98 x10 ⁻⁴ | 1.42 x10 ⁻⁵ – 2.13x10 ⁻³ | 0/0 | 0 | 1741 | GGTGGTCAGAA AATCCCTCA | GCAGGCCAATAT TCACCATT | TD 70-50 | 26 | [29] |
| DYS559 | (TAAA) _{7,9} | 9.63 x10 ⁻⁴ | 1.40 x10 ⁻⁴ – 3.19x10 ⁻³ | 1/0 | 1 | 1750 | AGCCAAGGTCA TACCACTGC | TCGGTGAAGGC ACCAATAAT | TD 70-50 | 25 | [29] |

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| DYS561 | (GATA) ₉₋₁₃ (GACA) ₄ | 9.41 x10 ⁻⁴ | 1.36 x10 ⁻⁴ – 3.11x10 ⁻³ | 0/1 | 1 | 1783 | GCCTGATGCCAT CTGAAAAT | TGATCCCAACAA CTGCACTC | TD 60-50 | 17 | [29] |
| DYS565 | (ATAA) ₉₋₁₄ | 2.09 x10 ⁻³ | 6.20 x10 ⁻⁴ – 4.95x10 ⁻³ | 1/2 | 3 | 1757 | AAACCCAGGAA GCAGTGTG | CCTGGCTCAGCA CATGAATA | TD 70-50 | 11 | [29] |
| DYS567 | (ATAA) ₇₋₁₃ | 4.08 x10 ⁻⁴ | 1.48 x10 ⁻⁵ – 2.14x10 ⁻³ | 0/0 | 0 | 1713 | GGAAAGCTGAGG AAGGAGGAG | TTATGACCGGGA TCAAGTGC | TD 70-50 | 10 | [29] |
| DYS568 | (AAAT) ₉₋₁₃ | 1.08 x10 ⁻³ | 1.56 x10 ⁻⁴ – 3.60x10 ⁻³ | 0/1 | 1 | 1547 | GTGGCAGACAA AACCCAGTT | TTGAAAAGGGA TGGGACTCA | TD 70-50 | 3 | [29] |
| DYS569 | (ATTT) _{8,2-13} | 1.58 x10 ⁻³ | 3.66 x10 ⁻⁴ – 4.24x10 ⁻³ | 0/2 | 2 | 1696 | TCCATGGGATAT GATGAGCA | GGCAGCCTGTAG GACAGAGA | TD 70-50 | 12 | [29] |
| DYS570 | (TTTC) ₁₄₋₂₄ | 1.24 x10 ⁻² | 7.52 x10 ⁻³ – 1.91x10 ⁻² | 8/9 | 17 | 1426 | GAACTGTCTACA ATGGCTCACG | TCAGCATAGTCA AGAAACCAGAC A | TD 70-50 | 1 | [29] |
| DYS571 | (TTTT) ₄ N ₇ (TTTA) ₉₋₁₂ | 4.13 x10 ⁻⁴ | 1.51 x10 ⁻⁵ – 2.20x10 ⁻³ | 0/0 | 0 | 1682 | AGCCTTCAGCGA CTGCTTTA | AGCTGAGATCAT CCCATTGC | TD 70-50 | 47 | [29] |
| DYS572 | (AAAT) ₈₋₁₃ | 2.07 x10 ⁻³ | 6.17 x10 ⁻⁴ – 4.96x10 ⁻³ | 0/3 | 3 | 1770 | CTAAGGACGCCT CCCATAACA | CTCATTCCCTAT GGTTTGCAC | TD 70-50 | 9 | [29] |
| DYS573 | (TTTA) ₈₋₁₃ | 4.10 x10 ⁻⁴ | 1.51 x10 ⁻⁵ – 2.17x10 ⁻³ | 0/0 | 0 | 1698 | GGGGGAGAAAA AGTTTGGTG | AAAAATGGGGA GGTGGAAT | TD 70-50 | 14 | [29] |
| DYS574 | (TTAT) ₈₋₁₂ | 9.77 x10 ⁻⁴ | 1.43 x10 ⁻⁴ – 3.25x10 ⁻³ | 0/1 | 1 | 1721 | GGTGGGGCTTCC ATATTTTT | AATGTAGACGA CGGGTTGATG | TD 60-50 | 43 | [29] |
| DYS575 | (AAAT) ₈₋₁₁ | 3.91 x10 ⁻⁴ | 1.47 x10 ⁻⁵ – 2.09x10 ⁻³ | 0/0 | 0 | 1764 | GGTGGTGGACA TCCGTAATC | AGTAATGGGAT GCTGGGTCA | TD 70-50 | 11 | [29] |
| DYS576 | (AAAG) ₁₃₋₂₂ | 1.43 x10 ⁻² | 9.41 x10 ⁻³ – 2.07x10 ⁻² | 12/12 | 24 | 1727 | TTGGGCTGAGG AGTTCAATC | GGCAGTCTCATT TCCTGGAG | TD 70-50 | 12 | [29] |
| DYS577 | (ATTC) ₆₋₁₀ | 4.11 x10 ⁻⁴ | 1.51 x10 ⁻⁵ – 2.19x10 ⁻³ | 0/0 | 0 | 1691 | TCAATGCATGTT TTTCTACGTG | GGAGGATGGTTT GAACCTGA | TD 60-50 | 40 | [29] |
| DYS578 | (AAAT) ₇₋₁₀ | 9.95 x10 ⁻⁴ | 1.43 x10 ⁻⁴ – 3.30x10 ⁻³ | 1/0 | 1 | 1686 | GAGGCGGAACT TTCAGTGAG | GCTTCAACAACC CTGGACAT | TD 70-50 | 4 | [29] |
| DYS579 | (TATT) ₇₋₁₀ | 3.94 x10 ⁻⁴ | 1.40 x10 ⁻⁵ – 2.10x10 ⁻³ | 0/0 | 0 | 1755 | GCCAGCAGTAG ACCCAGACT | AGGCAGAGGTT GCAGTGAGT | TD 70-50 | 2 | [29] |
| DYS580 | (AATA) ₈₋₁₀ | 4.05 x10 ⁻⁴ | 1.47 x10 ⁻⁵ – 2.13x10 ⁻³ | 0/0 | 0 | 1725 | GCAGTGAGCCG AGATCAGG | GGAGCAAACAC TGCAATTTCC | TD 70-50 | 4 | [29] |
| DYS581 | (TAGG) _{7,9} | 3.84 x10 ⁻⁴ | 1.43 x10 ⁻⁵ – 2.04x10 ⁻³ | 0/0 | 0 | 1807 | GTAGGGTCTTGA ACAGCATACG | CGAGCCAAGCT GCTGTTAT | TD 70-50 | 36 | [29] |
| DYS583 | (AAAC) _{7,9} | 3.99 x10 ⁻⁴ | 1.44 x10 ⁻⁵ – 2.12x10 ⁻³ | 0/0 | 0 | 1730 | GCAGGAAAATT GCTTGAACC | CCTCATCCAATA GCTCTTCCT | TD 70-50 | 2 | [29] |
| DYS584 | (CAAT) ₇₋₈ | 3.90 x10 ⁻⁴ | 1.43 x10 ⁻⁵ – 2.10x10 ⁻³ | 0/0 | 0 | 1777 | TGCAGAATGTAT GGTCTTTTTGA | CTGCCAGTCTAT TGCCCTTC | TD 60-50 | 45 | [29] |
| DYS585 | (TTATG) ₈₋₁₂ | 2.12 x10 ⁻³ | 6.33 x10 ⁻⁴ – 5.06x10 ⁻³ | 2/1 | 3 | 1734 | TGGAAGTATTCC ACTCACTTGCT | CTCAAGTGGGG AAGTCAAGG | TD 60-50 | 42 | [29] |
| DYS587 | (CAATA) ₈₋₁₆ [(CAGTA) ₁ (CAATA) ₁] ₃ | 2.62 x10 ⁻³ | 9.16 x10 ⁻⁴ – 5.75x10 ⁻³ | 2/2 | 4 | 1782 | CCTAAAGCGAA GAGACCATGA | TGAAGGCCAAA GAGTGAAAGA | TD 60-50 | 18 | [29] |
| DYS588 | (GCATT) ₉₋₁₆ | 3.92 x10 ⁻⁴ | 1.47 x10 ⁻⁵ – 2.10x10 ⁻³ | 0/0 | 0 | 1747 | GAATGCAGAAC CCTCAAGGA | AGCCTGGGTGAC AGAAACAC | TD 60-50 | 42 | [29] |
| DYS590 | (TTTTG) _{5,9} | 3.91 x10 ⁻⁴ | 1.45 x10 ⁻⁵ – 2.06x10 ⁻³ | 0/0 | 0 | 1780 | GGGAACATAGT CGGGCTGTA | GGGTGACAGAG CAAGAATCC | TD 70-50 | 5 | [29] |

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| DYS593 | (AAAAC) ₄ (AAAAT) ₇₋₁₀ | 1.51 x 10 ⁻³ | 3.47 x 10 ⁻⁴ – 4.06x10 ⁻³ | 1/1 | 2 | 1775 | CTTGAACCCAGG AAGCAGAC | TTATGCCCAAGT GACTGTGC | TD 70-50 | 29 | [29] |
| DYS594 | (AAATA) ₈₋₁₃ | 1.03 x 10 ⁻³ | 1.46 x 10 ⁻⁴ – 3.41x10 ⁻³ | 1/0 | 1 | 1635 | GATGTGCCTAAT GCCACAGA | CCCTGGTGTTAA TCGTGTCC | TD 70-50 | 5 | [29] |
| DYS595 | (ATTTA) ₆₋₉ | 3.96 x 10 ⁻⁴ | 1.46 x 10 ⁻⁵ – 2.10x10 ⁻³ | 0/0 | 0 | 1750 | TGTTTTCGGTTC CTCTGTCC | AGGGGAACAAC ACACACTGG | TD 70-50 | 28 | [29] |
| DYS596 | (GGA) ₅ (GTA) ₁ (GGA) ₃ (GAA) ₃ [(GGA) ₁ (GAA) ₁] ₈₋₁₀ | 4.24 x 10 ⁻⁴ | 1.52 x 10 ⁻⁵ – 2.28x10 ⁻³ | 0/0 | 0 | 1630 | ATAACCGTGCCC TTTACTGC | TTTTGACAAGCC CAAAGTTCT | TD 60-50 | 44 | [29] |
| DYS611 | (TTC) ₅ N ₉ (TTC) ₄ (CTC) ₁ (TTC) ₃ N ₉ (TTC) ₅ (CTC) ₁ (TTC) ₃ N ₁₅ (TTC) ₄ (CT) ₁ (TTC) ₃ (CTC) ₁ (TTC) ₃ N ₂₀ (TTC) ₃ T(TTC) ₄ N ₇ (TTC) ₃ N ₉ (TTC) ₄ (TCC) ₁ (TTC) ₇₋₂₁ N ₂₃ (TTC) ₄ N ₄ [(TTC) ₁ (CTC) ₁] ₂ [(CTC) ₁ (TTC) ₁] ₃ | 8.89 x 10 ⁻³ | 4.86 x 10 ⁻³ – 1.47x10 ⁻² | 3/9 | 12 | 1426 | TACAGGTGTGCA CCATGAGG | CTTGGCAACATA GCAGATCC | TD 70-50 | 35 | [29] |
| DYS612 | (CCT) ₅ (CTT) ₁ (TCT) ₃ (CCT) ₁ (TCT) ₁₉₋₃₁ | 1.45 x 10 ⁻² | 9.61 x 10 ⁻³ – 2.09x10 ⁻² | 11/14 | 25 | 1767 | CCCCATGCCAG TAAGAATA | TGAGGGAAGGC AAAAGAAAA | TD 60-50 | 42 | [29] |
| DYS613 | (ATG) ₈ (ATA) ₁ (ATG) ₈₋₉ | 4.35 x 10 ⁻⁴ | 1.60 x 10 ⁻⁵ – 2.32x10 ⁻³ | 0/0 | 0 | 1588 | ATAGAAGGCAA ATTCCTTATCAA | AAAGTTAATGAC GCCTTGTC | TD 60-50 | 23 | [29] |
| DYS614 | (CTT) ₄ (CCT) ₁ (CTT) ₃ N ₁₅ (CCT) ₄ (CTT) ₄ (CCT) ₁ (CTT) ₃ N ₁₈ (CCT) ₃ (CTT) ₅ N ₂₀ [(CTT) ₁ (CTG) ₁] ₃ (CT) ₁ (CTT) ₁₂₋₂₂ N ₈ (CTT) ₄ [(CTC) ₁ (CTT) ₁] ₃ [(CTC) ₁ (TTT) ₁] ₁ (CTT) ₅ | 4.32 x 10 ⁻³ | 1.94 x 10 ⁻³ – 8.14x10 ⁻³ | 2/5 | 7 | 1776 | GTGGCGATGTTG TGAGTGTT | GCCACAAAAG GTTTTCAGA | TD 70-50 | 33 | [29] |
| DYS615 | (TTG) ₇₋₈ | 3.91 x 10 ⁻⁴ | 1.43 x 10 ⁻⁵ – 2.09x10 ⁻³ | 0/0 | 0 | 1766 | GGTCGAAGAAG GTGTCACAGA | TGATTCTGCTAA TTCCCATGC | TD 70-50 | 25 | [29] |
| DYS616 | (TAT) ₈₋₁₆ (CAT) ₁ (TAT) ₃ | 1.72 x 10 ⁻³ | 4.03 x 10 ⁻⁴ – 4.64x10 ⁻³ | 1/1 | 2 | 1564 | GGCAAACAGAT AGCAATTTACA | TTGTTCTGCCCA GCAGTAT | TD 70-50 | 35 | [29] |
| DYS617 | (TTA) ₁₁₋₁₅ | 4.13 x 10 ⁻⁴ | 1.53 x 10 ⁻⁵ – 2.21x10 ⁻³ | 0/0 | 0 | 1684 | AGCATGTGCCCT TCAGCTTT | GGATTGGGGAG TGATAGCAT | TD 70-50 | 5 | [29] |
| DYS618 | (TAT) ₈₋₁₄ | 3.95 x 10 ⁻⁴ | 1.46 x 10 ⁻⁵ – 2.09x10 ⁻³ | 0/0 | 0 | 1766 | CCCATACCCTTG GTGTTGTC | GAGGGCTATGG GAGGGATAG | TD 70-50 | 13 | [29] |
| DYS619 | (AAT) ₆₋₁₀ | 4.69 x 10 ⁻⁴ | 1.70 x 10 ⁻⁵ – 2.50x10 ⁻³ | 0/0 | 0 | 1479 | GGCGACAGAGC GAGACTCTA | GGCATGTGAGTT GAGGAACA | TD 70-50 | 16 | [29] |
| DYS620 | (ATA) ₈₋₉ | 4.11 x 10 ⁻⁴ | 1.47 x 10 ⁻⁵ – 2.22x10 ⁻³ | 0/0 | 0 | 1678 | TGACGAGTTAAT GGGTGCAG | TGAGTTTGCTCC TCTAGCTTTC | TD 60-50 | 40 | [29] |
| DYS621 | (TTA) ₇₋₉ | 4.44 x 10 ⁻⁴ | 1.70 x 10 ⁻⁵ – 2.38x10 ⁻³ | 0/0 | 0 | 1543 | GCCCAAATTA AAGGCACAA | TGACGAGTTAGT GGGTGCAG | TD 70-50 | 35 | [29] |
| DYS622 | (GAAA) ₆ (AGAAG) ₁ (GAAA) ₈₋₁₆ | 3.40 x 10 ⁻³ | 1.32 x 10 ⁻³ – 7.01x10 ⁻³ | 2/3 | 5 | 1663 | TCCAGCCTCGGT GATAAGAG | GGCTGAAGTGG GTTGTGTTA | TD 60-50 | 44 | [29] |
| DYS624 | (GGAT) ₈₋₁₀ (G/AGAT) ₁ N ₃₅ (GGAT) ₃ | 4.06 x 10 ⁻⁴ | 1.55 x 10 ⁻⁵ – 2.18x10 ⁻³ | 0/0 | 0 | 1699 | GCATCTCAAATC CTTTGTGGA | TCCACCTGCTTT TCTCTCA | TD 60-50 | 43 | [29] |
| DYS625 | (CTTT) ₄ (TTCT) ₁ (CTTT) ₃ (TTT) ₁ (CTTT) ₄ (TT) ₁ (CTTT) ₃ N ₄₇ (CTTT) ₃₋₅ (CT) ₁ (CTTT) ₄ (CCTT) ₁ (CTTT) ₃ N ₁₀ (CTTT) ₃ | 9.58 x 10 ⁻⁴ | 1.40 x 10 ⁻⁴ – 3.18x10 ⁻³ | 0/1 | 1 | 1746 | TCATCACACATG GCCTAATTG | GGCAAGTCACAT GCATTACAA | TD 60-50 | 22 | [29] |
| DYS626 | (GAAA) ₁₄₋₂₃ N ₂₄ (GAAA) ₃ N ₆ | 1.22 x 10 ⁻² | 7.70 x 10 ⁻³ – | 13/7 | 20 | 1689 | GCAAGACCCCA | AAGAAGAATTTT | TD | 23 | [29] |

| | | | | | | | | | | | |
|------------|--|------------------------------|--|-------------|-----------|-------------|---|---|---------------------|-----------|-------------|
| | (GAAA)₅(AAA)₁(GAAA)₂₋₃ (GAAG)₁(GAAA)₃ | | 1.82x10⁻² | | | | TAGCAAAAG | GGGACATGTTT | 60-50 | | |
| DYS627 | (AGAA)₃N₁₆(AGAG)₃ (AAAG)₁₂₋₂₄N₈₁(AAGG)₃ | 1.23 x10⁻² | 7.80 x10⁻³– 1.81x10⁻² | 12/9 | 21 | 1766 | CTAGGTGACAG CGCAGGATT | GGATAATGAGC AAATGGCAAG | TD 60-50 | 21 | [29] |
| DYS629 | (TATC)₅₋₁₂ | 9.91 x10⁻⁴ | 1.41 x10⁻⁴– 3.31x10⁻³ | 1/0 | 1 | 1689 | GGGATTATTACA ATTC AAGGTC | TATGGGTA AATG GCAAAAGT | TD 60-50 | 19 | [29] |
| DYS630 | (AAAG)₄(AGAG)₃N₁₈ (AAAG)₁₂₋₂₁ | 4.86 x10⁻³ | 2.31 x10⁻³– 8.85x10⁻³ | 5/3 | 8 | 1784 | GCCTTTGGACAG AGCAAGAC | AGCCATGGAAA GCTGTGAGT | TD 70-50 | 25 | [29] |
| DYS631 | (AATA)₄(CATA)₁(AATA)₇₋₁₁ | 9.77 x10⁻⁴ | 1.40 x10⁻⁴– 3.25x10⁻³ | 0/1 | 1 | 1721 | CACTCCAGCCTC GGAGATAG | GCGCTCTGTGGA CATTATCA | TD 60-50 | 43 | [29] |
| DYS632 | (CATT)₈₋₁₀ | 3.97 x10⁻⁴ | 1.47 x10⁻⁵– 2.13x10⁻³ | 0/0 | 0 | 1745 | GGCCGTTGCAA AATAAACTG | TCTGGGCAACAG AAGGAGAC | TD 70-50 | 24 | [29] |
| DYS633 | (AAAT)₅N₁₆(AAAT)₇₋₉ | 3.81 x10⁻⁴ | 1.38 x10⁻⁵– 2.02x10⁻³ | 0/0 | 0 | 1814 | GGCAACAAGAG CAAAACTCC | CCACCAGGGAA GTGTCTTTC | TD 60-50 | 46 | [29] |
| DYS634 | (GGAA)₆N₁₀(AAGG)₇₋₁₀ N₁₂(AGGG)₃ | 4.20 x10⁻⁴ | 1.51 x10⁻⁵– 2.25x10⁻³ | 0/0 | 0 | 1646 | TCAGAAGCATG CTAGAACCCTA | TTGCTCCTTACA GAAGAGGTGA | TD 60-50 | 19 | [29] |
| DYS635 | (TCTA)₄(TGTA)₂(TCTA)₂ (TGTA)₂(TCTA)₂(TATG)₀₋₂ (TCTA)₄₋₁₇ | 3.85 x10⁻³ | 1.63 x10⁻³– 7.55x10⁻³ | 1/5 | 6 | 1732 | Yfiler | Yfiler | Yfiler | Yfiler | [29] |
| DYS637 | (AAAT)₄(ACAT)₈₋₁₄ | 1.04 x10⁻³ | 1.53 x10⁻⁴– 3.42x10⁻³ | 0/1 | 1 | 1623 | AAGCCAGTCAA CCAAACACA | TGCTGGGGTTGA AGGTAAAA | TD 60-50 | 41 | [29] |
| DYS638 | (TTTA)₈₋₁₃ | 1.04 x10⁻³ | 1.47 x10⁻⁴– 3.45x10⁻³ | 1/0 | 1 | 1617 | ACAATTTCCCTT GGGGCTAC | CATGGTGGTAGG CACCTGTA | TD 70-50 | 6 | [29] |
| DYS640 | (AAAT)₉₋₁₃ | 3.98 x10⁻⁴ | 1.41 x10⁻⁵– 2.16x10⁻³ | 0/0 | 0 | 1716 | TGGGAAAAACC ATGAGATCC | TAGGGTCAAGCC CGTTCATA | TD 70-50 | 15 | [29] |
| DYS641 | (TAAA)₈₋₁₂ | 3.90 x10⁻⁴ | 1.41 x10⁻⁵– 2.09x10⁻³ | 0/0 | 0 | 1768 | CTTGAGCCCAGG AAGCATAG | CCACACGATGCA ATTTTGTC | TD 70-50 | 6 | [29] |
| DYS642 | (TAAA)₆₋₁₀ | 3.91 x10⁻⁴ | 1.45 x10⁻⁵– 2.07x10⁻³ | 0/0 | 0 | 1785 | CATTGTGCACGT GTACCCTAA | AAAGGTTGTGC TGCATGAT | TD 70-50 | 33 | [29] |
| DYS643 | (CTTTT)₆₋₁₅ | 1.50 x10⁻³ | 3.49 x10⁻⁴– 4.05x10⁻³ | 2/0 | 2 | 1773 | AAGCCATGCCTG GTTAAACT | TGTAACCAAAACA CCACCCATT | TD 70-50 | 14 | [29] |
| DYS644 | (TTTTA)₁₀₋₁₁(TTTA)₀₋₁ (TTTTA)₀₋₁₃ | 3.22 x10⁻³ | 1.25 x10⁻³– 6.62x10⁻³ | 3/2 | 5 | 1761 | TGACTTCGGGGT AGTTCCAG | CCTGGGCAAAA GAGTGAGAC | TD 70-50 | 8 | [29] |
| DYS645 | (TGTTT)₇₋₉ | 4.07 x10⁻⁴ | 1.49 x10⁻⁵– 2.14x10⁻³ | 0/0 | 0 | 1698 | GGTTACGGGTG GCAATCATA | ACTGCCAGACTC ACACATGG | TD 60-50 | 40 | [29] |
| Y-GATA-A10 | (ATCT)₁₁₋₁₆ | 3.32 x10⁻³ | 1.25 x10⁻³– 6.80x10⁻³ | 3/2 | 5 | 1713 | CCTGCCATCTCT ATTTATCTTGCA TATA | ATAAATGGAGA TAGTGGGTGGAT T | TD 60-50 | 41 | [29] |
| Y-GATA-H4 | (TAGA)₃N₁₂(TAGG)₃(TAGA)₈₋₁₅ N₂₂(TAGA)₄ | 3.22 x10⁻³ | 1.28 x10⁻³– 6.62x10⁻³ | 1/4 | 5 | 1755 | Yfiler | Yfiler | Yfiler | Yfiler | [29] |

Markers with median mutation rates above 10⁻² (the RM Y-STR set) are highlighted in red. Additionally included are PCR primers, PCR annealing temperature and locus assignment to the 54 multiplexes used for genotyping.

Notes: The repeat nomenclature used is in accordance with rules defined by Kayser *et al.* (2004). Separation of repeat blocks by "N" represents the break point between the separate loci present within complex markers (as described in Methods and Materials). Repeat arrays observed to be variable through sequence analysis are highlighted in bold. The allele ranges given are those seen with the 1966 father-son pairs of European origin.

