

ANNUAL REPORT SUMMARY FOR TESTING IN 2006
Prepared by the Relationship Testing Standards and Accreditation Program Units

PREFACE

The 2006 Annual Report includes respondents from accredited laboratories in the United States, Canada and Europe. Many of the laboratories report testing a broad range of cases, including relationship tests for routine paternity testing, immigration, as well as prenatal and postmortem evaluations. Almost all of the laboratories responding to the survey performed immigration testing and reconstruction (family study) cases. Two of the reporting laboratories indicated that they sent their cases to other laboratories for testing and, as such, the data presented here are from the remaining laboratories that performed the testing.

In this report, AABB provides some commentary for laymen on common misconceptions in paternity testing. Some of the commentary is from the previous year's report, as the commentary remains relevant to issues raised this year.

On January 1, 2008, the 8th edition of *Standards for Relationship Testing Laboratories* went into effect. The Relationship Testing Standards Program Unit would also like to remind readers that the *Guidance for Standards for Relationship Testing Laboratories*, published in 2008, discusses the *Standards* in detail and provides suggestions on how to comply with the standards and contains explanations of the requirements, the various types of calculations used, and addresses other issues in relationship testing.

ANNUAL VOLUME OF TESTING

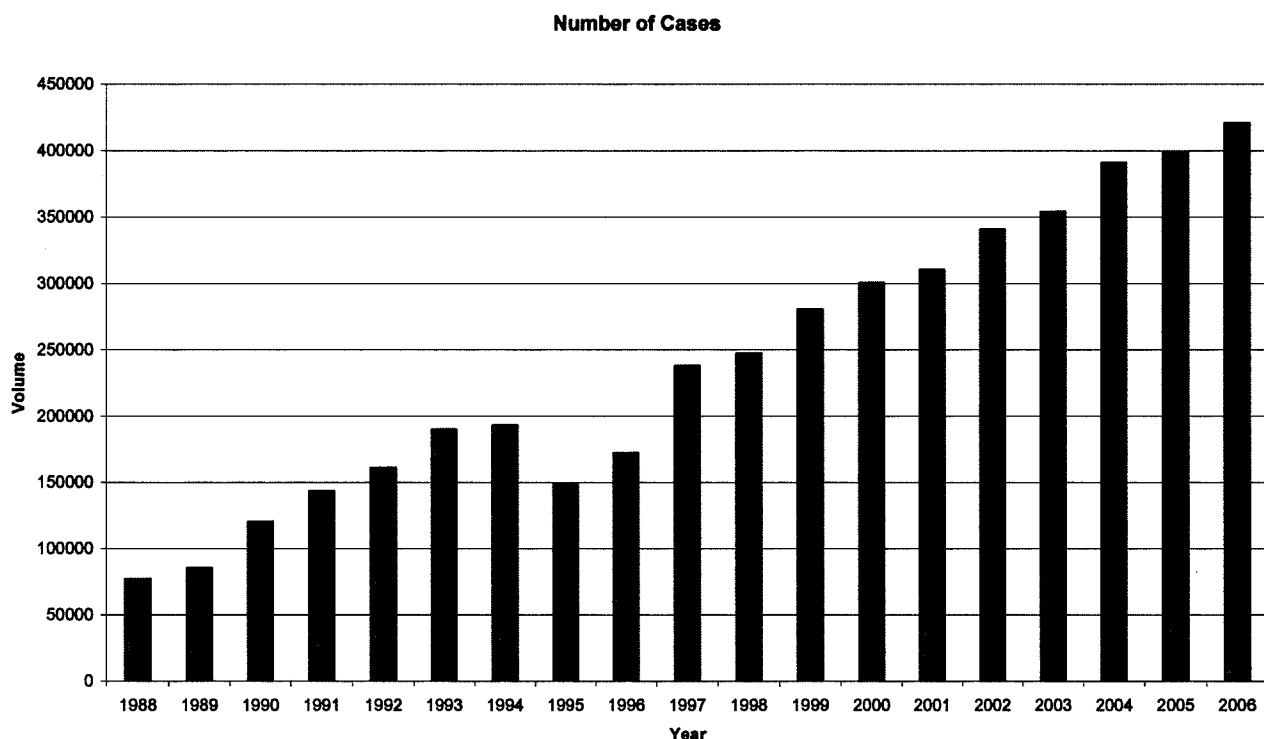
The volume reported for cases tested in 2006 was 420,740. Note that, for some laboratories, only the total number of cases reported is available. The statistics in this report therefore are based on a smaller number of laboratories, as indicated above. The total volume of cases represents an increase of 21,860 cases (or 5.48% increase) from the 2005 volume. A summary of the totals of all years since 1988 is shown in Table 1 and Figure 1.

Table 1. The Number of Relationship Cases Reported for 1988-2006.

Year	No. of Cases	Year	No. of Cases
1988	77000	1998	247317
1989	85231	1999	280510
1990	120436	2000	300626

1991	143459	2001	310490
1992	161000	2002	340798
1993	189904	2003	354011
1994	193000	2004	390928
1995	149100	2005	398880
1996	172316	2006	420740
1997	237981		

Figure 1. Graph of the Case Volume for 1988-2006.



In this report, laboratories were asked if they were testing cases where the chain of custody did not meet the requirements of the *Standards for Relationship Testing*. These so-called “non-legal” tests are generally collected by the individuals being tested, and are not “witnessed by a competent person with no interest in the test outcome” (as required by standard 5.2.2 in the *Standards for Relationship Testing Laboratories*, 8th edition.)

The *Standards* does not prohibit accredited laboratories from performing these types of tests, but because “non-legal” tests do not meet the requirements of *Standards*, laboratories cannot claim or advertise that their “non-legal” testing is encompassed by their AABB accreditation – regardless of whether the testing of the samples conforms to *Standards*. Particularly in relationship testing, the quality of the results depends on the testing just as much as it does on the integrity of the sample collection process. Of the laboratories reporting, over half (53%) reported that they performed “non-legal” testing. Those laboratories reported 19,582 non-legal cases or 5.55% of the total cases reported. Some

laboratories did not track the number of non-legal cases they evaluated, but it seems appropriate to estimate that no more than 10% of all cases were of a “non-legal” type.

Note that during 2007, the Department of Homeland Security, United States Citizenship and Immigration Service (USCIS) met with the committee to discuss this and other issues. Because the sample collection in “non-legal” tests is not controlled, there is the potential for fraud. Accordingly, “non-legal” testing is not acceptable for immigration purposes.

LABORATORIES BY SIZE

Table 2 indicates the size of the various responding laboratories by volume of cases reported. Note that this breakdown reflects individual laboratories, but that a single corporation may own several laboratories. There appears to be a decrease in the number of laboratories reporting 1-500 cases per year.

Table 2. Laboratories by the Volume of Cases Reported.

Case Volumes	1994	1995	1996	1997	1998	1999	2000	2001	2002	2003	2004	2005	2006
1-500	40	26	25	20	19	19	13	17	14	18	16	16	8
501-1,000	6	4	8	7	6	5	6	6	2	3	2	4	3
1,001-5,000	7	9	6	10	11	9	11	11	13	11	7	8	8
5,001-10,000	6	4	3	5	0	3	3	5	1	3	7	7	5
10,001-50,000	1	2	3	5	5	7	8	6	7	7	6	5	5
50,001-100,000	2	1	1	1	2	1	1	1	0	0	1	1	0
>100,000	0	0	0	0	0	0	0	0	1	1	1	1	1
Total Laboratories	62	46	46	48	43	44	42	46	38	43	40	42	30

EXCLUSION RATE

In 2006, two laboratories did not track the number of exclusions. For the laboratories tracking exclusions, there were 347,719 cases completed, and 89,890 of those (25.85%) were reported as exclusions. The average exclusion rate for the laboratories reporting exclusions is 23.13% with a standard deviation of 5.64. The median exclusion rate is 25.06% with a range of 10.99% to 31.83%. The explanation for the range of exclusion rates is complex but appears to be related to the laboratory’s volume and client base. Anecdotal explanations for the various exclusion rates include differences with the type of case (private versus public contracts), and the geographic source of the case (rural versus

metropolitan areas). For the non-legal testing laboratories, there were 4,579 exclusions from laboratories reporting exclusion data (total of 15,082 cases) or an exclusion rate of 30.36%, a higher percentage than the 26% seen for legal testing. The range for non-legal laboratories is 10.99 to 35.04%.

MISCONCEPTIONS IN PATERNITY TESTING – EXCLUSION RATE

AABB has seen the exclusion rate misused by several organizations trying to claim that 30% of men are misled into believing they are biological fathers of children. This claim is incorrect. The exclusion rate includes a number of factors. One is that a case may include several men as alleged fathers because the mother was sexually active with these individuals. These are not men who were misled into believing they were fathers and then later discover they are not. The testing merely determines which man is the likely biological father while excluding the others. Another factor is that the unexcluded alleged father, as part of his defense, may allege that the mother had multiple sexual partners during the time of conception. These other men are subsequently tested. Sometimes testing of a man is required because of a legal presumption. This is when the mother properly names the correct biological father but because the child is the product of a marriage (for example, the mother was married to someone other than the biological father at the time of conception) there is a legal presumption that the husband is the father. The husband is then tested to rebut the legal presumption even though no one believes he is the biological father of the child. There is no evidence that a large number of the men excluded in the testing were misled into believing they are the biological father of a given child.

COMBINED PATERNITY INDEX

The laboratories were asked to indicate what combined paternity index (CPI) they considered acceptable for cases with a standard trio (mother, child, father), single parent cases (mother, or father, not tested cases), and reconstruction cases (disputed parent is missing and other relatives are used to evaluate parentage). Some laboratories reported using different CPIs for different classes of clients (private versus public contracts, or for different technologies). For these laboratories, the higher CPI was used for this report.

The results for the laboratories that responded are shown in Table 3. The most common minimum CPI for a standard trio is 100, with 55% of laboratories using this value, with a range of 100 to 100,000. For cases where the mother is not tested, the most common minimum CPI is 100 with 62% of laboratories using this value, with a range of 100 to 10,000. For the family study or reconstruction cases, 58% indicated that they report “whatever was obtained” and 81% considered a combined paternity index of 101 or less reportable. For sibling studies about 90% of the laboratories considered a combined paternity index of 101 or less reportable.

A common issue is the significance of the paternity index and the reliability of the AABB standard requiring a CPI of 100 to 1. At least one laboratory has claimed that AABB is only concerned about how the testing is performed, but not the meaning of the test. In fact, the Relationship Testing Standards Program Unit is concerned about the meaning of the tests and thus chose a CPI of 100 to 1. First and foremost, this level was chosen because it provides reasonable evidence of paternity in a standard case where a trio is tested. Generally, when a laboratory tests a case, if the disputed person is not excluded and does not reach the laboratory's minimum value, additional testing is performed to evaluate this person. This additional testing may result in non-exclusion, exclusion, or inconclusive reports.

Another issue arises with regard to performing other relationship analyses, such as reconstruction cases, trios with genetic anomalies, and samples from exhumations, coroners, and other postmortem testing. Importantly, note that a CPI of less than 100 is not an indicator of no relationship, unless the CPI is 0 (or much less than 1), and may still in fact be a strong indicator of a relationship. Practical difficulties exist with the ability to obtain results from degraded samples, which are typically used in postmortem testing, and in the mathematical analysis of the relationships in reconstruction cases. Understanding this is particularly important for legislators who establish presumption levels based on paternity calculations, and contract administrators, since testing is often performed in conditions that are not ideal. Another important concept is that a laboratory's minimum combined paternity index, which may reflect scientific reality, is not necessarily the laboratory's testing goal or median combined paternity index.

Table 3. The Number of Laboratories Using Various Minimum Combined Likelihood Ratios for Standard Trios, One Parent (Mother, or father, Not Tested (MNT)) and Reconstruction Cases (Note: not all laboratories indicated a CPI for each type of case).

CPI	Trio	One Parent	Family Study (Reconstruction)	Full Sib	Half Sib
Whatever Is Obtained	0	0	15	12	12
10	0	0	0	3	1
25	0	0	0	0	1
100	16	18	5	2	1
101	0	0	1	0	0
150	1	1	0	0	0
200	2	2	2	1	1
400	0	0	0	0	0
500	2	1	0	0	0
1000	3	3	2	1	0

1001	1	1	0	0	0
2500	2	1	1	0	0
10000	1	2	0	0	0
100000	1	0	0	0	0
Sum	29	29	26	19	16

TESTING WITHOUT THE MOTHER

There is still a strong concern about the apparent increase in the number of clients submitting disputed paternity cases without the mother. Testing without the mother presents a number of problems. First, the paternity index, on average, is cut in half. This also greatly reduces the ability to detect a falsely accused man, and in some cases, such as incest, can easily produce false inclusions. When an apparent inconsistency (mutation) is present, it may not be possible to render an opinion of paternity without obtaining a sample from the mother. Obtaining a sample from the mother is also an important quality control step, because results that exclude the mother may indicate a problem with the testing. The testing of the mother may also allow for the detection of fraud, such as welfare fraud on the part of the mother or cases where the alleged father brings a child he knows is his, but who is not the child of the mother. Thus, the testing of the mother, even if maternity is not disputed, is important in evaluating the questioned relationship, because it improves the chance of obtaining clear results and is a quality control check for both the scientific and legal community. Testing without the mother should only be done when the mother's location is unknown or she is deceased. Every effort should be made to test the mother.

TECHNOLOGY USE

In 2006, the survey showed a continued trend toward the increased use of polymerase chain reaction (PCR) technology (STR analysis) with a decrease in the use of restriction fragment length polymorphism (RFLP) methods. PCR technology was used in 98.53% of reported cases and RFLP was used in 1.12% of reported cases. This is also the first year that no cases were evaluated using serological HLA testing.

Table 4 provides a breakdown of the technology used to resolve the reported paternity cases. The three laboratories using HLA molecular methods were asked to identify the source of the frequencies. Laboratories using HLA molecular for Class I HLA methods reported using serologic tables for calculating paternity indices.

