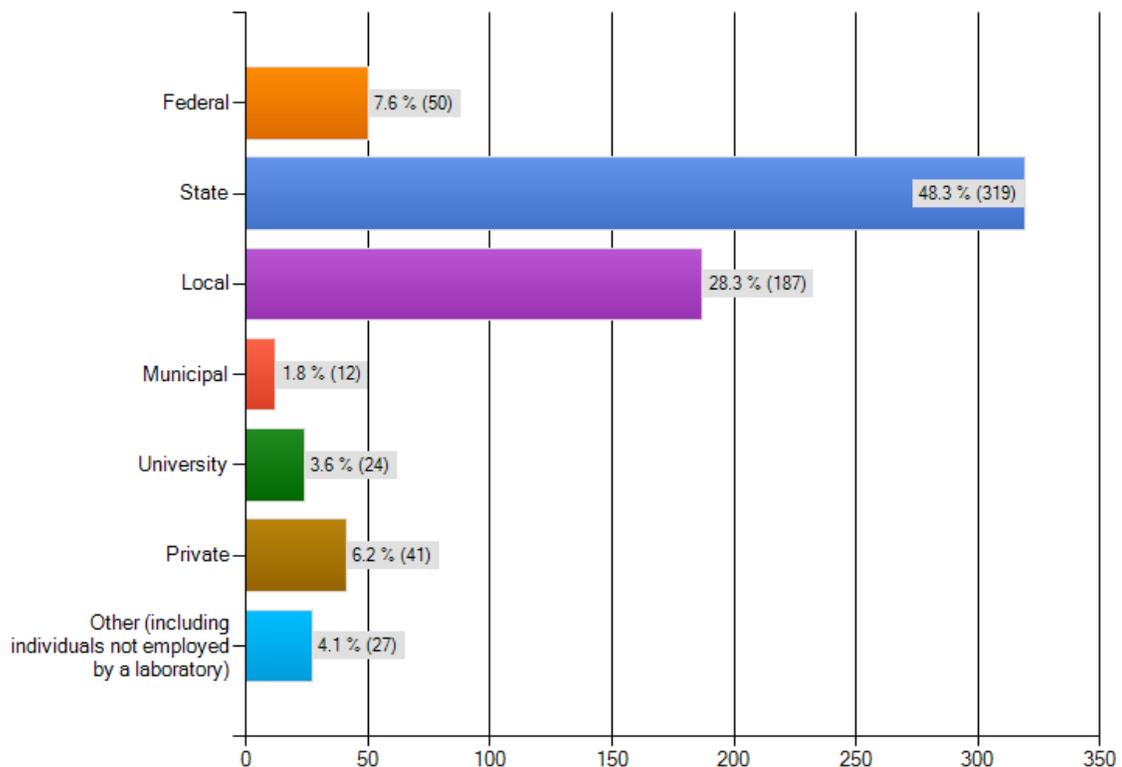


NIST DNA Mixture Interpretation Workshop

Poll Results from April 12, 2013

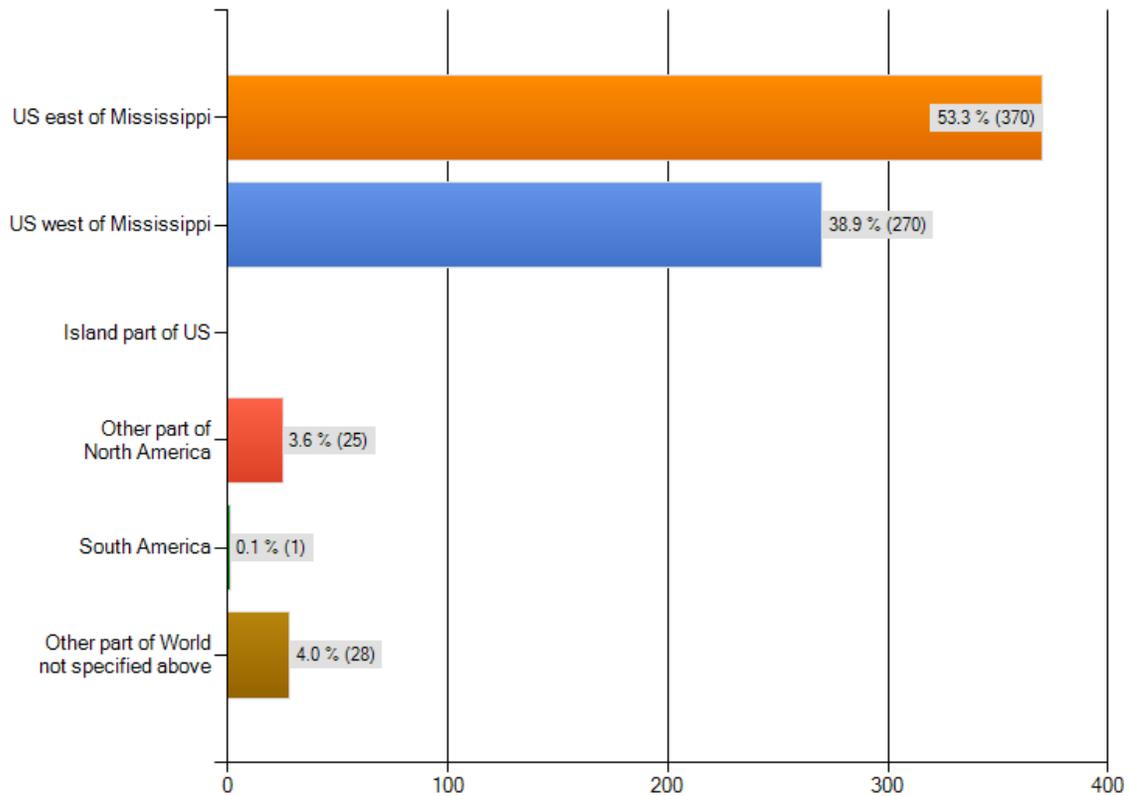
During the NIST DNA Mixture Interpretation Workshop and Webcast on April 12, 2013, a total of 20 poll questions were made available for workshop attendees and webcast viewers to answer. The responses were used as discussion points by the presenters during the event. Responses were received from individual viewers using computers or web browsers on smart phones. These