The "Total number of Meioses" reported is the number of meioses for the marker, and is not the number of allele transmissions for multicopy markers.

The mutations reported for the Yfiler loci have been previously reported in Reference 28.

PCR Protocols:

| TD60-50 | | | TD65-50 | | | TD65-55 | | | TD70-50 | | |
|---------|---------|-----|---------|---------|-----|---------|---------|-----|---------|---------|-----|
| 95C | 10 min | | 95C | 15 min | | 95C | 10 min | | 95C | 15 min | |
| 94C | 30s | X10 | 94C | 30s | X20 | 94C | 30s | X10 | 94C | 30s | X20 |
| 60-1C | 30s | | 65-1C | 45s | | 65-1C | 30s | | 70-1C | 45s | |
| 72C | 45s | | 72C | 1 min | | 72C | 45s | | 72C | 1 min | |
| 94C | 30s | X25 | 94C | 30s | X15 | 94C | 30s | X25 | 94C | 30s | X15 |
| 50C | 30s | | 50C | 30s | | 55C | 30s | | 50C | 30s | |
| 72C | 45s | | 72C | 45s | | 72C | 45s | | 72C | 45s | |
| 60C | 45min | | 60C | 45min | | 60C | 45min | | 60C | 45min | |
| 15C | forever | | 15C | forever | | 15C | forever | | 15C | forever | |

PCR Protocols for Extra Genotyping of RM Y-STRs:

| RM 1 (DYF387S1, DYF399S1, DYS570, DYS576) | | RM 2 (DYS518, DYS526a/b, DYS626, DYS627) | | RM 3 (DYF403S1a/b, DYF404S1, DYS449, DYS547, DYS612) | |
|--|---------|---|---------|---|---------|
| PCR Buffer | 1x | PCR Buffer | 1x | PCR Buffer | 1x |
| MgCl ₂ | 2.27mM | MgCl ₂ | 1.5mM | MgCl ₂ | 2.0mM |
| dNTPs | 220µM | dNTPs | 250µM | dNTPs | 250µM |
| DYF387S1 Primer | 0.09µM | DYS518 Primer | 0.5µM | DYF403S1a/b Primer | 0.6µM |
| DYF399S1 Primer | 0.36µM | DYS526a/b Primer | 0.35µM | DYSF404S1 Primer | 0.1µM |
| DYS570 Primer | 0.09µM | DYS626 Primer | 0.2µM | DYS449 Primer | 0.1µM |
| DYS576 Primer | 0.09µM | DYS627Primer | 0.15µM | DYS547 Primer | 0.6µM |
| | | | | DYS612 Primer | 0.2µM |
| Taq | 0.25U | Taq | 0.35U | Taq | 0.5U |
| DNA | 2ng | DNA | 2ng | DNA | 2ng |
| PCR Protocol | TD70-50 | PCR Protocol | TD65-55 | PCR Protocol | TD65-55 |

Table S2. Details of the 924 Y-STR Mutations Observed among DNA-Confirmed Father-Son Pairs (See Footnote for Explanation)

| Locus | Father Allele | Son Allele | Father's Age | Father Reference # |
|----------|---|---|--------------|--------------------|
| DYF382S1 | (GGAT) ₁₃ (AGAT) ₁ (GGAT) ₃ N ₈ (GGAC) ₃ | (GGAT) ₁₄ (AGAT) ₁ (GGAT) ₃ N ₈ (GGAC) ₃ | 59 | 1953 |
| DYF386S1 | (AAT) ₁₂ | (AAT) ₁₃ | 27 | 20 |
| DYF386S1 | (AAT) ₁₄ | (AAT) ₁₃ | 32 | 84 |
| DYF386S1 | (AAT) ₁₃ | (AAT) ₁₂ | 38 | 94 |
| DYF386S1 | (AAT) ₁₅ | (AAT) ₁₃ | 20 | 208 |
| DYF386S1 | (AAT) ₁₄ | (AAT) ₁₃ | 25 | 317 |
| DYF386S1 | (AAT) ₁₁ | (AAT) ₁₂ | 19 | 641 |
| DYF386S1 | (AAT) ₁₄ | (AAT) ₁₃ | 27 | 1195 |
| DYF386S1 | (AAT) ₁₄ | (AAT) ₁₅ | 40 | 1558 |
| DYF386S1 | (AAT) ₁₄ | (AAT) ₁₃ | 42 | 1644 |
| DYF386S1 | (AAT) ₁₄ | (AAT) ₁₃ | 52 | 1864 |
| DYF387S1 | 21 | 24 | 21 | 46 |
| DYF387S1 | 23 | 22 | 48 | 74 |
| DYF387S1 | 23 | 24 | 36 | 150 |
| DYF387S1 | 20 | 21 | 29 | 155 |
| DYF387S1 | 24 | 23 | 24 | 259 |
| DYF387S1 | 22 | 21 | 28 | 677 |
| DYF387S1 | 24 | 23 | 40 | 738 |
| DYF387S1 | 25 | 24 | 22 | 817 |
| DYF387S1 | 25 | 26 | 36 | 830 |
| DYF387S1 | 25 | 26 | 18 | 852 |
| DYF387S1 | 23 | 24 | 28 | 880 |
| DYF387S1 | 24 | 23 | 19 | 916 |
| DYF387S1 | 21 | 22 | 33 | 955 |
| DYF387S1 | 22 | 23 | 43 | 1159 |
| DYF387S1 | 23 | 24 | Unknown | 1202 |

| | | | | |
|----------|--|--|---------|------|
| DYF387S1 | 24 | 23 | 30 | 1274 |
| DYF387S1 | 23 | 22 | 37 | 1319 |
| DYF387S1 | 23 | 21 | Unknown | 1328 |
| DYF387S1 | 24 | 25 | 31 | 1390 |
| DYF387S1 | 23 | 24 | 39 | 1451 |
| DYF387S1 | 22 | 21 | 24 | 1469 |
| DYF387S1 | 22 | 23 | 38 | 1494 |
| DYF387S1 | 21 | 22 | 39 | 1552 |
| DYF387S1 | 21 | 22 | 20 | 1592 |
| DYF387S1 | 23 | 22 | 42 | 1644 |
| DYF387S1 | 23 | 22 | 23 | 1710 |
| DYF387S1 | 23 | 22 | 18 | 1750 |
| DYF387S1 | 23 | 25 | 54 | 1911 |
| DYF388S1 | (CTTC) ₆ (CTTT) ₁₂ N ₁₈ (CTTC) ₃ (TTTC) ₁ (CTTC) ₃ N ₈ (CTTC) ₃ N ₃₂ (CTTC) ₃ | (CTTC) ₆ (CTTT) ₁₃ N ₁₈ (CTTC) ₃ (TTTC) ₁ (CTTC) ₃ N ₈ (CTTC) ₃ N ₃₂ (CTTC) ₃ | 37 | 36 |
| DYF388S1 | (CTTC) ₆ (CTTT) ₁₄ N ₁₈ (CTTC) ₃ (TTTC) ₁ (CTTC) ₃ N ₈ (CTTC) ₃ N ₃₂ (CTTC) ₃ | (CTTC) ₆ (CTTT) ₁₂ N ₁₈ (CTTC) ₃ (TTTC) ₁ (CTTC) ₃ N ₈ (CTTC) ₃ N ₃₂ (CTTC) ₃ | 34 | 372 |
| DYF388S1 | (CTTC) ₆ (CTTT) ₁₂ N ₁₈ (CTTC) ₃ (TTTC) ₁ (CTTC) ₃ N ₈ (CTTC) ₃ N ₃₂ (CTTC) ₃ | (CTTC) ₆ (CTTT) ₁₃ N ₁₈ (CTTC) ₃ (TTTC) ₁ (CTTC) ₃ N ₈ (CTTC) ₃ N ₃₂ (CTTC) ₃ | 28 | 674 |
| DYF388S1 | (CTTC) ₆ (CTTT) ₁₂ N ₁₈ (CTTC) ₃ (TTTC) ₁ (CTTC) ₃ N ₈ (CTTC) ₃ N ₃₂ (CTTC) ₃ | (CTTC) ₆ (CTTT) ₁₃ N ₁₈ (CTTC) ₃ (TTTC) ₁ (CTTC) ₃ N ₈ (CTTC) ₃ N ₃₂ (CTTC) ₃ | 21 | 769 |
| DYF388S1 | (CTTC) ₆ (CTTT) ₁₁ N ₁₈ (CTTC) ₃ (TTTC) ₁ (CTTC) ₃ N ₈ (CTTC) ₃ N ₃₂ (CTTC) ₃ | (CTTC) ₆ (CTTT) ₁₂ N ₁₈ (CTTC) ₃ (TTTC) ₁ (CTTC) ₃ N ₈ (CTTC) ₃ N ₃₂ (CTTC) ₃ | 49 | 945 |
| DYF388S1 | (CTTC) ₆ (CTTT) ₁₂ N ₁₈ (CTTC) ₃ (TTTC) ₁ (CTTC) ₃ N ₈ (CTTC) ₃ N ₃₂ (CTTC) ₃ | (CTTC) ₆ (CTTT) ₁₁ N ₁₈ (CTTC) ₃ (TTTC) ₁ (CTTC) ₃ N ₈ (CTTC) ₃ N ₃₂ (CTTC) ₃ | 28 | 1035 |
| DYF388S1 | (CTTC) ₆ (CTTT) ₁₂ N ₁₈ (CTTC) ₃ (TTTC) ₁ (CTTC) ₃ N ₈ (CTTC) ₃ N ₃₂ (CTTC) ₃ | (CTTC) ₆ (CTTT) ₁₃ N ₁₈ (CTTC) ₃ (TTTC) ₁ (CTTC) ₃ N ₈ (CTTC) ₃ N ₃₂ (CTTC) ₃ | 30 | 1177 |
| DYF388S1 | (CTTC) ₆ (CTTT) ₁₃ N ₁₈ (CTTC) ₃ (TTTC) ₁ (CTTC) ₃ N ₈ (CTTC) ₃ N ₃₂ (CTTC) ₃ | (CTTC) ₆ (CTTT) ₁₂ N ₁₈ (CTTC) ₃ (TTTC) ₁ (CTTC) ₃ N ₈ (CTTC) ₃ N ₃₂ (CTTC) ₃ | 56 | 1272 |
| DYF388S1 | (CTTC) ₆ (CTTT) ₁₃ N ₁₈ (CTTC) ₃ (TTTC) ₁ (CTTC) ₃ N ₈ (CTTC) ₃ N ₃₂ (CTTC) ₃ | (CTTC) ₆ (CTTT) ₁₂ N ₁₈ (CTTC) ₃ (TTTC) ₁ (CTTC) ₃ N ₈ (CTTC) ₃ N ₃₂ (CTTC) ₃ | Unknown | 1352 |
| DYF388S1 | (CTTC) ₆ (CTTT) ₁₃ N ₁₈ (CTTC) ₃ (TTTC) ₁ (CTTC) ₃ N ₈ (CTTC) ₃ N ₃₂ (CTTC) ₃ | (CTTC) ₆ (CTTT) ₁₂ N ₁₈ (CTTC) ₃ (TTTC) ₁ (CTTC) ₃ N ₈ (CTTC) ₃ N ₃₂ (CTTC) ₃ | 31 | 1518 |
| DYF388S1 | (CTTC) ₆ (CTTT) ₁₄ N ₁₈ (CTTC) ₃ (TTTC) ₁ (CTTC) ₃ N ₈ (CTTC) ₃ N ₃₂ (CTTC) ₃ | (CTTC) ₆ (CTTT) ₁₂ N ₁₈ (CTTC) ₃ (TTTC) ₁ (CTTC) ₃ N ₈ (CTTC) ₃ N ₃₂ (CTTC) ₃ | 28 | 1734 |

| | | | | |
|----------|---|---|----|------|
| DYF390S1 | (TTTA) ₁₁ | (TTTA) ₁₀ | 21 | 1667 |
| DYF393S1 | (AAG) ₄ (AA) ₁ (AAG) ₂₆ (CAG) ₁ | (AAG) ₄ (AA) ₁ (AAG) ₂₅ (CAG) ₁ | 32 | 19 |
| DYF393S1 | (AAG) ₄ (AA) ₁ (AAG) ₂₈ (CAG) ₁ | (AAG) ₄ (AA) ₁ (AAG) ₂₉ (CAG) ₁ | 32 | 84 |
| DYF393S1 | (AAG) ₄ (AA) ₁ (AAG) ₁₉ (CAG) ₁ | (AAG) ₄ (AA) ₁ (AAG) ₂₀ (CAG) ₁ | 34 | 183 |
| DYF393S1 | (AAG) ₄ (AA) ₁ (AAG) ₂₆ (CAG) ₁ | (AAG) ₄ (AA) ₁ (AAG) ₂₅ (CAG) ₁ | 32 | 213 |
| DYF393S1 | (AAG) ₄ (AA) ₁ (AAG) ₂₅ (CAG) ₁ | (AAG) ₄ (AA) ₁ (AAG) ₂₄ (CAG) ₁ | 26 | 303 |
| DYF393S1 | (AAG) ₄ (AA) ₁ (AAG) ₂₂ (CAG) ₁ | (AAG) ₄ (AA) ₁ (AAG) ₂₃ (CAG) ₁ | 31 | 927 |
| DYF393S1 | (AAG) ₄ (AA) ₁ (AAG) ₂₆ (CAG) ₂ | (AAG) ₄ (AA) ₁ (AAG) ₂₇ (CAG) ₂ | 23 | 941 |
| DYF393S1 | (AAG) ₄ (AA) ₁ (AAG) ₂₂ (CAG) ₁ | (AAG) ₄ (AA) ₁ (AAG) ₂₃ (CAG) ₁ | 64 | 951 |
| DYF393S1 | (AAG) ₄ (AA) ₁ (AAG) ₂₇ (CAG) ₁ | (AAG) ₄ (AA) ₁ (AAG) ₂₆ (CAG) ₁ | 21 | 1207 |
| DYF393S1 | (AAG) ₄ (AA) ₁ (AAG) ₂₂ (CAG) ₁ | (AAG) ₄ (AA) ₁ (AAG) ₂₄ (CAG) ₁ | 17 | 1406 |
| DYF393S1 | (AAG) ₄ (AA) ₁ (AAG) ₂₈ (CAG) ₁ | (AAG) ₄ (AA) ₁ (AAG) ₂₉ (CAG) ₁ | 19 | 1530 |
| DYF393S1 | (AAG) ₄ (AA) ₁ (AAG) ₂₇ (CAG) ₁ | (AAG) ₄ (AA) ₁ (AAG) ₂₈ (CAG) ₁ | 26 | 1551 |
| DYF393S1 | (AAG) ₄ (AA) ₁ (AAG) ₂₃ (CAG) ₁ | (AAG) ₄ (AA) ₁ (AAG) ₂₄ (CAG) ₁ | 36 | 1672 |
| DYF393S1 | (AAG) ₄ (AA) ₁ (AAG) ₂₅ (CAG) ₁ | (AAG) ₄ (AA) ₁ (AAG) ₂₄ (CAG) ₁ | 55 | 1928 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₈ (GAAA) ₂₀ | 22 | 14 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₆ | (GAAA) ₃ N ₈ (GAAA) ₁₇ | 46 | 21 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₈ (GAAA) ₂₀ | 36 | 22 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₁₈ | (GAAA) ₃ N ₈ (GAAA) ₁₉ | 30 | 25 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₂₁ | (GAAA) ₃ N ₈ (GAAA) ₂₀ | 32 | 32 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₂₀ | (GAAA) ₃ N ₈ (GAAA) ₁₉ | 16 | 55 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₁₈ | (GAAA) ₃ N ₈ (GAAA) ₁₉ | 32 | 59 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₂₀ | (GAAA) ₃ N ₈ (GAAA) ₁₉ | 30 | 62 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₇ (GAAA) ₁₈ | 46 | 72 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₁₈ | (GAAA) ₃ N ₈ (GAAA) ₁₉ | 46 | 72 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₈ (GAAA) ₂₀ | 25 | 79 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₈ (GAAA) ₂₀ | 25 | 80 |

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| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₂₀ | (GAAA) ₃ N ₈ (GAAA) ₁₉ | 28 | 91 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₂₁ | (GAAA) ₃ N ₈ (GAAA) ₂₀ | 32 | 95 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₇ | (GAAA) ₃ N ₈ (GAAA) ₁₆ | 30 | 99 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₂₀ | (GAAA) ₃ N ₈ (GAAA) ₁₉ | 27 | 119 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₈ (GAAA) ₂₀ | 48 | 122 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₇ (GAAA) ₁₈ | 33 | 126 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₇ | (GAAA) ₃ N ₈ (GAAA) ₁₆ | 36 | 136 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₈ (GAAA) ₂₀ | 28 | 153 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₁₈ | (GAAA) ₃ N ₈ (GAAA) ₁₇ | 33 | 189 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₇ | (GAAA) ₃ N ₇ (GAAA) ₁₈ | 39 | 200 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₆ | (GAAA) ₃ N ₈ (GAAA) ₁₅ | 22 | 203 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₇ (GAAA) ₁₈ | 22 | 229 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₆ | (GAAA) ₃ N ₈ (GAAA) ₁₅ | 50 | 270 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₈ (GAAA) ₂₀ | 28 | 287 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₂₁ | (GAAA) ₃ N ₈ (GAAA) ₂₀ | 32 | 290 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₆ | (GAAA) ₃ N ₈ (GAAA) ₁₇ | 21 | 299 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₂₂ | (GAAA) ₃ N ₇ (GAAA) ₂₁ | 27 | 302 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₅ | (GAAA) ₃ N ₈ (GAAA) ₁₆ | 38 | 336 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₁₈ | (GAAA) ₃ N ₈ (GAAA) ₁₇ | 50 | 356 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₂₁ | (GAAA) ₃ N ₈ (GAAA) ₂₂ | 28 | 367 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₆ | (GAAA) ₃ N ₈ (GAAA) ₁₇ | 34 | 372 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₂₁ | (GAAA) ₃ N ₈ (GAAA) ₂₂ | 28 | 373 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₁₈ | (GAAA) ₃ N ₈ (GAAA) ₁₇ | 35 | 389 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₆ | (GAAA) ₃ N ₈ (GAAA) ₁₇ | 32 | 401 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₁₈ | (GAAA) ₃ N ₈ (GAAA) ₁₇ | 53 | 453 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₇ | (GAAA) ₃ N ₇ (GAAA) ₁₈ | 34 | 459 |

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| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₇ | (GAAA) ₃ N ₈ (GAAA) ₁₆ | 26 | 480 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₁₈ | (GAAA) ₃ N ₈ (GAAA) ₁₉ | 28 | 484 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₇ (GAAA) ₁₈ | 26 | 488 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₂₁ | (GAAA) ₃ N ₈ (GAAA) ₂₀ | 39 | 492 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₈ (GAAA) ₂₀ | 27 | 494 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₈ (GAAA) ₂₀ | 35 | 500 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₆ | (GAAA) ₃ N ₈ (GAAA) ₁₅ | 19 | 546 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₈ (GAAA) ₂₀ | 34 | 559 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₂₁ | (GAAA) ₃ N ₈ (GAAA) ₂₂ | 30 | 586 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₂₀ | (GAAA) ₃ N ₈ (GAAA) ₁₉ | 29 | 603 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₁₈ | (GAAA) ₃ N ₈ (GAAA) ₁₇ | 56 | 608 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₇ (GAAA) ₁₈ | 32 | 614 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₇ | (GAAA) ₃ N ₈ (GAAA) ₁₆ | 43 | 624 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₂₀ | (GAAA) ₃ N ₈ (GAAA) ₁₉ | 25 | 630 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₂₁ | (GAAA) ₃ N ₈ (GAAA) ₂₀ | 30 | 640 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₇ (GAAA) ₁₈ | 43 | 657 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₂₂ | (GAAA) ₃ N ₇ (GAAA) ₂₁ | 25 | 666 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₈ (GAAA) ₂₀ | 23 | 675 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₇ (GAAA) ₁₈ | 18 | 680 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₁₈ | (GAAA) ₃ N ₈ (GAAA) ₁₉ | 23 | 687 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₇ | (GAAA) ₃ N ₇ (GAAA) ₁₈ | 20 | 706 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₈ (GAAA) ₂₀ | 33 | 718 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₅ | (GAAA) ₃ N ₈ (GAAA) ₁₆ | 21 | 720 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₅ | (GAAA) ₃ N ₈ (GAAA) ₁₄ | 22 | 747 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₂₀ | (GAAA) ₃ N ₈ (GAAA) ₁₉ | 28 | 772 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₆ | (GAAA) ₃ N ₈ (GAAA) ₁₅ | 28 | 805 |

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| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₂₂ | (GAAA) ₃ N ₈ (GAAA) ₂₀ | 22 | 817 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₂₁ | (GAAA) ₃ N ₈ (GAAA) ₂₀ | 20 | 824 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₂₀ | (GAAA) ₃ N ₈ (GAAA) ₁₉ | 20 | 827 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₇ (GAAA) ₁₈ | 25 | 877 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₂₀ | (GAAA) ₃ N ₈ (GAAA) ₁₉ | 31 | 881 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₇ (GAAA) ₁₈ | 19 | 900 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₆ | (GAAA) ₃ N ₈ (GAAA) ₁₇ | 37 | 904 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₅ | (GAAA) ₃ N ₈ (GAAA) ₁₄ | 34 | 911 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₁₈ | (GAAA) ₃ N ₈ (GAAA) ₁₇ | 19 | 916 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₁₈ | (GAAA) ₃ N ₈ (GAAA) ₁₆ | 24 | 926 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₇ (GAAA) ₁₈ | 22 | 986 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₂₀ | (GAAA) ₃ N ₇ (GAAA) ₂₁ | 36 | 989 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₁₈ | (GAAA) ₃ N ₈ (GAAA) ₁₇ | 35 | 1001 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₂₁ | (GAAA) ₃ N ₈ (GAAA) ₂₀ | 41 | 1008 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₂₀ | (GAAA) ₃ N ₈ (GAAA) ₁₉ | 41 | 1008 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₆ | (GAAA) ₃ N ₈ (GAAA) ₁₅ | 33 | 1046 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₅ | (GAAA) ₃ N ₈ (GAAA) ₁₆ | 33 | 1046 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₇ (GAAA) ₁₈ | 31 | 1072 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₂₀ | (GAAA) ₃ N ₈ (GAAA) ₁₉ | 22 | 1088 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₁₈ | (GAAA) ₃ N ₈ (GAAA) ₁₉ | 36 | 1091 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₅ | (GAAA) ₃ N ₈ (GAAA) ₁₆ | 23 | 1095 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₇ | (GAAA) ₃ N ₇ (GAAA) ₁₈ | 25 | 1104 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₂₀ | (GAAA) ₃ N ₈ (GAAA) ₁₉ | 43 | 1138 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₁₈ | (GAAA) ₃ N ₈ (GAAA) ₁₇ | 20 | 1154 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₇ | (GAAA) ₃ N ₈ (GAAA) ₁₆ | 36 | 1155 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₆ | (GAAA) ₃ N ₈ (GAAA) ₁₇ | 34 | 1158 |