Table 4. The Technology Used in Cases Reported in 2006

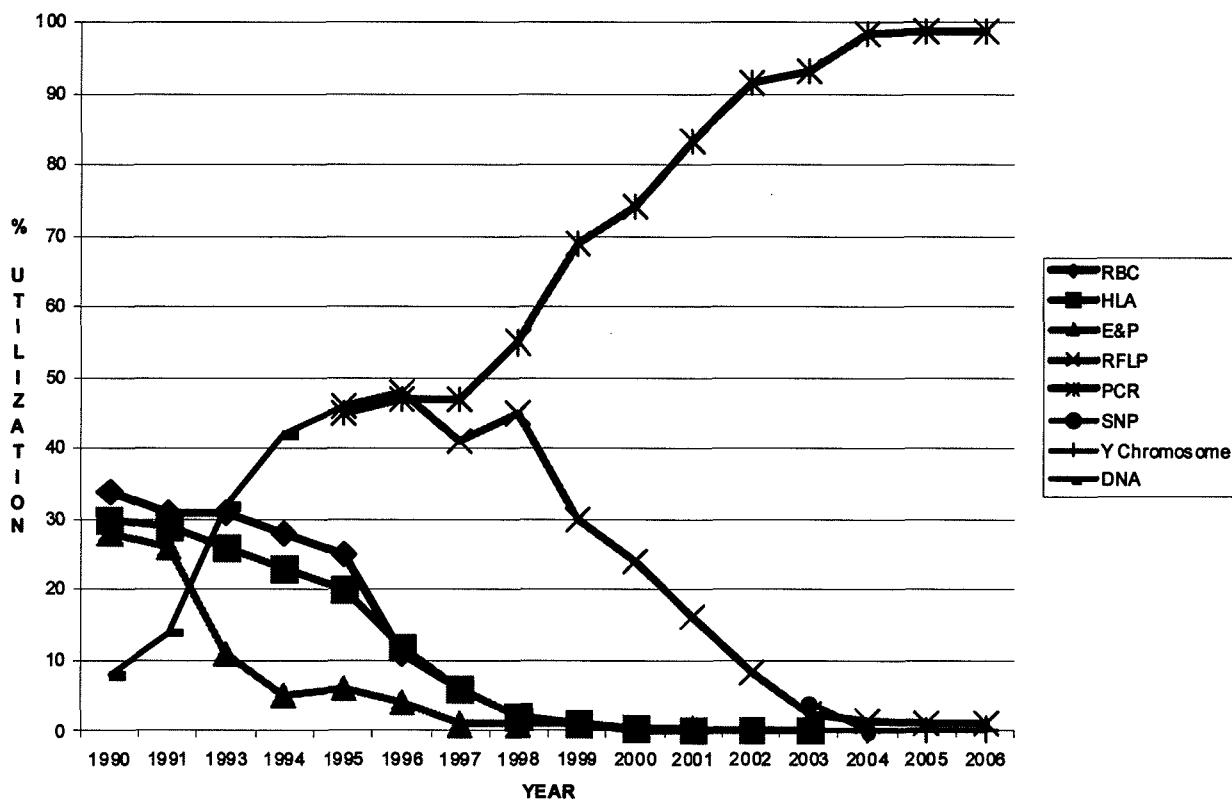
Technology	Number of Cases	Utilization (%)
STR	344616	98.53

RFLP	3906	1.12
HLA Class II Molecular	35	0.01
Y Chromosome	1079	0.31
HLA Class I Molecular	139	0.04
SNP	0	0
HLA Serology	0	0
Red Cell Antigens	0	0
Red Cell Enzymes/	0	0
Serum Proteins		
Allotyping	0	0

*Note that some cases used more than one technology. Not all laboratories responded to this question.

Figure 2 shows the use of various technologies since 1990. As indicated above, the most commonly used technologies in 1990 (red cell antigens, HLA, and red cell enzymes and serum proteins) now account for less than 1% of all casework. The change in DNA technologies from RFLP to PCR technology is also obvious. Prior to 1995, the survey only asked about the use of DNA testing but not about which DNA technology was used (PCR verses RFLP). Note that in some instances, multiple technologies were used in the same case.

Figure 2. The Use of Various Technologies Since 1990.



SAMPLE SOURCE

Laboratories reported approximately 812,177 samples used for the casework in 2006. Not all laboratories reported the samples they used. Of these samples, buccal swabs accounted for 98.05% of the samples. Whole blood samples accounted for .64%. The use of blood spot cards decreased to 1.16% of samples. Various other samples were also reported (See Table 5).

Table 5. Sample Source in 2006

Sample	Number	Percent of Total
Buccal Swabs	812,177	98.0501
Blood	5,278	0.6372
Blood Spot Cards	9,641	1.1639
Amniotic Fluid	693	0.0837

Misc. Tissues	184	0.0222
Paraffin Blocks	10	0.0012
Hair	93	0.0112
CVS	225	0.0272
Products of Conception	15	0.0018
Bone	13	0.0016
Total	828,329	100

AMELOGENIN

The amelogenin locus is now used in a number of laboratories to test for the gender of the sample. A number of males lacking the Y or X amelogenin allele have been observed. Laboratories were asked to measure the apparent X males observed in their laboratory. Like other DNA loci, amelogenin is subject to mutations. Therefore, occasionally normal males have a female amelogenin phenotype or a Y phenotype. The X male phenotype was most commonly seen in the Hispanic populations, in about 1/1392 men. The Y male phenotype was most commonly seen in the Black population in about 1/1688 Black males.

Table 6. A Summary of Data on Apparent X and Y Males Seen with ABI Primers

	Black	White	Hispanic
Number X Males Observed	6	12	16
%	0.0099	0.0254	0.0718
Number Y Males Observed	36	6	4
%	0.0592	0.0127	0.0180

MUTATION REPORTS

One area of concern is the number of inconsistencies necessary to render an opinion of non-paternity. The laboratories were asked if they had seen any case where, in the opinion of the expert, the inconsistencies were double or triple "mutations" and not sufficient to render an opinion of non-paternity. Twelve laboratories stated they had reported cases with double or triple mutations. Eleven laboratories did not observe any mutations. The

laboratories reported 47 cases with double mutations and no cases with triple mutations as inclusions. Most laboratories report these “double mutation” cases with the inconsistencies noted and statistically considered. This illustrates the importance of accurate assessments of potential mutations and null alleles.

MUTATION CALCULATION AND FREQUENCIES

Single inconsistencies are routinely seen in the testing of paternity cases. If a laboratory comes to the conclusion that the inconsistency is a mutation, then the mutation result must be incorporated into the reported results. Laboratories were asked how they calculated the paternity index (PI) for these loci. The laboratories all appear to be using one of several calculation methods. Some laboratories are using the mutation rate as the PI, while others, most commonly, used the mutation rate divided by the average probability of exclusion. Some laboratories used the mutation rate as a transmission frequency and some of the laboratories used Brenner’s method in looking at the repeat length difference between STR alleles.

A summary of the mutation frequencies for each STR locus is provided in Appendix 1. Note that these frequencies incorporate the indeterminate findings. The calculations are summarized at the bottom of the table for each paternal allele as shown in Appendix 2. The frequencies for changes from one allele to another are presented in Appendix 2 Appendix 3.

A continuing objective of this year’s report is to begin to collect data on STR loci to provide laboratories with frequencies to use in the mutation calculation. The guidance document for the 7th and 8th edition of standards contains a discussion of two methods that might be useful. One limitation of this data set is if the laboratory did not see any mutations, the laboratory did not provide data on the maternal and paternal meiosis. Many laboratories did not provide any data so the data presented is from a few laboratories.

If one wished to determine the specific mutation frequency at locus D3S1358 for the apparent paternal mutation event of the alleged father’s allele 16 changing to an allele 17 in the child. Using the attached table specific calculations could be made. Suppose that there are 16 instances where, simply, 16 changed to 17 out of 79247 meiosis reported or a frequency of 0.000202. However there are several other opportunities for this change. If there were five instances where the alleged father’s 16 could have changed to either a 15 or 17 (child is a clone of the mother or mother was not tested). To incorporate this data one approach is to calculate the relative chance that the change was 16 to 17 rather than 16 to 15. Note the clear changes and calculate the relative chance of each change. Multiply the relative chance times the number of changes where the allele is 16 to 15 or 17, which is 5 in this data set, to obtain the relative portion attributable to a 16 to 17 change.

Table 6. Relative Chance of allele 16 changing to 15 or 17.

Change	Observed	Relative Chance	Portion of 5
16 to 17	16	$16/31 = 0.516$	$5 * 0.516 = 2.58$
16 to 15	15	$15/31 = 0.484$	$5 * 0.484 = 2.42$
Total	31	1	5

From this data, add 2.56 to the 16 observed potential changes from 16 to 17 to get the total of 18.56. Similarly, there were seven observations where the alleged father has alleles 16 and 18 either of which could mutate to a 17.

Table 7. Relative Chance of allele 16 or 18 changing 17.

Change	Observed	Relative Chance	Portion of 7
16 to 17	16	$16/26 = 0.615$	$7 * 0.615 = 4.305$
18 to 17	10	$10/26 = 0.385$	$7 * 0.385 = 2.695$
Total	26	1	5

From this data, add 4.305 to the 18.56 potential changes (paragraph above) from 16 to 17 to get the total of 22.865.

Hypothetically, there were instances where the father's alleles 16 and 19 could have changed to a 17 or 18. To incorporate this data a similar approach is used.

Table 8. Relative Chance of allele 16 or 19 changing to 17 or 18.

Change	Observed	Relative Chance	Portion of 1
16 to 17	16	$16/21 = 0.762$	$1 * 0.762 = 0.762$
16 to 18	0	$0/21 = 0$	$1 * 0 = 0$
19 to 17	0	$0/21 = 0$	$1 * 0 = 0$
19 to 18	5	$5/21 = 0.238$	$1 * 0.238 = 0.238$
Total	31	1	1

From this data, add 0.762 to the 22.865 above yielding 23.627.

Lastly, data from those cases where the mutation is either maternal or paternal may be incorporated (indeterminate). Hypothetically there were 7 instances where the mutation to a 17 could have been from a paternal 16. The approach to incorporate these data is similar to the above. First look to the data to determine the frequency of the changes.

Table 9. Relative Chance of allele 16 changing to 17.

Change	Observed	Relative Chance	Portion of 7

16 to 17 Maternal	$1 / 67521 = 1.481\text{e-}5$	$1.481\text{e-}5 / 2.167\text{e-}4 = 0.0683$	$7 * 0.0683 = 0.478$
16 to 17 Paternal	$16 / 79247 = 2.019\text{e-}4$	$2.019\text{e-}4 / 2.167\text{e-}4 = 0.9317$	$7 * 0.9317 = 6.522$
Total	$2.167\text{e-}4$	1	7

Finally add 6.522 to the 23.627 yielding 30.149. Thus for this hypothetical population the frequency of paternal mutation from a 16 to a 17 is $30.149 / 79247 = 0.00038$ as compared to the 0.000202 without incorporating all possible mutation events. The committee invites comments on alternative methods of determining the mutation frequencies.

Appendix 1. Summary of Apparent Mutations at various Loci analyzed by PCR in 2006.

	Caucasian		Black		Hispanic	
	Paternal	Maternal	Paternal	Maternal	Paternal	Maternal
D8S1179	0.0024079	0.0002582	0.0027566	0.0004389	0.0021734	0.0005536
D21S11	0.0018992	0.0015278	0.0020646	0.0016688	0.0016599	0.0021119
D7S820	0.0019085	0.0001861	0.0014712	0.0002058	0.0017771	0.0001206
CSF1PO	0.0022249	0.0003272	0.0025281	0.0003962	0.0022446	0.0006719
D3S1358	0.0023268	0.0002622	0.0022779	0.0002555	0.0019073	0.0001514
THO1	0.0000676	0.0000785	0.0000818	0.0001146	<0.0000570	0.0000445
D13S317	0.0021998	0.0001748	0.0023244	0.0007426	0.0016911	0.0003627
D16S539	0.0021727	0.0003588	0.0019858	0.0005468	0.0012631	0.0003971
D2S1338	0.0022049	0.0003617	0.0015539	0.0004469	0.0014671	0.0012259
D19S453	0.0017659	0.0004243	0.0020252	0.0007177	0.0011208	<0.0002909
VWA	0.0044028	0.0004994	0.0036031	0.0006894	0.0026509	0.0003165
TPOX	0.0001878	0.0000702	0.0001548	0.0001012	0.0001143	<0.0000539
D18S51	0.0023140	0.0007041	0.0044855	0.0008289	0.0019579	0.0008343
D5S818	0.0021447	0.0003833	0.0023605	0.0003969	0.0019594	0.0002712
FGA	0.0038970	0.0007061	0.0039352	0.0008705	0.0041331	0.0008344



Appendix 2. Paternal & Maternal Mutation Data

CSF Paternal Mutations

Paternal Mutation – Allele:		Caucasian		Black		Hispanic		American Indian	Asian (Oriental)
From	To	Number	Frequency	Number	Frequency	Number	Frequency		
7	8			1	0.0000229				
7	9								
8	9								
9	8	1	0.0000279						
9	10			2	0.0000458				
10	8								
10	9	1	0.0000279	2	0.0000458	1	0.0000588		
10	11	1	0.0000279	6	0.0001373	4	0.0002354		
11	9					1	0.0000588		
11	10	5	0.0001396	6	0.0001373			1	
11	12	4	0.0001117	3	0.0000687	2	0.0001177		2
11	15								
11	10 or 12			2	0.0000458	1	0.0000588		
12	10			1	0.0000229				
12	11	1	0.0000279	10	0.0002289	3	0.0001765		
12	13	11	0.0003072	18	0.0004120	6	0.0003531		1
12	11 or 13	2	0.0000558		0.0000000				
13	12	6	0.0001675	15	0.0003434	1	0.0000588		
13	14	4	0.0001117	6	0.0001373	1	0.0000588		
13	12 or 14	1	0.0000279						
14	13	5	0.0001396	2	0.0000458	2	0.0001177		1
14	15	1	0.0000279						
15	14	3	0.0000838			2	0.0001177		
10 or 11	9 or 12								

10 or 12	11	4	0.0001117	1	0.0000229	2	0.0001177
10 or 12	11 or 13	1	0.0000279				
10 or 12	9 or 11			1	0.0000229		
10 or 13	11 or 12	1	0.0000279	1	0.0000229		
10 or 13	9 or 12			1	0.0000229		
11 or 12	10 or 13	1	0.0000279				
11 or 13	12	4	0.0001117	4	0.0000916		
11 or 13	10 or 12	1	0.0000279				
12 or 14	13	2	0.0000558	2	0.0000458		
9 or 11	10						
	Sum	60	0.0016754	84	0.0019228	26	0.0015300
Total Paternal Meiosis by Race			43,686		16,993		229
Paternal Frequency	0.0016754		0.0019228		0.0015300		0.0218341
Maternal Frequency	0.0002464		0.0003013		0.0004580		
Sum	0.0019218		0.0022241		0.0019881		
Relative Frequency Paternal	0.8717879		0.8645264		0.7696148		
Relative Frequency Maternal	0.1282121		0.1354736		0.2303852		
Indeterminate	0.0006303		0.0007001		0.0009284		
Frequency Paternal Indeterminate	0.0005495		0.0006053		0.0007145		
Frequency Maternal Indeterminate	0.0000808		0.0000948		0.0002139		
Total Paternal Mutation Frequency	0.0022249		0.0025281		0.0022446		
Total Maternal Mutation Frequency	0.0003272		0.0003962		0.0006719		

CSF Maternal Mutations

Maternal Mutation – Allele:		Caucasian		Black		Hispanic		American Indian	Asian (Oriental)
From	To	Number	Frequency	Number	Frequency	Number	Frequency		
7	8								
8	9			1	0.0000215				
9	10								
10	8								
10	8			1	0.0000215				
10	9	1	0.0000224	1	0.0000215				
10	11							1	
11	10	1	0.0000224	1	0.0000215				
11	12	1	v	2	0.0000430				
11	15								
11	10 or 12								
12	11	2	0.0000448	1	0.0000215			1	
12	13	3	0.0000672	3	0.0000646	2	0.0001018		
12	14					1	0.0000509		
12	11 or 13	1	0.0000224						
13	12			1	0.0000215	1	0.0000509		
13	14	1	0.0000224			1	0.0000509		
14	13					1	0.0000509		
14	15							1	
15	16	1	0.0000224						
10 or 11	9 or 12								
10 or 12	11					1	0.0000509		
10 or 12	11 or 13					1	0.0000509		
10 or 12	9 or 11								
10 or 12	9 or 13					1	0.0000509		

11 or 12	10 or 13		1	0.0000215	
11 or 13	12		2	0.0000430	
11 or 13	10 or 12				
12 or 14	13				
9 or 11	10				
	sum	11	0.0002464	14	0.0003013
Total Maternal Meiosis by Race		44,642	46,464	19,650	0.0004580
	Freq	0.0002464		0.0003013	0.0004580
					0.0060241