anonymous responses were only presented in aggregate form as seen in this document. We are unable to determine the identity, occupation, or associated agency of the individuals that answered these poll questions. This document includes the aggregated responses to each of the 20 questions.

Poll Question 1: Please tell us what type of laboratory you work for (select the best single answer):



Total Responses: 660

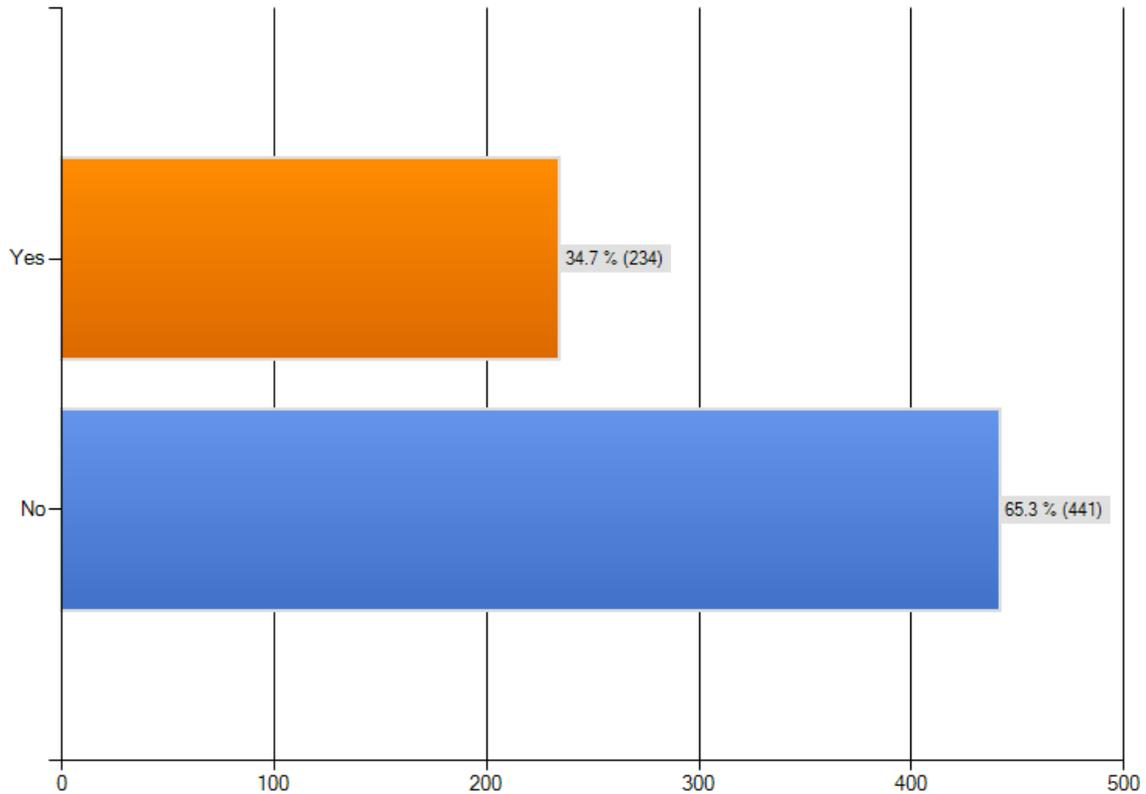
Poll Question 2: What part of the world are you watching from?