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| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₂₂ | (GAAA) ₃ N ₇ (GAAA) ₂₁ | 34 | 1158 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₁₈ | (GAAA) ₃ N ₈ (GAAA) ₁₇ | 43 | 1159 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₃ | (GAAA) ₃ N ₈ (GAAA) ₁₄ | 21 | 1163 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₇ (GAAA) ₁₈ | 21 | 1207 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₂₁ | (GAAA) ₃ N ₈ (GAAA) ₂₀ | Unknown | 1263 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₁₈ | (GAAA) ₃ N ₈ (GAAA) ₁₉ | Unknown | 1265 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₂₀ | (GAAA) ₃ N ₈ (GAAA) ₁₉ | Unknown | 1268 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₇ | (GAAA) ₃ N ₈ (GAAA) ₁₆ | 39 | 1327 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₂₁ | (GAAA) ₃ N ₈ (GAAA) ₂₀ | 27 | 1397 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₄ | (GAAA) ₃ N ₈ (GAAA) ₁₃ | 42 | 1407 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₇ (GAAA) ₁₈ | 42 | 1411 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₂₂ | (GAAA) ₃ N ₇ (GAAA) ₂₁ | 40 | 1443 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₇ | (GAAA) ₃ N ₈ (GAAA) ₁₆ | 17 | 1446 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₇ | (GAAA) ₃ N ₈ (GAAA) ₁₆ | 24 | 1455 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₁₈ | (GAAA) ₃ N ₈ (GAAA) ₁₉ | 18 | 1456 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₅ | (GAAA) ₃ N ₈ (GAAA) ₁₆ | 24 | 1466 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₂₀ | (GAAA) ₃ N ₇ (GAAA) ₂₁ | 24 | 1466 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₇ | (GAAA) ₃ N ₇ (GAAA) ₁₈ | 39 | 1471 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₄ | (GAAA) ₃ N ₈ (GAAA) ₁₅ | Unknown | 1479 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₇ (GAAA) ₁₈ | 28 | 1554 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₈ (GAAA) ₂₀ | 37 | 1577 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₇ | (GAAA) ₃ N ₇ (GAAA) ₁₈ | 40 | 1606 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₇ (GAAA) ₁₈ | 25 | 1612 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₇ | (GAAA) ₃ N ₈ (GAAA) ₁₆ | 17 | 1620 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₂₀ | (GAAA) ₃ N ₈ (GAAA) ₁₉ | Unknown | 1650 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₁₈ | (GAAA) ₃ N ₈ (GAAA) ₁₉ | 24 | 1651 |

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| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₇ (GAAA) ₁₈ | 24 | 1658 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₄ | (GAAA) ₃ N ₈ (GAAA) ₁₅ | 37 | 1663 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₂₂ | (GAAA) ₃ N ₇ (GAAA) ₂₁ | 32 | 1664 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₆ | (GAAA) ₃ N ₈ (GAAA) ₂₀ | 17 | 1665 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₅ | (GAAA) ₃ N ₈ (GAAA) ₁₄ | 20 | 1670 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₁₈ | (GAAA) ₃ N ₈ (GAAA) ₁₉ | 34 | 1691 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₂₀ | (GAAA) ₃ N ₇ (GAAA) ₂₁ | 28 | 1696 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₇ (GAAA) ₁₈ | 23 | 1722 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₆ | (GAAA) ₃ N ₈ (GAAA) ₁₇ | 30 | 1733 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₂₀ | (GAAA) ₃ N ₇ (GAAA) ₂₁ | 26 | 1760 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₇ | (GAAA) ₃ N ₇ (GAAA) ₁₈ | 29 | 1773 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₂₀ | (GAAA) ₃ N ₈ (GAAA) ₁₉ | 31 | 1798 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₅ | (GAAA) ₃ N ₈ (GAAA) ₁₆ | 51 | 1813 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₂₁ | (GAAA) ₃ N ₈ (GAAA) ₂₀ | 19 | 1841 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₂₂ | (GAAA) ₃ N ₇ (GAAA) ₂₁ | 55 | 1844 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₂₂ | (GAAA) ₃ N ₈ (GAAA) ₂₃ | 59 | 1867 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₇ | (GAAA) ₃ N ₈ (GAAA) ₁₆ | 53 | 1869 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₆ | (GAAA) ₃ N ₈ (GAAA) ₁₅ | 73 | 1884 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₁₈ | (GAAA) ₃ N ₈ (GAAA) ₁₉ | 52 | 1891 |
| DYF399S1 | (GAAA) ₃ N ₇ (GAAA) ₂₁ | (GAAA) ₃ N ₈ (GAAA) ₂₀ | 52 | 1891 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₆ | (GAAA) ₃ N ₈ (GAAA) ₁₅ | 55 | 1909 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₉ | (GAAA) ₃ N ₇ (GAAA) ₁₈ | 60 | 1913 |
| DYF399S1 | (GAAA) ₃ N ₈ (GAAA) ₁₇ | (GAAA) ₃ N ₈ (GAAA) ₁₆ | 50 | 1941 |
| DYF401S1 | (AAGG) ₃ (AAGC) ₁ (AAGG) ₃ N ₃₉ (AAGG) ₃ N ₈ (AAGG) ₃ (AAAG) ₁ (AAGG) ₃ N ₁₃ (AAAG) ₁₅ G(AAGG) ₆ | (AAGG) ₃ (AAGC) ₁ (AAGG) ₃ N ₃₉ (AAGG) ₃ N ₈ (AAGG) ₃ (AAAG) ₁ (AAGG) ₃ N ₁₃ (AAAG) ₁₆ G(AAGG) ₆ | 37 | 36 |
| DYF401S1 | (AAGG) ₃ (AAGC) ₁ (AAGG) ₃ N ₃₉ (AAGG) ₃ N ₈ (AAGG) ₃ (AAAG) ₁ (AAGG) ₃ N ₁₃ (AAAG) ₁₇ G(AAGG) ₆ | (AAGG) ₃ (AAGC) ₁ (AAGG) ₃ N ₃₉ (AAGG) ₃ N ₈ (AAGG) ₃ (AAAG) ₁ (AAGG) ₃ N ₁₃ (AAAG) ₁₅ G(AAGG) ₆ | 34 | 372 |
| DYF401S1 | (AAGG) ₃ (AAGC) ₁ (AAGG) ₃ N ₃₉ (AAGG) ₃ N ₈ (AAGG) ₃ | (AAGG) ₃ (AAGC) ₁ (AAGG) ₃ N ₃₉ (AAGG) ₃ N ₈ (AAGG) ₃ | 28 | 674 |

| | (AAAG) ₁ (AAGG) ₃ N ₁₃ (AAAG) ₁₅ G(AAGG) ₆ | (AAAG) ₁ (AAGG) ₃ N ₁₃ (AAAG) ₁₆ G(AAGG) ₆ | | |
|-----------|---|---|---------|------|
| DYF401S1 | (AAGG) ₃ (AAGC) ₁ (AAGG) ₃ N ₃₉ (AAGG) ₃ N ₈ (AAGG) ₃ (AAAG) ₁ (AAGG) ₃ N ₁₃ (AAAG) ₁₅ G(AAGG) ₆ | (AAGG) ₃ (AAGC) ₁ (AAGG) ₃ N ₃₉ (AAGG) ₃ N ₈ (AAGG) ₃ (AAAG) ₁ (AAGG) ₃ N ₁₃ (AAAG) ₁₆ G(AAGG) ₆ | 21 | 769 |
| DYF401S1 | (AAGG) ₃ (AAGC) ₁ (AAGG) ₃ N ₃₉ (AAGG) ₃ N ₈ (AAGG) ₃ (AAAG) ₁ (AAGG) ₃ N ₁₃ (AAAG) ₁₅ G(AAGG) ₆ | (AAGG) ₃ (AAGC) ₁ (AAGG) ₃ N ₃₉ (AAGG) ₃ N ₈ (AAGG) ₃ (AAAG) ₁ (AAGG) ₃ N ₁₃ (AAAG) ₁₄ G(AAGG) ₆ | Unknown | 1241 |
| DYF401S1 | (AAGG) ₃ (AAGC) ₁ (AAGG) ₃ N ₃₉ (AAGG) ₃ N ₈ (AAGG) ₃ (AAAG) ₁ (AAGG) ₃ N ₁₃ (AAAG) ₁₆ G(AAGG) ₆ | (AAGG) ₃ (AAGC) ₁ (AAGG) ₃ N ₃₉ (AAGG) ₃ N ₈ (AAGG) ₃ (AAAG) ₁ (AAGG) ₃ N ₁₃ (AAAG) ₁₅ G(AAGG) ₆ | 56 | 1272 |
| DYF401S1 | (AAGG) ₃ (AAGC) ₁ (AAGG) ₃ N ₃₉ (AAGG) ₃ N ₈ (AAGG) ₃ (AAAG) ₁ (AAGG) ₃ N ₁₃ (AAAG) ₁₆ G(AAGG) ₆ | (AAGG) ₃ (AAGC) ₁ (AAGG) ₃ N ₃₉ (AAGG) ₃ N ₈ (AAGG) ₃ (AAAG) ₁ (AAGG) ₃ N ₁₃ (AAAG) ₁₅ G(AAGG) ₆ | 31 | 1518 |
| DYF401S1 | (AAGG) ₃ (AAGC) ₁ (AAGG) ₃ N ₃₉ (AAGG) ₃ N ₈ (AAGG) ₃ (AAAG) ₁ (AAGG) ₃ N ₁₃ (AAAG) ₁₇ G(AAGG) ₆ | (AAGG) ₃ (AAGC) ₁ (AAGG) ₃ N ₃₉ (AAGG) ₃ N ₈ (AAGG) ₃ (AAAG) ₁ (AAGG) ₃ N ₁₃ (AAAG) ₁₅ G(AAGG) ₆ | 28 | 1734 |
| DYF403S1a | 342 | 338 | 20 | 73 |
| DYF403S1a | 321 | 316 | 32 | 128 |
| DYF403S1a | 316 | 321 | 27 | 132 |
| DYF403S1a | 316,329,338 | 321,325,334 | 17 | 175 |
| DYF403S1a | 346 | 350 | 37 | 186 |
| DYF403S1a | 342 | 346 | 22 | 201 |
| DYF403S1a | 354 | 350 | 37 | 406 |
| DYF403S1a | 350 | 354 | 20 | 423 |
| DYF403S1a | 342 | 346 | 19 | 546 |
| DYF403S1a | 325 | 329 | 24 | 681 |
| DYF403S1a | 325 | 321 | 39 | 749 |
| DYF403S1a | 334 | 338 | 22 | 817 |
| DYF403S1a | 321 | 342 | 28 | 841 |
| DYF403S1a | 338 | 342 | 37 | 904 |
| DYF403S1a | 308 | 312 | 34 | 911 |
| DYF403S1a | 342 | 346 | 18 | 943 |
| DYF403S1a | 312 | 316 | 20 | 977 |
| DYF403S1a | 342 | 346 | 46 | 1033 |
| DYF403S1a | 321 | 316 | 21 | 1053 |
| DYF403S1a | 350 | 354 | 39 | 1071 |
| DYF403S1a | 342 | 346 | 27 | 1085 |

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|-----------|---------|---------|---------|------|
| DYF403S1a | 325 | 329 | 44 | 1110 |
| DYF403S1a | 342 | 338 | 22 | 1323 |
| DYF403S1a | 346 | 350 | 36 | 1336 |
| DYF403S1a | 312 | 316 | Unknown | 1353 |
| DYF403S1a | 354 | 350 | 45 | 1364 |
| DYF403S1a | 342 | 338 | 44 | 1378 |
| DYF403S1a | 342 | 338 | 42 | 1411 |
| DYF403S1a | 346 | 350 | 42 | 1441 |
| DYF403S1a | 338 | 342 | 24 | 1480 |
| DYF403S1a | 325 | 329 | 40 | 1531 |
| DYF403S1a | 346 | 342 | 28 | 1554 |
| DYF403S1a | 312 | 316 | 33 | 1561 |
| DYF403S1a | 321 | 316 | 25 | 1634 |
| DYF403S1a | 342 | 334 | 36 | 1643 |
| DYF403S1a | 329 | 334 | 24 | 1704 |
| DYF403S1a | 312 | 316 | 26 | 1725 |
| DYF403S1a | 312 | 316 | 40 | 1785 |
| DYF403S1a | 312 | 316 | 31 | 1798 |
| DYF403S1a | 329 | 334 | 43 | 1818 |
| DYF403S1a | 346 | 342 | 27 | 1840 |
| DYF403S1a | 346 | 350 | 43 | 1883 |
| DYF403S1a | 329 | 334 | 57 | 1896 |
| DYF403S1a | 334 | 321 | 64 | 1912 |
| DYF403S1a | 346 | 342 | 50 | 1941 |
| DYF403S1a | 325,346 | 329,339 | 56 | 1947 |
| DYF403S1b | 50 | 49 | 40 | 28 |
| DYF403S1b | 46.1 | 45.1 | 32 | 115 |
| DYF403S1b | 50 | 49 | 33 | 137 |
| DYF403S1b | 49 | 47 | 17 | 175 |
| DYF403S1b | 51 | 50 | 53 | 453 |

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|-----------|--|--|---------|------|
| DYF403S1b | 51 | 50 | 18 | 470 |
| DYF403S1b | 50 | 49 | 39 | 749 |
| DYF403S1b | 53 | 52 | 19 | 916 |
| DYF403S1b | 52 | 53 | 21 | 1090 |
| DYF403S1b | 50 | 49 | Unknown | 1288 |
| DYF403S1b | 52 | 51 | 25 | 1381 |
| DYF403S1b | 46.1 | 47.1 | 54 | 1447 |
| DYF403S1b | 49 | 50 | Unknown | 1479 |
| DYF403S1b | 48 | 49 | 23 | 1761 |
| DYF403S1b | 49 | 48 | 56 | 1947 |
| DYF403S1b | 46.1 | 47.1 | 54 | 1951 |
| DYF404S1 | (TTTC) ₁₅ N ₄₂ (TTTC) ₃ | (TTTC) ₁₆ N ₄₂ (TTTC) ₃ | 28 | 9 |
| DYF404S1 | (TTTC) ₁₅ N ₄₂ (TTTC) ₃ | (TTTC) ₁₄ N ₄₂ (TTTC) ₃ | 26 | 376 |
| DYF404S1 | (TTTC) ₁₆ N ₄₂ (TTTC) ₃ | (TTTC) ₁₇ N ₄₂ (TTTC) ₃ | 29 | 415 |
| DYF404S1 | (TTTC) ₁₆ N ₄₂ (TTTC) ₃ | (TTTC) ₁₇ N ₄₂ (TTTC) ₃ | 29 | 481 |
| DYF404S1 | (TTTC) ₁₆ N ₄₂ (TTTC) ₃ | (TTTC) ₁₇ N ₄₂ (TTTC) ₃ | 20 | 706 |
| DYF404S1 | (TTTC) ₁₅ N ₄₂ (TTTC) ₃ | (TTTC) ₁₆ N ₄₂ (TTTC) ₃ | 22 | 757 |
| DYF404S1 | (TTTC) ₁₅ N ₄₂ (TTTC) ₃ | (TTTC) ₁₆ N ₄₂ (TTTC) ₃ | 18 | 910 |
| DYF404S1 | (TTTC) ₁₆ N ₄₂ (TTTC) ₃ | (TTTC) ₁₇ N ₄₂ (TTTC) ₃ | 32 | 1007 |
| DYF404S1 | (TTTC) ₁₄ N ₄₂ (TTTC) ₃ | (TTTC) ₁₅ N ₄₂ (TTTC) ₃ | 23 | 1012 |
| DYF404S1 | (TTTC) ₁₅ N ₄₂ (TTTC) ₃ | (TTTC) ₁₆ N ₄₂ (TTTC) ₃ | 27 | 1049 |
| DYF404S1 | (TTTC) ₁₈ N ₄₂ (TTTC) ₃ | (TTTC) ₁₇ N ₄₂ (TTTC) ₃ | 31 | 1084 |
| DYF404S1 | (TTTC) ₁₄ N ₄₂ (TTTC) ₃ | (TTTC) ₁₅ N ₄₂ (TTTC) ₃ | 38 | 1114 |
| DYF404S1 | (TTTC) ₁₆ N ₄₂ (TTTC) ₃ | (TTTC) ₁₅ N ₄₂ (TTTC) ₃ | 21 | 1396 |
| DYF404S1 | (TTTC) ₁₆ N ₄₂ (TTTC) ₃ | (TTTC) ₁₇ N ₄₂ (TTTC) ₃ | 23 | 1546 |
| DYF404S1 | (TTTC) ₁₅ N ₄₂ (TTTC) ₃ | (TTTC) ₁₄ N ₄₂ (TTTC) ₃ | 40 | 1578 |
| DYF404S1 | (TTTC) ₁₇ N ₄₂ (TTTC) ₃ | (TTTC) ₁₈ N ₄₂ (TTTC) ₃ | 36 | 1655 |
| DYF404S1 | (TTTC) ₁₇ N ₄₂ (TTTC) ₃ | (TTTC) ₁₆ N ₄₂ (TTTC) ₃ | 26 | 1657 |
| DYF404S1 | (TTTC) ₁₆ N ₄₂ (TTTC) ₃ | (TTTC) ₁₇ N ₄₂ (TTTC) ₃ | 36 | 1687 |
| DYF404S1 | (TTTC) ₁₅ N ₄₂ (TTTC) ₃ | (TTTC) ₁₄ N ₄₂ (TTTC) ₃ | 19 | 1739 |

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|----------|---|---|---------|------|
| DYF404S1 | (TTTC) ₁₈ N ₄₂ (TTTC) ₃ | (TTTC) ₁₇ N ₄₂ (TTTC) ₃ | 50 | 1881 |
| DYF404S1 | (TTTC) ₁₇ N ₄₂ (TTTC) ₃ | (TTTC) ₁₈ N ₄₂ (TTTC) ₃ | 60 | 1913 |
| DYF405S1 | (GGAA) ₁₁ N ₁₁₅ (GGAA) ₃ (GAAA) ₁ (GGAA) ₃ | (GGAA) ₁₂ N ₁₁₅ (GGAA) ₃ (GAAA) ₁ (GGAA) ₃ | 31 | 438 |
| DYF405S1 | (GGAA) ₁₄ N ₁₁₅ (GGAA) ₃ (GAAA) ₁ (GGAA) ₃ | (GGAA) ₁₃ N ₁₁₅ (GGAA) ₃ (GAAA) ₁ (GGAA) ₃ | 34 | 629 |
| DYF406S1 | (TATC) ₁₂ | (TATC) ₁₁ | 26 | 204 |
| DYF406S1 | (TATC) ₁₁ | (TATC) ₁₀ | 47 | 347 |
| DYF406S1 | (TATC) ₁₂ | (TATC) ₁₃ | 35 | 518 |
| DYF406S1 | (TATC) ₁₁ | (TATC) ₁₀ | 33 | 1011 |
| DYF406S1 | (TATC) ₁₂ | (TATC) ₁₃ | 41 | 1122 |
| DYF406S1 | (TATC) ₁₂ | (TATC) ₁₃ | Unknown | 1263 |
| DYF410S1 | (AAAT) ₉ | (AAAT) ₁₀ | Unknown | 1457 |
| DYF410S1 | (AAAT) ₉ | (AAAT) ₁₀ | 21 | 1667 |
| DYS19 | (TAGA) ₃ (TAGG) ₁ (TAGA) ₁₃ | (TAGA) ₃ (TAGG) ₁ (TAGA) ₁₄ | 46 | 21 |
| DYS19 | (TAGA) ₃ (TAGG) ₁ (TAGA) ₁₁ | (TAGA) ₃ (TAGG) ₁ (TAGA) ₁₂ | 43 | 472 |
| DYS19 | (TAGA) ₃ (TAGG) ₁ (TAGA) ₁₄ | (TAGA) ₃ (TAGG) ₁ (TAGA) ₁₃ | 24 | 726 |
| DYS19 | (TAGA) ₃ (TAGG) ₁ (TAGA) ₁₄ | (TAGA) ₃ (TAGG) ₁ (TAGA) ₁₅ | 31 | 927 |
| DYS19 | (TAGA) ₃ (TAGG) ₁ (TAGA) ₁₂ | (TAGA) ₃ (TAGG) ₁ (TAGA) ₁₃ | Unknown | 1224 |
| DYS19 | (TAGA) ₃ (TAGG) ₁ (TAGA) ₁₄ | (TAGA) ₃ (TAGG) ₁ (TAGA) ₁₃ | 29 | 1257 |
| DYS19 | (TAGA) ₃ (TAGG) ₁ (TAGA) ₁₄ | (TAGA) ₃ (TAGG) ₁ (TAGA) ₁₃ | 24 | 1767 |
| DYS385a | (AAGG) ₄ N ₁₄ (AAAG) ₃ N ₁₂ (AAAG) ₃ N ₂₉ (AAGG) ₆ (GAAA) ₁₃ | (AAGG) ₄ N ₁₄ (AAAG) ₃ N ₁₂ (AAAG) ₃ N ₂₉ (AAGG) ₅ (GAAA) ₁₃ | 22 | 602 |
| DYS385a | (AAGG) ₄ N ₁₄ (AAAG) ₃ N ₁₂ (AAAG) ₃ N ₂₉ (AAGG) ₆ (GAAA) ₁₃ | (AAGG) ₄ N ₁₄ (AAAG) ₃ N ₁₂ (AAAG) ₃ N ₂₉ (AAGG) ₆ (GAAA) ₁₄ | 30 | 1000 |
| DYS385a | (AAGG) ₄ N ₁₄ (AAAG) ₃ N ₁₂ (AAAG) ₃ N ₂₉ (AAGG) ₆ (GAAA) ₁₄ | (AAGG) ₄ N ₁₄ (AAAG) ₃ N ₁₂ (AAAG) ₃ N ₂₉ (AAGG) ₆ (GAAA) ₁₅ | 60 | 1695 |
| DYS385b | (AAGG) ₄ N ₁₄ (AAAG) ₃ N ₁₂ (AAAG) ₃ N ₂₉ (AAGG) ₆ (GAAA) ₁₅ | (AAGG) ₄ N ₁₄ (AAAG) ₃ N ₁₂ (AAAG) ₃ N ₂₉ (AAGG) ₆ (GAAA) ₁₆ | 39 | 501 |
| DYS385b | (AAGG) ₄ N ₁₄ (AAAG) ₃ N ₁₂ (AAAG) ₃ N ₂₉ (AAGG) ₆ (GAAA) ₁₅ | (AAGG) ₄ N ₁₄ (AAAG) ₃ N ₁₂ (AAAG) ₃ N ₂₉ (AAGG) ₆ (GAAA) ₁₆ | 48 | 781 |
| DYS385b | (AAGG) ₄ N ₁₄ (AAAG) ₃ N ₁₂ (AAAG) ₃ N ₂₉ (AAGG) ₆ (GAAA) ₁₄ | (AAGG) ₄ N ₁₄ (AAAG) ₃ N ₁₂ (AAAG) ₃ N ₂₉ (AAGG) ₆ (GAAA) ₁₅ | 24 | 835 |
| DYS385b | (AAGG) ₄ N ₁₄ (AAAG) ₃ N ₁₂ (AAAG) ₃ N ₂₉ (AAGG) ₆ (GAAA) ₁₄ | (AAGG) ₄ N ₁₄ (AAAG) ₃ N ₁₂ (AAAG) ₃ N ₂₉ (AAGG) ₆ (GAAA) ₁₅ | 21 | 896 |