D2S1338 Paternal Mutations

Paternal Mutation – Allele:		Caucasian	Black	Hispanic	American Indian	Asian (Oriental)
From	To	Number	Frequency	Number	Frequency	Number
16	17			2	0.0000914	
17	16			1	0.0000457	
17	18	4	0.0002125			
17	20			1	0.0000457	
18	17			1	0.0000457	
18	19			1	0.0000457	
18	17 or 19					
19	17	1	0.0000531			
19	18	2	0.0001062			
19	20			2	0.0000914	1
20	19					0.0002934
20	21	1	0.0000531	1	0.0000457	1
21	20			1	0.0000457	1
21	22	1	0.0000531	3	0.0001371	1
						0.0002934

22	21	2	0.0001062	1	0.0000457		
22	23	2	0.0001062	3	0.0001371		1
23	22	2	0.0001062	2	0.0000914	1	0.0002934
23	24	4	0.0002125				1
23	25						
24	23	3	0.0001593				
24	25	2	0.0001062				
24	26			1	0.0000457		
25	24	5	0.0002656	5	0.0002285		1
25	26	2	0.0001062	2	0.0000914	1	0.0002934
26	25	2	0.0001062	5	0.0002285		
26	27	3	0.0001593				1
27	26			2	0.0000914		
20 or 24	19 or 23						
21 or 23	22						
23 or 25	24						
23 or 25	24						
	Sum	36	0.0019121	34	0.0015539	5	0.0014671
Total Paternal Meiosis by Race				3408		0	5
		18827					229
Paternal Frequency			0.0019121	0.0015539	0.0014671		0.0218341
Maternal Frequency			0.0003137	0.0004469	0.0012259		
	Sum	0.0022258		0.0020008	0.0026930		
Relative Frequency							
Paternal		0.8590726		0.7766339	0.5447958		
Relative Frequency							
Maternal		0.1409274		0.2233661	0.4552042		
Indeterminate		0.0003408		0.0000000	0.0000000		
Frequency Paternal							
Indeterminate		0.0002928		0.0000000	0.0000000		

Frequency Maternal				
Indeterminate	0.0000480		0.0000000	
Total Paternal			0.0000000	0.0000000
Mutation Frequency	0.0022049		0.0015539	0.0014671
Total Maternal			0.0004469	0.0012259
Mutation Frequency	0.0003617			

D2S1338 Maternal Mutations

Maternal Mutation – Allele:		Caucasian	Black	Hispanic	American Indian	Asian (Oriental)
From	To	Number	Frequency	Number	Frequency	Number
17	16					1
17	18	1	0.0000627			1
18	19	1	0.0000627	1	0.0000559	
18	22			1	0.0000559	
18	17 or 19					
19	18					
19	20			1	0.0000559	
20	19					
20	21					
21	20					
21	22			1	0.0000559	1
22	21	1	0.0000627	1	0.0000559	
22	23					
23	22					
23	24					
23	25					
24	23	1	0.0000627	2	0.0001117	
24	25	1	0.0000627			
25	24					

D3S1358 Paternal Mutations

17	16	11	0.0002057	20	0.0002934	1	0.0000425
17	18	9	0.0001683	18	0.0002640	7	0.0002976
17	19			2	0.0000293		1
17	16 or 18						
18	17	13	0.0002431	8	0.0001173	4	0.0001701
18	19	10	0.0001870	8	0.0001173	3	0.0001275
18	16 or 17						1
19	18	11	0.0002057	2	0.0000293		
19	20						
20	19	2	0.0000374	2	0.0000293		
20	21						
22	23					1	0.0000425
23	22	1	0.0000187				
14 or 16	15	1	0.0000187	1	0.0000147		
14 or 17	15 or 16			3	0.0000440		
15 or 16	14 or 17			1	0.0000147		
15 or 17	16	1	0.0000187	5	0.0000733	1	0.0000425
15 or 17	14 or 16						
15 or 17	16 or 18	1	0.0000187				
15 or 18	16 or 17			1	0.0000147	1	0.0000425
15 or 19	14 or 18			1	0.0000147		
15 or 19	16 or 18	1	0.0000187				
16 or 17	15 or 18	1	0.0000187				
16 or 18	17	4	0.0000748	5	0.0000733	2	0.0000850
16 or 19	15 or 18					1	0.0000425
16 or 19	17 or 18						
17 or 18	16						
17 or 19	18						
	Sum	92	0.0017201	113	0.0016575	31	0.0013180
							5

Total Paternal Meiosis by Race	53,486		68,176		23,521		229
Paternal Frequency	0.0017201		0.0016575		0.0013180		0.0218341
Maternal Frequency	0.0001938		0.0001859		0.0001046		
Sum	0.0019139		0.0018434		0.0014226		
Relative Frequency Paternal	0.8987395		0.8991524		0.9264712		
Relative Frequency Maternal	0.1012605		0.1008476		0.0735288		
Indeterminate	0.0006751		0.0006900		0.0006361		
Frequency Paternal Indeterminate	0.0006067		0.0006204		0.0005893		
Frequency Maternal Indeterminate	0.0000684		0.0000696		0.0000468		
Total Paternal Mutation Frequency	0.0023268		0.0022779		0.0019073		
Total Maternal Mutation Frequency	0.0002672		0.0002555		0.0001514		

D3S1358 Maternal Mutations

Maternal Mutation – Allele:		Caucasian		Black		Hispanic		American Indian		Asian (Oriental)	
From	To	Number	Frequency	Number	Frequency	Number	Frequency	Number	Frequency	Number	Frequency
12	14	1	0.0000194								
14	15										
15	12										
15	14										
15	16										

Total Maternal Meiosis by Race Trios Only	51,611	59,157	19,126	
Frequency	0.0001938	0.0001859	0.0001046	

D5S818 Paternal Mutations

Paternal Mutation – Allele:		Caucasian		Black		Hispanic		American Indian	Asian (Oriental)
From	To	Number	Frequency	Number	Frequency	Number	Frequency		
9	10								
10	11	2	0.0000516	1	0.00000208				
11	10	1	0.0000258	3	0.00000623				
11	12	5	0.0001291	7	0.0001455	3	0.0001682	1	
11	13	5	0.0001291						
11	9							1	0.0000561
12	10								
12	11	1	0.0000258	6	0.0001247	1	0.0000561		
12	13	12	0.0003098	17	0.0003533	1	0.0000561		
12	10 or 11								
12	11 or 13								
13	12	8	0.0002065	6	0.0001247	1	0.0000561		
13	14	7	0.0001807	19	0.0003949	7	0.0003925	1	
13	12 or 14								
14	13	11	0.0002840	11	0.0002286	3	0.0001682		
14	15			2	0.0000416				
15	14								
17	18								
10 or 12	11							1	0.0000561
10 or 13	11 or 12			1	0.0000208				

10 or 14	9 or 13							
11 or 12	13 or 14							
11 or 13	12	2	0.0000516	9	0.0001870			
11 or 13	12 or 14			1	0.0000208			
11 or 14	12 or 15			1	0.0000208			
11 or 14	12 or 13					1	0.0000561	
12 or 14	13	2	0.0000516	3	0.0000623	1	0.0000561	1
12 or 14	11 or 13	1	0.0000258					
9 or 11	10							
	Sum	57	0.0014715	88	0.0018288	20	0.0011213	0
Total Paternal Meiosis by Race								
			38,736	48,119	17,836		581	
Paternal Frequency		0.0014715		0.0018288		0.0011213		0.0051635
Maternal Frequency		0.0002630		0.0003075		0.0001552		
Sum		0.0017345		0.0021363		0.0012765		
Relative Frequency Paternal								
Relative Frequency Maternal		0.8483712		0.8560507		0.8784097		
Indeterminate		0.0007935		0.0006211		0.0009541		
Frequency Paternal Indeterminate		0.0006732		0.0005317		0.0008381		
Frequency Maternal Indeterminate		0.0001203		0.0000894		0.0001160		
Total Paternal Mutation Frequency		0.0021447		0.0023605		0.0019594		
Total Maternal Mutation Frequency		0.0003833		0.0003969		0.0002712		

D5S18 Maternal Mutations

Maternal Mutation – Allele:		Caucasian		Black		Hispanic		Asian (Oriental)	American Indian
From	To	Number	Frequency	Number	Frequency	Number	Frequency		
9	10								
10	11								
11	10	2	0.0000405						
11	12	1	0.0000202	2	0.0000410				
11	13								
11	13					1	0.0000517		
12	10								
12	11	3	0.0000607	1	0.0000205				
12	13			4	0.0000820	1	0.0000517		
12	10 or 11								
12	11 or 13								
13	12	1	0.0000202	1	0.0000205				
13	14	2	0.0000405	3	0.0000615			1	
13	12 or 14					1	0.0000205		
14	13					1	0.0000205		
14	15	1	0.0000202	1	0.0000205				
15	14	2	0.0000405						
17	18								
10 or 12	11								
10 or 14	9 or 13								
11 or 12	13 or 14								
11 or 13	12	1	0.0000202	1	0.0000205	1	0.0000517		
12 or 14	13								
12 or 14	11 or 13								
9 or 11	10								

	Sum	13	0.0002630	15	0.0003075	3	0.0001552	0	1
Total Maternal Meiosis by Race	49,428		48,777		19,328				234
Frequency	0.0002630		0.0003075		0.0001552				0.0042735

D7S820 Paternal Mutations

15	14		1	0.0000212		
10 or 12	11		1	0.0000212	2	0.0001030
10 or 12	11 or 13	1	0.0000263			
10 or 12	9 or 11					
10 or 12	9 or 13					
10 or 13	11 or 12					
10 or 13	11 or 12		1	0.0000212		
11 or 13	12	4	0.0001054			
11 or 13	10 or 12		1	0.0000212	1	0.0000515
11 or 14	10 or 15	1	0.0000263			
12 or 14	13					
8 or 10	9		2	0.0000423		
8 or 11	9 or 10		1	0.0000212		
8 or 11	9 or 12	1	0.0000263			
8 or 12	9 or 11		1	0.0000212		
9 or 11	10	2	0.0000527	2	0.0000423	2
9 or 12	10 or 11					0.0001030
9 or 12	10 or 13	1	0.0000263			
9 or 13	10 or 12					
	Sum	55	0.0014490	57	0.0012065	31
Total Paternal Meiosis by Race		37,957	47,244	19,422	352	229
Paternal Frequency		0.0014490	0.0012065	0.0015961	0.0028409	0.0131004
Maternal Frequency		0.0001413	0.0001688	0.0001083		
	Sum	0.0015903	0.0013753	0.0017044		
Relative Frequency		0.9111493	0.8772634	0.9364596		
Relative Frequency		0.0888507	0.1227366	0.0635404		
Maternal						

Indeterminate	0.0005043		0.0003017		0.0001932	
Frequency Paternal Indeterminate	0.0004595		0.0002647		0.0001809	
Frequency Maternal Indeterminate	0.0000448		0.0000370		0.0000123	
Total Paternal						
Mutation Frequency	0.0019085		0.0014712		0.0017771	
Total Maternal						
Mutation Frequency	0.0001861		0.0002058		0.0001206	

D7S820 Maternal Mutations

Maternal Mutation – Allele:		Caucasian	Black	Hispanic	American Indian	Asian (Oriental)
From	To	Number	Frequency	Number	Frequency	
7	6					
8	7					
8	9					
9	8					
9	10					
10	9	1	0.0000202	1	0.0000211	
10	11	1	0.0000202	1	0.0000211	
10.1	9.1			1	0.0000211	
11	10			1	0.0000211	
11	12			1	0.0000211	1
11	12 or 13					
12	11					
12	13	1	0.0000202			
13	12					

13	14	1	0.0000202
14	13	1	0.0000202
14	15	1	0.0000202
10 or 12	11		
10 or 12	9 or 11		
10 or 12	9 or 13		
10 or 13	11 or 12		
11 or 13	12	1	0.0000211
12 or 14	13		
8 or 11	9 or 10	1	0.0000211
9 or 10	8 or 11	1	0.0000211
9 or 11	10		
9 or 11	8 or 10		
9 or 12	10 or 11		
9 or 13	10 or 12		
9 or 13	8 or 12	1	0.0000202
	Sum	7	0.0001413
Total Maternal Meiosis by Race		8	0.0001688
Frequency	0.0001413		0.0001688
			0.0001083
			2
			0.0001083
			0
			1
			264
			0.0037879

D8S1179 Paternal Mutations

Paternal Mutation – Allele:		Caucasian		Black		Hispanic		Asian American Indian (Oriental)	
From	To	Number	Frequency	Number	Frequency	Number	Frequency	American Indian	Asian (Oriental)
9	10								
10	11	2	0.0000370	2	0.0000291	1	0.0000411	1	
11	7								

11 or 13	12 or 14			1	0.0000145	
12 or 13	11 or 14					
12 or 14	13	1	0.0000185	6	0.0000872	1
12 or 14	11 or 13					
12 or 14	13 or 15			1	0.0000145	
12 or 15	13 or 14					
12 or 18	13 or 17					
13 or 14	15					
13 or 14	12 or 15			2	0.0000291	
13 or 15	14	4	0.0000740	7	0.0001018	2
13 or 15	12 or 14				0.0000823	
13 or 15	14 or 16			1	0.0000145	
13 or 16	12 or 17					
13 or 16	14 or 15					
14 or 16	15			3	0.0000436	1
14 or 16	13 or 15				0.0000411	
15 or 16	14 or 17			1	0.0000145	
15 or 17	16			2	0.0000291	
15 or 18	14 or 17					
	Sum	106	0.0019619	153	0.0022240	38
Total Paternal Meiosis by Race					0.0015633	1
						6
Paternal Frequency	0.0019619					
Maternal Frequency	0.0002104					
Sum	0.0021723					
Relative Frequency Paternal	0.9031462					

Relative Frequency Maternal	0.0968538		0.1373510		0.2030109	
Indeterminate	0.0004938		0.0006174		0.0007656	
Frequency Paternal Indeterminate	0.0004460		0.0005326		0.0006102	
Frequency Maternal Indeterminate	0.0000478		0.0000848		0.0001554	
Total Paternal Mutation Frequency	0.0024079		0.0027566		0.0021734	
Total Maternal Mutation Frequency	0.0002582		0.0004389		0.0005536	

D8S1179 Maternal Mutations

Maternal Mutation – Allele:	Caucasian	Black	Hispanic	American Indian	Asian (Oriental)
From	To	Number	Frequency	Number	Frequency
9	10	2	0.0000301		
10	11	7		1	0.0000498
11	10	1	0.0000150		
11	12	1	0.0000150	1	0.0000169
12	11			1	0.0000169
12	13	1	0.0000150	2	0.0000337
12	11 or 13			2	0.0000996
13	12			1	0.0000498
13	14	1	0.0000150	2	0.0000337
13	15			3	0.0001493
13	12 or 14				

14	13	3	0.0000451	4	0.0000675	1	0.0000498
14	15	1	0.0000150	2	0.0000337		
14	13 or 15						
15	14	2	0.0000301	2	0.0000337		
15	16			2	0.0000337		
16	15			3	0.0000506		
16	17						
16	18						
16	15 or 17						
17	16						
17	18						
18	17			1	0.0000169		
10 or 12	11						
10 or 13	11 or 14						
11 or 13	12						
11 or 13	10 or 12						
12 or 13	11 or 14						
12 or 14	13						
12 or 14	11 or 13						
12 or 15	13 or 14						
13 or 14	15						
13 or 14	12 or 15						
13 or 15	14	1	0.0000150				
13 or 15	12 or 14						
13 or 15	14 or 16						
13 or 16	12 or 17						
13 or 16	14 or 15						
13 or 16	14 or 15	1	0.0000150				
14 or 16	15						
14 or 16	13 or 15						
15 or 17	16						

15 or 18	14 or 17	Sum	14	0.0002104	21	0.0003541	8	0.0003982
Total Maternal Meiosis by Race			66,529	66,529	59,299		20,090	
	Frequency	0.0002104		0.0003541		0.0003982		