Total Responses: 694

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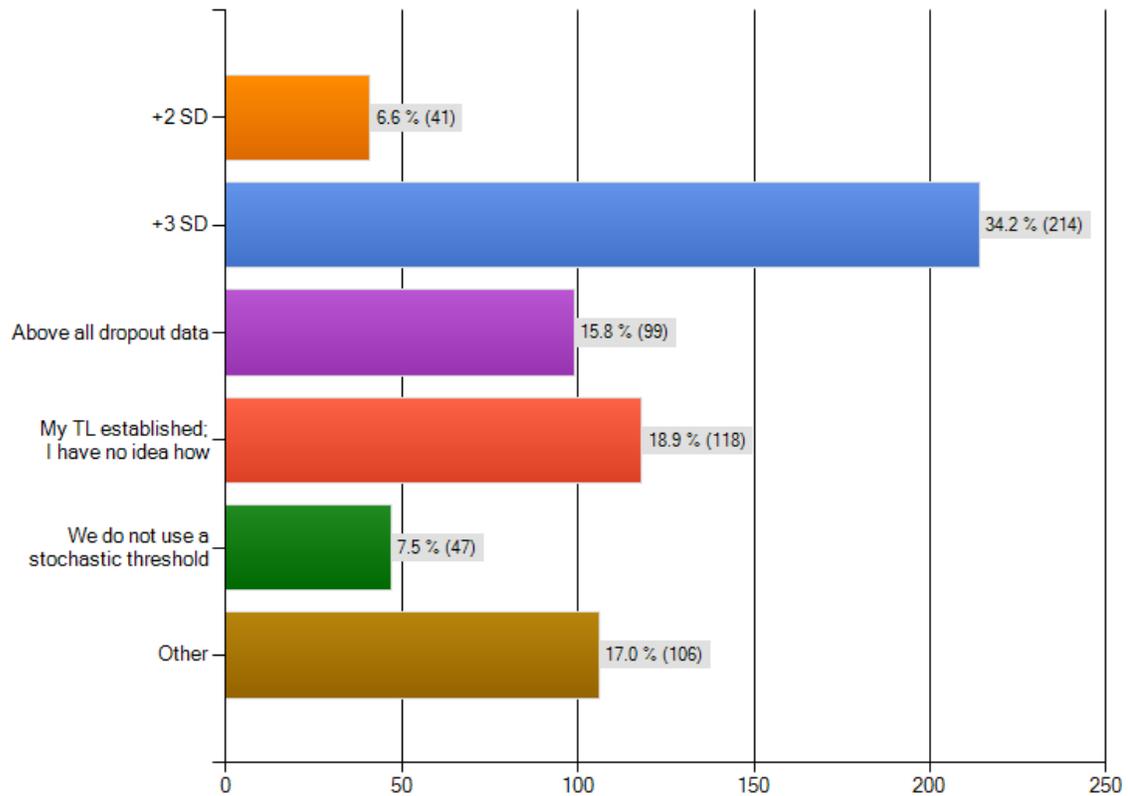
Poll Question 3: Would you prefer specific protocols for every detail in your lab?