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| DYS385b | (AAGG) ₄ N ₁₄ (AAAG) ₃ N ₁₂ (AAAG) ₃ N ₂₉ (AAGG) ₆ (GAAA) ₁₄ | (AAGG) ₄ N ₁₄ (AAAG) ₃ N ₁₂ (AAAG) ₃ N ₂₉ (AAGG) ₆ (GAAA) ₁₅ | 23 | 1080 |
| DYS385b | (AAGG) ₄ N ₁₄ (AAAG) ₃ N ₁₂ (AAAG) ₃ N ₂₉ (AAGG) ₆ (GAAA) ₁₁ | (AAGG) ₄ N ₁₄ (AAAG) ₃ N ₁₂ (AAAG) ₃ N ₂₉ (AAGG) ₆ (GAAA) ₁₂ | 52 | 1527 |
| DYS389I | (TCTG) ₃ (TCTA) ₁₂ | (TCTG) ₃ (TCTA) ₁₁ | 32 | 12 |
| DYS389I | (TCTG) ₃ (TCTA) ₁₁ | (TCTG) ₃ (TCTA) ₁₀ | 23 | 96 |
| DYS389I | (TCTG) ₃ (TCTA) ₁₀ | (TCTG) ₃ (TCTA) ₁₁ | 23 | 207 |
| DYS389I | (TCTG) ₃ (TCTA) ₁₁ | (TCTG) ₃ (TCTA) ₁₀ | 26 | 214 |
| DYS389I | (TCTG) ₃ (TCTA) ₉ | (TCTG) ₃ (TCTA) ₁₀ | 23 | 1119 |
| DYS389I | (TCTG) ₃ (TCTA) ₁₀ | (TCTG) ₃ (TCTA) ₁₁ | Unknown | 1265 |
| DYS389I | (TCTG) ₃ (TCTA) ₁₀ | (TCTG) ₃ (TCTA) ₉ | 39 | 1700 |
| DYS389I | (TCTG) ₃ (TCTA) ₁₀ | (TCTG) ₃ (TCTA) ₁₁ | 46 | 1946 |
| DYS389I | (TCTG) ₃ (TCTA) ₁₁ | (TCTG) ₃ (TCTA) ₁₀ | 33 | 1966 |
| DYS389II | (TCTG) ₅ (TCTA) ₁₂ N ₂₈ (TCTG) ₃ (TCTA) ₁₀ | (TCTG) ₅ (TCTA) ₁₁ N ₂₈ (TCTG) ₃ (TCTA) ₁₀ | 32 | 401 |
| DYS389II | (TCTG) ₅ (TCTA) ₁₄ N ₂₈ (TCTG) ₃ (TCTA) ₁₁ | (TCTG) ₅ (TCTA) ₁₃ N ₂₈ (TCTG) ₃ (TCTA) ₁₁ | 36 | 519 |
| DYS389II | (TCTG) ₅ (TCTA) ₁₃ N ₂₈ (TCTG) ₃ (TCTA) ₁₁ | (TCTG) ₅ (TCTA) ₁₂ N ₂₈ (TCTG) ₃ (TCTA) ₁₁ | 21 | 721 |
| DYS389II | (TCTG) ₅ (TCTA) ₁₂ N ₂₈ (TCTG) ₃ (TCTA) ₁₀ | (TCTG) ₅ (TCTA) ₁₃ N ₂₈ (TCTG) ₃ (TCTA) ₁₀ | Unknown | 1221 |
| DYS389II | (TCTG) ₅ (TCTA) ₁₂ N ₂₈ (TCTG) ₃ (TCTA) ₉ | (TCTG) ₅ (TCTA) ₁₃ N ₂₈ (TCTG) ₃ (TCTA) ₉ | 24 | 1312 |
| DYS389II | (TCTG) ₅ (TCTA) ₁₃ N ₂₈ (TCTG) ₃ (TCTA) ₉ | (TCTG) ₅ (TCTA) ₁₂ N ₂₈ (TCTG) ₃ (TCTA) ₉ | 54 | 1942 |
| DYS390 | (TCTG) ₈ (TCTA) ₁₂ (TCTG) ₁ (TCTG) ₄ | (TCTG) ₈ (TCTA) ₁₁ (TCTG) ₁ (TCTG) ₄ | 30 | 975 |
| DYS390 | (TCTG) ₈ (TCTA) ₁₂ (TCTG) ₁ (TCTG) ₄ | (TCTG) ₈ (TCTA) ₁₁ (TCTG) ₁ (TCTG) ₄ | 24 | 1148 |
| DYS391 | (TCTG) ₃ (TCTA) ₁₁ | (TCTG) ₃ (TCTA) ₁₀ | 20 | 240 |
| DYS391 | (TCTG) ₃ (TCTA) ₁₀ | (TCTG) ₃ (TCTA) ₁₁ | 22 | 689 |
| DYS391 | (TCTG) ₃ (TCTA) ₁₁ | (TCTG) ₃ (TCTA) ₁₀ | 31 | 881 |
| DYS391 | (TCTG) ₃ (TCTA) ₁₁ | (TCTG) ₃ (TCTA) ₁₂ | 50 | 884 |
| DYS391 | (TCTG) ₃ (TCTA) ₁₁ | (TCTG) ₃ (TCTA) ₁₂ | 42 | 1411 |
| DYS392 | (TAT) ₁₃ | (TAT) ₁₄ | 60 | 1802 |
| DYS393 | (AGAT) ₁₄ | (AGAT) ₁₃ | 22 | 1028 |
| DYS393 | (AGAT) ₁₃ | (AGAT) ₁₄ | 42 | 1411 |
| DYS393 | (AGAT) ₁₄ | (AGAT) ₁₅ | 51 | 1852 |
| DYS425 | (TGT) ₁₃ | (TGT) ₁₂ | 31 | 1084 |

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| DYS425 | (TGT) ₁₂ | (TGT) ₁₃ | 41 | 1476 |
| DYS435 | (TGGA) ₁₁ | (TGGA) ₁₀ | 19 | 916 |
| DYS437 | (TCTA) ₈ (TCTG) ₂ (TCTA) ₄ | (TCTA) ₉ (TCTG) ₂ (TCTA) ₄ | 32 | 1025 |
| DYS437 | (TCTA) ₉ (TCTG) ₂ (TCTA) ₄ | (TCTA) ₁₀ (TCTG) ₂ (TCTA) ₄ | 53 | 1869 |
| DYS438 | (TTTTC) ₁₂ | (TTTTC) ₁₀ , (TTTTC) ₁₂ | 46 | 23 |
| DYS439 | (GATA) ₃ N ₃₂ (GATA) ₁₀ | (GATA) ₃ N ₃₂ (GATA) ₁₁ | 24 | 516 |
| DYS439 | (GATA) ₃ N ₃₂ (GATA) ₁₃ | (GATA) ₃ N ₃₂ (GATA) ₁₄ | 36 | 617 |
| DYS439 | (GATA) ₃ N ₃₂ (GATA) ₁₃ | (GATA) ₃ N ₃₂ (GATA) ₁₂ | 23 | 620 |
| DYS439 | (GATA) ₃ N ₃₂ (GATA) ₁₃ | (GATA) ₃ N ₃₂ (GATA) ₁₂ | 40 | 1204 |
| DYS439 | (GATA) ₃ N ₃₂ (GATA) ₁₄ | (GATA) ₃ N ₃₂ (GATA) ₁₃ | 37 | 1211 |
| DYS439 | (GATA) ₃ N ₃₂ (GATA) ₁₃ | (GATA) ₃ N ₃₂ (GATA) ₁₂ | 21 | 1463 |
| DYS441 | (TTCC) ₁₄ | (TTCC) ₁₅ | 38 | 589 |
| DYS442 | (GATA) ₁₃ (GACA) ₃ | (GATA) ₁₂ (GACA) ₃ | 32 | 213 |
| DYS442 | (GATA) ₁₃ (GACA) ₃ | (GATA) ₁₂ (GACA) ₃ | 26 | 354 |
| DYS442 | (GATA) ₁₃ (GACA) ₃ | (GATA) ₁₂ (GACA) ₃ | 45 | 409 |
| DYS442 | (GATA) ₁₅ (GACA) ₃ | (GATA) ₁₄ (GACA) ₃ | 28 | 425 |
| DYS442 | (GATA) ₁₆ (GACA) ₃ | (GATA) ₁₅ (GACA) ₃ | 30 | 533 |
| DYS442 | (GATA) ₁₄ (GACA) ₃ | (GATA) ₁₃ (GACA) ₃ | 26 | 775 |
| DYS442 | (GATA) ₁₅ (GACA) ₃ | (GATA) ₁₄ (GACA) ₃ | 27 | 953 |
| DYS442 | (GATA) ₁₄ (GACA) ₃ | (GATA) ₁₅ (GACA) ₃ | 30 | 1181 |
| DYS442 | (GATA) ₁₃ (GACA) ₃ | (GATA) ₁₂ (GACA) ₃ | 16 | 1238 |
| DYS442 | (GATA) ₁₂ (GACA) ₃ | (GATA) ₁₁ (GACA) ₃ | 39 | 1239 |
| DYS442 | (GATA) ₁₄ (GACA) ₃ | (GATA) ₁₃ (GACA) ₃ | Unknown | 1245 |
| DYS442 | (GATA) ₁₂ (GACA) ₃ | (GATA) ₁₃ (GACA) ₃ | 28 | 1537 |
| DYS442 | (GATA) ₁₂ (GACA) ₃ | (GATA) ₁₁ (GACA) ₃ | 73 | 1884 |
| DYS442 | (GATA) ₁₃ (GACA) ₃ | (GATA) ₁₂ (GACA) ₃ | 51 | 1888 |
| DYS443 | (TTCC) ₁₅ (CTT) ₃ | (TTCC) ₁₄ (CTT) ₃ | 18 | 69 |
| DYS443 | (TTCC) ₁₄ (CTT) ₃ | (TTCC) ₁₅ (CTT) ₃ | 35 | 97 |
| DYS443 | (TTCC) ₁₄ (CTT) ₃ | (TTCC) ₁₅ (CTT) ₃ | 28 | 473 |
| DYS444 | (TAGA) ₁₂ | (TAGA) ₁₁ | 46 | 23 |

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| DYS444 | (TAGA) ₁₃ | (TAGA) ₁₂ | 20 | 73 |
| DYS444 | (TAGA) ₁₃ | (TAGA) ₁₂ | 30 | 448 |
| DYS444 | (TAGA) ₁₄ | (TAGA) ₁₅ | 25 | 673 |
| DYS444 | (TAGA) ₁₅ | (TAGA) ₁₄ | 30 | 1067 |
| DYS444 | (TAGA) ₁₅ | (TAGA) ₁₄ | 22 | 1131 |
| DYS444 | (TAGA) ₁₅ | (TAGA) ₁₄ | 28 | 1294 |
| DYS444 | (TAGA) ₁₂ | (TAGA) ₁₃ | 33 | 1680 |
| DYS444 | (TAGA) ₁₃ | (TAGA) ₁₂ | 51 | 1813 |
| DYS445 | (TTTA) ₁₂ | (TTTA) ₁₁ | 57 | 464 |
| DYS445 | (TTTA) ₁₁ | (TTTA) ₁₂ | 22 | 886 |
| DYS445 | (TTTA) ₁₃ | (TTTA) ₁₄ | 21 | 1448 |
| DYS446 | (TCTCT) ₁₃ | (TCTCT) ₁₄ | 41 | 313 |
| DYS446 | (TCTCT) ₁₁ | (TCTCT) ₁₀ | 51 | 1442 |
| DYS446 | (TCTCT) ₁₂ | (TCTCT) ₁₁ | 55 | 1844 |
| DYS446 | (TCTCT) ₁₂ | (TCTCT) ₁₃ | 53 | 1893 |
| DYS447 | (TTATA) ₆ (TTATT) ₁ (TTATA) ₈ (TTATT) ₁ (TTATA) ₇ | (TTATA) ₆ (TTATT) ₁ (TTATA) ₉ (TTATT) ₁ (TTATA) ₇ | 33 | 690 |
| DYS447 | (TTATA) ₇ (TTATT) ₁ (TTATA) ₁₀ (TTATT) ₁ (TTATA) ₇ | (TTATA) ₇ (TTATT) ₁ (TTATA) ₉ (TTATT) ₁ (TTATA) ₇ | 47 | 1302 |
| DYS447 | (TTATA) ₆ (TTATT) ₁ (TTATA) ₉ (TTATT) ₁ (TTATA) ₉ | (TTATA) ₆ (TTATT) ₁ (TTATA) ₉ (TTATT) ₁ (TTATA) ₈ | 56 | 1677 |
| DYS449 | (TTCT) ₁₅ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₆ | (TTCT) ₁₅ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₅ | 29 | 78 |
| DYS449 | (TTCT) ₁₅ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₉ | (TTCT) ₁₄ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₉ | 27 | 170 |
| DYS449 | (TTCT) ₁₆ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₇ | (TTCT) ₁₇ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₇ | 23 | 251 |
| DYS449 | (TTCT) ₁₆ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₇ | (TTCT) ₁₅ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₇ | 38 | 449 |
| DYS449 | (TTCT) ₁₅ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₈ | (TTCT) ₁₅ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₉ | 39 | 492 |
| DYS449 | (TTCT) ₁₇ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₃ | (TTCT) ₁₇ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₄ | 35 | 531 |
| DYS449 | (TTCT) ₁₄ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₄ | (TTCT) ₁₄ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₅ | 34 | 568 |
| DYS449 | (TTCT) ₁₆ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₆ | (TTCT) ₁₆ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₇ | 21 | 786 |
| DYS449 | (TTCT) ₁₆ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₇ | (TTCT) ₁₆ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₈ | 25 | 840 |
| DYS449 | (TTCT) ₁₅ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₃ | (TTCT) ₁₆ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₃ | 22 | 894 |
| DYS449 | (TTCT) ₁₅ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₈ | (TTCT) ₁₅ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₇ | 37 | 904 |
| DYS449 | (TTCT) ₁₅ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₇ | (TTCT) ₁₅ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₆ | 33 | 966 |

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| DYS449 | (TTCT) ₁₆ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₆ | (TTCT) ₁₅ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₆ | 24 | 1167 |
| DYS449 | (TTCT) ₁₅ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₄ | (TTCT) ₁₆ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₄ | 22 | 1349 |
| DYS449 | (TTCT) ₁₅ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₄ | (TTCT) ₁₆ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₄ | 45 | 1364 |
| DYS449 | (TTCT) ₁₈ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₆ | (TTCT) ₁₉ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₆ | 37 | 1418 |
| DYS449 | (TTCT) ₁₄ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₄ | (TTCT) ₁₄ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₅ | 21 | 1505 |
| DYS449 | (TTCT) ₁₅ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₅ | (TTCT) ₁₄ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₅ | 22 | 1526 |
| DYS449 | (TTCT) ₁₄ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₅ | (TTCT) ₁₄ N ₂₂ (TTCT) ₃ N ₁₂ (TTCT) ₁₆ | 54 | 1845 |
| DYS450 | (TTTTA) ₉ N ₁₂ (TTTTA) ₃ | (TTTTA) ₈ N ₁₂ (TTTTA) ₃ | 44 | 1619 |
| DYS452 | (TATAC) ₁₂ [(CATAC) ₁ (TATAC) ₁] ₂ N ₂₀ (TATAC) ₃ (CATAC) ₁ (TATAC) ₃ | (TATAC) ₁₃ [(CATAC) ₁ (TATAC) ₁] ₂ N ₂₀ (TATAC) ₃ (CATAC) ₁ (TATAC) ₃ | 17 | 967 |
| DYS452 | (TATAC) ₁₁ [(CATAC) ₁ (TATAC) ₁] ₂ N ₂₀ (TATAC) ₃ (CATAC) ₁ (TATAC) ₃ | (TATAC) ₁₂ [(CATAC) ₁ (TATAC) ₁] ₂ N ₂₀ (TATAC) ₃ (CATAC) ₁ (TATAC) ₃ | 26 | 971 |
| DYS452 | (TATAC) ₁₁ [(CATAC) ₁ (TATAC) ₁] ₂ N ₂₀ (TATAC) ₃ (CATAC) ₁ (TATAC) ₃ | (TATAC) ₁₀ [(CATAC) ₁ (TATAC) ₁] ₂ N ₂₀ (TATAC) ₃ (CATAC) ₁ (TATAC) ₃ | 33 | 1046 |
| DYS452 | (TATAC) ₁₀ [(CATAC) ₁ (TATAC) ₁] ₂ N ₂₀ (TATAC) ₃ (CATAC) ₁ (TATAC) ₃ | (TATAC) ₉ [(CATAC) ₁ (TATAC) ₁] ₂ N ₂₀ (TATAC) ₃ (CATAC) ₁ (TATAC) ₃ | 41 | 1453 |
| DYS452 | (TATAC) ₈ [(CATAC) ₁ (TATAC) ₁] ₄ N ₂₀ (TATAC) ₃ (CATAC) ₁ (TATAC) ₃ | (TATAC) ₈ [(CATAC) ₁ (TATAC) ₁] ₃ N ₂₀ (TATAC) ₃ (CATAC) ₁ (TATAC) ₃ | 55 | 1858 |
| DYS456 | (AGAT) ₁₅ | (AGAT) ₁₆ | 34 | 308 |
| DYS456 | (AGAT) ₁₆ | (AGAT) ₁₇ | 32 | 401 |
| DYS456 | (AGAT) ₁₅ | (AGAT) ₁₆ | 28 | 525 |
| DYS456 | (AGAT) ₁₆ | (AGAT) ₁₇ | 24 | 560 |
| DYS456 | (AGAT) ₁₅ | (AGAT) ₁₆ | 36 | 830 |
| DYS456 | (AGAT) ₁₇ | (AGAT) ₁₆ | 20 | 1037 |
| DYS456 | (AGAT) ₁₇ | (AGAT) ₁₆ | Unknown | 1333 |
| DYS456 | (AGAT) ₁₆ | (AGAT) ₁₇ | 29 | 1790 |
| DYS458 | (GAAA) ₁₈ | (GAAA) ₁₇ | 25 | 307 |
| DYS458 | (GAAA) ₁₈ | (GAAA) ₁₉ | 41 | 313 |
| DYS458 | (GAAA) ₁₇ | (GAAA) ₁₈ | 28 | 466 |
| DYS458 | (GAAA) ₁₇ | (GAAA) ₁₈ | 27 | 734 |
| DYS458 | (GAAA) ₁₆ | (GAAA) ₁₅ | 24 | 771 |
| DYS458 | (GAAA) ₁₇ | (GAAA) ₁₈ | 29 | 826 |

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| DYS458 | (GAAA) ₁₅ | (GAAA) ₁₆ | Unknown | 1063 |
| DYS458 | (GAAA) ₁₆ | (GAAA) ₁₅ | 19 | 1132 |
| DYS458 | (GAAA) ₁₇ | (GAAA) ₁₆ | 34 | 1428 |
| DYS458 | (GAAA) ₁₆ | (GAAA) ₁₇ | 37 | 1454 |
| DYS458 | (GAAA) ₁₇ | (GAAA) ₁₆ | 19 | 1508 |
| DYS458 | (GAAA) ₁₇ | (GAAA) ₁₆ | Unknown | 1613 |
| DYS458 | (GAAA) ₁₈ | (GAAA) ₁₇ | 64 | 1912 |
| DYS458 | (GAAA) ₁₈ | (GAAA) ₁₉ | 65 | 1920 |
| DYS459 | (ATTT) ₁₀ | (ATTT) ₉ | 29 | 246 |
| DYS459 | (ATTT) ₉ | (ATTT) ₁₀ | 26 | 573 |
| DYS459 | (ATTT) ₁₀ | (ATTT) ₉ | 43 | 928 |
| DYS459 | (ATTT) ₁₀ | (ATTT) ₁₁ | 27 | 1195 |
| DYS460 | (TAGA) ₁₃ | (TAGA) ₁₂ | 27 | 41 |
| DYS460 | (TAGA) ₉ | (TAGA) ₁₀ | 29 | 481 |
| DYS460 | (TAGA) ₁₁ | (TAGA) ₁₀ | 35 | 522 |
| DYS460 | (TAGA) ₁₂ | (TAGA) ₁₁ | 24 | 560 |
| DYS460 | (TAGA) ₁₁ | (TAGA) ₁₀ | 27 | 777 |
| DYS460 | (TAGA) ₁₁ | (TAGA) ₁₀ | 30 | 1062 |
| DYS460 | (TAGA) ₁₂ | (TAGA) ₁₁ | 33 | 1112 |
| DYS460 | (TAGA) ₁₁ | (TAGA) ₁₀ | 29 | 1601 |
| DYS460 | (TAGA) ₁₀ | (TAGA) ₁₁ | 21 | 1728 |
| DYS460 | (TAGA) ₁₃ | (TAGA) ₁₂ | 28 | 1734 |
| DYS461 | (TAGA) ₁₁ | (TAGA) ₁₀ | 23 | 1676 |
| DYS462 | (CATA) ₁₁ | (CATA) ₁₀ | 18 | 692 |
| DYS462 | (CATA) ₁₃ | (CATA) ₁₂ | 30 | 1425 |
| DYS462 | (CATA) ₁₁ | (CATA) ₁₀ | 39 | 1502 |
| DYS462 | (CATA) ₁₁ | (CATA) ₁₂ | 64 | 1912 |
| DYS463 | (AAAGG) ₆ (AAGGG) ₁₅ | (AAAGG) ₆ (AAGGG) ₁₆ | 36 | 327 |
| DYS463 | (AAAGG) ₆ (AAGGG) ₁₃ | (AAAGG) ₆ (AAGGG) ₁₄ | 54 | 1447 |
| DYS464 | (CCTT) ₁₆ N ₄₆ (CCTT) ₃ N ₈ (CCTT) ₄ | (CCTT) ₁₇ N ₄₆ (CCTT) ₃ N ₈ (CCTT) ₄ | 25 | 4 |

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| DYS464 | (CCTT) ₁₅ N ₄₆ (CCTT) ₃ N ₈ (CCTT) ₄ | (CCTT) ₁₄ N ₄₆ (CCTT) ₃ N ₈ (CCTT) ₄ | 18 | 470 |
| DYS464 | (CCTT) ₁₇ N ₄₆ (CCTT) ₃ N ₈ (CCTT) ₄ | (CCTT) ₁₆ N ₄₆ (CCTT) ₃ N ₈ (CCTT) ₄ | 25 | 512 |
| DYS464 | (CCTT) ₁₆ N ₄₆ (CCTT) ₃ N ₈ (CCTT) ₄ | (CCTT) ₁₉ N ₄₆ (CCTT) ₃ N ₈ (CCTT) ₄ | 53 | 760 |
| DYS464 | (CCTT) ₁₃ N ₄₆ (CCTT) ₃ N ₈ (CCTT) ₄ | (CCTT) ₁₄ N ₄₆ (CCTT) ₃ N ₈ (CCTT) ₄ | 18 | 847 |
| DYS464 | (CCTT) ₁₈ N ₄₆ (CCTT) ₃ N ₈ (CCTT) ₄ | (CCTT) ₁₆ N ₄₆ (CCTT) ₃ N ₈ (CCTT) ₄ | 19 | 900 |
| DYS464 | (CCTT) ₁₆ N ₄₆ (CCTT) ₃ N ₈ (CCTT) ₄ | (CCTT) ₁₅ N ₄₆ (CCTT) ₃ N ₈ (CCTT) ₄ | 20 | 1185 |
| DYS464 | (CCTT) ₁₇ N ₄₆ (CCTT) ₃ N ₈ (CCTT) ₄ | (CCTT) ₁₆ N ₄₆ (CCTT) ₃ N ₈ (CCTT) ₄ | 31 | 1339 |
| DYS464 | (CCTT) ₁₅ N ₄₆ (CCTT) ₃ N ₈ (CCTT) ₄ | (CCTT) ₁₆ N ₄₆ (CCTT) ₃ N ₈ (CCTT) ₄ | 22 | 1526 |
| DYS464 | (CCTT) ₁₈ N ₄₆ (CCTT) ₃ N ₈ (CCTT) ₄ | (CCTT) ₁₉ N ₄₆ (CCTT) ₃ N ₈ (CCTT) ₄ | 43 | 1540 |
| DYS464 | (CCTT) ₁₇ N ₄₆ (CCTT) ₃ N ₈ (CCTT) ₄ | (CCTT) ₁₆ N ₄₆ (CCTT) ₃ N ₈ (CCTT) ₄ | 19 | 1784 |
| DYS464 | (CCTT) ₁₈ N ₄₆ (CCTT) ₃ N ₈ (CCTT) ₄ | (CCTT) ₁₇ N ₄₆ (CCTT) ₃ N ₈ (CCTT) ₄ | 56 | 1892 |
| DYS468 | (CTG) ₄ N ₄₄ (CCT) ₃ N ₄₀ (CTT) ₃ N ₃₅ (CCT) ₄ N ₈ (CTC) ₄ (CTT) ₈ (ATTCAT) ₈ | (CTG) ₄ N ₄₄ (CCT) ₃ N ₄₀ (CTT) ₃ N ₃₅ (CCT) ₄ N ₈ (CTC) ₄ (CTT) ₉ (ATTCAT) ₈ | 29 | 1743 |
| DYS468 | (CTG) ₄ N ₄₄ (CCT) ₃ N ₄₀ (CTT) ₃ N ₃₅ (CCT) ₄ N ₈ (CTC) ₄ (CTT) ₉ (ATTCAT) ₉ | (CTG) ₄ N ₄₄ (CCT) ₃ N ₄₀ (CTT) ₃ N ₃₅ (CCT) ₄ N ₈ (CTC) ₄ (CTT) ₉ (ATTCAT) ₈ | 60 | 1802 |
| DYS469 | (CTT) ₃ N ₃₉ (CTT) ₄ (GTT) ₁ (CTT) ₂₀ T(CTT) ₃ N ₁₇ (CTT) ₅ N ₃₇ (CTT) ₃ N ₁₂ (CTT) ₄ N ₁₂ (CTT) ₃ N ₁₂ (CTT) ₅ (CCT) ₄ N ₉ (CTT) ₃ (CCT) ₃ | (CTT) ₃ N ₃₉ (CTT) ₄ (GTT) ₁ (CTT) ₂₁ T(CTT) ₃ N ₁₇ (CTT) ₅ N ₃₇ (CTT) ₃ N ₁₂ (CTT) ₄ N ₁₂ (CTT) ₃ N ₁₂ (CTT) ₅ (CCT) ₄ N ₉ (CTT) ₃ (CCT) ₃ | 24 | 107 |
| DYS469 | (CTT) ₃ N ₃₉ (CTT) ₄ (GTT) ₁ (CTT) ₁₅ T(CTT) ₃ N ₁₇ (CTT) ₅ N ₃₇ (CTT) ₃ N ₁₂ (CTT) ₄ N ₁₂ (CTT) ₃ N ₁₂ (CTT) ₅ (CCT) ₄ N ₉ (CTT) ₃ (CCT) ₃ | (CTT) ₃ N ₃₉ (CTT) ₄ (GTT) ₁ (CTT) ₁₆ T(CTT) ₃ N ₁₇ (CTT) ₅ N ₃₇ (CTT) ₃ N ₁₂ (CTT) ₄ N ₁₂ (CTT) ₃ N ₁₂ (CTT) ₅ (CCT) ₄ N ₉ (CTT) ₃ (CCT) ₃ | 21 | 769 |
| DYS469 | (CTT) ₃ N ₃₉ (CTT) ₄ (GTT) ₁ (CTT) ₁₅ T(CTT) ₃ N ₁₇ (CTT) ₅ N ₃₇ (CTT) ₃ N ₁₂ (CTT) ₄ N ₁₂ (CTT) ₃ N ₁₂ (CTT) ₅ (CCT) ₄ N ₉ (CTT) ₃ (CCT) ₃ | (CTT) ₃ N ₃₉ (CTT) ₄ (GTT) ₁ (CTT) ₁₆ T(CTT) ₃ N ₁₇ (CTT) ₅ N ₃₇ (CTT) ₃ N ₁₂ (CTT) ₄ N ₁₂ (CTT) ₃ N ₁₂ (CTT) ₅ (CCT) ₄ N ₉ (CTT) ₃ (CCT) ₃ | 29 | 1491 |
| DYS469 | (CTT) ₃ N ₃₉ (CTT) ₄ (GTT) ₁ (CTT) ₁₆ T(CTT) ₃ N ₁₇ (CTT) ₅ N ₃₇ (CTT) ₃ N ₁₂ (CTT) ₄ N ₁₂ (CTT) ₃ N ₁₂ (CTT) ₅ (CCT) ₄ N ₉ (CTT) ₃ (CCT) ₃ | (CTT) ₃ N ₃₉ (CTT) ₄ (GTT) ₁ (CTT) ₁₅ T(CTT) ₃ N ₁₇ (CTT) ₅ N ₃₇ (CTT) ₃ N ₁₂ (CTT) ₄ N ₁₂ (CTT) ₃ N ₁₂ (CTT) ₅ (CCT) ₄ N ₉ (CTT) ₃ (CCT) ₃ | 29 | 1693 |
| DYS476 | (TGA) ₁₁ | (TGA) ₁₂ | 59 | 1867 |
| DYS481 | (CTT) ₂₈ | (CTT) ₂₉ | 30 | 99 |
| DYS481 | (CTT) ₂₂ | (CTT) ₂₃ | 32 | 655 |
| DYS481 | (CTT) ₃₀ | (CTT) ₂₉ | 24 | 778 |
| DYS481 | (CTT) ₃₂ | (CTT) ₃₁ | 17 | 845 |
| DYS481 | (CTT) ₃₁ | (CTT) ₃₀ | 24 | 1370 |
| DYS481 | (CTT) ₂₃ | (CTT) ₂₁ | 42 | 1411 |