D13S317 Paternal Mutations

Paternal Mutation – Allele:		Caucasian	Black	Hispanic	American Indian	Asian (Oriental)
From	To	Number	Frequency	Number	Frequency	Number
8	9	1	0.0000245			
9	10			1	0.0000209	
10	8			1	0.0000209	1
10	9			1	0.0000209	1
10	11			1	0.0000209	
10	15					
11	9					
11	10	3	0.0000736	2	0.0000418	1
11	12	6	0.0001472	9	0.0001883	3
11	15					
11	10 or 12	1	0.0000245			
12	9					
12	11	6	0.0001472	9	0.0001883	
12	13	8	0.0001963	14	0.0002929	4
12	11 or 13	1	0.0000245	4	0.0000837	2
13	12	3	0.0000736	6	0.0001255	1
13	14	8	0.0001963	6	0.0001255	6
						0.0003040

13	12 or 14		2	0.0000418			
14	13	10	0.0002453	7	0.0001464	3	0.0001520
14	15	2	0.0000491	2	0.0000418	2	0.0001013
15	14	3	0.0000736	4	0.0000837	1	0.0000507
16	15			1	0.0000209	1	0.0000507
16	17			1	0.0000209		
10 or 12	11						
10 or 13	11 or 12						
11 or 12	10 or 13						
11 or 13	12	5	0.0001227	7	0.0001464	1	0.0000507
11 or 13	12 or 14			2	0.0000418		
11 or 14	12 or 13			1	0.0000209	1	0.0000507
11 or 15	13			1	0.0000209		
12 or 13	11 or 14						
12 or 14	13		4	0.0000837			
12 or 14	11 or 13	1	0.0000245				
12 or 14	11 or 15						
12 or 15	13 or 14	1	0.0000245				
13 or 14	12 or 15			1	0.0000209		
13 or 15	14	3	0.0000736	1	0.0000209		
16 or 18	15 or 17	1	0.0000245				
8 or 12	11 or 9						
8 or 12	9 or 13						
	Sum	63	0.0015455	87	0.0018200	25	0.0012667
Total Paternal Meiosis by Race						0	4
Paternal Frequency		40,763		47,801		19,736	229
Maternal Frequency		0.0015455		0.0018200		0.0012667	0.0174672
Sum		0.0001228		0.0005815		0.0002717	
		0.0016683		0.0024015		0.0015384	

Relative Frequency Paternal	0.9263930		0.7578643		0.8233903	
Relative Frequency Maternal	0.0736070		0.2421357		0.1766097	
Indeterminate	0.0007063		0.0006655		0.0005154	
Frequency Paternal	0.0006543		0.0005044		0.0004244	
Indeterminate						
Frequency Maternal	0.0000520		0.0001611		0.0000910	
Indeterminate						
Total Paternal	0.0021998		0.0023244		0.0016911	
Mutation Frequency						
Total Maternal			0.0007426		0.0003627	
Mutation Frequency	0.0001748					

D13S317 Maternal Mutations

Maternal Mutation – Allele:		Caucasian	Black	Hispanic	American Indian	Asian (Oriental)
From	To	Number	Frequency	Number	Frequency	Number
8	9					
9	10					
10	9					
10	11					
10	15					
11	9					
11	10			1	0.0000208	
11	12	2	0.0000409	6	0.0001246	
11	15					
12	9					
12	11			1	0.0000208	1
12	13	1	0.0000205	7	0.0001454	1
						0.0000543

12	11or13		1	0.0000208
13	12		4	0.0000831
13	14		3	0.0000623
13	12 or 14		1	0.0000543
14	13	1	0.0000205	
14	15	1	0.0000205	1
14	16		1	0.0000208
15	14		1	0.0000208
10 or 12	11		1	0.0000208
10 or 13	11 or 12		1	0.0000208
11 or 12	10 or 13		1	0.0000208
11 or 13	12		1	0.0000208
11 or 13	12 or 14		1	0.0000208
11 or 14	12 or 13		1	0.0000208
12 or 13	11 or 14		1	0.0000208
12 or 14	13	1	0.0000205	1
12 or 14	11 or 13		1	0.0000208
12 or 14	11 or 15		1	0.0000208
8 or 12	11 or 9		1	0.0000208
8 or 12	9 or 13		1	0.0000208
9 or 12	10 or 11		1	0.0000208
Sum		6	0.0001228	28
Total Maternal Meiosis by Race Trios Only			0.0005815	5
Frequency		48,852	48,154	18,406
		0.0001228	0.0005815	0.0002717
				234
				0.0042735

D16S539 Paternal Mutations

Paternal Mutation – Allele:	Caucasian	Black	Hispanic	American Indian	Asian (Oriental)

From	To	Number	Frequency	Number	Frequency	Number	Frequency
8	9	1	0.0000237				
9	8	1	0.0000237				
9	10	2	0.0000473	2	0.0000367		
9	11			1	0.0000184		
10	11			5	0.0000918	1	0.0000510
10	9 or 11						
11	9	1	0.0000237				
11	10	2	0.0000473	1	0.0000184	1	0.0000510
11	12	6	0.0001420	17	0.0003120	1	0.0000510
11	13			1	0.0000184		
11	14					1	0.0000510
11	10 or 12	1	0.0000237				
12	10	1	0.0000237				
12	11	13	0.0003076	8	0.0001468	2	0.0001019
12	13	10	0.0002366	12	0.0002202	3	0.0001529
12	11 or 13	2	0.0000473				
13	9						
13	11						
13	12	7	0.0001656	8	0.0001468	1	0.0000510
13	14	12	0.0002839	13	0.0002386	2	0.0001019
13	12 or 14					1	0.0000510
14	13	7	0.0001656	9	0.0001652	2	0.00001019
14	15	3	0.0000710	3	0.0000551		
15	14			1	0.0000184	1	0.0000510
15	16					1	0.0000510
10 or 12	11	1	0.0000237	2	0.0000367		
10 or 12	11 or 13						
10 or 12	9 or 13						
11 or 13	12	3	0.0000710	3	0.0000551	1	0.0000510

11 or 13	12 or 14		1	0.0000184
11 or 14	10 or 13			
11 or 14	12 or 13		1	0.0000184
11 or 15	12 or 14			
12 or 14	13	3	0.0000710	1
12 or 15	13 or 14			
13 or 15	14			
9 or 11	10	1	0.0000237	
9 or 12	10 or 11			
9 or 13	10 or 12			
	Sum	77	0.0018219	89
Total Paternal Meiosis by Race		42,264	54,494	19,622
Paternal Frequency	0.0018219		0.0016332	0.0010193
Maternal Frequency	0.0003009		0.0004497	0.0003204
Sum	0.0021228		0.0020829	0.0013397
Relative Frequency				
Paternal	0.8582520		0.7840919	0.7608132
Relative Frequency Maternal	0.1417480		0.2159081	0.2391868
Indeterminate	0.0004088		0.0004497	0.0003204
Frequency Paternal Indeterminate	0.0003509		0.0003526	0.0002438
Frequency Maternal Indeterminate	0.0000579		0.0000971	0.0000766
Total Paternal Mutation Frequency	0.0021727		0.0019858	0.0012631
Total Maternal Mutation Frequency	0.0003588		0.0005468	0.0003971

D16S539 Maternal Mutations

Maternal Mutation – Allele:	Caucasian			Black			Hispanic			American Indian	Asian (Oriental)
From	To	Number	Frequency	Number	Frequency	Number	Frequency	Number	Frequency		
8	9										
9	10	1	0.00000158								
10	9			1	0.00000150						
10	11			1	0.00000150						1
10	9 or 11										
11	9			1	0.00000150						
11	10							1	0.00000458		
11	12										
12	11	3	0.00000475	5	0.00000750	1	0.00000458				
12	13										
12	11 or 13	1	0.00000158								
13	9										
13	11										
13	12	2	0.00000317	10	0.0001499						
13	14	1	0.00000158	2	0.00000300						
13	12 or 14	1	0.00000158								
14	13	4	0.00000534	5	0.00000750	1	0.00000458				1
14	15			1	0.00000150	1	0.00000458				
15	14	2	0.00000317	1	0.00000150	2	0.00000916				
15	16										
10 or 12	11			1	0.00000150						
10 or 12	11 or 13										
10 or 12	9 or 13										
10 or 13	11 or 12	1	0.00000158								
11 or 13	12	1	0.00000158	1	0.00000150	1	0.00000458				
11 or 14	10 or 13										

11 or 15	12 or 14							
12 or 14	13	2	0.0000317	1	0.0000150			
12 or 15	13 or 14							
13 or 15	14							
9 or 12	10 or 11							
9 or 13	10 or 12							
	sum	19	0.0003009	30	0.0004497	7	0.0003204	0
Total Maternal Meiosis by Race								2
	Frequency	0.0003009		0.0004497		0.0003204		0.0085470

D19S453 Paternal Mutations

Paternal Mutation – Allele:		Caucasian		Black		Hispanic		American Indian	Asian (Oriental)
From	To	Number	Frequency	Number	Frequency	Number	Frequency		
12	10								
12	11	2	0.0001028						
12	13								
12	14	1	0.0000514						
12.2	13.2								
13	14	1	0.0000514	3	0.0001298	1	0.0002802		
13	12	4	0.0002057	1	0.0000433				
13	15			1	0.0000433				
13.2	11.2								
13.2	12.2					1	0.0000433		
14	13	1	0.0000514	3	0.0001298	1	0.0002802		
14	15	3	0.0001543	3	0.0001298			1	
14	13 or 15								

14	14.2			1	0.0000433		
14.2	13.2	1	0.0000514	1	0.0000433		
15	14	2	0.0001028	2	0.0000865		
15	16	2	0.0001028	3	0.0001298	1	0.0002802
15.2	16.2			1	0.0000433	1	0.0002802
15.2	14.2	4	0.0002057	1	0.0000433		
16	15	3	0.0001543	1	0.0000433		
16	17	2	0.0001028				
16.2	15.2	1	0.0000514	1	0.0000433		
16.2	17.2	1	0.0000514				
17	16						
17	18						
17.2	16.2	1	0.0000514				
13 or 15	14						
14 or 16	15						
	Sum	29	0.0014912	24	0.0010381	4	0.0011208
Total Paternal Meiosis by Race			19448	23119	3569		229
Paternal Frequency		0.0014912		0.0010381		0.0011208	
Maternal Frequency		0.0003583		0.0003679		0.0000000	
Sum		0.0018495		0.0014060		0.0011208	
Relative Frequency		0.8062673		0.7383370		1.0000000	
Paternal							
Relative Frequency		0.1937327		0.2616630		0.0000000	
Maternal							
Indeterminate		0.0003408		0.0013369		0.0000000	
Frequency Paternal		0.0002748		0.0009871		0.0000000	
Indeterminate							
Frequency Maternal		0.0000660		0.0003498		0.0000000	
Indeterminate							

Total Paternal Mutation Frequency	0.0017659		0.0020252		0.0011208	
Total Maternal Mutation Frequency	0.0004243		0.0007177		<0.0002909	

D19S453 Maternal Mutations

Maternal Mutation – Allele:		Caucasian		Black		Hispanic	American Indian	Asian (Oriental)
From	To	Number	Frequency	Number	Frequency	Number	Frequency	
12	10							
12.2	13.2							
13	12	3	0.0001535					
13	14							
13.2	11.2							
14	13					2	0.0001051	
14	15							
14	13 or 15							
14.2	13.2							
14.2	15.2					1	0.0000526	
15	11					1	0.0000526	
15	14	2	0.0001024	2	0.0001051			
15	16							
15.2	14.2					1	0.0000526	
15.2	16.2							
16	15	2	0.0001024					
16	17							
16.2	15.2							
16.2	17.2							
17	16							
17	18							

13 or 15	14						
14 or 16	15						
	Sum	7	0.0003583	7	0.0003679	0	0
Total Maternal Meiosis by Race	19539			19027		3438	0.00029087
	Frequency	0.0003583		0.0003679		0.0000000	

D21S51 Paternal Mutations

Paternal Mutation – Allele:		Caucasian			Black			Hispanic		American Indian	Asian (Oriental)
From	To	Number	Frequency	Number	Frequency	Number	Frequency	Number	Frequency		
23	29										
26	27										
27	28			2	0.0000292	1	0.0000427				
28	26										
28	27	1	0.0000187	2	0.0000292						
28	27.2										
28	29	4	0.0000747	8	0.0001168	3	0.0001280				
28	30	1	0.0000187	1	0.0000146						
28	27 or 29										
29	28	2	0.0000374	4	0.0000584	1	0.0000427				
29	30	6	0.0001121	12	0.0001752					1	
29	30.2					1	0.0000146				
29	33										
29	28 or 30										
30	28										
30	29	6	0.0001121	7	0.0001022	2	0.0000854				

30	29.2	1	0.0000187				
30	30.2	1	0.0000187				
30	31	16	0.0002990	16	0.0002336	5	0.0002134
30	32						
30	32.2	1	0.0000187				
30	29 or 31						
30.2	31.2	3	0.0000561	1	0.0000146		
31	30	6	0.0001121	2	0.0000292		1
31	31.2						
31	32	4	0.0000747	7	0.0001022	2	0.0000854
31.2	30.2			2	0.0000292		2
31.2	31	1	0.0000187				
31.2	32.2	6	0.0001121	3	0.0000438	1	0.0000427
32	31	4	0.0000747				
32	33	2	0.0000374	3	0.0000438		
32.2	30					1	0.0000427
32.2	31.2			2	0.0000292	4	0.0001707
32.2	33.2	8	0.0001495	7	0.0001022	3	0.0001280
32.2	31.2 or 33.2						
33	32	1	0.0000187				
33	34	1	0.0000187				
33.2	31.2	1	0.0000187				
33.2	32.2	3	0.0000561	4	0.0000584	1	0.0000427
33.2	34.2	3	0.0000561	7	0.0001022	2	0.0000854
33.2	32.2 or 34.2					1	0.0000427
34	33			2	0.0000292		
34	35			2	0.0000292		
34.2	33.2			1	0.0000146		
34.2	35.2	1	0.0000187	1	0.0000146		

35	34		2	0.0000292
35	36		5	0.0000730
35.2	34.2			
36	35		2	0.0000292
36	37		3	0.0000438
36.2	35.2			
37	36		1	0.0000146
37	38		1	0.0000146
16 or 19	14 or 20			
27 or 29	28		1	0.0000146
27 or 29	28 or 30			
27 or 30	28 or 29			
27 or 32	28 or 31			
28 or 29	29 or 30			
28 or 30	29		2	0.0000292
28 or 30	27 or 29			
28 or 30	27 or 31			
28 or 30	29 or 31			
28 or 31	29 or 30			
28 or 31.2	29 or 32.2	1	0.0000187	
28 or 32.2	29 or 31.2			
29 or 30	28 or 31		1	0.0000146
29 or 31	30	1	0.0000187	2
29 or 31	28 or 30			
29 or 32	30 or 31			
29 or 32.2	28 or 31.2			
30 or 31	29 or 32	1	0.0000187	
30 or 31.2	31			