Total Responses: 675

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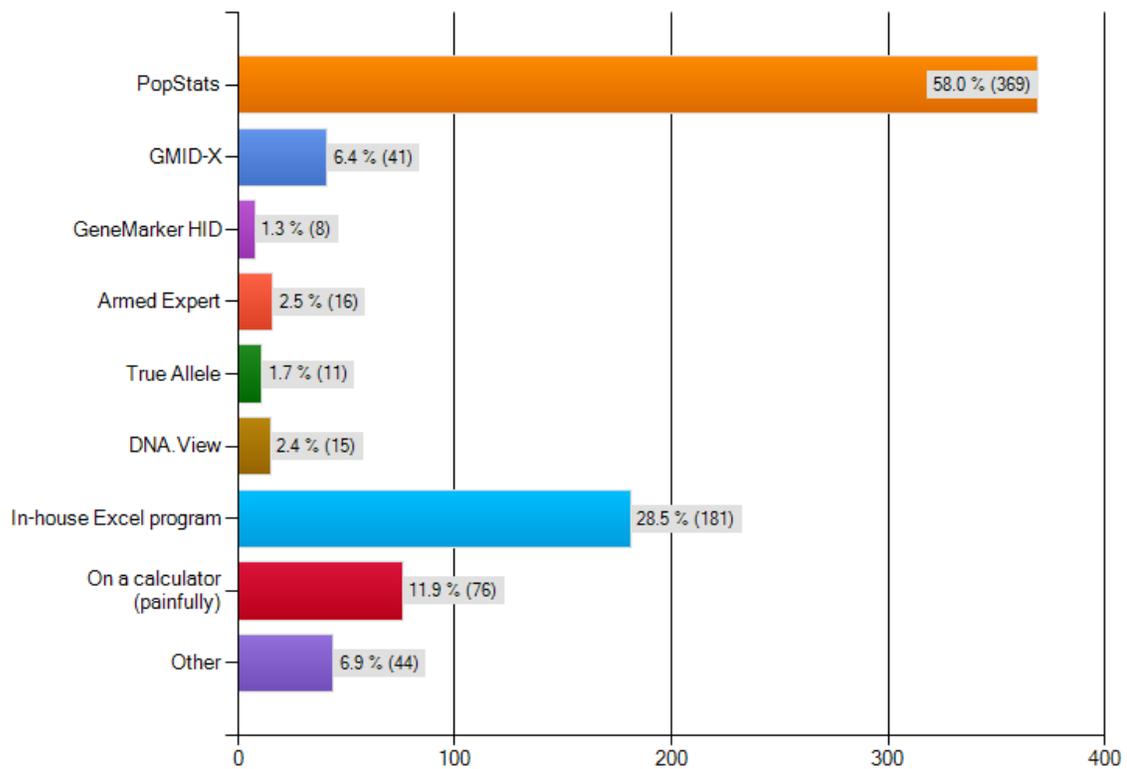
Poll Question 4: How were the RFU levels set for your laboratory stochastic threshold?



Total Responses: 625

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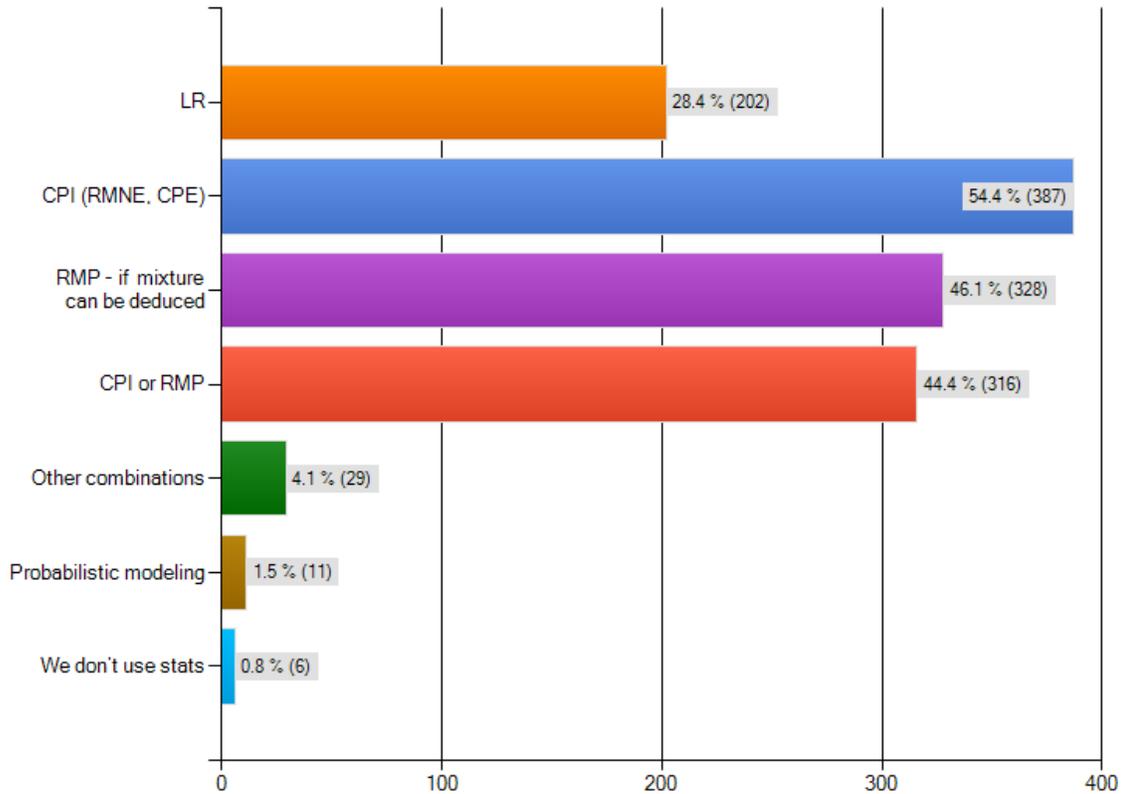
Poll Question 5: Does your lab use any software to help calculate mixture stats? Check all that apply:



Total Responses: 636

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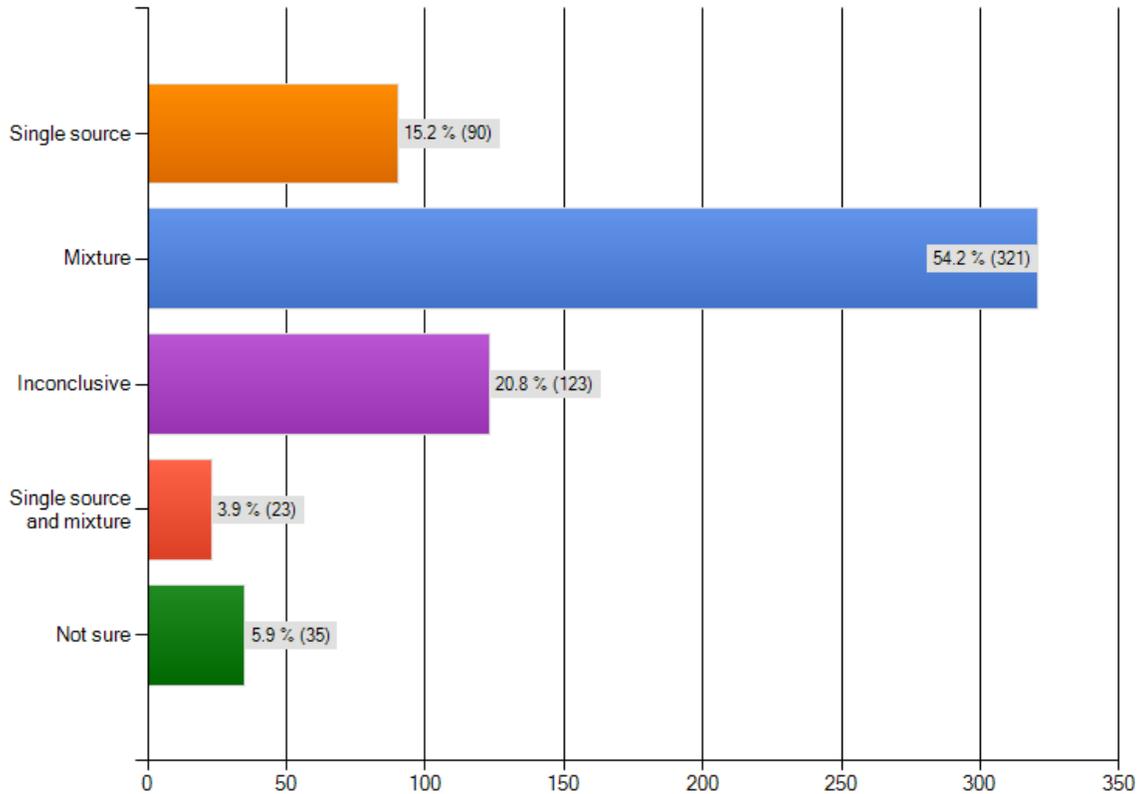
Poll Question 6: What kind of mixture statistic does your lab use? Check all that apply:



Total Responses: 711

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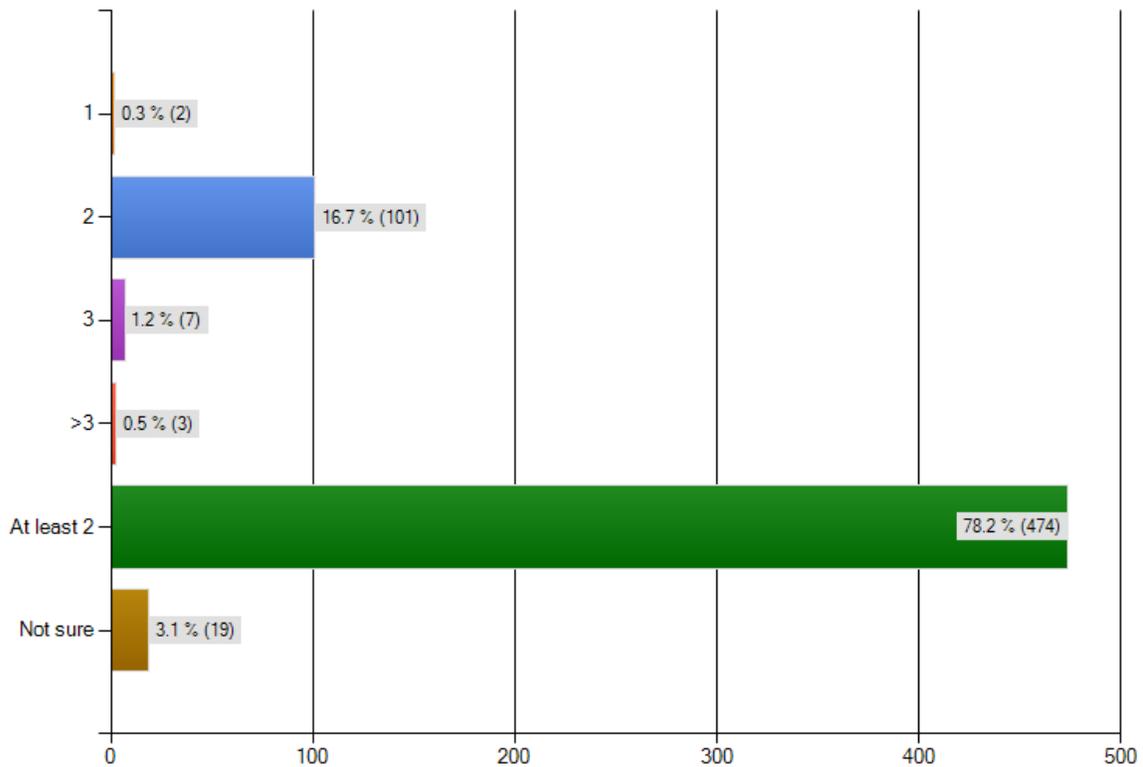
Poll Question 7: I would report Profile #4 (see Example Profiles) as:



Total Responses: 592

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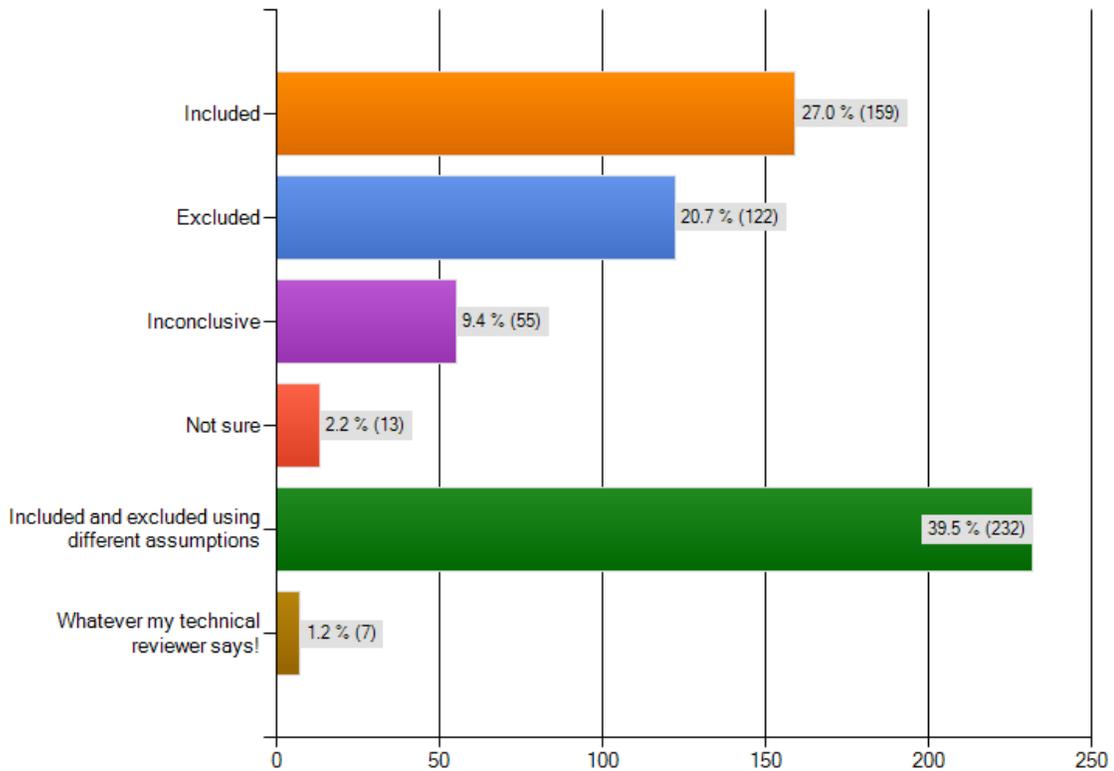
Poll Question 8: I am confident that the number of contributors in Profile 5 (see Example Profiles) is...