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| DYS481 | (CTT) ₂₅ | (CTT) ₂₄ | 36 | 1672 |
| DYS481 | (CTT) ₂₂ | (CTT) ₂₃ | 54 | 1895 |
| DYS484 | (AAT) ₁₃ N ₁₂ (AAT) ₃ (TAT) ₃ | (AAT) ₁₁ N ₁₂ (AAT) ₃ (TAT) ₃ | 22 | 45 |
| DYS484 | (AAT) ₁₃ N ₁₂ (AAT) ₃ (TAT) ₃ | (AAT) ₁₄ N ₁₂ (AAT) ₃ (TAT) ₃ | 35 | 875 |
| DYS484 | (AAT) ₁₂ N ₁₂ (AAT) ₃ (TAT) ₃ | (AAT) ₁₃ N ₁₂ (AAT) ₃ (TAT) ₃ | 25 | 1213 |
| DYS484 | (AAT) ₁₂ N ₁₂ (AAT) ₃ (TAT) ₃ | (AAT) ₁₀ N ₁₂ (AAT) ₃ (TAT) ₃ | 26 | 1653 |
| DYS487 | (AAT) ₁₃ | (AAT) ₁₄ | 21 | 172 |
| DYS487 | (AAT) ₁₄ | (AAT) ₁₃ | 28 | 1410 |
| DYS495 | (AAT) ₁₅ | (AAT) ₁₆ | 26 | 775 |
| DYS495 | (AAT) ₁₅ | (AAT) ₁₆ | 30 | 1274 |
| DYS495 | (AAT) ₁₇ | (AAT) ₁₆ | 31 | 1596 |
| DYS497 | (TTA) ₁₅ | (TTA) ₁₄ | 25 | 58 |
| DYS497 | (TTA) ₁₅ | (TTA) ₁₆ | 40 | 305 |
| DYS504 | (CCTT) ₁₇ N ₇ (CCCT) ₃ | (CCTT) ₁₆ N ₇ (CCCT) ₃ | 25 | 275 |
| DYS504 | (CCTT) ₁₉ N ₇ (CCCT) ₃ | (CCTT) ₁₈ N ₇ (CCCT) ₃ | Unknown | 1223 |
| DYS504 | (CCTT) ₁₇ N ₇ (CCCT) ₃ | (CCTT) ₁₈ N ₇ (CCCT) ₃ | 39 | 1502 |
| DYS504 | (CCTT) ₁₈ N ₇ (CCCT) ₃ | (CCTT) ₁₇ N ₇ (CCCT) ₃ | 30 | 1796 |
| DYS504 | (CCTT) ₁₈ N ₇ (CCCT) ₃ | (CCTT) ₁₇ N ₇ (CCCT) ₃ | 54 | 1895 |
| DYS505 | (TCCT) ₁₃ | (TCCT) ₁₂ | 64 | 951 |
| DYS505 | (TCCT) ₁₂ | (TCCT) ₁₃ | 25 | 1070 |
| DYS508 | (TATC) ₁₀ | (TATC) ₁₁ | 32 | 1369 |
| DYS508 | (TATC) ₁₁ | (TATC) ₁₂ | 21 | 1635 |
| DYS508 | (TATC) ₁₁ | (TATC) ₁₀ | 20 | 1862 |
| DYS508 | (TATC) ₁₄ | (TATC) ₁₃ | 67 | 1954 |
| DYS509 | (AAAT) ₁₀ (AATAA) ₁ (AAAT) ₃ | (AAAT) ₉ (AATAA) ₁ (AAAT) ₃ | 18 | 680 |
| DYS510 | (GATA) ₃ N ₁₂ (GATA) ₁₂ N ₁₃ (GGAT) ₄ N ₉ (GATA) ₃ | (GATA) ₃ N ₁₂ (GATA) ₁₃ N ₁₃ (GGAT) ₄ N ₉ (GATA) ₃ | 24 | 17 |
| DYS510 | (GATA) ₃ N ₁₂ (GATA) ₁₂ N ₁₃ (GGAT) ₄ N ₉ (GATA) ₃ | (GATA) ₃ N ₁₂ (GATA) ₁₁ N ₁₃ (GGAT) ₄ N ₉ (GATA) ₃ | 64 | 166 |
| DYS510 | (GATA) ₃ N ₁₂ (GATA) ₁₂ N ₁₃ (GGAT) ₄ N ₉ (GATA) ₃ | (GATA) ₃ N ₁₂ (GATA) ₁₁ N ₁₃ (GGAT) ₄ N ₉ (GATA) ₃ | 25 | 185 |
| DYS510 | (GATA) ₃ N ₁₂ (GATA) ₁₅ N ₁₃ (GGAT) ₄ N ₉ (GATA) ₃ | (GATA) ₃ N ₁₂ (GATA) ₁₄ N ₁₃ (GGAT) ₄ N ₉ (GATA) ₃ | 19 | 761 |
| DYS510 | (GATA) ₃ N ₁₂ (GATA) ₁₁ N ₁₃ (GGAT) ₄ N ₉ (GATA) ₃ | (GATA) ₃ N ₁₂ (GATA) ₁₀ N ₁₃ (GGAT) ₄ N ₉ (GATA) ₃ | 19 | 916 |

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| DYS510 | (GATA) ₃ N ₁₂ (GATA) ₁₄ N ₁₃ (GGAT) ₄ N ₉ (GATA) ₃ | (GATA) ₃ N ₁₂ (GATA) ₁₅ N ₁₃ (GGAT) ₄ N ₉ (GATA) ₃ | 43 | 928 |
| DYS510 | (GATA) ₃ N ₁₂ (GATA) ₁₂ N ₁₃ (GGAT) ₄ N ₉ (GATA) ₃ | (GATA) ₃ N ₁₂ (GATA) ₁₁ N ₁₃ (GGAT) ₄ N ₉ (GATA) ₃ | 33 | 1112 |
| DYS510 | (GATA) ₃ N ₁₂ (GATA) ₁₃ N ₁₃ (GGAT) ₄ N ₉ (GATA) ₃ | (GATA) ₃ N ₁₂ (GATA) ₁₂ N ₁₃ (GGAT) ₄ N ₉ (GATA) ₃ | 34 | 1118 |
| DYS510 | (GATA) ₃ N ₁₂ (GATA) ₁₄ N ₁₃ (GGAT) ₄ N ₉ (GATA) ₃ | (GATA) ₃ N ₁₂ (GATA) ₁₅ N ₁₃ (GGAT) ₄ N ₉ (GATA) ₃ | 45 | 1364 |
| DYS510 | (GATA) ₃ N ₁₂ (GATA) ₁₂ N ₁₃ (GGAT) ₄ N ₉ (GATA) ₃ | (GATA) ₃ N ₁₂ (GATA) ₁₃ N ₁₃ (GGAT) ₄ N ₉ (GATA) ₃ | 28 | 1758 |
| DYS511 | (GATA) ₁₃ | (GATA) ₁₂ | 22 | 418 |
| DYS511 | (GATA) ₁₁ | (GATA) ₁₂ | 52 | 1804 |
| DYS513 | (TCTA) ₄ (TCCA) ₁ (TATC) ₃ (CGTA) ₁ (TCTA) ₁₂ | (TCTA) ₄ (TCCA) ₁ (TATC) ₃ (CGTA) ₁ (TCTA) ₁₃ | 33 | 29 |
| DYS513 | (TCTA) ₄ (TCCA) ₁ (TATC) ₃ (CGTA) ₁ (TCTA) ₁₄ | (TCTA) ₄ (TCCA) ₁ (TATC) ₃ (CGTA) ₁ (TCTA) ₁₃ | 23 | 134 |
| DYS513 | (TCTA) ₄ (TCCA) ₁ (TATC) ₃ (CGTA) ₁ (TCTA) ₁₃ | (TCTA) ₄ (TCCA) ₁ (TATC) ₃ (CGTA) ₁ (TCTA) ₁₂ | 33 | 187 |
| DYS513 | (TCTA) ₄ (TCCA) ₁ (TATC) ₃ (CGTA) ₁ (TCTA) ₁₃ | (TCTA) ₄ (TCCA) ₁ (TATC) ₃ (CGTA) ₁ (TCTA) ₁₄ | 30 | 1181 |
| DYS513 | (TCTA) ₄ (TCCA) ₁ (TATC) ₃ (CGTA) ₁ (TCTA) ₁₃ | (TCTA) ₄ (TCCA) ₁ (TATC) ₃ (CGTA) ₁ (TCTA) ₁₄ | 21 | 1216 |
| DYS513 | (TCTA) ₄ (TCCA) ₁ (TATC) ₃ (CGTA) ₁ (TCTA) ₁₃ | (TCTA) ₄ (TCCA) ₁ (TATC) ₃ (CGTA) ₁ (TCTA) ₁₄ | Unknown | 1308 |
| DYS513 | (TCTA) ₄ (TCCA) ₁ (TATC) ₃ (CGTA) ₁ (TCTA) ₁₂ | (TCTA) ₄ (TCCA) ₁ (TATC) ₃ (CGTA) ₁ (TCTA) ₁₁ | 22 | 1323 |
| DYS513 | (TCTA) ₄ (TCCA) ₁ (TATC) ₃ (CGTA) ₁ (TCTA) ₁₂ | (TCTA) ₄ (TCCA) ₁ (TATC) ₃ (CGTA) ₁ (TCTA) ₁₃ | 36 | 1421 |
| DYS513 | (TCTA) ₄ (TCCA) ₁ (TATC) ₃ (CGTA) ₁ (TCTA) ₁₃ | (TCTA) ₄ (TCCA) ₁ (TATC) ₃ (CGTA) ₁ (TCTA) ₁₂ | 36 | 1672 |
| DYS513 | (TCTA) ₄ (TCCA) ₁ (TATC) ₃ (CGTA) ₁ (TCTA) ₁₁ | (TCTA) ₄ (TCCA) ₁ (TATC) ₃ (CGTA) ₁ (TCTA) ₁₂ | 60 | 1695 |
| DYS516 | (TTCT) ₄ N ₃₀ (TTCT) ₁₆ | (TTCT) ₄ N ₃₀ (TTCT) ₁₅ | 28 | 106 |
| DYS516 | (TTCT) ₄ N ₃₀ (TTCT) ₁₄ | (TTCT) ₄ N ₃₀ (TTCT) ₁₃ | 37 | 904 |
| DYS516 | (TTCT) ₄ N ₃₀ (TTCT) ₁₂ | (TTCT) ₄ N ₃₀ (TTCT) ₁₃ | 44 | 973 |
| DYS516 | (TTCT) ₄ N ₃₀ (TTCT) ₁₅ | (TTCT) ₄ N ₃₀ (TTCT) ₁₆ | 34 | 1030 |
| DYS516 | (TTCT) ₄ N ₃₀ (TTCT) ₁₃ | (TTCT) ₄ N ₃₀ (TTCT) ₁₄ | Unknown | 1241 |
| DYS516 | (TTCT) ₄ N ₃₀ (TTCT) ₁₂ | (TTCT) ₄ N ₃₀ (TTCT) ₁₃ | 38 | 1524 |
| DYS516 | (TTCT) ₄ N ₃₀ (TTCT) ₁₄ | (TTCT) ₄ N ₃₀ (TTCT) ₁₅ | 42 | 1628 |
| DYS516 | (TTCT) ₄ N ₃₀ (TTCT) ₁₅ | (TTCT) ₄ N ₃₀ (TTCT) ₁₆ | 22 | 1778 |
| DYS516 | (TTCT) ₄ N ₃₀ (TTCT) ₁₅ | (TTCT) ₄ N ₃₀ (TTCT) ₁₆ | 24 | 1782 |
| DYS516 | (TTCT) ₄ N ₃₀ (TTCT) ₁₂ | (TTCT) ₄ N ₃₀ (TTCT) ₁₁ | 53 | 1869 |
| DYS516 | (TTCT) ₄ N ₃₀ (TTCT) ₁₇ | (TTCT) ₄ N ₃₀ (TTCT) ₁₅ | 50 | 1881 |
| DYS517 | (AAAG) ₁₆ N ₈ (AAAG) ₃ | (AAAG) ₁₅ N ₈ (AAAG) ₃ | 26 | 802 |
| DYS517 | (AAAG) ₁₄ N ₈ (AAAG) ₃ | (AAAG) ₁₅ N ₈ (AAAG) ₃ | 30 | 1796 |

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| DYS517 | (AAAG) ₁₆ N ₈ (AAAG) ₃ | (AAAG) ₁₅ N ₈ (AAAG) ₃ | 63 | 1807 |
| DYS517 | (AAAG) ₁₄ N ₈ (AAAG) ₃ | (AAAG) ₁₅ N ₈ (AAAG) ₃ | 54 | 1860 |
| DYS517 | (AAAG) ₁₄ N ₈ (AAAG) ₃ | (AAAG) ₁₅ N ₈ (AAAG) ₃ | 53 | 1869 |
| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₆ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₇ N ₂₇ (AAGG) ₄ | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₅ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₇ N ₂₇ (AAGG) ₄ | 33 | 29 |
| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₄ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₄ N ₂₇ (AAGG) ₄ | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₄ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₅ N ₂₇ (AAGG) ₄ | 32 | 56 |
| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₇ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₂ N ₂₇ (AAGG) ₄ | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₆ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₂ N ₂₇ (AAGG) ₄ | 48 | 74 |
| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₅ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₅ N ₂₇ (AAGG) ₄ | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₄ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₅ N ₂₇ (AAGG) ₄ | 30 | 87 |
| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₈ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₆ N ₂₇ (AAGG) ₄ | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₇ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₆ N ₂₇ (AAGG) ₄ | 34 | 88 |
| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₇ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₃ N ₂₇ (AAGG) ₄ | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₆ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₃ N ₂₇ (AAGG) ₄ | 29 | 89 |
| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₉ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₇ N ₂₇ (AAGG) ₄ | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₈ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₇ N ₂₇ (AAGG) ₄ | 50 | 270 |
| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₈ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₅ N ₂₇ (AAGG) ₄ | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₉ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₅ N ₂₇ (AAGG) ₄ | 25 | 426 |
| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₂₂ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₃ N ₂₇ (AAGG) ₄ | (AAAG) ₃ (GAAG) ₁ (AAAG) ₂₁ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₃ N ₂₇ (AAGG) ₄ | 26 | 433 |
| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₅ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₂ N ₂₇ (AAGG) ₄ | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₅ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₁ N ₂₇ (AAGG) ₄ | 28 | 525 |
| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₅ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₇ N ₂₇ (AAGG) ₄ | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₅ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₆ N ₂₇ (AAGG) ₄ | 24 | 571 |
| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₈ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₄ N ₂₇ (AAGG) ₄ | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₇ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₄ N ₂₇ (AAGG) ₄ | 21 | 593 |
| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₆ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₁ N ₂₇ (AAGG) ₄ | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₅ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₇ N ₂₇ (AAGG) ₄ | 23 | 687 |
| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₆ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₄ N ₂₇ (AAGG) ₄ | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₆ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₅ N ₂₇ (AAGG) ₄ | 20 | 703 |
| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₇ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₅ N ₂₇ (AAGG) ₄ | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₆ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₅ N ₂₇ (AAGG) ₄ | 20 | 741 |
| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₅ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₆ N ₂₇ (AAGG) ₄ | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₅ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₇ N ₂₇ (AAGG) ₄ | 22 | 747 |
| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₅ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₆ N ₂₇ (AAGG) ₄ | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₆ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₆ N ₂₇ (AAGG) ₄ | 15 | 763 |

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| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₇ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₄ N ₂₇ (AAGG) ₄ | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₆ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₆ N ₂₇ (AAGG) ₄ | 22 | 817 |
| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₆ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₈ N ₂₇ (AAGG) ₄ | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₆ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₇ N ₂₇ (AAGG) ₄ | 23 | 888 |
| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₅ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₆ N ₂₇ (AAGG) ₄ | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₄ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₆ N ₂₇ (AAGG) ₄ | 19 | 916 |
| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₇ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₆ N ₂₇ (AAGG) ₄ | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₇ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₅ N ₂₇ (AAGG) ₄ | 56 | 1043 |
| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₇ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₇ N ₂₇ (AAGG) ₄ | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₇ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₆ N ₂₇ (AAGG) ₄ | 37 | 1107 |
| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₈ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₇ N ₂₇ (AAGG) ₄ | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₈ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₈ N ₂₇ (AAGG) ₄ | 19 | 1115 |
| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₇ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₇ N ₂₇ (AAGG) ₄ | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₇ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₆ N ₂₇ (AAGG) ₄ | 21 | 1273 |
| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₅ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₃ N ₂₇ (AAGG) ₄ | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₆ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₃ N ₂₇ (AAGG) ₄ | 45 | 1364 |
| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₆ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₇ N ₂₇ (AAGG) ₄ | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₅ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₅ N ₂₇ (AAGG) ₄ | 42 | 1411 |
| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₆ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₈ N ₂₇ (AAGG) ₄ | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₆ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₉ N ₂₇ (AAGG) ₄ | 32 | 1545 |
| DYS518 | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₆ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₅ N ₂₇ (AAGG) ₄ | (AAAG) ₃ (GAAG) ₁ (AAAG) ₁₅ (GGAG) ₁ (AAAG) ₄ N ₆ (AAAG) ₁₅ N ₂₇ (AAGG) ₄ | 29 | 1790 |
| DYS520 | (GATA) ₁₂ (CATA) ₁₁ | (GATA) ₁₁ (CATA) ₁₁ | 32 | 56 |
| DYS520 | (GATA) ₁₂ (CATA) ₁₁ | (GATA) ₁₁ (CATA) ₁₁ | 34 | 88 |
| DYS520 | (GATA) ₁₂ (CATA) ₁₀ | (GATA) ₁₂ (CATA) ₁₁ | 31 | 141 |
| DYS520 | (GATA) ₁₁ (CATA) ₁₁ | (GATA) ₁₂ (CATA) ₁₁ | 31 | 434 |
| DYS521 | (CTTT) ₅ (TCTT) ₃ (TTTT) ₁ (CTTT) ₅ T(CTTT) ₁₃ | (CTTT) ₅ (TCTT) ₃ (TTTT) ₁ (CTTT) ₅ T(CTTT) ₁₂ | 25 | 133 |
| DYS522 | (ATAG) ₁₀ | (ATAG) ₁₁ | 22 | 1088 |
| DYS525 | (AGAT) ₁₁ | (AGAT) ₁₀ | 24 | 571 |
| DYS526a | (CCTT) ₁₆ | (CCTT) ₁₅ | 64 | 166 |
| DYS526a | (CCTT) ₁₃ | (CCTT) ₁₄ | 53 | 453 |
| DYS526a | (CCTT) ₁₄ | (CCTT) ₁₃ | 31 | 1315 |
| DYS526a | (CCTT) ₁₁ | (CCTT) ₁₂ | 31 | 1652 |
| DYS526b | (CCCT) ₃ N ₂₀ (CTTT) ₁₄ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₁ | (CCCT) ₃ N ₂₀ (CTTT) ₁₃ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₁ | 24 | 17 |
| DYS526b | (CCCT) ₃ N ₂₀ (CTTT) ₁₆ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₄ | (CCCT) ₃ N ₂₀ (CTTT) ₁₅ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₄ | 50 | 42 |