30 or 32	31	1	0.00000187			
30 or 32.2	31 or 33.2		1	0.00000146		
31.2 or 32	31					
31.2 or 33.2	32.2					
	Sum	87	0.0016256	118	0.0017232	31
Total Paternal Meiosis by Race		53,520	68,479	23,432		229
Paternal Frequency		0.0016256	0.0017232	0.0013230		0.0174672
Maternal Frequency		0.0013077	0.0013928	0.0016832		
	Sum	0.0029333	0.0031160	0.0030062		
Relative Frequency Paternal	Paternal	0.5541821	0.5530104	0.4400862		
Relative Frequency Maternal	Maternal	0.4458179	0.4469896	0.5599138		
Indeterminate		0.0004938	0.0006174	0.0007656		
Frequency Paternal Indeterminate		0.0002737	0.0003414	0.0003369		
Frequency Maternal Indeterminate		0.0002201	0.0002760	0.0004287		
Total Paternal Mutation Frequency		0.0018992	0.0020646	0.0016599		
Total Maternal Mutation Frequency		0.0015278	0.0016688	0.0021119		

D21S51 Maternal Mutations

Maternal Mutation – Allele:	Caucasian			Black			Hispanic		American Indian	Asian (Oriental)
From	To	Number	Frequency	Number	Frequency	Number	Frequency			
23	29									
26	27									
27	28									
28	26									
28	27	1	0.0000147							
28	27.2									
28	29	1	0.0000147	5	0.00000718					
28	30									
28	27 or 29									
29	27	1	0.0000147							
29	28	4	0.0000588	2	0.00000287	4	0.0001772	1		
29	30	2	0.0000294	1	0.00000144	2	0.00000886			
29	31.2	1	0.0000147							
29	33									
29	28 or 30									
30	28									
30	29	12	0.0001763	14	0.00002010	2	0.00000886			
30	31	1	0.0000147	1	0.00000144	2	0.00000886			
30	32									
30	29 or 31									
30.2	31.2									
31	30	18	0.0002645	17	0.0002441	5	0.0002215			
31	31.2									
31	32	2	0.0000294	4	0.0000574					
31	30 or 32			1	0.0000144					

31.2	30.2	1	0.0000147	1	0.0000144
31.2	31	1	0.0000147		
31.2	32.2	1	0.0000147	1	0.0000144
32	31	3	0.0000441	8	0.0001149
32	33				3 0.0001329
32.2	31.2	3	0.0000441	4	0.0000574
32.2	33.2	3	0.0000441	3	0.0000431
32.2	31.2 or 33.2				1 0.0000443
33	32	2	0.0000294	1	0.0000144
33	34	2	0.0000294		0.0000000
33.2	32.2	8	0.0001175	10	0.0001436
33.2	34.2	1	0.0000147	1	0.0000144
33.2	32.2 or 34.2				1 0.0000443
34	33				
34	35			1	0.0000144
34.2	33.2	6	0.0000882	5	0.0000718
34.2	35.2			4	0.0001772
35	34			1	0.0000144
35	36				
35.2	34.2				
36	35				
36	37			2	0.0000287
36.2	35.2				
37	36			2	0.0000287
16 or 19	14 or 20			1	0.0000443
26 or 32.2	27 or 33.2	1	0.0000147		
27 or 29	28				
27 or 29	28 or 30				

27 or 32	28 or 31							
27 or 34.2	28 or 33.2			1	0.0000144			
28 or 29	29 or 30							
28 or 30	29	2	0.0000294	2	0.0000287			
28 or 30	27 or 29			1	0.0000144			
28 or 30	27 or 31							
28 or 30	29 or 31	1	0.0000147					
28 or 31	29 or 30							
28 or 32.2	29 or 31.2							
29 or 30	28 or 31							
29 or 31	30	5	0.0000735	7	0.0001005	1	0.0000443	1
29 or 31	28 or 30	1	0.0000147					
29 or 32	30 or 31							
29 or 32.2	28 or 33.2	1	0.0000147					
30 or 31.2	31							
30 or 32	31	2	0.0000294					
30 or 32.2	31 or 33.2							
30 or 33	31 or 32	1	0.0000147					
31.2 or 32	31							
31.2 or 33.2	32.2	1	0.0000147	1	0.0000144	1	0.0000443	
	Sum	89	0.0013077	97	0.0013928	38	0.0016832	2
Total Maternal Meiosis by Race			68,057	69,644	22,576			
	Frequency	0.0013077		0.0013928		0.0016832		

FGA Paternal Mutations

Paternal Mutation – Allele:	From	To	Caucasian Number	Frequency	Black Number	Frequency	Hispanic Number	Frequency	American Indian	Asian (Oriental)
17.2	18.2									
18	19									
18.2	19.2				1	0.0000148				
19	17	1	0.0000188							
19	18	3	0.0000565				1	0.0000426		
19	20	4	0.0000753		2	0.0000297	1	0.0000426		
19	21	1	0.0000188							
19.2	20.2									
20	18	1	0.0000188							
20	19	4	0.0000753		2	0.0000297				
20	21	6	0.0001129		5	0.0000742	4	0.0001703		
20	22				1	0.0000148				
20	19 or 21									
20.2	21.2				1	0.0000148				
21	18									
21	19	1	0.0000188		1	0.0000148				
21	20	2	0.0000376		5	0.0000742				
21	20.2				1	0.0000148				
21	22	7	0.0001318		16	0.0002374	4	0.0001703	1	
21.2	21									
21.2	22.2								1	
22	20	1	0.0000188		1	0.0000148				
22	21	2	0.0000376		11	0.0001632	1	0.0000426	1	
22	23	10	0.0001882		12	0.0001780	6	0.0002555	1	
22	24				1	0.0000148				

22	21 or 23		1	0.0000148				
22.2	21.2							
22.2	22							
22.2	23.2	1	0.0000188	1	0.0000148			
23	21	1	0.0000188					
23	22	11	0.0002071	6	0.0000890	1	0.0000426	2
23	24	10	0.0001882	18	0.0002670	3	0.0001277	2
23	22 or 24	1	0.0000188					
23.2	22.2		1	0.0000148				
23.2	24.2	1	0.0000188	1	0.0000148			
24	19		1	0.0000148				
24	22	1	0.0000188	1	0.0000148			
24	23	4	0.0000753	15	0.0002225	6	0.0002555	1
24	25	20	0.0003765	27	0.0004005	2	0.0000852	1
24	26	1	0.0000188					
24	27							
24	23 or 25				1	0.0000426		
24	25 or 26							
24.2	23.2	1	0.0000188					
24.3	25.3							
25	24	16	0.0003012	17	0.0002522	2	0.0000852	2
25	26	17	0.0003200	18	0.0002670	5	0.0002129	1
25	27							
25	24 or 26	1	0.0000188					
25.2	24.2							
26	23							
26	25	11	0.0002071	18	0.0002670	15	0.0006387	
26	27	8	0.0001506	7	0.0001038	8	0.0003406	1
26	25 or 27							
27	24		1	0.0000148				
27	25	1	0.0000188					

27	26	6	0.0001129	8	0.0001187	5	0.0002129
27	28	1	0.0000188	2	0.0000297	4	0.0001703
28	26					1	0.0000426
28	27	1	0.0000188	4	0.0000593	1	0.0000426
28	29			3	0.0000445		1
29	28	1	0.0000188				
29	30			1	0.0000148		
30	29			2	0.0000297		
30.2	31.2						
33.2	34.2						
43.2	42.2			1	0.0000148		
43.2	44.2				1	0.0000148	
45.2	46.2						
46.2	45.2				1	0.0000148	
19 or 21	20						
19 or 23	20 or 24					1	0.0000426
19 or 24	21 or 25						
19 or 25	20 or 24	1	0.0000188				
20 or 22	21	5	0.0000941			1	0.0000426
20 or 22	21 or 23						
20 or 24	19 or 25						
20 or 24	21 or 23					1	0.0000426
21 or 23	22				1	0.0000148	1
21 or 23	22 or 24						
21 or 25	20 or 26			1	0.0000148		
21 or 25	22 or 24	1	0.0000188				
21 or 26	20 or 25					1	0.0000426
22 or 24	23	3	0.0000565	4	0.0000593		
22 or 24	21 or 23			1	0.0000148		
22 or 24	21 or 25			1	0.0000148		
22 or 24	22 or 25						

22 or 24	23 or 25	1	0.0000188				
22 or 25	23 or 24			1	0.0000426		
22 or 26	21 or 25						
22 or 26	23 or 25	1	0.0000188				
22 or 29	23 or 28						
23 or 24	22 or 25						
23 or 25	24	2	0.0000376	1	0.0000148	2	0.0000852
23 or 25	22 or 24						
23 or 25	24 or 26	1	0.0000188				
23 or 26	22 or 25						
23 or 26	24 or 25			1	0.0000148		
24 or 26	25	3	0.0000565	1	0.0000148	3	0.0001277
24 or 26	23 or 25						
24 or 26	25 or 27				1	0.0000426	
25 or 27	26			1	0.0000148		
25 or 28	24 or 27			1	0.0000148		
26 or 28	27						
	Sum	176	0.0033132	229	0.0033972	83	0.0035340
Total Paternal Meiosis by Race		53,121	67,408	23,486		352	523
Paternal Frequency		0.0033132	0.0033972	0.0035340		0.0113636	0.0248566
Maternal Frequency		0.0006003	0.0007515	0.0007135			
	Sum	0.0039135	0.0041487	0.0042475			
Relative Frequency Paternal		0.8466075	0.8188586	0.8320279			
Relative Frequency Maternal		0.1533925	0.1811414	0.1679721			
Indeterminate		0.0006896	0.0006570	0.0007200			
Frequency Paternal Indeterminate		0.0005838	0.0005380	0.0005991			

Frequency Maternal Indeterminate	0.0001058		0.0001190		0.0001209	
Total Paternal	0.0038970		0.0039352		0.0041331	
Mutation Frequency						
Total Maternal	0.0007061		0.0008705		0.0008344	

FGA Maternal Mutations

Maternal Mutation – Allele:		Caucasian		Black		Hispanic		American Indian	Asian (Oriental)
From	To	Number	Frequency	Number	Frequency	Number	Frequency		
17.2	18.2								
18	19								
18	24			1	0.0000150				
18.2	19.2			1	0.0000150				
19	20			1	0.0000150	1	0.0000446		
19.2	20.2								
20	19	2	0.00000308						
20	21	1	0.00000154						
20	19 or 21								
21	18								
21	19			1	0.0000150				
21	20	1	0.00000154						
21	22	2	0.00000308	1	0.0000150				
21.2	21								
21.2	22.2	1	0.00000154						
22	20								
22	21	2	0.00000308	2	0.00000301	2	0.0000892		
22	23	3	0.00000462	3	0.00000451	2	0.0000892		

22	24		1	0.0000150
22	29		1	0.0000150
22	21 or 23		1	
22.2	21.2	1	0.0000154	
22.2	22			
22.2	23.2			
23	21		1	0.0000150
23	22	3	0.0000462	7
23	24	2	0.0000308	5
23.2	24.2			
24	19		1	0.0000150
24	22		1	0.0000150
24	23	5	0.0000770	7
24	25	2	0.0000308	2
24	27			
24	23 or 25			
24	25 or 26			
24.3	25.3			
25	24	4	0.0000616	2
25	26	2	0.0000308	4
25	27			
25	24 or 26			
26	23			
26	25	2	0.0000308	3
26	27			
26	25 or 27			
27	26	1	0.0000154	2
27	28	1	0.0000154	
28	27			
28	29			
29	28			

30	29		
30.2	31.2		
33.2	34.2		
45.2	46.2		
19 or 21	20		
19 or 24	21 or 25		
19 or 25	20 or 24		
20 or 22	21	0.00000154	
20 or 24	19 or 25		
20 or 24	21 or 23		1 0.0000446
21 or 23	22		
21 or 23	22 or 24		
21 or 24	20 or 25		1 0.0000150
21 or 25	22 or 24		
21 or 26	22 or 25		1 0.0000446
22 or 24	23	1 0.00000154	
22 or 24	21 or 25		
22 or 24	22 or 25		
22 or 24	23 or 25		
22 or 26	21 or 25		
22 or 26	23 or 25		
22 or 29	23 or 28		
23 or 24	22 or 25		
23 or 25	24	1 0.00000154	1 0.00000150
23 or 25	22 or 24		
23 or 25	22 or 26		1 0.00000150
23 or 25	24 or 26		
23 or 26	22 or 25		
24 or 26	25	1 0.00000154	
24 or 26	23 or 25		
25 or 27	26		

26 or 28	27							
	Sum	39	0.0006003	50	0.0007515	16	0.0007135	
Total Maternal Meiosis by Race		64,971		66,533		22,426		
Frequency	0.0006003		0.0007515		0.0007135			

THO Paternal Mutations

Paternal Mutation – Allele:		Caucasian		Black		Hispanic		American Indian	Asian (Oriental)
From	To	Number	Frequency	Number	Frequency	Number	Frequency		
6	7								
7	6								
7	8								
8	7			1	0.0000153				
8	9			1	0.0000153				
8	9.3								
9	8	1	0.0000225	1	0.0000153				
9	10	1	0.0000225			1	0.0000153		
9.3	7								
10	8								
10	11	1	0.0000225						
6 or 8	7 or 9								
8 or 10	9								
	Sum	3	0.0000676	4	0.0000612	0			
Total Paternal Meiosis by Race				65,369		17,554			
Paternal Frequency	0.0000676			0.0000612		0.0000000			

Maternal Frequency	0.0000785		0.0000858		0.0000445	
Sum	0.0001461		0.0001470		0.0000445	
Relative Frequency Paternal	0.4628688		0.4162911		0.0000000	
Relative Frequency Maternal	0.5371312		0.5837089		1.0000000	
Indeterminate	0.0000000		0.0000494		0.0000000	
Frequency Paternal Indeterminate	0.0000000		0.0000206		0.0000000	
Frequency Maternal Indeterminate	0.0000000		0.0000288		0.0000000	
Total Paternal Mutation Frequency	0.0000676		0.0000818		<0.0000570	
Total Maternal Mutation Frequency	0.0000785		0.0001146		0.0000445	

THO Maternal Mutations

Maternal Mutation – Allele:		Caucasian	Black	Hispanic	American Indian	Asian (Oriental)
From	To	Number	Frequency	Number	Frequency	Number
6	7					1
7	6					
7	8					
8	7	2	0.0000392			
8	9.3					
9	7	1	0.0000196			
9	8			5	0.0000858	
9.3	8	1	0.0000196			
10	8					
6 or 8	7 or 9					

8 or 10	9	Sum	4	0.0000785	5	0.0000858	1	0.0000445
Total Maternal Meiosis by Race			50,961		58,261		22,448	
	Frequency	0.0000785		0.0000858		0.0000445		

TPOX Paternal Mutations

Paternal Mutation – Allele:		Caucasian		Black		Hispanic		American Indian		Asian (Oriental)	
From	To	Number	Frequency	Number	Frequency	Number	Frequency	Number	Frequency	Number	Frequency
8	9										
9	10					2	0.0000442				
9	11					1	0.0000221				
10	9										
10	11	1	0.0000271 ^r								
11	10	1	0.0000271	2	0.0000442						
11	12	3	0.0000814								
11	16										
12	11	1	0.0000271					1	0.0000572		
12	13							1	0.0000572		
13	12										
10 or 12	11					1	0.0000221				
9 or 10	8 or 11					1	0.0000221				
9 or 11	10										
	Sum	6	0.0001628	7	0.0001548	2	0.0001143				
Total Paternal Meiosis by Race		36,845		45,229		17,496					

Paternal Frequency	0.0001628		0.0001548		0.0001143
Maternal Frequency	0.0000609		0.0001012		0.0000000
Sum	0.0002237		0.0002559		0.0001143
Relative Frequency Paternal	0.7278144		0.6047342		1.0000000
Relative Frequency Maternal	0.2721856		0.3952658		0.0000000
Indeterminate	0.0000343		0.0000000		0.0000000
Frequency Paternal Indeterminate	0.0000250		0.0000000		0.0000000
Frequency Maternal Indeterminate	0.0000093		0.0000000		0.0000000
Total Paternal	0.0001878		0.0001548		0.0001143
Mutation Frequency					
Total Maternal	0.0000702		0.0001012		<0.0000539
Mutation Frequency					

