NIST has not submitted comments or rebuttals. The Editor's response is the only set of statements from NIST

| Comment Number | Comment | Editor's Response |
|---|---|---|
| 1 | We are looking at using 40 STR loci (the CODIS core 13 loci, 25 additional NIST loci, and the 2 sex-typing loci) to ensure that we have strong enough likelihood ratios that fully automated match results will be sufficient for kinship determinations. | All of these loci are included in the list |
| 2 | I would be very concerned if a DNA standard was published that was limited to only 13 loci or implied that 13 is sufficient | The table is not limited to the CODIS 13 loci |
| 3 | The DNA list should include markers for familial relationship checking | The X-STR and Y-STR loci have been cross-checked against several commerical family geneology sites to ensure that the markers that they test for are included |
| 4 | The DNA record should include all of the information in the ISO draft standard | The record contains that information and more, such as X-STR and a more extensive list of loci, as well as distinctions for lab and Rapid-DNA types of testing |