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| DYS526b | (CCCT) ₃ N ₂₀ (CTTT) ₁₇ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₄ | (CCCT) ₃ N ₂₀ (CTTT) ₁₆ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₄ | 34 | 88 |
| DYS526b | (CCCT) ₃ N ₂₀ (CTTT) ₁₆ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₂ | (CCCT) ₃ N ₂₀ (CTTT) ₁₅ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₂ | 25 | 185 |
| DYS526b | (CCCT) ₃ N ₂₀ (CTTT) ₁₇ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₄ | (CCCT) ₃ N ₂₀ (CTTT) ₁₆ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₄ | 41 | 298 |
| DYS526b | (CCCT) ₃ N ₂₀ (CTTT) ₁₆ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₄ | (CCCT) ₃ N ₂₀ (CTTT) ₁₅ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₄ | 41 | 386 |
| DYS526b | (CCCT) ₃ N ₂₀ (CTTT) ₁₁ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₄ | (CCCT) ₃ N ₂₀ (CTTT) ₁₂ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₄ | 31 | 505 |
| DYS526b | (CCCT) ₃ N ₂₀ (CTTT) ₁₆ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₀ | (CCCT) ₃ N ₂₀ (CTTT) ₁₇ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₀ | 32 | 523 |
| DYS526b | (CCCT) ₃ N ₂₀ (CTTT) ₁₇ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₀ | (CCCT) ₃ N ₂₀ (CTTT) ₁₆ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₀ | 39 | 654 |
| DYS526b | (CCCT) ₃ N ₂₀ (CTTT) ₁₆ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₄ | (CCCT) ₃ N ₂₀ (CTTT) ₁₇ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₄ | 25 | 918 |
| DYS526b | (CCCT) ₃ N ₂₀ (CTTT) ₁₅ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₄ | (CCCT) ₃ N ₂₀ (CTTT) ₁₅ (CCTT) ₈ N ₁₁₃ (CCTT) ₁₄ | 36 | 983 |
| DYS526b | (CCCT) ₃ N ₂₀ (CTTT) ₁₅ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₄ | (CCCT) ₃ N ₂₀ (CTTT) ₁₆ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₄ | 37 | 1107 |
| DYS526b | (CCCT) ₃ N ₂₀ (CTTT) ₁₄ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₄ | (CCCT) ₃ N ₂₀ (CTTT) ₁₅ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₄ | 41 | 1122 |
| DYS526b | (CCCT) ₃ N ₂₀ (CTTT) ₁₅ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₄ | (CCCT) ₃ N ₂₀ (CTTT) ₁₄ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₄ | 22 | 1161 |
| DYS526b | (CCCT) ₃ N ₂₀ (CTTT) ₁₃ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₄ | (CCCT) ₃ N ₂₀ (CTTT) ₁₄ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₄ | 45 | 1171 |
| DYS526b | (CCCT) ₃ N ₂₀ (CTTT) ₁₆ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₄ | (CCCT) ₃ N ₂₀ (CTTT) ₁₅ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₄ | 48 | 1250 |
| DYS526b | (CCCT) ₃ N ₂₀ (CTTT) ₁₃ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₂ | (CCCT) ₃ N ₂₀ (CTTT) ₁₄ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₂ | 54 | 1447 |
| DYS526b | (CCCT) ₃ N ₂₀ (CTTT) ₁₃ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₃ | (CCCT) ₃ N ₂₀ (CTTT) ₁₂ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₃ | 21 | 1555 |
| DYS526b | (CCCT) ₃ N ₂₀ (CTTT) ₁₄ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₃ | (CCCT) ₃ N ₂₀ (CTTT) ₁₅ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₃ | 29 | 1662 |
| DYS526b | (CCCT) ₃ N ₂₀ (CTTT) ₁₅ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₄ | (CCCT) ₃ N ₂₀ (CTTT) ₁₆ (CCTT) ₉ N ₁₁₃ (CCTT) ₁₄ | 50 | 1881 |
| DYS531 | (AAAT) ₁₁ | (AAAT) ₉ | 22 | 747 |
| DYS532 | (TCCC) ₃ N ₅ (TTCC) ₅ N ₉ (TTCT) ₃ (TTCC) ₁ (TTCT) ₁₂ N ₁₇ (TTCT) ₃ N ₁₃ (TTCC) ₄ N ₇₀ (TTCT) ₃ N ₆ (TTCT) ₃ | (TCCC) ₃ N ₅ (TTCC) ₅ N ₉ (TTCT) ₃ (TTCC) ₁ (TTCT) ₁₃ N ₁₇ (TTCT) ₃ N ₁₃ (TTCC) ₄ N ₇₀ (TTCT) ₃ N ₆ (TTCT) ₃ | 26 | 759 |
| DYS532 | (TCCC) ₃ N ₅ (TTCC) ₅ N ₉ (TTCT) ₃ (TTCC) ₁ (TTCT) ₁₂ N ₁₇ (TTCT) ₃ N ₁₃ (TTCC) ₄ N ₇₀ (TTCT) ₃ N ₆ (TTCT) ₃ | (TCCC) ₃ N ₅ (TTCC) ₅ N ₉ (TTCT) ₃ (TTCC) ₁ (TTCT) ₁₁ N ₁₇ (TTCT) ₃ N ₁₃ (TTCC) ₄ N ₇₀ (TTCT) ₃ N ₆ (TTCT) ₃ | 30 | 1101 |
| DYS532 | (TCCC) ₃ N ₅ (TTCC) ₅ N ₉ (TTCT) ₃ (TTCC) ₁ (TTCT) ₁₂ N ₁₇ (TTCT) ₃ N ₁₃ (TTCC) ₄ N ₇₀ (TTCT) ₃ N ₆ (TTCT) ₃ | (TCCC) ₃ N ₅ (TTCC) ₅ N ₉ (TTCT) ₃ (TTCC) ₁ (TTCT) ₁₃ N ₁₇ (TTCT) ₃ N ₁₃ (TTCC) ₄ N ₇₀ (TTCT) ₃ N ₆ (TTCT) ₃ | 29 | 1255 |
| DYS532 | (TCCC) ₃ N ₅ (TTCC) ₅ N ₉ (TTCT) ₃ (TTCC) ₁ (TTCT) ₁₂ N ₁₇ (TTCT) ₃ N ₁₃ (TTCC) ₄ N ₇₀ (TTCT) ₃ N ₆ (TTCT) ₃ | (TCCC) ₃ N ₅ (TTCC) ₅ N ₉ (TTCT) ₃ (TTCC) ₁ (TTCT) ₁₃ N ₁₇ (TTCT) ₃ N ₁₃ (TTCC) ₄ N ₇₀ (TTCT) ₃ N ₆ (TTCT) ₃ | 30 | 1347 |
| DYS533 | (TATC) ₁₃ | (TATC) ₁₂ | 34 | 27 |
| DYS533 | (TATC) ₁₃ | (TATC) ₁₂ | 29 | 892 |
| DYS533 | (TATC) ₁₃ | (TATC) ₁₂ | 37 | 905 |
| DYS533 | (TATC) ₁₃ | (TATC) ₁₄ | 18 | 1039 |
| DYS533 | (TATC) ₁₂ | (TATC) ₁₃ | 42 | 1054 |

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| DYS533 | (TATC) ₁₄ | (TATC) ₁₅ | 34 | 1158 |
| DYS533 | (TATC) ₁₂ | (TATC) ₁₃ | 21 | 1166 |
| DYS533 | (TATC) ₁₃ | (TATC) ₁₂ | 40 | 1281 |
| DYS534 | (CTTT) ₃ N ₈ (CTTT) ₁₆ N ₉ (CTTT) ₃ | (CTTT) ₃ N ₈ (CTTT) ₁₇ N ₉ (CTTT) ₃ | 37 | 135 |
| DYS534 | (CTTT) ₃ N ₈ (CTTT) ₁₆ N ₉ (CTTT) ₃ | (CTTT) ₃ N ₈ (CTTT) ₁₇ N ₉ (CTTT) ₃ | 27 | 167 |
| DYS534 | (CTTT) ₃ N ₈ (CTTT) ₁₅ N ₉ (CTTT) ₃ | (CTTT) ₃ N ₈ (CTTT) ₁₆ N ₉ (CTTT) ₃ | 17 | 235 |
| DYS534 | (CTTT) ₃ N ₈ (CTTT) ₁₄ N ₉ (CTTT) ₃ | (CTTT) ₃ N ₈ (CTTT) ₁₅ N ₉ (CTTT) ₃ | 41 | 250 |
| DYS534 | (CTTT) ₃ N ₈ (CTTT) ₁₄ N ₉ (CTTT) ₃ | (CTTT) ₃ N ₈ (CTTT) ₁₅ N ₉ (CTTT) ₃ | 34 | 308 |
| DYS534 | (CTTT) ₃ N ₈ (CTTT) ₁₇ N ₉ (CTTT) ₃ | (CTTT) ₃ N ₈ (CTTT) ₁₆ N ₉ (CTTT) ₃ | 39 | 419 |
| DYS534 | (CTTT) ₃ N ₈ (CTTT) ₁₈ N ₉ (CTTT) ₃ | (CTTT) ₃ N ₈ (CTTT) ₁₉ N ₉ (CTTT) ₃ | 28 | 674 |
| DYS534 | (CTTT) ₃ N ₈ (CTTT) ₁₃ N ₉ (CTTT) ₃ | (CTTT) ₃ N ₈ (CTTT) ₁₄ N ₉ (CTTT) ₃ | 23 | 1205 |
| DYS534 | (CTTT) ₃ N ₈ (CTTT) ₁₆ N ₉ (CTTT) ₃ | (CTTT) ₃ N ₈ (CTTT) ₁₇ N ₉ (CTTT) ₃ | 45 | 1364 |
| DYS534 | (CTTT) ₃ N ₈ (CTTT) ₁₄ N ₉ (CTTT) ₃ | (CTTT) ₃ N ₈ (CTTT) ₁₃ N ₉ (CTTT) ₃ | 58 | 1808 |
| DYS534 | (CTTT) ₃ N ₈ (CTTT) ₁₇ N ₉ (CTTT) ₃ | (CTTT) ₃ N ₈ (CTTT) ₁₈ N ₉ (CTTT) ₃ | 61 | 1836 |
| DYS536 | (TCCT) ₁₂ N ₈ (TTCT) ₄ | (TCCT) ₁₃ N ₈ (TTCT) ₄ | 20 | 1092 |
| DYS537 | (TCTA) ₁₂ | (TCTA) ₁₃ | 29 | 609 |
| DYS537 | (TCTA) ₁₃ | (TCTA) ₁₂ | 27 | 1248 |
| DYS537 | (TCTA) ₁₁ | (TCTA) ₁₂ | 40 | 1427 |
| DYS539 | (TAGA) ₁₁ | (TAGA) ₁₀ | 63 | 1902 |
| DYS540 | (TTAT) ₁₂ | (TTAT) ₁₃ | 31 | 141 |
| DYS540 | (TTAT) ₁₂ | (TTAT) ₁₁ | 59 | 152 |
| DYS540 | (TTAT) ₁₁ | (TTAT) ₁₀ | 19 | 682 |
| DYS540 | (TTAT) ₁₁ | (TTAT) ₁₂ | 38 | 1020 |
| DYS540 | (TTAT) ₁₂ | (TTAT) ₁₁ | 31 | 1134 |
| DYS541 | (TATC) ₁₂ (TTC) ₁ (TATC) ₃ | (TATC) ₁₃ (TTC) ₁ (TATC) ₃ | 34 | 151 |
| DYS541 | (TATC) ₁₂ (TTC) ₁ (TATC) ₃ | (TATC) ₁₁ (TTC) ₁ (TATC) ₃ | 34 | 239 |
| DYS541 | (TATC) ₁₄ (TTC) ₁ (TATC) ₃ | (TATC) ₁₃ (TTC) ₁ (TATC) ₃ | 33 | 339 |
| DYS541 | (TATC) ₁₃ (TTC) ₁ (TATC) ₃ | (TATC) ₁₂ (TTC) ₁ (TATC) ₃ | 36 | 733 |
| DYS541 | (TATC) ₁₄ (TTC) ₁ (TATC) ₃ | (TATC) ₁₃ (TTC) ₁ (TATC) ₃ | 25 | 1415 |
| DYS541 | (TATC) ₁₂ (TTC) ₁ (TATC) ₃ | (TATC) ₁₃ (TTC) ₁ (TATC) ₃ | 64 | 1843 |

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| DYS543 | (AGAT) ₃ (GATA) ₁₁ N ₄₂ (ATGT) ₄ (ATGG) ₂ N ₃₅ (GAAA) ₃ | (AGAT) ₃ (GATA) ₁₂ N ₄₂ (ATGT) ₄ (ATGG) ₂ N ₃₅ (GAAA) ₃ | 23 | 16 |
| DYS543 | (AGAT) ₃ (GATA) ₁₅ N ₄₂ (ATGT) ₃ (ATGG) ₃ N ₃₅ (GAAA) ₃ | (AGAT) ₃ (GATA) ₁₄ N ₄₂ (ATGT) ₃ (ATGG) ₃ N ₃₅ (GAAA) ₃ | 40 | 774 |
| DYS543 | (AGAT) ₃ (GATA) ₁₅ N ₄₂ (ATGT) ₃ (ATGG) ₃ N ₃₅ (GAAA) ₃ | (AGAT) ₃ (GATA) ₁₄ N ₄₂ (ATGT) ₃ (ATGG) ₃ N ₃₅ (GAAA) ₃ | 33 | 844 |
| DYS543 | (AGAT) ₃ (GATA) ₁₂ N ₄₂ (ATGT) ₄ (ATGG) ₂ N ₃₅ (GAAA) ₃ | (AGAT) ₃ (GATA) ₁₁ N ₄₂ (ATGT) ₄ (ATGG) ₂ N ₃₅ (GAAA) ₃ | 43 | 939 |
| DYS543 | (AGAT) ₃ (GATA) ₁₃ N ₄₂ (ATGT) ₃ (ATGG) ₃ N ₃₅ (GAAA) ₃ | (AGAT) ₃ (GATA) ₁₄ N ₄₂ (ATGT) ₃ (ATGG) ₃ N ₃₅ (GAAA) ₃ | 42 | 1054 |
| DYS543 | (AGAT) ₃ (GATA) ₁₅ N ₄₂ (ATGT) ₃ (ATGG) ₃ N ₃₅ (GAAA) ₃ | (AGAT) ₃ (GATA) ₁₆ N ₄₂ (ATGT) ₃ (ATGG) ₃ N ₃₅ (GAAA) ₃ | Unknown | 1063 |
| DYS543 | (AGAT) ₃ (GATA) ₁₃ N ₄₂ (ATGT) ₄ (ATGG) ₂ N ₃₅ (GAAA) ₃ | (AGAT) ₃ (GATA) ₁₂ N ₄₂ (ATGT) ₄ (ATGG) ₂ N ₃₅ (GAAA) ₃ | Unknown | 1223 |
| DYS543 | (AGAT) ₃ (GATA) ₁₅ N ₄₂ (ATGT) ₃ (ATGG) ₃ N ₃₅ (GAAA) ₃ | (AGAT) ₃ (GATA) ₁₄ N ₄₂ (ATGT) ₃ (ATGG) ₃ N ₃₅ (GAAA) ₃ | 31 | 1229 |
| DYS543 | (AGAT) ₃ (GATA) ₁₃ N ₄₂ (ATGT) ₃ (ATGG) ₃ N ₃₅ (GAAA) ₃ | (AGAT) ₃ (GATA) ₁₂ N ₄₂ (ATGT) ₃ (ATGG) ₃ N ₃₅ (GAAA) ₃ | 31 | 1315 |
| DYS543 | (AGAT) ₃ (GATA) ₁₁ N ₄₂ (ATGT) ₄ (ATGG) ₂ N ₃₅ (GAAA) ₃ | (AGAT) ₃ (GATA) ₁₂ N ₄₂ (ATGT) ₄ (ATGG) ₂ N ₃₅ (GAAA) ₃ | Unknown | 1591 |
| DYS543 | (AGAT) ₃ (GATA) ₁₃ N ₄₂ (ATGT) ₄ (ATGG) ₂ N ₃₅ (GAAA) ₃ | (AGAT) ₃ (GATA) ₁₂ N ₄₂ (ATGT) ₄ (ATGG) ₂ N ₃₅ (GAAA) ₃ | 28 | 1712 |
| DYS546 | (TTCC) ₃ N ₂₃ (TTCT) ₃ N ₃₃ (TTCC) ₃ N ₁₆ (TTCT) ₁₇ | (TTCC) ₃ N ₂₃ (TTCT) ₃ N ₃₃ (TTCC) ₃ N ₁₆ (TTCT) ₁₈ | 34 | 371 |
| DYS546 | (TTCC) ₃ N ₂₃ (TTCT) ₃ N ₃₃ (TTCC) ₃ N ₁₆ (TTCT) ₁₄ | (TTCC) ₃ N ₂₃ (TTCT) ₃ N ₃₃ (TTCC) ₃ N ₁₆ (TTCT) ₁₃ | 20 | 706 |
| DYS546 | (TTCC) ₃ N ₂₃ (TTCT) ₃ N ₃₃ (TTCC) ₃ N ₁₆ (TTCT) ₁₆ | (TTCC) ₃ N ₂₃ (TTCT) ₃ N ₃₃ (TTCC) ₃ N ₁₆ (TTCT) ₁₇ | 20 | 756 |
| DYS546 | (TTCC) ₃ N ₂₃ (TTCT) ₃ N ₃₃ (TTCC) ₃ N ₁₆ (TTCT) ₁₇ | (TTCC) ₃ N ₂₃ (TTCT) ₃ N ₃₃ (TTCC) ₃ N ₁₆ (TTCT) ₁₆ | 31 | 878 |
| DYS546 | (TTCC) ₃ N ₂₃ (TTCT) ₃ N ₃₃ (TTCC) ₃ N ₁₆ (TTCT) ₁₈ | (TTCC) ₃ N ₂₃ (TTCT) ₃ N ₃₃ (TTCC) ₃ N ₁₆ (TTCT) ₁₇ | 23 | 1119 |
| DYS546 | (TTCC) ₃ N ₂₃ (TTCT) ₃ N ₃₃ (TTCC) ₃ N ₁₆ (TTCT) ₁₆ | (TTCC) ₃ N ₂₃ (TTCT) ₃ N ₃₃ (TTCC) ₃ N ₁₆ (TTCT) ₁₄ | 55 | 1840 |
| DYS547 | (CCTT) ₁₀ T(C TTC) ₅ N ₅₆ (TTTC) ₁₆ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₁ N ₁₄ (TTTC) ₃ | (CCTT) ₁₀ T(C TTC) ₅ N ₅₆ (TTTC) ₁₇ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₁ N ₁₄ (TTTC) ₃ | 28 | 1 |
| DYS547 | (CCTT) ₁₂ T(C TTC) ₄ N ₅₆ (TTTC) ₁₅ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₂ N ₁₄ (TTTC) ₃ | (CCTT) ₁₂ T(C TTC) ₄ N ₅₆ (TTTC) ₁₅ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₃ N ₁₄ (TTTC) ₃ | 46 | 10 |
| DYS547 | (CCTT) ₁₁ T(C TTC) ₅ N ₅₆ (TTTC) ₁₇ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₄ N ₁₄ (TTTC) ₃ | (CCTT) ₁₁ T(C TTC) ₅ N ₅₆ (TTTC) ₁₈ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₄ N ₁₄ (TTTC) ₃ | 59 | 152 |
| DYS547 | (CCTT) ₁₃ T(C TTC) ₅ N ₅₆ (TTTC) ₁₆ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₃ N ₁₄ (TTTC) ₃ | (CCTT) ₁₂ T(C TTC) ₅ N ₅₆ (TTTC) ₁₆ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₃ N ₁₄ (TTTC) ₃ | 34 | 243 |
| DYS547 | (CCTT) ₁₃ T(C TTC) ₅ N ₅₆ (TTTC) ₁₄ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₂ N ₁₄ (TTTC) ₃ | (CCTT) ₁₃ T(C TTC) ₅ N ₅₆ (TTTC) ₁₄ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₁ N ₁₄ (TTTC) ₃ | 31 | 268 |
| DYS547 | (CCTT) ₁₂ T(C TTC) ₄ N ₅₆ (TTTC) ₁₇ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₂ N ₁₄ (TTTC) ₃ | (CCTT) ₁₂ T(C TTC) ₄ N ₅₆ (TTTC) ₁₇ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₁ N ₁₄ (TTTC) ₃ | 50 | 270 |
| DYS547 | (CCTT) ₁₀ T(C TTC) ₅ N ₅₆ (TTTC) ₁₈ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₀ N ₁₄ (TTTC) ₃ | (CCTT) ₁₀ T(C TTC) ₅ N ₅₆ (TTTC) ₁₉ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₀ N ₁₄ (TTTC) ₃ | 33 | 339 |
| DYS547 | (CCTT) ₁₂ T(C TTC) ₅ N ₅₆ (TTTC) ₁₇ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₁ N ₁₄ (TTTC) ₃ | (CCTT) ₁₂ T(C TTC) ₅ N ₅₆ (TTTC) ₁₇ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₀ N ₁₄ (TTTC) ₃ | 36 | 378 |
| DYS547 | (CCTT) ₁₃ T(C TTC) ₄ N ₅₆ (TTTC) ₁₆ N ₁₀ (CCTT) ₄ | (CCTT) ₁₃ T(C TTC) ₄ N ₅₆ (TTTC) ₁₇ N ₁₀ (CCTT) ₄ | 28 | 425 |

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|--------|--|--|----|------|
| | $(TCTC)_1(TTTC)_{12}N_{14}(TTTC)_3$ | $(TCTC)_1(TTTC)_{12}N_{14}(TTTC)_3$ | | |
| DYS547 | $(CCTT)_{13}T(CTTC)_5N_{56}(TTTC)_{16}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{11}N_{14}(TTTC)_3$ | $(CCTT)_{13}T(CTTC)_5N_{56}(TTTC)_{18}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{11}N_{14}(TTTC)_3$ | 28 | 484 |
| DYS547 | $(CCTT)_{12}T(CTTC)_4N_{56}(TTTC)_{17}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{12}N_{14}(TTTC)_3$ | $(CCTT)_{12}T(CTTC)_4N_{56}(TTTC)_{16}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{12}N_{14}(TTTC)_3$ | 31 | 613 |
| DYS547 | $(CCTT)_{12}T(CTTC)_4N_{56}(TTTC)_{15}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{12}N_{14}(TTTC)_3$ | $(CCTT)_{12}T(CTTC)_4N_{56}(TTTC)_{15}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{11}N_{14}(TTTC)_3$ | 39 | 654 |
| DYS547 | $(CCTT)_{12}T(CTTC)_5N_{56}(TTTC)_{15}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{12}N_{14}(TTTC)_3$ | $(CCTT)_{12}T(CTTC)_5N_{56}(TTTC)_{14}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{12}N_{14}(TTTC)_3$ | 27 | 710 |
| DYS547 | $(CCTT)_{13}T(CTTC)_5N_{56}(TTTC)_{15}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{11}N_{14}(TTTC)_3$ | $(CCTT)_{13}T(CTTC)_5N_{56}(TTTC)_{14}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{11}N_{14}(TTTC)_3$ | 25 | 711 |
| DYS547 | $(CCTT)_{10}T(CTTC)_5N_{56}(TTTC)_{18}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{10}N_{14}(TTTC)_3$ | $(CCTT)_{10}T(CTTC)_5N_{56}(TTTC)_{17}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{10}N_{14}(TTTC)_3$ | 59 | 846 |
| DYS547 | $(CCTT)_{12}T(CTTC)_4N_{56}(TTTC)_{16}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{12}N_{14}(TTTC)_3$ | $(CCTT)_{12}T(CTTC)_4N_{56}(TTTC)_{17}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{12}N_{14}(TTTC)_3$ | 37 | 904 |
| DYS547 | $(CCTT)_{12}T(CTTC)_4N_{56}(TTTC)_{18}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{13}N_{14}(TTTC)_3$ | $(CCTT)_{12}T(CTTC)_4N_{56}(TTTC)_{19}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{13}N_{14}(TTTC)_3$ | 22 | 986 |
| DYS547 | $(CCTT)_{12}T(CTTC)_5N_{56}(TTTC)_{22}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_9N_{14}(TTTC)_3$ | $(CCTT)_{12}T(CTTC)_5N_{56}(TTTC)_{21}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_9N_{14}(TTTC)_3$ | 55 | 1022 |
| DYS547 | $(CCTT)_{12}T(CTTC)_4N_{56}(TTTC)_{16}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{13}N_{14}(TTTC)_3$ | $(CCTT)_{12}T(CTTC)_4N_{56}(TTTC)_{16}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{14}N_{14}(TTTC)_3$ | 43 | 1153 |
| DYS547 | $(CCTT)_{12}T(CTTC)_4N_{56}(TTTC)_{19}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{12}N_{14}(TTTC)_3$ | $(CCTT)_{12}T(CTTC)_4N_{56}(TTTC)_{18}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{12}N_{14}(TTTC)_3$ | 31 | 1229 |
| DYS547 | $(CCTT)_{12}T(CTTC)_4N_{56}(TTTC)_{16}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{13}N_{14}(TTTC)_3$ | $(CCTT)_{12}T(CTTC)_4N_{56}(TTTC)_{16}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{14}N_{14}(TTTC)_3$ | 29 | 1276 |
| DYS547 | $(CCTT)_{12}T(CTTC)_4N_{56}(TTTC)_{18}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{12}N_{14}(TTTC)_3$ | $(CCTT)_{12}T(CTTC)_4N_{56}(TTTC)_{19}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{12}N_{14}(TTTC)_3$ | 28 | 1294 |
| DYS547 | $(CCTT)_{11}T(CTTC)_4N_{56}(TTTC)_{16}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{12}N_{14}(TTTC)_3$ | $(CCTT)_{11}T(CTTC)_4N_{56}(TTTC)_{16}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{13}N_{14}(TTTC)_3$ | 30 | 1297 |
| DYS547 | $(CCTT)_{11}T(CTTC)_5N_{56}(TTTC)_{15}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{11}N_{14}(TTTC)_3$ | $(CCTT)_{11}T(CTTC)_5N_{56}(TTTC)_{15}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{12}N_{14}(TTTC)_3$ | 32 | 1321 |
| DYS547 | $(CCTT)_{12}T(CTTC)_5N_{56}(TTTC)_{15}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{12}N_{14}(TTTC)_3$ | $(CCTT)_{12}T(CTTC)_5N_{56}(TTTC)_{15}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{13}N_{14}(TTTC)_3$ | 41 | 1379 |
| DYS547 | $(CCTT)_{10}T(CTTC)_5N_{56}(TTTC)_{18}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{10}N_{14}(TTTC)_3$ | $(CCTT)_{10}T(CTTC)_5N_{56}(TTTC)_{19}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{10}N_{14}(TTTC)_3$ | 24 | 1466 |
| DYS547 | $(CCTT)_{10}T(CTTC)_5N_{56}(TTTC)_{16}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{10}N_{14}(TTTC)_3$ | $(CCTT)_{10}T(CTTC)_5N_{56}(TTTC)_{16}N_{10}(CCTT)_4$ $(TCTC)_1(TTTC)_{11}N_{14}(TTTC)_3$ | 42 | 1485 |
| DYS547 | $(CCTT)_{12}T(CTTC)_5N_{56}(TTTC)_{15}N_{10}(CCTT)_4$ | $(CCTT)_{12}T(CTTC)_5N_{56}(TTTC)_{16}N_{10}(CCTT)_4$ | 29 | 1491 |