TPOX Maternal Mutations

Maternal Mutation – Allele:		Caucasian	Black	Hispanic	American Indian	Asian (Oriental)
From	To	Number	Frequency	Number	Frequency	Number
8	9	1	0.0000203			
9	10			1	0.0000202	
10	9			1	0.0000202	
11	10	1	0.0000203	1	0.0000202	
11	12	1	0.0000203	1	0.0000202	
11	16					
12	11					
12	13					

	13	12						
8 or 10	9				1	0.0000202		
9 or 11	10							
	Sum	3	0.0000609	5	0.0001012	0		
Total Maternal Meiosis by Race	49,229		49,427		18,539			
	Frequency	0.0000609		0.0001012		0.0000000		

VWA Paternal Mutations

Paternal Mutation – Allele:	Caucasian	Black	Hispanic	American Indian	Asian (Oriental)
Paternal Mutation – Allele:	Number	Frequency	Number	Frequency	Number
From	To				
14	11				
14	12			1	0.0000414
14	13			1	0.0000414
14	15				
15	14	2	0.0000375	4	0.0000583
15	16	6	0.0001126	13	0.0001895
15	14 or 16	1	0.0000188		
16	13	1	0.0000188		
16	15	5	0.0000938	12	0.0001749
16	17	9	0.0001689	10	0.0001457
16	18			2	0.0000291
16	15 or 17			2	0.0000291
17	16	18	0.0003379	11	0.0001603
17	18	20	0.0003754	21	0.0003061
17	16 or 18				6
					0.0002486

18	16	2	0.0000375	1	0.0000146		
18	17	22	0.0004129	19	0.0002769	3	0.0001243
18	19	16	0.0003003	18	0.0002623	6	0.0002486
18	20						
18	21			1	0.0000146		
18	17 or 19	4	0.0000751				
19	18	29	0.0005443	29	0.0004226	4	0.0001658
19	20	15	0.0002815	14	0.0002040	6	0.0002486
19	18 or 20			1	0.0000146		
20	19	17	0.0003191	17	0.0002478	3	0.0001243
20	21	2	0.0000375	1	0.0000146		1
21	19			1	0.0000146		
21	20	5	0.0000938	2	0.0000291	1	0.0000414
21	22						
22	21	1	0.0000188	2	0.0000291		
23	22			1	0.0000146		
10 or 12	11						
13 or 15	14			1	0.0000146		
14 or 16	15	1	0.0000188				
14 or 16	15 or 17						
14 or 18	16						
14 or 18	15 or 17			1	0.0000146		
15 or 17	16	1	0.0000188	4	0.0000583	1	0.0000414
15 or 17	14 or 18						
15 or 17	16 or 18						
15 or 18	16 or 17			1	0.0000146	1	0.0000414
15 or 18	16 or 19					1	0.0000414
15 or 19	16						
15 or 19	17						
15 or 19	14 or 20			1	0.0000146		
15 or 20	16 or 19					1	0.0000414

16 or 17	15 or 18						
16 or 18	17	3	0.0000563	4	0.0000583	1	0.0000414
16 or 18	15 or 17			2	0.0000291		
16 or 18	17 or 19	1	0.0000188				
16 or 19	15 or 18						
16 or 19	17 or 18	1	0.0000188	1	0.0000146		
17 or 18	16 or 19	1	0.0000188				
17 or 19	18	4	0.0000751	4	0.0000583	1	0.0000414
17 or 19	16 or 18			1	0.0000146		
17 or 19	18 or 20	1	0.0000188				
17 or 20	16 or 19						
17 or 20	18 or 19						
18 or 20	19	2	0.0000375	3	0.0000437	2	0.0000829
18 or 20	17 or 19						
18 or 20	19 or 21	1	0.0000188				
19 or 21	20						
	Sum	191	0.0035850	208	0.0030314	53	0.0021963
Total Paternal Meiosis by Race		53,278	68,616		24,132		523
Paternal Frequency	0.0035850		0.0030314		0.0021963		0.0133843
Maternal Frequency	0.0004066		0.0005800		0.0002622		
Sum	0.0039916		0.0036114		0.0024585		
Relative Frequency	0.8981353		0.8393870		0.8933355		
Paternal							
Relative Frequency	0.1018647		0.1606130		0.1066645		
Maternal							
Indeterminate	0.0009106		0.0006811		0.0005090		
Frequency Paternal Indeterminate	0.0008178		0.0005717		0.0004547		

Frequency Maternal	0.0000928		0.0001094		0.0000543	
Indeterminate						
Total Paternal						
Mutation Frequency	0.0044028		0.0036031		0.0026509	
Total Maternal						
Mutation Frequency	0.0004994		0.0006894		0.0003165	

VWA Maternal Mutations

Maternal Mutation – Allele:		Caucasian		Black		Hispanic		American Indian		Asian (Oriental)	
From	To	Number	Frequency	Number	Frequency	Number	Frequency	Number	Frequency	Number	Frequency
14	15										
15	14										
15	16	2	0.0000301	3	0.0000435	1	0.0000524				
16	15			3	0.0000435	1	0.0000524				
16	17	4	0.0000602	3	0.0000435	1	0.0000524				
16	18										
16	15 or 17			1	0.0000145						
17	15			1	0.0000145						
17	16										
17	18	3	0.0000452	5	0.0000725	1	0.0000524				
17	16 or 18										
18	17	1	0.0000151	4	0.0000580						
18	19	6	0.0000904	2	0.0000290	1	0.0000524				
18	20	1	0.0000151								
18	17 or 19										
19	18					2	0.0000290				
19	20	3	0.0000452	2	0.0000290						
20	19	1	0.0000151	1	0.0000145						
20	21			4	0.0000580						

21	20	1	0.0000151	1	0.0000145
21	22				
14 or 16	15			1	0.0000145
14 or 16	15 or 17				
14 or 18	16				
14 or 18	15 or 17	1	0.0000151		
14 or 20	15 or 19			1	0.0000145
15 or 17	16				
15 or 17	16 or 18	1	0.0000151		
15 or 18	16				
15 or 18	16 or 17			1	0.0000145
15 or 18	16 or 19				
16 or 17	15 or 18				
16 or 18	17	2	0.0000301	1	0.0000145
16 or 18	15 or 17				
16 or 18	15 or 19			1	0.0000145
16 or 18	17 or 19			1	0.0000145
16 or 19	15 or 18				
16 or 19	17 or 18				
17 or 19	18			1	0.0000145
17 or 20	16 or 19				
17 or 20	18 or 19				
17 or 21	16 or 22			1	0.0000145
18 or 20	19	1	0.0000151		
18 or 20	17 or 19				
19 or 21	20				
	Sum	27	0.0004066	40	0.0005800
Total Maternal Meiosis by Race		66,397	68,961	5	0.0002622
Frequency		0.0004066	0.0005800	0.0002622	

D16S539 Paternal Mutations

Paternal Mutation – Allele:		Caucasian		Black		Hispanic		American Indian	Asian (Oriental)
From	To	Number	Frequency	Number	Frequency	Number	Frequency		
8	9	1	0.0000237						
9	8	1	0.0000237						
9	10	2	0.0000473	2	0.0000367				
9	11			1	0.0000184				
10	11			5	0.0000918	1	0.00000510		
10	9 or 11								
11	9	1	0.0000237						
11	10	2	0.0000473	1	0.0000184	1	0.00000510		
11	12	6	0.0001220	17	0.0003120	1	0.00000510	1	
11	13			1	0.0000184				
11	14					1	0.00000510		
11	10 or 12	1	0.0000237						
12	10	1	0.0000237						
12	11	13	0.0003076	8	0.0001468	2	0.00001019	1	
12	13	10	0.0002366	12	0.0002202	3	0.00001529	1	
12	11 or 13	2	0.0000473						
13	9								
13	11								
13	12	7	0.0001656	8	0.0001468	1	0.00000510		
13	14	12	0.0002839	13	0.0002386	2	0.0001019	1	
13	12 or 14					1	0.00000510		
14	13	7	0.0001656	9	0.0001652	2	0.0001019		
14	15	3	0.0000710	3	0.0000551				
15	14			1	0.0000184	1	0.00000510		
15	16					1	0.00000510		

10 or 12	11	1	0.0000237	2	0.0000367
10 or 12	11 or 13				
10 or 12	9 or 13				
11 or 13	12	3	0.0000710	3	0.0000551
11 or 13	12 or 14			1	0.0000184
11 or 14	10 or 13				
11 or 14	12 or 13			1	0.0000184
11 or 15	12 or 14				
12 or 14	13	3	0.0000710	1	0.0000184
12 or 15	13 or 14				
13 or 15	14				
9 or 11	10	1	0.0000237		
9 or 12	10 or 11				
9 or 13	10 or 12				
	Sum	77	0.0018219	89	0.0016332
Total Paternal Meiosis by Race		42,264	54,494	19,622	229
Paternal Frequency		0.0018219	0.0016332	0.0010193	0.0174672
Maternal Frequency		0.0003009	0.0004497	0.0003204	
Sum		0.0021228	0.0020829	0.0013397	
Relative Frequency					
Paternal	Paternal	0.8582520	0.7840919	0.7608132	
Relative Frequency	Maternal	0.1417480	0.2159081	0.2391868	
Indeterminate	Indeterminate	0.0004088	0.0004497	0.0003204	
Frequency Paternal					
Indeterminate	Indeterminate	0.0003509	0.0003526	0.0002438	
Frequency Maternal					
Indeterminate	Indeterminate	0.0000579	0.0000971	0.0000766	
Total Paternal					
Mutation Frequency		0.0021727	0.0019858	0.0012631	

Total Maternal Mutation Frequency	0.0003588		0.0005468		0.0003971	
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D16S539 Maternal Mutations

Maternal Mutation – Allele:		Caucasian			Black			Hispanic			American Indian			Asian (Oriental)		
From	To	Number	Frequency	Number	Frequency	Number	Frequency	Number	Frequency	Number	Frequency	Number	Frequency	Number	Frequency	
8	9															
9	10	1	0.0000158			1	0.0000150									
10	9														1	
10	11					1	0.0000150									
10	9 or 11															
11	9					1	0.0000150									
11	10											1	0.0000458			
11	12															
12	11	3	0.0000475	5	0.0000750	5	0.0000750	1	0.0000458							
12	13															
12	11 or 13	1	0.0000158													
13	9															
13	11															
13	12	2	0.0000317	10	0.0001499											
13	14	1	0.0000158	2	0.0000300											
13	12 or 14	1	0.0000158													
14	13	4	0.0000634	5	0.0000750	1	0.0000458								1	
14	15					1	0.0000150	1	0.0000458							
15	14	2	0.0000317	1	0.0000150	2	0.0000916									
15	16															
10 or 12	11					1	0.0000150									
10 or 12	11 or 13															

10 or 12	9 or 13						
10 or 13	11 or 12	1	0.0000158				
11 or 13	12	1	0.0000158	1	0.0000150	1	0.0000458
11 or 14	10 or 13						
11 or 15	12 or 14						
12 or 14	13	2	0.0000317	1	0.0000150		
12 or 15	13 or 14						
13 or 15	14						
9 or 12	10 or 11						
9 or 13	10 or 12						
	sum	19	0.0003009	30	0.0004497	7	0.0003204
						0	2
Total Maternal Meiosis by Race		63,137	66,708		21,845		234
Frequency		0.0003009	0.0004497	0.0003204			0.0085470

Appendix 3. Mutations that are Indeterminate as to their parental origin.

CSF1PO Mutations that are Indeterminate as to the Parental Origin

MOTHER'S ALLELES	CHILD'S ALLELES	FATHER'S ALLELES	Caucasian	Black	Hispanic	American Indian	East Asian (Oriental)
11	10, 11	11					
10, 12	10, 11	10	1				
10, 12	10, 11	10, 12	1	1			
11, 12	10, 11	11, 12			2		
11, 12	10, 11	11, 13	1	1			
10, 11	10, 12	10, 11		1			
10, 11	10, 12	10, 13			1		
10, 13	10, 12	10, 11					
11, 12	10, 12	11					
10, 12	10, 13	10, 12					
10, 12	10, 13	10, 12					
10	10, 11	10			1		
10	10, 11	10, 12		1			
11	10, 11	11, 13		1			
11, 12	10, 11	11		1		1	
11, 12	10, 11	9, 11		1			
9, 10	10, 11	10		1			
10, 13	10, 12	10, 13		1			
11, 12	10, 12	11, 12			1		
11	11, 12	10, 11			1		
11	11, 12	11, 13					
11	11, 12	7, 11					
12	11, 12	10, 12		1			
10, 11	11, 12	11					
10, 11	11, 12	11, 13			2		
10, 12	11, 12	10, 12		1	3		1

11, 13		11, 12	11, 13	
8, 11		11, 12	11	1
8, 11		11, 12	9, 11	
9, 11		11, 12	11, 13	
9, 12		11, 12	12, 13	1
11, 12		11, 13	11, 12	2
11		11, 12	11	1
12		11, 12	12	2
10, 12		11, 12	12	1
10, 12		11, 12	12, 13	1
7, 12		11, 12	12	1
12		12, 13	10, 12	1
12		12, 13	11, 12	1
12		12, 13	8, 12	
13		12, 13	10, 13	
10, 12		12, 13	12	
10, 12		12, 13	10, 12	1
10, 13		12, 13	10, 13	1
11, 12		12, 13	12	1
11, 12		12, 13	10, 13	
11, 12		12, 13	11, 12	1
8, 12		12, 13	12	
8, 12		12, 13	8, 12	
12		12, 13	12	
11, 12		12, 13	12, 14	1
11, 13		12, 13	11, 13	1
9, 13		12, 13	11, 13	1
10, 13		13, 14	8, 13	2
10, 12		8, 10	10, 12	
10		8, 10	10, 11	1
8, 10		8, 11	8, 12	1

10	9, 10	10, 12						
10, 12	9, 12	11, 12						
10, 13	9, 10	10	1					
		Sum	20	27	14	0	2	
Indicate the number of cases.								
	31729	38566	15079	998			229	
Frequency								
	0.0006303	0.0007001	0.0000284	0.0000000	0.0087336			

D3S1358 Mutations that are Indeterminate as to the Parental Origin

MOTHER'S ALLELES	CHILD'S ALLELES	FATHER'S ALLELES	Caucasian	Black	Hispanic	American Indian	East Asian (Oriental)
15, 17	13, 15	15, 16					
14, 16	14, 15	14, 16					
14, 16	14, 15	14, 18					
15, 17	14, 15	15					
15, 18	14, 15	15, 17					
16	14, 16	15, 16					
14	14, 15	14, 16	1				
15	14, 15	15, 16	1				
14, 19	14, 15	14, 17	1				
14, 16	14, 17	14, 16		1			
15	15, 16	15, 17	1	2			
15	15, 16	15, 18			1		
16	15, 16	13, 16					
16	15, 16	14, 16					
16	15, 16	16, 17	1				
14, 15	15, 16	15					
15, 17	15, 16	15			2		
15, 17	15, 16	15, 17	1		1		
15, 18	15, 16	15					

