

1 Minutes (Draft)  
2 Scientific Advisory Committee  
3 Subcommittee on Y-STR analysis  
4 May 5, 2008 at 9:00 A.M.  
5 DFS Central Laboratory, Classroom 1  
6  
7

8 Subcommittee Members Present:

9 Dr. Robin Cotton  
10 Ms. Ann Marie Gross  
11 Dr. Dan Krane, Chair  
12 Mr. Barry Fisher  
13

14 Staff Members Present:

15 Mr. Jeff Ban, Central Laboratory Director  
16 Mr. Brad Jenkins, Forensic Biology Section Chief  
17 Ms. Beth Ballard, Forensic Biologist  
18 Ms. Angie Cunningham, Forensic Biologist  
19 Ms. Katie Hall, Forensic Biologist  
20

21 Call to Order:

22 Angie Cunningham called the meeting to order at 9:29 A.M. and asked the subcommittee  
23 to appoint a Chairperson. Dr. Krane was elected Chair.  
24

25 Validation Review:

26 Ms. Gross noted that some of the summaries were missing from the documentation that  
27 she was provided beforehand and Mr. Ban pointed out that all of the summaries were  
28 there in the binders presently provided including the Y-STR decision tree. The  
29 subcommittee members spent 15 minutes reviewing the four summaries not previously  
30 provided.  
31

32 Dr. Cotton asked Mr. Ban to explain the quantitative PCR process (Plexor<sup>®</sup>) that DFS  
33 was planning to implement. Mr. Ban described the mechanism of Plexor<sup>®</sup> analysis and  
34 Ms. Ballard provided an example of how the data would appear. Dr. Krane asked about  
35 the cost and Mr. Ban said that the Stratagene instrument is half of the price of an Applied  
36 Biosystems instrument. Some discussion ensued regarding the autosomal to Y DNA  
37 ratio that Plexor<sup>®</sup> calculates. Dr. Krane asked if the decision tree could be amended to  
38 include the possibility of going straight to Y-STR testing if the Plexor<sup>®</sup> data indicated an  
39 excess of female to male DNA. Mr. Ban maintained that autosomal DNA analysis would  
40 still be attempted before considering Y-STR analysis at this point; however the decision  
41 tree will be amended to include the option of Y-STR analysis in lieu of autosomal  
42 analysis.  
43

44 Dr. Krane asked if there was an idea of the percentage of Y-STR cases vs. autosomal that  
45 the lab would be analyzing. Mr. Ban said that there was no good sense of how many the  
46 lab may analyze. Dr. Cotton added that it is not unusual for the lab to be unsure of the

47 number of cases when going online. Ms. Gross agreed that the longer the lab is online;  
48 the number of cases could grow.

49  
50 Dr. Krane commended the lab on choosing values for the limit of detection and minimum  
51 peak height thresholds based on empirical data and not an arbitrary number. Since a  
52 threshold has been determined for casework, and some of the previous validation data  
53 had been analyzed with a lower threshold, he would like the previous data re-analyzed  
54 with the new thresholds. After discussion, it was decided that the current data could be  
55 reviewed to determine if the validation summaries needed to be updated based upon the  
56 new thresholds.

57  
58 Dr. Krane would like the limit of detection evaluated with each electrophoresis run since  
59 noise can change with time, as indicated in his publication. Ms. Gross disagrees with  
60 evaluating it with each run and thinks it's not necessary. Dr. Cotton also believes that  
61 each run is not necessary and would rather see consistency. Dr. Krane agrees with using  
62 the current values derived, but suggests the consideration of using run specific values in  
63 the future.

64  
65 Suggestions were made to improve and clarify the results in Table 1 for the precision  
66 validation. Ms. Gross asked how long we can use the capillary. Ms. Ballard indicated  
67 that the manufacturer's recommendations will be used until more runs were evaluated.  
68 Ms. Gross would like that procedure mentioned in the conclusions and Dr. Krane agreed  
69 that language needed to be worked up to that effect.

70  
71 Ms. Gross mentioned that there was no validation summary of female/male mixture  
72 samples in the specificity validation. Ms. Ballard explained that male/male mixtures and  
73 female samples were tested separately; although one female/male mixture was included  
74 with the non-probative casework study. Although the literature could be cited for other  
75 female/male mixture studies, Ms. Ballard offered to include a female/male mixture study  
76 in the validation.

77  
78 Ms. Gross asked why the stutter study did not include quantitation data and Ms. Ballard  
79 explained that those samples were extracted with QIAamp<sup>®</sup> and no quantitation was  
80 needed. Mr. Ban offered to add that statement to the summary. Dr. Cotton asked if a  
81 scatter plot was used in evaluating the data and Mr. Ban said that it was not used. She  
82 recommended that the lab be allowed to use the current stutter data in the interpretation  
83 guidelines, but recommended a plot for a training aid. Dr. Krane also recommends using  
84 a scatter plot to minimize the opportunity for the analyst to invoke her/his experience and  
85 expertise by going outside of the protocol.

86  
87 Some discussion ensued regarding the male/male mixture study. Suggestions were made  
88 to clarify the tables in the summary. Dr. Krane commented that the data does not support  
89 calling a major or minor contributor by using a 2-fold difference in peak heights.

90  
91 Ms. Ballard explained the process in making the environmental samples. Some  
92 discussion was held regarding Table 2 of the summary.

93

94 Mr. Ban explained how the tissue samples were acquired from the medical examiner's  
95 office and the possibilities for introduced contamination in one of the samples.

96

97 Ms. Ballard and Mr. Ban explained how the non-probative samples were obtained. It was  
98 suggested to add more detail to the procedure in how the samples were quantitated and  
99 the volume used.

100

101 The results of the concordance study were not evaluated pending verification by Ms.  
102 Gross.

103

104 Protocol Review:

105

106 Dr. Krane recommended including guidance for amplification of excess female to male  
107 DNA in the sample.

108

109 Ms. Gross asked if the flags in the GeneMapper<sup>®</sup> ID software would be used and if so,  
110 which flags. Ms. Ballard explained that the flags could be used for possible additional  
111 interpretation. Some discussion ensued regarding what is determined to be an actual peak  
112 versus an artificial one.

113

114 Regarding chapter 5, Dr. Krane believes it would be good to safeguard against contextual  
115 effect, if discretion was invoked, by shielding the analyst from the reference sample  
116 results. Mr. Ban clarified that the manual does not indicate that the reference sample will  
117 be used for that purpose, but Dr. Krane would like to see that the reference samples are  
118 excluded in print. A discussion was held regarding the need to express in the manual that  
119 reference samples will not be used for interpretation of the results and/or statistical  
120 calculations and exactly where in the manual that statement would be placed, should it be  
121 agreed upon.

122

123 The lack of statistical results for inclusion in a mixture was discussed. Dr. Krane referred  
124 to NRC I when discussing the need for a calculation and also the need for a theta  
125 correction. Discussion ensued regarding a calculation for mixtures and whether or not to  
126 include individuals if stats cannot be provided.

127

128 More discussion was needed and more validation work was advised, so the committee  
129 decided to adjourn without resolution regarding Y-STR endorsement.

130

131 Public Comment:

132

133 Public Comment was taken.

134

135 Adjourn:

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137 The meeting was adjourned at 8:30 P.M.