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| | (TCTC) ₁ (TTTC) ₁₃ N ₁₄ (TTTC) ₃ | (TCTC) ₁ (TTTC) ₁₃ N ₁₄ (TTTC) ₃ | | |
| DYS547 | (CCTT) ₁₂ T(CTTC) ₄ N ₅₆ (TTTC) ₁₆ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₂ N ₁₄ (TTTC) ₃ | (CCTT) ₁₂ T(CTTC) ₄ N ₅₆ (TTTC) ₁₇ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₂ N ₁₄ (TTTC) ₃ | 23 | 1510 |
| DYS547 | (CCTT) ₁₂ T(CTTC) ₅ N ₅₆ (TTTC) ₁₆ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₁ N ₁₄ (TTTC) ₃ | (CCTT) ₁₂ T(CTTC) ₅ N ₅₆ (TTTC) ₁₆ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₀ N ₁₄ (TTTC) ₃ | 38 | 1524 |
| DYS547 | (CCTT) ₁₂ T(CTTC) ₅ N ₅₆ (TTTC) ₁₀ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₇ N ₁₄ (TTTC) ₃ | (CCTT) ₁₂ T(CTTC) ₅ N ₅₆ (TTTC) ₁₀ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₆ N ₁₄ (TTTC) ₃ | 36 | 1582 |
| DYS547 | (CCTT) ₁₂ T(CTTC) ₅ N ₅₆ (TTTC) ₁₆ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₃ N ₁₄ (TTTC) ₃ | (CCTT) ₁₂ T(CTTC) ₅ N ₅₆ (TTTC) ₁₇ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₃ N ₁₄ (TTTC) ₃ | 28 | 1640 |
| DYS547 | (CCTT) ₁₂ T(CTTC) ₄ N ₅₆ (TTTC) ₁₅ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₂ N ₁₄ (TTTC) ₃ | (CCTT) ₁₂ T(CTTC) ₄ N ₅₆ (TTTC) ₁₄ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₂ N ₁₄ (TTTC) ₃ | 22 | 1648 |
| DYS547 | (CCTT) ₁₂ T(CTTC) ₅ N ₅₆ (TTTC) ₁₆ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₂ N ₁₄ (TTTC) ₃ | (CCTT) ₁₂ T(CTTC) ₅ N ₅₆ (TTTC) ₁₇ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₂ N ₁₄ (TTTC) ₃ | 37 | 1663 |
| DYS547 | (CCTT) ₁₂ T(CTTC) ₄ N ₅₆ (TTTC) ₁₆ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₂ N ₁₄ (TTTC) ₃ | (CCTT) ₁₂ T(CTTC) ₄ N ₅₆ (TTTC) ₁₇ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₂ N ₁₄ (TTTC) ₃ | Unknown | 1723 |
| DYS547 | (CCTT) ₁₂ T(CTTC) ₅ N ₅₆ (TTTC) ₁₇ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₂ N ₁₄ (TTTC) ₃ | (CCTT) ₁₂ T(CTTC) ₅ N ₅₆ (TTTC) ₁₆ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₂ N ₁₄ (TTTC) ₃ | 54 | 1860 |
| DYS547 | (CCTT) ₁₀ T(CTTC) ₅ N ₅₆ (TTTC) ₁₆ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₃ N ₁₄ (TTTC) ₃ | (CCTT) ₁₀ T(CTTC) ₅ N ₅₆ (TTTC) ₁₆ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₂ N ₁₄ (TTTC) ₃ | 53 | 1871 |
| DYS547 | (CCTT) ₁₂ T(CTTC) ₅ N ₅₆ (TTTC) ₁₆ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₁ N ₁₄ (TTTC) ₃ | (CCTT) ₁₂ T(CTTC) ₅ N ₅₆ (TTTC) ₁₅ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₁ N ₁₄ (TTTC) ₃ | 55 | 1910 |
| DYS547 | (CCTT) ₁₂ T(CTTC) ₅ N ₅₆ (TTTC) ₁₄ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₁ N ₁₄ (TTTC) ₃ | (CCTT) ₁₂ T(CTTC) ₅ N ₅₆ (TTTC) ₁₄ N ₁₀ (CCTT) ₄ (TCTC) ₁ (TTTC) ₁₂ N ₁₄ (TTTC) ₃ | 65 | 1920 |
| DYS549 | (GATA) ₁₃ | (GATA) ₁₂ | 30 | 87 |
| DYS549 | (GATA) ₁₃ | (GATA) ₁₂ | 47 | 113 |
| DYS549 | (GATA) ₁₂ | (GATA) ₁₁ | 27 | 119 |
| DYS549 | (GATA) ₁₂ | (GATA) ₁₃ | 38 | 336 |
| DYS549 | (GATA) ₁₄ | (GATA) ₁₃ | 43 | 472 |
| DYS549 | (GATA) ₁₂ | (GATA) ₁₁ | 25 | 517 |
| DYS549 | (GATA) ₁₂ | (GATA) ₁₁ | 22 | 625 |
| DYS551 | (AGAT) ₁₅ N ₈ (AGAC) ₃ (AGGT) ₁ (AGAT) ₄ | (AGAT) ₁₄ N ₈ (AGAC) ₃ (AGGT) ₁ (AGAT) ₄ | 21 | 436 |
| DYS551 | (AGAT) ₁₄ N ₈ (AGAC) ₃ (AGGT) ₁ (AGAT) ₄ | (AGAT) ₁₃ N ₈ (AGAC) ₃ (AGGT) ₁ (AGAT) ₄ | 22 | 604 |
| DYS551 | (AGAT) ₁₅ N ₈ (AGAC) ₃ (AGGT) ₁ (AGAT) ₄ | (AGAT) ₁₄ N ₈ (AGAC) ₃ (AGGT) ₁ (AGAT) ₄ | 41 | 862 |
| DYS551 | (AGAT) ₁₄ N ₈ (AGAC) ₃ (AGGT) ₁ (AGAT) ₄ | (AGAT) ₁₃ N ₈ (AGAC) ₃ (AGGT) ₁ (AGAT) ₄ | 51 | 1212 |
| DYS551 | (AGAT) ₁₃ N ₈ (AGAC) ₃ (AGGT) ₁ (AGAT) ₄ | (AGAT) ₁₄ N ₈ (AGAC) ₃ (AGGT) ₁ (AGAT) ₄ | 27 | 1671 |

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| DYS552 | (TCTA) ₃ (TCTG) ₁ (TCTA) ₉ N ₄₀ (TCTA) ₁₅ | (TCTA) ₃ (TCTG) ₁ (TCTA) ₇ N ₄₀ (TCTA) ₁₅ | 25 | 471 |
| DYS552 | (TCTA) ₃ (TCTG) ₁ (TCTA) ₁₀ N ₄₀ (TCTA) ₁₅ | (TCTA) ₃ (TCTG) ₁ (TCTA) ₁₀ N ₄₀ (TCTA) ₁₆ | 59 | 846 |
| DYS552 | (TCTA) ₃ (TCTG) ₁ (TCTA) ₁₀ N ₄₀ (TCTA) ₁₄ | (TCTA) ₃ (TCTG) ₁ (TCTA) ₁₀ N ₄₀ (TCTA) ₁₅ | 18 | 847 |
| DYS552 | (TCTA) ₃ (TCTG) ₁ (TCTA) ₁₁ N ₄₀ (TCTA) ₁₄ | (TCTA) ₃ (TCTG) ₁ (TCTA) ₁₂ N ₄₀ (TCTA) ₁₄ | 31 | 1486 |
| DYS554 | (TAAA) ₁₀ | (TAAA) ₁₁ | 37 | 1277 |
| DYS556 | (AAAT) ₁₂ | (AAAT) ₁₁ | 30 | 1101 |
| DYS556 | (AAAT) ₁₁ | (AAAT) ₁₂ | 21 | 1448 |
| DYS557 | (TTTC) ₄ (TTCTC) ₁ (TTTC) ₄ (TTC) ₁ (TTTC) ₁₆ | (TTTC) ₄ (TTCTC) ₁ (TTTC) ₄ (TTC) ₁ (TTTC) ₁₅ | 24 | 17 |
| DYS557 | (TTTC) ₄ (TTCTC) ₁ (TTTC) ₄ (TTC) ₁ (TTTC) ₁₆ | (TTTC) ₄ (TTCTC) ₁ (TTTC) ₄ (TTC) ₁ (TTTC) ₁₅ | 23 | 52 |
| DYS557 | (TTTC) ₄ (TTCTC) ₁ (TTTC) ₄ (TTC) ₁ (TTTC) ₁₅ | (TTTC) ₄ (TTCTC) ₁ (TTTC) ₄ (TTC) ₁ (TTTC) ₁₆ | 34 | 394 |
| DYS557 | (TTTC) ₄ (TTCTC) ₁ (TTTC) ₄ (TTC) ₁ (TTTC) ₁₅ | (TTTC) ₄ (TTCTC) ₁ (TTTC) ₄ (TTC) ₁ (TTTC) ₁₆ | 38 | 589 |
| DYS557 | (TTTC) ₄ (TTCTC) ₁ (TTTC) ₄ (TTC) ₁ (TTTC) ₁₇ | (TTTC) ₄ (TTCTC) ₁ (TTTC) ₄ (TTC) ₁ (TTTC) ₁₆ | 38 | 1494 |
| DYS557 | (TTTC) ₄ (TTCTC) ₁ (TTTC) ₄ (TTC) ₁ (TTTC) ₁₅ | (TTTC) ₄ (TTCTC) ₁ (TTTC) ₄ (TTC) ₁ (TTTC) ₁₆ | 17 | 1517 |
| DYS559 | (TAAA) ₉ | (TAAA) ₈ | 27 | 1357 |
| DYS561 | (GATA) ₁₃ (GACA) ₄ | (GATA) ₁₂ (GACA) ₄ | 34 | 359 |
| DYS565 | (ATAA) ₁₃ | (ATAA) ₁₂ | 34 | 101 |
| DYS565 | (ATAA) ₁₃ | (ATAA) ₁₂ | 43 | 624 |
| DYS565 | (ATAA) ₁₃ | (ATAA) ₁₄ | 27 | 1673 |
| DYS568 | (AAAT) ₁₂ | (AAAT) ₁₃ | 35 | 1547 |
| DYS569 | (ATTT) ₁₂ | (ATTT) ₁₁ | 31 | 598 |
| DYS569 | (ATTT) ₁₂ | (ATTT) ₁₁ | 21 | 1053 |
| DYS570 | (TTTC) ₁₇ | (TTTC) ₁₆ | 34 | 92 |
| DYS570 | (TTTC) ₁₉ | (TTTC) ₂₀ | 39 | 112 |
| DYS570 | (TTTC) ₁₉ | (TTTC) ₁₈ | 20 | 240 |
| DYS570 | (TTTC) ₂₀ | (TTTC) ₁₉ | 37 | 293 |
| DYS570 | (TTTC) ₁₇ | (TTTC) ₁₈ | 41 | 313 |
| DYS570 | (TTTC) ₁₉ | (TTTC) ₁₇ | 16 | 316 |
| DYS570 | (TTTC) ₁₉ | (TTTC) ₁₈ | 25 | 317 |
| DYS570 | (TTTC) ₂₀ | (TTTC) ₂₁ | 32 | 614 |
| DYS570 | (TTTC) ₂₀ | (TTTC) ₂₁ | 24 | 855 |

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|--------|----------------------|----------------------|---------|------|
| DYS570 | (TTTC) ₂₀ | (TTTC) ₁₉ | 30 | 867 |
| DYS570 | (TTTC) ₂₁ | (TTTC) ₂₂ | 22 | 922 |
| DYS570 | (TTTC) ₁₈ | (TTTC) ₁₉ | 36 | 1061 |
| DYS570 | (TTTC) ₁₉ | (TTTC) ₂₀ | 20 | 1253 |
| DYS570 | (TTTC) ₁₆ | (TTTC) ₁₅ | 41 | 1256 |
| DYS570 | (TTTC) ₂₁ | (TTTC) ₂₀ | 45 | 1364 |
| DYS570 | (TTTC) ₁₇ | (TTTC) ₁₈ | 54 | 1895 |
| DYS570 | (TTTC) ₁₉ | (TTTC) ₁₈ | 66 | 1901 |
| DYS572 | (AAAT) ₁₁ | (AAAT) ₁₀ | 35 | 735 |
| DYS572 | (AAAT) ₁₁ | (AAAT) ₁₀ | 24 | 1342 |
| DYS572 | (AAAT) ₁₁ | (AAAT) ₁₀ | 28 | 1696 |
| DYS574 | (TTAT) ₁₀ | (TTAT) ₉ | 43 | 1818 |
| DYS576 | (AAAG) ₁₇ | (AAAG) ₁₈ | 28 | 153 |
| DYS576 | (AAAG) ₁₉ | (AAAG) ₁₈ | 34 | 236 |
| DYS576 | (AAAG) ₁₇ | (AAAG) ₁₈ | 34 | 243 |
| DYS576 | (AAAG) ₁₈ | (AAAG) ₁₇ | 47 | 347 |
| DYS576 | (AAAG) ₁₉ | (AAAG) ₁₈ | 17 | 635 |
| DYS576 | (AAAG) ₁₉ | (AAAG) ₁₈ | 37 | 715 |
| DYS576 | (AAAG) ₂₀ | (AAAG) ₂₁ | 23 | 716 |
| DYS576 | (AAAG) ₁₈ | (AAAG) ₁₇ | 38 | 719 |
| DYS576 | (AAAG) ₂₀ | (AAAG) ₂₁ | 29 | 789 |
| DYS576 | (AAAG) ₂₀ | (AAAG) ₁₉ | 50 | 884 |
| DYS576 | (AAAG) ₁₇ | (AAAG) ₁₈ | 42 | 1054 |
| DYS576 | (AAAG) ₁₉ | (AAAG) ₁₈ | 36 | 1061 |
| DYS576 | (AAAG) ₁₈ | (AAAG) ₁₉ | Unknown | 1200 |
| DYS576 | (AAAG) ₁₈ | (AAAG) ₁₉ | 40 | 1204 |
| DYS576 | (AAAG) ₁₈ | (AAAG) ₁₉ | 51 | 1212 |
| DYS576 | (AAAG) ₁₈ | (AAAG) ₁₇ | Unknown | 1224 |
| DYS576 | (AAAG) ₁₈ | (AAAG) ₁₉ | Unknown | 1265 |
| DYS576 | (AAAG) ₁₈ | (AAAG) ₁₉ | 38 | 1278 |

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|--------|---|---|---------|------|
| DYS576 | (AAAG) ₁₇ | (AAAG) ₁₆ | 22 | 1403 |
| DYS576 | (AAAG) ₁₉ | (AAAG) ₁₈ | 42 | 1411 |
| DYS576 | (AAAG) ₁₈ | (AAAG) ₁₇ | 26 | 1440 |
| DYS576 | (AAAG) ₁₈ | (AAAG) ₁₉ | 47 | 1675 |
| DYS576 | (AAAG) ₁₈ | (AAAG) ₁₉ | 55 | 1844 |
| DYS576 | (AAAG) ₂₀ | (AAAG) ₁₉ | 54 | 1939 |
| DYS578 | (AAAT) ₈ | (AAAT) ₉ | Unknown | 1353 |
| DYS585 | (TTATG) ₉ | (TTATG) ₁₀ | 35 | 794 |
| DYS585 | (TTATG) ₉ | (TTATG) ₈ | 26 | 1105 |
| DYS585 | (TTATG) ₉ | (TTATG) ₁₀ | 17 | 1109 |
| DYS587 | (CAATA) ₁₁ [(CAGTA) ₁ (CAATA) ₁] ₃ | (CAATA) ₁₀ [(CAGTA) ₁ (CAATA) ₁] ₃ | 24 | 83 |
| DYS587 | (CAATA) ₁₁ [(CAGTA) ₁ (CAATA) ₁] ₃ | (CAATA) ₁₂ [(CAGTA) ₁ (CAATA) ₁] ₃ | 59 | 152 |
| DYS587 | (CAATA) ₁₂ [(CAGTA) ₁ (CAATA) ₁] ₃ | (CAATA) ₁₁ [(CAGTA) ₁ (CAATA) ₁] ₃ | 25 | 260 |
| DYS587 | (CAATA) ₁₂ [(CAGTA) ₁ (CAATA) ₁] ₃ | (CAATA) ₁₃ [(CAGTA) ₁ (CAATA) ₁] ₃ | 63 | 917 |
| DYS593 | (AAAAC) ₄ (AAAAT) ₈ | (AAAAC) ₄ (AAAAT) ₇ | 26 | 1082 |
| DYS593 | (AAAAC) ₄ (AAAAT) ₈ | (AAAAC) ₄ (AAAAT) ₉ | Unknown | 1353 |
| DYS594 | (AAATA) ₁₀ | (AAATA) ₁₁ | 31 | 1134 |
| DYS611 | (TTC) ₅ N ₉ (TTC) ₄ (CTC) ₁ (TTC) ₃ N ₉ (TTC) ₅ (CTC) ₁ (TTC) ₃ N ₁₅ (TTC) ₄ (CT) ₁ (TTC) ₃ (CTC) ₁ (TTC) ₃ N ₂₀ (TTC) ₃ T (TTC) ₄ N ₇ (TTC) ₃ N ₉ (TTC) ₄ (TCC) ₁ (TTC) ₁₉ N ₂₃ (TTC) ₄ N ₄ [(TTC) ₁ (CTC) ₁] ₂ [(CTC) ₁ (TTC) ₁] ₃ | (TTC) ₅ N ₉ (TTC) ₄ (CTC) ₁ (TTC) ₃ N ₉ (TTC) ₅ (CTC) ₁ (TTC) ₃ N ₁₅ (TTC) ₄ (CT) ₁ (TTC) ₃ (CTC) ₁ (TTC) ₃ N ₂₀ (TTC) ₃ T (TTC) ₄ N ₇ (TTC) ₃ N ₉ (TTC) ₄ (TCC) ₁ (TTC) ₂₀ N ₂₃ (TTC) ₄ N ₄ [(TTC) ₁ (CTC) ₁] ₂ [(CTC) ₁ (TTC) ₁] ₃ | 30 | 99 |
| DYS611 | (TTC) ₅ N ₉ (TTC) ₄ (CTC) ₁ (TTC) ₃ N ₉ (TTC) ₅ (CTC) ₁ (TTC) ₃ N ₁₅ (TTC) ₄ (CT) ₁ (TTC) ₃ (CTC) ₁ (TTC) ₃ N ₂₀ (TTC) ₃ T (TTC) ₄ N ₇ (TTC) ₃ N ₉ (TTC) ₄ (TCC) ₁ (TTC) ₁₇ N ₂₃ (TTC) ₄ N ₄ [(TTC) ₁ (CTC) ₁] ₂ [(CTC) ₁ (TTC) ₁] ₃ | (TTC) ₅ N ₉ (TTC) ₄ (CTC) ₁ (TTC) ₃ N ₉ (TTC) ₅ (CTC) ₁ (TTC) ₃ N ₁₅ (TTC) ₄ (CT) ₁ (TTC) ₃ (CTC) ₁ (TTC) ₃ N ₂₀ (TTC) ₃ T (TTC) ₄ N ₇ (TTC) ₃ N ₉ (TTC) ₄ (TCC) ₁ (TTC) ₁₆ N ₂₃ (TTC) ₄ N ₄ [(TTC) ₁ (CTC) ₁] ₂ [(CTC) ₁ (TTC) ₁] ₃ | 31 | 164 |
| DYS611 | (TTC) ₅ N ₉ (TTC) ₄ (CTC) ₁ (TTC) ₃ N ₉ (TTC) ₅ (CTC) ₁ (TTC) ₃ N ₁₅ (TTC) ₄ (CT) ₁ (TTC) ₃ (CTC) ₁ (TTC) ₃ N ₂₀ (TTC) ₃ T (TTC) ₄ N ₇ (TTC) ₃ N ₉ (TTC) ₄ (TCC) ₁ (TTC) ₁₈ N ₂₃ (TTC) ₄ N ₄ [(TTC) ₁ (CTC) ₁] ₂ [(CTC) ₁ (TTC) ₁] ₃ | (TTC) ₅ N ₉ (TTC) ₄ (CTC) ₁ (TTC) ₃ N ₉ (TTC) ₅ (CTC) ₁ (TTC) ₃ N ₁₅ (TTC) ₄ (CT) ₁ (TTC) ₃ (CTC) ₁ (TTC) ₃ N ₂₀ (TTC) ₃ T (TTC) ₄ N ₇ (TTC) ₃ N ₉ (TTC) ₄ (TCC) ₁ (TTC) ₁₆ N ₂₃ (TTC) ₄ N ₄ [(TTC) ₁ (CTC) ₁] ₂ [(CTC) ₁ (TTC) ₁] ₃ | 20 | 254 |
| DYS611 | (TTC) ₅ N ₉ (TTC) ₄ (CTC) ₁ (TTC) ₃ N ₉ (TTC) ₅ (CTC) ₁ (TTC) ₃ N ₁₅ (TTC) ₄ (CT) ₁ (TTC) ₃ (CTC) ₁ (TTC) ₃ N ₂₀ (TTC) ₃ T (TTC) ₄ N ₇ (TTC) ₃ N ₉ (TTC) ₄ (TCC) ₁ (TTC) ₁₅ N ₂₃ (TTC) ₄ N ₄ [(TTC) ₁ (CTC) ₁] ₂ [(CTC) ₁ (TTC) ₁] ₃ | (TTC) ₅ N ₉ (TTC) ₄ (CTC) ₁ (TTC) ₃ N ₉ (TTC) ₅ (CTC) ₁ (TTC) ₃ N ₁₅ (TTC) ₄ (CT) ₁ (TTC) ₃ (CTC) ₁ (TTC) ₃ N ₂₀ (TTC) ₃ T (TTC) ₄ N ₇ (TTC) ₃ N ₉ (TTC) ₄ (TCC) ₁ (TTC) ₁₄ N ₂₃ (TTC) ₄ N ₄ [(TTC) ₁ (CTC) ₁] ₂ [(CTC) ₁ (TTC) ₁] ₃ | 25 | 517 |
| DYS611 | (TTC) ₅ N ₉ (TTC) ₄ (CTC) ₁ (TTC) ₃ N ₉ (TTC) ₅ (CTC) ₁ (TTC) ₃ N ₁₅ | (TTC) ₅ N ₉ (TTC) ₄ (CTC) ₁ (TTC) ₃ N ₉ (TTC) ₅ (CTC) ₁ (TTC) ₃ N ₁₅ | 18 | 956 |