16, 17	15, 16	16	2	1
16, 18	15, 16	16		
15, 16	15, 17	15, 16		
15, 16	15, 17	15, 18	2	
15, 18	15, 17	15, 16	1	
15, 18	15, 17	15, 18		
16, 17	15, 17	16, 17		
18	15, 18	17, 18		
14, 18	15, 18	14, 18		
15, 16	15, 18	15		
15, 16	15, 19	15		
16	15, 16	16	1	2
16	15, 16	16, 18	1	
14, 16	15, 16	14, 16		1
14, 16	15, 16	16, 17	1	
15, 18	15, 16	15, 17	2	
16	16, 17	16, 18		1
17	16, 17	15, 17	4	
17	16, 17	17, 18	1	1
14, 16	16, 17	15, 16		
14, 16	16, 17	16, 18	1	
15, 16	16, 17	15, 16		1
15, 16	16, 17	16, 18		
15, 17	16, 17	17	2	3
16, 18	16, 17	16		
16, 19	16, 17	16		
15, 18	16, 18	17, 18		
16, 17	16, 18	16		
17, 18	16, 18	17, 18	1	

		16	16, 17	16, 17	16	14, 16	1	2	
		16	16, 17	16, 17	16	15, 16	1	1	
16, 18	16, 18	16, 17	16, 17	15, 16	15, 16	15, 16	1	1	
16, 18	16, 18	16, 17	16, 17	16, 18	16, 18	16, 18	2	1	
17, 18	17, 18	16, 17	16, 17	15, 17	15, 17	15, 17	1	1	
16, 17	16, 17	16, 19	16, 19	16	16	16	1	1	
17	17	17, 18	17, 18	17	17	17			
17	17	17, 18	17, 18	16, 17	16, 17	16, 17			
14, 18	14, 18	17, 18	17, 18	15, 18	15, 18	15, 18			
15, 17	15, 17	17, 18	17, 18	15, 17	15, 17	15, 17	1	1	
16, 17	16, 17	17, 18	17, 18	16, 17	16, 17	16, 17			
16, 17	16, 17	17, 18	17, 18	17, 19	17, 19	17, 19			
16, 18	16, 18	17, 18	17, 18	16, 18	16, 18	16, 18			
15, 17	15, 17	17, 19	17, 19	16, 17	16, 17	16, 17			
18	18	17, 18	17, 18	16, 18	16, 18	16, 18	1	1	1
14, 18	14, 18	17, 18	17, 18	16, 18	16, 18	16, 18	1	1	
15, 18	15, 18	17, 18	17, 18	15, 18	15, 18	15, 18	1	1	
15, 18	15, 18	17, 18	17, 18	16, 18	16, 18	16, 18			
16, 17	16, 17	17, 18	17, 18	15, 17	15, 17	15, 17	1	1	
16, 18	16, 18	17, 18	17, 18	18	18	18	1	1	
18	18	18, 19	18, 19	18, 20	18, 20	18, 20			
14, 19	14, 19	18, 19	18, 19	19	19	19		1	
16, 18	16, 18	18, 19	18, 19	14, 18	14, 18	14, 18	1	1	
		Sum		22	28	28	10	0	3
		Indicate the number of cases.		32590	40581	15720	1060	229	
		Frequency		0.0006751	0.0006900	0.0006361	0.0000000	0.0131004	

D5S818 Mutations that are Indeterminate as to the Parental Origin

MOTHER'S ALLELES	CHILD'S ALLELES	FATHER'S ALLELES	Caucasian	Black	Hispanic	American Indian	East Asian (Oriental)
11, 12	10, 11	11	3		2		
11, 12	10, 11	11, 12			1		
10, 11	10, 12	10, 13					
10, 12	10, 13	10, 12					
9, 13	10, 13	11, 13					
11	10, 11	11, 12	1				
11, 12	10, 11	9, 11			1		
11, 12	10, 12	11, 12			1		
12, 13	10, 12	11, 12					
11, 13	10, 13	11, 13		1			
11	11, 12	11	1		1		
12	11, 12	12			1		
12	11, 12	8, 12		1			
10, 12	11, 12	12			1		
11, 13	11, 12	11, 13					
12, 13	11, 12	12		2			
12, 13	11, 12	10, 12					
9, 11	11, 12	11					
11	11, 13	11, 12					
10, 13	11, 13	12, 13					
11, 12	11, 13	11, 12		1	1		
11, 12	11, 13	11, 12					
11, 12	11, 13	11, 14	2		1		
12, 13	11, 13	12, 13					
11	11, 12	11, 13	2				
12	11, 12	10, 12	1				
11, 13	11, 12	11	4				
11, 13	11, 12	9, 11			1		
12, 13	11, 12	12, 13	1				

8,11	11,12	10,11		1
8,11	11,12	11,13		1
11,14	11,13	11,12	1	
12	12,13	11,12	2	1
12	12,13	12,14	4	
12	12,13	9,12	1	
13	12,13	13	1	
11,12	12,13	12		
11,12	12,13	12	1	1
11,12	12,13	10,12		
11,12	12,13	11,12	1	1
11,12	12,13	12,14	2	1
11,12	12,13	7,12		
11,12	12,13	9,12	1	
11,13	12,13	9,13	1	
7,13	12,13	13		
8,12	12,13	10,12	1	
8,12	12,13	12,14		
8,13	12,13	13		
11,12	12,14	12,13		
12	12,13	12	3	2
12,14	12,13	12	1	
11,13	13,14	13		
12,13	13,14	11,13	2	
12,13	13,14	11,13		
12,13	13,14	12,13	3	
12,13	13,14	12,13		
13	13,14	13	1	
7,11	7,12	7,11		

9, 12	9, 13	9, 12	Sum	27	25	15	0	0
Indicate the number of cases.								
				34028	40251	15721	1059	
				Frequency	0.0007935	0.0006211	0.0009541	

D7S820 Mutations that are Indeterminate as to the Parental Origin

MOTHER'S ALLELES	CHILD'S ALLELES	FATHER'S ALLELES	Caucasian	Black	Hispanic	American Indian	East Asian (Oriental)
10	10, 11	10					
10	10, 11	10, 12		3			
10	10, 11	8, 10		1			
11	10, 11	11					
10, 12	10, 11	10					
10, 12	10, 11	9, 10					
11, 12	10, 11	11		1			
11, 12	10, 11	11, 12					
8, 11	10, 11	11					
9, 10	10, 11	10		1			
9, 10	10, 11	10, 12		1			
10, 11	10, 12	10, 11					
10, 11	10, 12	10, 13		1			
10, 11	11, 12	11					
10, 11	11, 12	10, 11					
10, 12	11, 12	10, 12			1		
10, 12	11, 12	12, 13					
10, 12	11, 12	8, 12					
10, 12	11, 12	9, 12					

11, 13	11, 12	11, 13	
12, 14	11, 12	10, 12	
8, 11	11, 12	11	
8, 12	11, 12	8, 12	
9, 11	11, 12	8, 11	1
9, 11	11, 12	9, 11	
9, 12	11, 12	12	1
9, 12	11, 12	8, 12	1
10, 14	11, 14	10, 14	1
12	12, 13	10, 12	2
10, 12	12, 13	10, 12	
10, 13	12, 13	10, 13	1
11, 12	12, 13	9, 12	
8, 12	12, 13	12	
8, 12	12, 13	8, 12	1
9, 13	12, 13	11, 13	1
8, 13	13, 14	10, 13	1
8, 10	7, 8	8	1
8, 10	8, 11	8, 12	1
8, 11	8, 12	8, 11	
8, 13	8, 12	8, 13	
8, 12	8, 13	8, 12	
8, 11	8, 9	8, 11	
9	9, 10	9	1
10	9, 10	10, 12	
10, 11	9, 10	10	1
10, 11	9, 10	8, 10	
8, 10	9, 10	8, 10	
8, 9	9, 10	9, 11	1
9, 10	9, 11	9, 10	1
9, 10	9, 11	9, 12	1

9, 11	9, 12	9, 13	Sum	1	17	12	3	0	1
Indicate the number of cases.									
	Frequency	0.0005043		0.0003017	39774	15530	1050	229	
					0.0000000	0.0001932	0.0000000	0.00043668	

D8S1179 Mutations that are Indeterminate as to the Parental Origin

MOTHER'S ALLELES	CHILD'S ALLELES	FATHER'S ALLELES	Caucasian	Black	Hispanic	American Indian	East Asian (Oriental)
10, 15	10, 11	10, 13					
10, 15	10, 16	10, 15					
11	11, 12	11, 14					
11, 14	11, 12	11, 15					
10, 13	11, 13	10, 13					
12, 13	11, 12	10, 12					
11, 13	11, 14	11, 15	1				
12, 14	11, 14	12, 14					
13	12, 13	13					
13	12, 13	11, 13					
13	12, 13	13, 14					
13	12, 13	13, 15					
10, 12	12, 13	12, 14					
11, 13	12, 13	13, 15					
12, 14	12, 13	12					
9, 13	12, 13	13, 14			1		
10, 13	12, 15	13, 15					
12, 14	12, 15	12, 14					
13	12, 13	11, 13			1		
10, 12	12, 13	8, 12	1	1			
12, 13	12, 14	12, 15			1		

12, 15	12, 14	12, 13	1
13	13, 14	13	
13	13, 14	13, 15	1
14	13, 14	14	
14	13, 14	12, 14	1
14	13, 14	14, 15	
10, 13	13, 14	13	
10, 14	13, 14	11, 14	1
11, 13	13, 14	13	1
11, 13	13, 14	13, 15	2
11, 14	13, 14	12, 14	
11, 14	13, 14	14	
12, 14	13, 14	12, 14	
13, 15	13, 14	12, 13	1
13, 15	13, 14	13, 15	2
14, 16	13, 14	11, 14	1
14, 16	13, 14	12, 14	
9, 14	13, 14	14	
11, 15	13, 15	14, 15	
13, 14	13, 15	13, 14	
13	13, 14	12, 13	1
11, 13	13, 14	8, 13	2
12, 14	13, 14	14, 15	1
14	14, 15	11, 14	1
14	14, 15	12, 14	1
14	14, 15	14, 16	
10, 14	14, 15	12, 14	
10, 14	14, 15	13, 14	1
11, 14	14, 15	14	
11, 14	14, 15	12, 14	1
11, 14	14, 15	13, 14	1

12, 14	14, 15	11, 14	
12, 14	14, 15	14, 16	1
13, 14	14, 15	13, 14	1
13, 14	14, 15	14, 16	
13, 15	14, 15	13, 15	1
13, 15	14, 15	15, 16	
14, 16	14, 15	12, 14	
14, 16	14, 15	13, 14	1
14, 17	14, 15	14, 16	
15, 16	14, 15	15	
15, 16	14, 15	12, 15	
15, 16	14, 15	15, 16	
9, 15	14, 15	8, 15	
14, 15	14, 16	14, 15	1
15	14, 15	15	1
12, 15	14, 15	13, 15	1
13, 15	14, 15	15	1
15, 17	14, 15	13, 15	1
14, 17	14, 16	14, 15	1
14, 17	14, 16	14, 17	1
15	15, 16	13, 15	
11, 15	15, 16	15	2
12, 15	15, 16	14, 15	
13, 15	15, 16	12, 15	
13, 15	15, 16	14, 15	1
14, 15	15, 16	15	
14, 15	15, 16	13, 15	1
14, 16	15, 16	12, 16	
13, 16	15, 16	14, 16	1
10, 16	16, 17	13, 16	1
13	9, 13	13, 14	1

		Sum	15	25	12	0
Indicate the number of cases.		30379	40492	15675	1058	
	Frequency	0.0004938	0.0006174	0.0007656	0.0000000	

D13S317 Mutations that are Indeterminate as to the Parental Origin

MOTHER'S ALLELES	CHILD'S ALLELES	FATHER'S ALLELES	Caucasian	Black	Hispanic	American Indian	East Asian (Oriental)
8, 10	8, 9	8, 11					
8, 10	8, 11	8, 12					
9, 11	9, 10	9, 12					
9, 11	9, 12	9, 10					
11	10, 11	9, 11					
11, 12	10, 11	11	1				
11, 13	10, 11	11, 12		2			
11, 12	10, 12	11, 12					
11	11, 12	11		1			
11	11, 12	10, 11	1				
12	11, 12	12		2			
12	11, 12	12, 14	1		1		
11, 13	11, 12	11					
11, 14	11, 12	11, 13	2		1		
8, 11	11, 12	11				1	
9, 12	11, 12	12	1		1		
9, 12	11, 12	10, 12					
11	11, 13	11, 14					
11, 12	11, 13	11, 12					
11, 12	11, 13	11, 14		1			
11, 12	11, 13	9, 11					

11, 14	11, 13	11, 12	11, 12	1
12, 13	11, 13	12, 13	12	1
12	12, 13	12, 13	10, 12	3
12	12, 13	12, 13	12, 14	2
12	12, 13	12, 13	9, 12	4
13	12, 13	12, 13	13, 14	
10, 12	12, 13	11, 12		
10, 13	12, 13	13		1
10, 13	12, 13	12, 13	13, 14	
10, 13	12, 13	12, 13	8, 13	
11, 12	12, 13	12, 14		
11, 13	12, 13	12, 13	13	
11, 13	12, 13	12, 13	11, 13	
12, 14	12, 13	11, 12	1	1
12, 14	12, 13	12, 14		
7, 12	12, 13	11, 12	3	1
8, 12	12, 13	12	1	
8, 13	12, 13	11, 13	1	
8, 13	12, 13	8, 13		
9, 13	12, 13	11, 13		
12	12, 14	12, 13		
12, 13	12, 14	12, 15		
9, 13	13, 14	12, 13	2	2
13, 14	14, 15	11, 14		2
11, 12	12, 13	12		1
11, 13	12, 13	9, 13	1	
11, 14	13, 14	12, 14	1	
10, 12	10, 11	10, 12	1	
11, 13	13, 14	13	1	
11, 14	11, 12	8, 11	1	

12	11, 12	10, 12	1
8, 10	10, 11	10, 12	1
12, 13	11, 12	8, 12	1
12, 14	13, 14	11, 14	1
12	11, 12	8, 12	1
11, 13	11, 12	11, 13	1
11, 12	12, 13	8, 12	1
			0
		Sum	23

Indicate the number of cases.

Frequency	0.0007063	0.0006655	0.0005154	0.0000000

D16S539 Mutations that are Indeterminate as to the Parental Origin

MOTHER'S ALLELES	CHILD'S ALLELES	FATHER'S ALLELES	Caucasian	Black	Hispanic	American Indian	East Asian (Oriental)
10, 12	10, 11	10, 12					
10, 11	10, 12	10, 12					1
10	10, 13	10, 12					
10, 13	10, 14	10, 15					
10, 12	10, 11	10, 13					1
10, 11	10, 12	10, 11				1	
10, 14	10, 13	10, 12				1	
11	11, 12	11					
11	11, 12	11, 13			1	1	
11	11, 12	9, 11		1	1		
12	11, 12	10, 12		1			
12	11, 12	12, 13		1			
12	11, 12	9, 12					
11, 13	11, 12	11				1	
12, 13	11, 12	12		1			1

12,13	11,12	9,12	1	1
8,11	11,12	10,11	1	1
9,11	11,12	11,13		
9,12	11,12	12,13		
11	11,13	11,12		
11,12	11,13	11,12	2	1
11,12	11,13	11,14		
12,13	11,13	10,13		
11	11,14	11,13		
12	11,12	12	2	
10,12	11,12	9,12	1	1
11,13	11,12	11,13	1	1
9,11	11,12	11	1	1
11,14	11,13	11,14	2	
12	12,13	12	1	1
12	12,13	11,12	1	
10,13	12,13	10,13		
11,12	12,13	12	1	1
11,12	12,13	10,12	1	1
11,12	12,13	12,14		
11,13	12,13	13		
13,14	12,13	9,13	1	
8,12	12,13	12		
9,12	12,13	12		
9,13	12,13	11,13		
13	12,13	10,13	1	
13	13,14	12,13		
13	13,14	9,13		
11,13	13,14	11,13		
11,13	13,14	12,13		
12,13	13,14	10,13	1	

12, 13	13, 14	13, 14	12, 13	1
11, 13	13, 14	13, 14	13	1
9, 13	9, 10	8, 9	8, 9	1
9, 10	9, 11	9, 10	9, 10	
9, 12	9, 11	9, 10	9, 10	
9, 12	9, 11	9, 12	9, 12	
9, 12	9, 13	9, 13	9, 14	
9, 11	9, 10	9, 11	1	
		Sum	13	17
		Indicate the number of cases.	31803	39988
		Frequency	0.0004088	0.0004251
			0.0006469	0.0000000
			0.0043668	0.0043668

D18S51 Mutations that are Indeterminate as to the Parental Origin

MOTHER'S ALLELES	CHILD'S ALLELES	FATHER'S ALLELES	Caucasian	Black	Hispanic	American Indian	East Asian (Oriental)
19	10, 19	19, 20		1			
13, 17	12, 13	10, 13	1				
14, 15	12, 15	14					
12, 15	12, 13	12	1				
14	13, 14	14, 18					
12, 16	13, 16	12, 16		1			
15	14, 15	13, 15					
13, 14	14, 15	14, 16					
14, 18	14, 15	14					
14, 19	14, 15	13, 14					
15, 18	14, 15	15					
15, 19	14, 19	19					
14	14, 15	14, 18	1				
15	14, 15	15		1			

14, 15	14, 16	14, 17	1
13, 15	15, 16	12, 15	
13, 15	15, 16	15, 17	1
13, 15	15, 16	15, 19	
13, 16	15, 16	16, 20	
15, 17	15, 16	15	
15, 18	15, 16	14, 15	
16, 17	15, 16	16	
16, 17	15, 16	12, 16	1
16, 17	15, 16	16, 17	
14, 17	15, 17	14, 17	
15, 16	15, 17	15, 16	
15, 17	15, 18	15, 17	1
15, 18	15, 26	15, 21	
14, 16	15, 16	16	1
15, 17	15, 16	15, 17	1
16, 17	15, 17	16, 17	1
17	16, 17	11, 17	
12, 16	16, 17	16, 20	
13, 17	16, 17	13, 17	1
14, 16	16, 17	14, 16	2
14, 16	16, 17	16, 20	
14, 17	16, 17	15, 17	
16, 18	16, 17	16	
16, 18	16, 17	16, 18	1
16, 19	16, 17	14, 16	
16, 20	16, 17	16	1
17, 18	16, 17	15, 17	
17, 20	16, 17	17, 20, 2	
16, 17	16, 18	16, 17	
16, 19	16, 18	16, 17	

16,20	16,19	16,20	
17	16,17	15,17	1
15,16	16,17	16,18	1
17	17,18	17	
17	17,18	10,2,17	1
12,17	17,18	15,17	
14,17	17,18	16,17	
16,17	17,18	15,17	1
16,17	17,18	16,17	
16,18	17,18	16,18	
14,2,17	17,19	17,20	
17	17,18	17,19	1
13,18	17,18	18,20	1
19	18,19	16,19	
12,18	18,19	14,18	1
12,18	18,19	18,20	1
12,19	18,19	17,19	
16,18	18,19	18,20	
18,21	18,19	12,18	
12,19	19,20	19	1
14,19	19,20	13,19	
15,19	19,20	15,19	
15,19	19,20	17,19	
16,19	19,20	16,19	
21	20,21	18,21	1
12,20	20,21	16,20	1
		Sum	0
		10	18
		2	0
		0	0
Indicate the number of cases.		31231	38315
Frequency		0.0003202	0.0004698
		0.0001362	14679
		1007	

D19S433 Mutations that are Indeterminate as to the Parental Origin

MOTHER'S ALLELES	CHILD'S ALLELES	FATHER'S ALLELES	Caucasian	Black	Hispanic	American Indian	East Asian (Oriental)
13, 14, 2	12, 13	13					1
13	13, 14	13, 13, 2					
14	13, 14	12, 14					
14	13, 14	14, 15					
12, 14	13, 14	14, 14, 2	1				
13, 15	13, 14	13, 15				1	
13, 15	13, 14	13, 15, 2					
14, 15	13, 14	14					
14, 15	13, 14	14, 15					
13, 16	13, 15	13, 14					
13, 14	14, 15	14					
13, 14	14, 15	13, 14					1
13, 15	14, 15	12, 15					
14, 17, 2	14, 15	14, 16	1				
14, 16	16, 17	13, 16					
		Sum	1	1	0	1	2
		Indicate the number of cases.	2934	748			229
		Frequency	0.0003408	0.0013369			0.0087336

D21S11 Mutations that are Indeterminate as to the Parental Origin

MOTHER'S ALLELES	CHILD'S ALLELES	FATHER'S ALLELES	Caucasian	Black	Hispanic	American Indian	East Asian (Oriental)
28	28, 29	28, 30					

28	28,29	28,29	28,32,2
29	28,29	29,30.2	
28,30	28,29	28,31	
28,30	28,29	28,31.2	1
28,31	28,29	28,30	1
29,30	28,29	29	
29,31.2	28,29	29,30	
28,31	28,30	28,29	
28,31	28,30	28,31.2	
29	29,30	29	
29	29,30	29,31.2	
29	29,30	29,32.2	
29	29,30	29,35	
30	29,30	30	
30	29,30	24.2,30	1
28,29	29,30	29,30.2	1
28,30	29,30	30	
28,30	29,30	27,30	
29,31	29,30	29	
29,31	29,30	29,31.2	1
30,31	29,30	30	2
30,32.2	29,30	30	
30,32.2	29,30	27,30	1
30,34.2	29,30	28,30	2
29,30	29,31	29,30	1
30,31	29,31	29,31	
30	30,31	30	
30	30,31	28,30	2
30	30,31	30,32.2	
28,31	30,31	31,32.2	1
29,30	30,31	29,30	4
			2

29,30	30,31	28,30		
30, 31.2	30, 31	30, 30.2		1
31, 31.2	30, 31	31, 31.2		1
31, 32.2	30, 32.2	28, 32.2		
30, 32.2	30, 33.2	30, 33		
28, 32	31, 32	30, 32		
31.2	31.2, 32.2	28, 31.2		
29, 31.2	31.2, 32.2	30, 31.2	1	
30, 33.2	32.2, 33.2	30.2, 33.2		
28, 29	29, 30	29	1	1
28, 30	28, 29	28	1	1
28, 31	30, 31	29, 31	1	1
28, 31	31, 32	30, 31	3	1
28, 32.2	28, 29	27, 28	2	
28, 32.2	32.2, 33.2	32, 32.2	1	2
29, 31	30, 31	29, 31	1	1
31.2, 33.2	31.2, 32.2	29, 31.2	1	
31.2, 32.2	30.2, 31.2	29, 31.2	1	
27, 31	30, 31	29, 31	2	
28, 31	28, 29	28	1	
28, 30	30, 31	30	1	
28, 32.2	28, 31.2	28, 32.2		2
31, 33.2	31, 32.2	31, 33.2		1
		Sum	20	21
			5	0
		Indicate the number of cases.	33312	40408
			0.0006004	0.0005197
	Frequency		15654	1057
			0.0000000	0.0000000
			229	229
			0.0087336	0.0087336

FGA Mutations that are Indeterminate as to the Parental Origin

MOTHER'S ALLELES	CHILD'S ALLELES	FATHER'S ALLELES	Caucasian	Black	Hispanic	American Indian	East Asian (Oriental)
19,24	19,20	19,22					
19,20	19,21	19,22		1			
20,23	20,22	20,23					
21,22	20,22	21,23		1			
22,25	20,22	22,23					
20,22	20,23	20,24					
24,25	20,24	23,24					
20,25	20,26	20,27					
20,25	20,26	25,27					
19,21	21,22	21,23					
20,21	21,22	21,25					
21,24	21,22	21					
22,25	21,22	22,24					
21,22	21,23	21,24					
21,24	21,25	21,23		1			
21,24	21,25	21,24			1		
21,26	21,25	20,21					
21,26	21,25	21,26				1	
21,23	21,26	21,25					
20,22	21,22	22		1			
21,23	21,22	21,23			1		
21,24	21,23	21,22			1		
22	22,23	22		1			
22	22,23	22,24			1		
22	22,23	22,25			1	1	
18,2,23	22,23	21,23					
21,22	22,23	22			1		
21,22	22,23	22,24		2		1	
21,22	22,23	22,25				1	

21, 23	22, 23	21, 23
22, 24	22, 23	22, 25
23, 24	22, 23	23
23, 24	22, 23	23, 46
23, 24	22, 24	21, 24
23, 26	22, 26	23, 26
20, 22	22, 23	20, 22
22, 23	22, 24	22, 23
22, 25	22, 24	22, 25
23	23, 24	23
23	23, 24	21, 23
23	23, 24	23, 23, 3
24	23, 24	20, 24
19, 23	23, 24	18, 23
19, 23	23, 24	22, 23
20, 24	23, 24	22, 24
21, 23	23, 24	20, 23
22, 23	23, 24	23, 25
22, 24	23, 24	24
22, 24	23, 24	20, 24
23, 25	23, 24	23, 25
23, 26	23, 24	23, 25
24, 25	23, 24	22, 24
24, 26	23, 24	21, 24
24, 26	23, 24	24, 28
24, 27	23, 24	21, 24
22, 25	23, 25	24, 25
24	23, 24	24
18, 2, 24	23, 24	24
22, 23	23, 24	23
23, 25	23, 24	22, 23

24	24,25	24,25	20,24	1	1
24	24,25	24,25	21,24		
18,25	24,25	24,25	21,25	1	
18,2,24	24,25	24,25	23,24		
20,24	24,25	24,25	21,24	1	
20,24	24,25	24,25	22,24		
20,24	24,25	24,25	23,24		
21,24	24,25	24,25	24	1	
21,24	24,25	24,25	23,24		
22,24	24,25	24,25	21,24	1	1
22,24	24,25	24,25	21,24		
22,25	24,25	24,25	22,25	1	
23,24	24,25	24,25	24,26	1	1
25,26	24,25	24,25	23,25		
24	24,25	24	1	1	
18,2,25	24,25	25	1		
23,25	24,25	20,25	1		1
25	25,26	23,25	1		
21,25	25,26	25			1
23,26	25,26	22,3,26			
23,26	25,26	23,26			
26,27	25,26	21,26			
22,25	25,26	21,25	1		
24,25	25,26	25,28	1	1	
21,27	27,28	24,27			
27	27,28	24,27	1		
29	28,29	23,29	1		
		Sum	20	26	11
				0	2
Indicate the number of cases.		29002	39572	15277	1038
Frequency	0.0006896	0.0006570	0.0007200	0.0000000	0.0087336

THO Mutations that are Indeterminate as to the Parental Origin

MOTHER'S ALLELES	CHILD'S ALLELES	FATHER'S ALLELES	Caucasian	Black	Hispanic	American Indian	East Asian (Oriental)
7,8	8,9	7,8		1			
9,9,3	9,10	7,9		1			
		Sum	0	2	0	0	0
Indicate the number of cases.		29270	40510	15665	1056		
Frequency		0.0000494					

D2S1338 Mutations that are Indeterminate as to the Parental Origin

MOTHER'S ALLELES	CHILD'S ALLELES	FATHER'S ALLELES	Caucasian	Black	Hispanic	American Indian	East Asian (Oriental)
17,19	17,18	17					
17,25	17,19	17,18					
17	17,25	17,18					
17,26	17,27	17,26					
16,21	20,21	19,21					
23,24	23	20,24					
19,23	22,23	20,23					
20,23	22,23	21,23					
22	22,23	21,22					
16,23	23,24	21,23					
19,23	23,24	23,25					
17,25	24,25	22,25		1			
24	24,26	24,25					
18,26	25,26	17,26					

	Sum	1		
Indicate the number of cases.	2934			
Frequency	0.0003408			

VWA Mutations that are Indeterminate as to the Parental Origin

MOTHER'S ALLELES	CHILD'S ALLELES	FATHER'S ALLELES	Caucasian	Black	Hispanic	American Indian	East Asian (Oriental)
15, 16	12, 16	16, 19					
15	13, 15	15, 17			1		
15	14, 15	15, 20					
14, 18	14, 15	14					
14, 17	14, 18	14, 17					
14, 17	14, 18	14, 19	1				
15	15, 16	15, 19					
16	15, 16	16, 18					
15, 17	15, 16	15, 17					
15, 19	15, 16	15, 17					
15, 19	15, 16	15, 18					
16, 17	15, 16	16, 18					
15, 18	15, 17	15, 18	1				
15, 17	15, 18	15, 17					
15	15, 16	15	1				
14, 15	15, 16	15		2			
16, 18	15, 16	14, 16		1			
17	15, 17	17, 18	1				
15, 18	15, 17	15, 16		1			
15, 18	15, 19	15, 18		1			
17	16, 17	17, 19			1		
16	16, 17	15, 16			1		

16	16,17	16,17	16,18	2	2	2	1
14,16	16,17	16,17	16				
14,16	16,17	14,16					1
14,17	16,17	17					
15,16	16,17	16	1				
15,17	16,17	17,18					
15,17	16,17	17,19					
16,18	16,17	15,16					
16,18	16,17	16,18	1				
17,18	16,17	17					
17,18	16,17	14,17	1				
17,19	16,17	17					
17,19	16,17	17,19					
16	16,18	16,17					
15,16	16,18	16,17					
16,17	16,18	16,17	1				
16,17	16,18	16,19		2			
16,19	16,18	16,17					
16,19	16,18	16,19					
17,18	16,18	17,18					
16,18	16,19	16,18					
16,18	16,19	16,20					
17	16,17	17	1				
14,17	16,17	17,19	1				
15,16	16,17	16,18	2	1			1
15,17	16,17	15,17		2			
16,19	16,17	16,19					
17,18	16,17	15,17	2				
17,18	16,17	17,18	1				
15,18	16,18	15,18	1				

15, 17	17, 18	17, 19	1
16, 18	17, 18	18	
17, 19	17, 18	17	
16, 17	17, 19	17, 20	
17	17, 18	17	
17	17, 18	13, 17	
17	17, 18	14, 17	1
17	17, 18	15, 17	1
17	17, 18	16, 17	
17	17, 18	17, 19	
18	17, 18	18	
18	17, 18	14, 18	
13, 18	17, 18	15, 18	1
16, 17	17, 18	16, 17	
16, 17	17, 18	17, 19	1
17, 19	17, 18	17	
18, 19	17, 18	14, 18	
18, 20	17, 18	18	
14, 17	17, 20	17, 19	
18	17, 18	16, 18	1
17, 19	17, 18	15, 17	1
16, 19	17, 19	18, 19	1
17, 18	17, 19	17, 20	1
18	18, 19	16, 18	
14, 19	18, 19	17, 19	
15, 18	18, 19	17, 18	
15, 19	18, 19	16, 19	
16, 18	18, 19	14, 18	3
16, 18	18, 19	17, 18	1
17, 18	18, 19	18	2
17, 18	18, 19	16, 18	1

17, 18	18, 19	18, 19	18, 20	1
17, 19	18, 19	19		
18, 21	18, 19	17, 18		
19, 10	18, 19	17, 19		
19	18, 19	16, 19	1	
16, 19	18, 19	19		1
17, 19	19, 20	16, 19		
16, 20	19, 20	17, 20	2	1
19, 20	20, 21	18, 20		
Sum		31	28	8
Indicate the number of cases.		34044	41110	15718
Frequency		0.00091059	0.0006811	0.00050897
			0.0000000	0.0008733624