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|--------|---|--|----|------|
| DYS614 | (CTT) ₄ (CCT) ₁ (CTT) ₃ N ₁₅ (CCT) ₄ (CTT) ₄ (CCT) ₁ (CTT) ₃ N ₁₈ (CCT) ₃ (CTT) ₅ N ₂₀ [(CTT) ₁ (CTG) ₁] ₃ (CT) ₁ (CTT) ₁₉ N ₈ (CTT) ₄ [(CTC) ₁ (CTT) ₁] ₃ [(CTC) ₁ (TTT) ₁] ₁ (CTT) ₅ | (CTT) ₄ (CCT) ₁ (CTT) ₃ N ₁₅ (CCT) ₄ (CTT) ₄ (CCT) ₁ (CTT) ₃ N ₁₈ (CCT) ₃ (CTT) ₅ N ₂₀ [(CTT) ₁ (CTG) ₁] ₃ (CT) ₁ (CTT) ₁₈ N ₈ (CTT) ₄ [(CTC) ₁ (CTT) ₁] ₃ [(CTC) ₁ (TTT) ₁] ₁ (CTT) ₅ | 19 | 1784 |
| DYS614 | (CTT) ₄ (CCT) ₁ (CTT) ₃ N ₁₅ (CCT) ₄ (CTT) ₄ (CCT) ₁ (CTT) ₃ N ₁₈ (CCT) ₃ (CTT) ₅ N ₂₀ [(CTT) ₁ (CTG) ₁] ₃ (CT) ₁ (CTT) ₁₈ N ₈ (CTT) ₄ [(CTC) ₁ (CTT) ₁] ₃ [(CTC) ₁ (TTT) ₁] ₁ (CTT) ₅ | (CTT) ₄ (CCT) ₁ (CTT) ₃ N ₁₅ (CCT) ₄ (CTT) ₄ (CCT) ₁ (CTT) ₃ N ₁₈ (CCT) ₃ (CTT) ₅ N ₂₀ [(CTT) ₁ (CTG) ₁] ₃ (CT) ₁ (CTT) ₁₆ N ₈ (CTT) ₄ [(CTC) ₁ (CTT) ₁] ₃ [(CTC) ₁ (TTT) ₁] ₁ (CTT) ₅ | 52 | 1965 |
| DYS616 | (TAT) ₁₄ (CAT) ₁ (TAT) ₃ | (TAT) ₁₅ (CAT) ₁ (TAT) ₃ | 25 | 40 |
| DYS616 | (TAT) ₁₅ (CAT) ₁ (TAT) ₃ | (TAT) ₁₄ (CAT) ₁ (TAT) ₃ | 41 | 417 |
| DYS622 | (GAAA) ₆ (AGAAG) ₁ (GAAA) ₁₂ | (GAAA) ₆ (AGAAG) ₁ (GAAA) ₁₃ | 33 | 187 |
| DYS622 | (GAAA) ₆ (AGAAG) ₁ (GAAA) ₁₄ | (GAAA) ₆ (AGAAG) ₁ (GAAA) ₁₅ | 34 | 308 |
| DYS622 | (GAAA) ₆ (AGAAG) ₁ (GAAA) ₁₄ | (GAAA) ₆ (AGAAG) ₁ (GAAA) ₁₃ | 30 | 842 |
| DYS622 | (GAAA) ₆ (AGAAG) ₁ (GAAA) ₁₁ | (GAAA) ₆ (AGAAG) ₁ (GAAA) ₁₀ | 19 | 1006 |
| DYS622 | (GAAA) ₆ (AGAAG) ₁ (GAAA) ₁₃ | (GAAA) ₆ (AGAAG) ₁ (GAAA) ₁₂ | 21 | 1436 |
| DYS625 | (CTTT) ₄ (TTCT) ₁ (CTTT) ₃ (TTT) ₁ (CTTT) ₄ (TT) ₁ (CTTT) ₃ N ₄₇ (CTTT) ₄ (CT) ₁ (CTTT) ₄ (CCTT) ₁ (CTTT) ₃ N ₁₀ (CTTT) ₃ | (CTTT) ₄ (TTCT) ₁ (CTTT) ₃ (TTT) ₁ (CTTT) ₄ (TT) ₁ (CTTT) ₃ N ₄₇ (CTTT) ₃ (CT) ₁ (CTTT) ₄ (CCTT) ₁ (CTTT) ₃ N ₁₀ (CTTT) ₃ | 38 | 445 |
| DYS626 | (GAAA) ₁₉ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | (GAAA) ₂₀ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | 42 | 383 |
| DYS626 | (GAAA) ₁₆ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | (GAAA) ₁₇ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | 37 | 388 |
| DYS626 | (GAAA) ₁₇ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | (GAAA) ₁₈ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | 35 | 500 |
| DYS626 | (GAAA) ₁₈ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | (GAAA) ₁₇ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | 38 | 529 |
| DYS626 | (GAAA) ₁₉ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | (GAAA) ₁₈ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | 24 | 571 |
| DYS626 | (GAAA) ₁₈ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | (GAAA) ₁₉ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | 40 | 612 |
| DYS626 | (GAAA) ₁₇ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | (GAAA) ₁₈ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | 29 | 650 |
| DYS626 | (GAAA) ₁₉ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | (GAAA) ₂₀ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | 36 | 733 |
| DYS626 | (GAAA) ₁₈ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | (GAAA) ₁₉ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | 41 | 779 |
| DYS626 | (GAAA) ₂₀ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | (GAAA) ₁₉ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | 29 | 901 |

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|--------|---|---|----|------|
| DYS626 | (GAAA) ₂₀ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | (GAAA) ₂₁ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | 27 | 953 |
| DYS626 | (GAAA) ₁₉ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | (GAAA) ₂₀ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | 39 | 1071 |
| DYS626 | (GAAA) ₁₇ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | (GAAA) ₁₈ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | 17 | 1109 |
| DYS626 | (GAAA) ₁₉ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | (GAAA) ₂₀ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | 33 | 1112 |
| DYS626 | (GAAA) ₁₉ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | (GAAA) ₂₀ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | 27 | 1389 |
| DYS626 | (GAAA) ₁₉ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | (GAAA) ₁₈ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | 25 | 1445 |
| DYS626 | (GAAA) ₁₆ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | (GAAA) ₁₅ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | 25 | 1514 |
| DYS626 | (GAAA) ₁₈ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | (GAAA) ₁₉ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | 19 | 1530 |
| DYS626 | (GAAA) ₁₉ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | (GAAA) ₂₀ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | 59 | 1823 |
| DYS626 | (GAAA) ₂₂ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | (GAAA) ₂₁ N ₂₄ (GAAA) ₃ N ₆ (GAAA) ₅ (AAA) ₁ (GAAA) ₂ (GAAG) ₁ (GAAA) ₃ | 56 | 1907 |
| DYS627 | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₂₁ N ₈₁ (AAGG) ₃ | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₂₀ N ₈₁ (AAGG) ₃ | 25 | 4 |
| DYS627 | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₂₂ N ₈₁ (AAGG) ₃ | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₂₁ N ₈₁ (AAGG) ₃ | 36 | 49 |
| DYS627 | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₁₈ N ₈₁ (AAGG) ₃ | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₁₉ N ₈₁ (AAGG) ₃ | 29 | 82 |
| DYS627 | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₁₈ N ₈₁ (AAGG) ₃ | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₁₉ N ₈₁ (AAGG) ₃ | 39 | 112 |
| DYS627 | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₁₉ N ₈₁ (AAGG) ₃ | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₁₈ N ₈₁ (AAGG) ₃ | 27 | 170 |
| DYS627 | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₂₀ N ₈₁ (AAGG) ₃ | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₂₁ N ₈₁ (AAGG) ₃ | 34 | 243 |
| DYS627 | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₁₉ N ₈₁ (AAGG) ₃ | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₁₈ N ₈₁ (AAGG) ₃ | 20 | 256 |
| DYS627 | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₁₉ N ₈₁ (AAGG) ₃ | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₂₀ N ₈₁ (AAGG) ₃ | 21 | 328 |
| DYS627 | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₁₉ N ₈₁ (AAGG) ₃ | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₂₀ N ₈₁ (AAGG) ₃ | 31 | 331 |
| DYS627 | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₁₈ N ₈₁ (AAGG) ₃ | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₁₇ N ₈₁ (AAGG) ₃ | 37 | 355 |
| DYS627 | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₂₀ N ₈₁ (AAGG) ₃ | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₂₁ N ₈₁ (AAGG) ₃ | 23 | 496 |
| DYS627 | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₂₃ N ₈₁ (AAGG) ₃ | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₂₂ N ₈₁ (AAGG) ₃ | 35 | 500 |
| DYS627 | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₂₀ N ₈₁ (AAGG) ₃ | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₁₉ N ₈₁ (AAGG) ₃ | 36 | 619 |
| DYS627 | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₂₀ N ₈₁ (AAGG) ₃ | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₁₉ N ₈₁ (AAGG) ₃ | 25 | 711 |

| | | | | |
|--------|---|--|---------|------|
| DYS627 | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₁₉ N ₈₁ (AAGG) ₃ | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₁₈ N ₈₁ (AAGG) ₃ | 20 | 742 |
| DYS627 | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₁₉ N ₈₁ (AAGG) ₃ | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₂₀ N ₈₁ (AAGG) ₃ | 29 | 789 |
| DYS627 | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₁₉ N ₈₁ (AAGG) ₃ | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₂₀ N ₈₁ (AAGG) ₃ | Unknown | 1310 |
| DYS627 | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₁₅ N ₈₁ (AAGG) ₃ | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₁₆ N ₈₁ (AAGG) ₃ | 22 | 1323 |
| DYS627 | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₁₈ N ₈₁ (AAGG) ₃ | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₁₉ N ₈₁ (AAGG) ₃ | 42 | 1407 |
| DYS627 | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₂₂ N ₈₁ (AAGG) ₃ | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₂₃ N ₈₁ (AAGG) ₃ | 17 | 1416 |
| DYS627 | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₁₉ N ₈₁ (AAGG) ₃ | (AGAA) ₃ N ₁₆ (AGAG) ₃ (AAAG) ₂₀ N ₈₁ (AAGG) ₃ | 54 | 1860 |
| DYS629 | (TATC) ₉ | (TATC) ₁₀ | 29 | 609 |
| DYS630 | (AAAG) ₄ (AGAG) ₃ N ₁₈ (AAAG) ₁₄ | (AAAG) ₄ (AGAG) ₃ N ₁₈ (AAAG) ₁₅ | 21 | 46 |
| DYS630 | (AAAG) ₄ (AGAG) ₃ N ₁₈ (AAAG) ₁₅ | (AAAG) ₄ (AGAG) ₃ N ₁₈ (AAAG) ₁₄ | 33 | 53 |
| DYS630 | (AAAG) ₄ (AGAG) ₃ N ₁₈ (AAAG) ₁₈ | (AAAG) ₄ (AGAG) ₃ N ₁₈ (AAAG) ₁₇ | 23 | 96 |
| DYS630 | (AAAG) ₄ (AGAG) ₃ N ₁₈ (AAAG) ₁₆ | (AAAG) ₄ (AGAG) ₃ N ₁₈ (AAAG) ₁₅ | 40 | 255 |
| DYS630 | (AAAG) ₄ (AGAG) ₃ N ₁₈ (AAAG) ₁₅ | (AAAG) ₄ (AGAG) ₃ N ₁₈ (AAAG) ₁₆ | 30 | 448 |
| DYS630 | (AAAG) ₄ (AGAG) ₃ N ₁₈ (AAAG) ₁₃ | (AAAG) ₄ (AGAG) ₃ N ₁₈ (AAAG) ₁₄ | 21 | 478 |
| DYS630 | (AAAG) ₄ (AGAG) ₃ N ₁₈ (AAAG) ₁₇ | (AAAG) ₄ (AGAG) ₃ N ₁₈ (AAAG) ₁₈ | 39 | 501 |
| DYS630 | (AAAG) ₄ (AGAG) ₃ N ₁₈ (AAAG) ₁₄ | (AAAG) ₄ (AGAG) ₃ N ₁₈ (AAAG) ₁₅ | 36 | 665 |
| DYS631 | (AATA) ₄ (CATA) ₁ (AATA) ₁₁ | (AATA) ₄ (CATA) ₁ (AATA) ₁₀ | 47 | 347 |
| DYS635 | (TCTA) ₄ (TGTA) ₂ (TCTA) ₂ (TGTA) ₂ (TCTA) ₂ (TCTA) ₁₂ | (TCTA) ₄ (TGTA) ₂ (TCTA) ₂ (TGTA) ₂ (TCTA) ₂ (TCTA) ₁₁ | 53 | 35 |
| DYS635 | (TCTA) ₄ (TGTA) ₂ (TCTA) ₂ (TGTA) ₂ (TCTA) ₂ (TCTA) _{13,14} | (TCTA) ₄ (TGTA) ₂ (TCTA) ₂ (TGTA) ₂ (TCTA) ₂ (TCTA) ₁₃ | 33 | 528 |
| DYS635 | (TCTA) ₄ (TGTA) ₂ (TCTA) ₂ (TGTA) ₂ (TCTA) ₂ (TCTA) ₁₃ | (TCTA) ₄ (TGTA) ₂ (TCTA) ₂ (TGTA) ₂ (TCTA) ₂ (TCTA) ₁₂ | 36 | 617 |
| DYS635 | (TCTA) ₄ (TGTA) ₂ (TCTA) ₂ (TGTA) ₂ (TCTA) ₂ (TCTA) ₁₂ | (TCTA) ₄ (TGTA) ₂ (TCTA) ₂ (TGTA) ₂ (TCTA) ₂ (TCTA) ₁₁ | 26 | 800 |
| DYS635 | (TCTA) ₄ (TGTA) ₂ (TCTA) ₂ (TGTA) ₂ (TCTA) ₂ (TCTA) ₁₂ | (TCTA) ₄ (TGTA) ₂ (TCTA) ₂ (TGTA) ₂ (TCTA) ₂ (TCTA) ₁₁ | 29 | 1674 |
| DYS635 | (TCTA) ₄ (TGTA) ₂ (TCTA) ₂ (TGTA) ₂ (TCTA) ₂ (TCTA) ₁₁ | (TCTA) ₄ (TGTA) ₂ (TCTA) ₂ (TGTA) ₂ (TCTA) ₂ (TCTA) ₁₂ | 52 | 1891 |
| DYS637 | (AAAT) ₄ (ACAT) ₁₁ | (AAAT) ₄ (ACAT) ₁₀ | 25 | 950 |
| DYS638 | (TTTA) ₁₁ | (TTTA) ₁₂ | 56 | 1677 |
| DYS643 | (AAAT) ₁₁ | (AAAT) ₁₂ | 32 | 95 |
| DYS643 | (AAAT) ₁₃ | (AAAT) ₁₄ | 19 | 1697 |
| DYS644 | (TTTTA) ₁₀ (TTTTA) ₇ | (TTTTA) ₁₀ (TTTTA) ₈ | 22 | 487 |
| DYS644 | (TTTTA) ₁₀ (TTTTA) ₆ | (TTTTA) ₁₀ (TTTTA) ₅ | 24 | 681 |
| DYS644 | (TTTTA) ₁₀ (TTTA) ₁ (TTTTA) ₁₃ | (TTTTA) ₁₁ (TTTA) ₁ (TTTTA) ₁₃ | 21 | 1667 |

| | | | | |
|------------|--|--|----|------|
| DYS644 | (TTTTA) ₁₀ (TTTFA) ₆ | (TTTTA) ₁₀ (TTTFA) ₇ | 19 | 1717 |
| DYS644 | (TTTTA) ₁₀ (TTTFA) ₇ | (TTTTA) ₁₀ (TTTFA) ₆ | 50 | 1832 |
| Y-GATA-A10 | (ATCT) ₁₃ | (ATCT) ₁₄ | 41 | 417 |
| Y-GATA-A10 | (ATCT) ₁₄ | (ATCT) ₁₅ | 35 | 735 |
| Y-GATA-A10 | (ATCT) ₁₃ | (ATCT) ₁₂ | 24 | 855 |
| Y-GATA-A10 | (ATCT) ₁₃ | (ATCT) ₁₂ | 46 | 1252 |
| Y-GATA-A10 | (ATCT) ₁₃ | (ATCT) ₁₄ | 40 | 1606 |
| Y-GATA-H4 | (TAGA) ₃ N ₁₂ (TAGG) ₃ (TAGA) ₁₂ N ₂₂ (TAGA) ₄ | (TAGA) ₃ N ₁₂ (TAGG) ₃ (TAGA) ₁₁ N ₂₂ (TAGA) ₄ | 23 | 251 |
| Y-GATA-H4 | (TAGA) ₃ N ₁₂ (TAGG) ₃ (TAGA) ₁₁ N ₂₂ (TAGA) ₄ | (TAGA) ₃ N ₁₂ (TAGG) ₃ (TAGA) ₁₂ N ₂₂ (TAGA) ₄ | 19 | 1004 |
| Y-GATA-H4 | (TAGA) ₃ N ₁₂ (TAGG) ₃ (TAGA) ₁₂ N ₂₂ (TAGA) ₄ | (TAGA) ₃ N ₁₂ (TAGG) ₃ (TAGA) ₁₁ N ₂₂ (TAGA) ₄ | 29 | 1051 |
| Y-GATA-H4 | (TAGA) ₃ N ₁₂ (TAGG) ₃ (TAGA) ₁₃ N ₂₂ (TAGA) ₄ | (TAGA) ₃ N ₁₂ (TAGG) ₃ (TAGA) ₁₂ N ₂₂ (TAGA) ₄ | 42 | 1411 |
| Y-GATA-H4 | (TAGA) ₃ N ₁₂ (TAGG) ₃ (TAGA) ₁₃ N ₂₂ (TAGA) ₄ | (TAGA) ₃ N ₁₂ (TAGG) ₃ (TAGA) ₁₂ N ₂₂ (TAGA) ₄ | 53 | 1799 |

The repeat structure of both the father and son's alleles at the mutated Y-STR are given where possible. In the case of multicopy markers with multiple variable segments within the STR, total repeat numbers or amplicon size is given in the absence of sequence information. The age of the father at the time of the son's birth is given, as is an individual pair reference.

Table S3. Ability of 13 Rapidly-Mutating RM Y-STRs and 17 Yfiler Y-STRs to Differentiate between Pairs of Male Relatives by One or More Mutations

| Number of Meioses Separating Pair | RM Y-STR Mutations | RM Y-STR Locus Comparisons | Yfiler Mutations | Yfiler Locus Comparisons |
|--|---------------------------|-----------------------------------|-------------------------|---------------------------------|
| 1 | 1 | 9 | 0 | 17 |
| 1 | 1 | 12 | 0 | 17 |
| 1 | 1 | 12 | 0 | 17 |
| 1 | 1 | 11 | 0 | 17 |
| 1 | 1 | 10 | 0 | 17 |
| 1 | 1 | 12 | 0 | 17 |
| 1 | 2 | 5 | 0 | 17 |
| 1 | 0 | 13 | 0 | 17 |
| 1 | 1 | 12 | 0 | 17 |
| 1 | 0 | 13 | 0 | 17 |
| 1 | 0 | 13 | 0 | 17 |
| 1 | 1 | 13 | 0 | 17 |
| 1 | 0 | 13 | 0 | 17 |
| 1 | 0 | 13 | 0 | 17 |
| 1 | 3 | 11 | 0 | 17 |
| 1 | 1 | 10 | 0 | 17 |
| 1 | 0 | 13 | 0 | 17 |
| 1 | 1 | 10 | 0 | 17 |
| 1 | 1 | 4 | 0 | 17 |
| 1 | 1 | 12 | 0 | 17 |
| 2 | 1 | 12 | 0 | 17 |
| 2 | 2 | 11 | 0 | 17 |
| 2 | 1 | 8 | 0 | 17 |
| 2 | 2 | 11 | 0 | 17 |
| 2 | 2 | 13 | 0 | 17 |
| 2 | 1 | 9 | 0 | 17 |
| 2 | 2 | 13 | 0 | 17 |
| 2 | 0 | 13 | 0 | 17 |
| 2 | 0 | 13 | 0 | 17 |
| 2 | 0 | 13 | 0 | 17 |
| 2 | 1 | 13 | 0 | 17 |
| 2 | 0 | 13 | 0 | 17 |
| 2 | 0 | 13 | 0 | 17 |
| 2 | 1 | 12 | 0 | 17 |
| 2 | 3 | 13 | 0 | 17 |
| 2 | 0 | 13 | 0 | 17 |

| | | | | |
|---|---|----|---|----|
| 2 | 0 | 13 | 0 | 17 |
| 2 | 0 | 13 | 1 | 17 |
| 2 | 0 | 13 | 0 | 17 |
| 2 | 0 | 13 | 0 | 17 |
| 2 | 3 | 13 | 0 | 17 |
| 2 | 0 | 12 | 0 | 17 |
| 2 | 4 | 13 | 0 | 17 |
| 2 | 1 | 13 | 0 | 17 |
| 2 | 3 | 13 | 0 | 17 |
| 2 | 0 | 13 | 0 | 17 |
| 2 | 0 | 13 | 0 | 17 |
| 2 | 0 | 13 | 0 | 17 |
| 2 | 1 | 13 | 0 | 17 |
| 2 | 0 | 13 | 0 | 17 |
| 2 | 1 | 10 | 1 | 17 |
| 2 | 1 | 13 | 0 | 17 |
| 2 | 1 | 13 | 0 | 17 |
| 2 | 2 | 3 | 0 | 17 |
| 3 | 0 | 13 | 0 | 17 |
| 3 | 0 | 13 | 0 | 17 |
| 3 | 0 | 13 | 0 | 17 |
| 3 | 2 | 12 | 0 | 17 |
| 3 | 2 | 12 | 0 | 17 |
| 3 | 2 | 13 | 0 | 17 |
| 3 | 3 | 13 | 0 | 17 |
| 4 | 0 | 13 | 0 | 17 |
| 4 | 1 | 13 | 0 | 17 |
| 4 | 1 | 13 | 0 | 17 |
| 5 | 1 | 5 | 0 | 17 |
| 5 | 1 | 13 | 0 | 17 |
| 5 | 1 | 12 | 0 | 17 |
| 5 | 2 | 12 | 0 | 17 |
| 6 | 3 | 9 | 0 | 17 |
| 6 | 1 | 10 | 0 | 17 |
| 6 | 1 | 13 | 2 | 17 |
| 6 | 5 | 12 | 1 | 17 |
| 6 | 3 | 13 | 0 | 17 |
| 6 | 4 | 13 | 0 | 17 |
| 6 | 3 | 13 | 0 | 17 |
| 6 | 0 | 13 | 0 | 17 |
| 6 | 2 | 13 | 0 | 17 |

| | | | | |
|----------------|-------------|--------------|-------------|-------------|
| 7 | 0 | 13 | 0 | 17 |
| 7 | 4 | 13 | 1 | 17 |
| 8 | 3 | 13 | 0 | 17 |
| 8 | 4 | 13 | 0 | 17 |
| 8 | 2 | 13 | 0 | 17 |
| 8 | 0 | 13 | 0 | 17 |
| 8 | 0 | 13 | 1 | 17 |
| 8 | 4 | 13 | 0 | 17 |
| 8 | 2 | 13 | 0 | 17 |
| 9 | 1 | 13 | 1 | 17 |
| 10 | 1 | 13 | 0 | 17 |
| 10 | 4 | 12 | 1 | 17 |
| 10 | 2 | 13 | 0 | 17 |
| 10 | 3 | 13 | 0 | 17 |
| 10 | 3 | 13 | 1 | 17 |
| 10 | 1 | 12 | 2 | 17 |
| 10 | 0 | 12 | 1 | 17 |
| 11 | 6 | 13 | 0 | 17 |
| 11 | 6 | 13 | 0 | 17 |
| 11 | 3 | 12 | 0 | 17 |
| 11 | 4 | 13 | 2 | 17 |
| 11 | 1 | 13 | 1 | 17 |
| 11 | 3 | 13 | 0 | 17 |
| 13 | 4 | 12 | 1 | 17 |
| 13 | 5 | 13 | 0 | 17 |
| 20 | 4 | 13 | 0 | 17 |
| Total | 158 | 1246 | 17 | 1751 |
| Average | 1.53 | 12.10 | 0.17 | 17 |