Total Responses: 606

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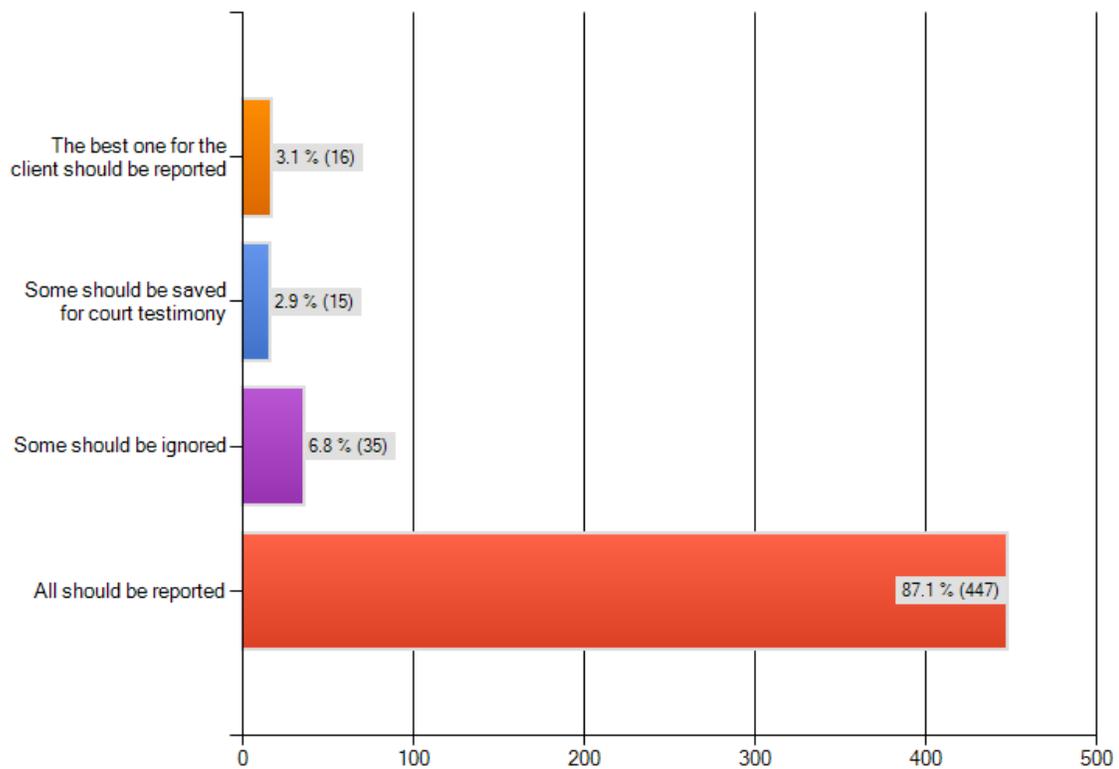
Poll Question 9: The known individual (genotypes 13,14 and 28, 30) compared to the two loci shown from Profile 5 (see Example Profiles) is



Total Responses: 588

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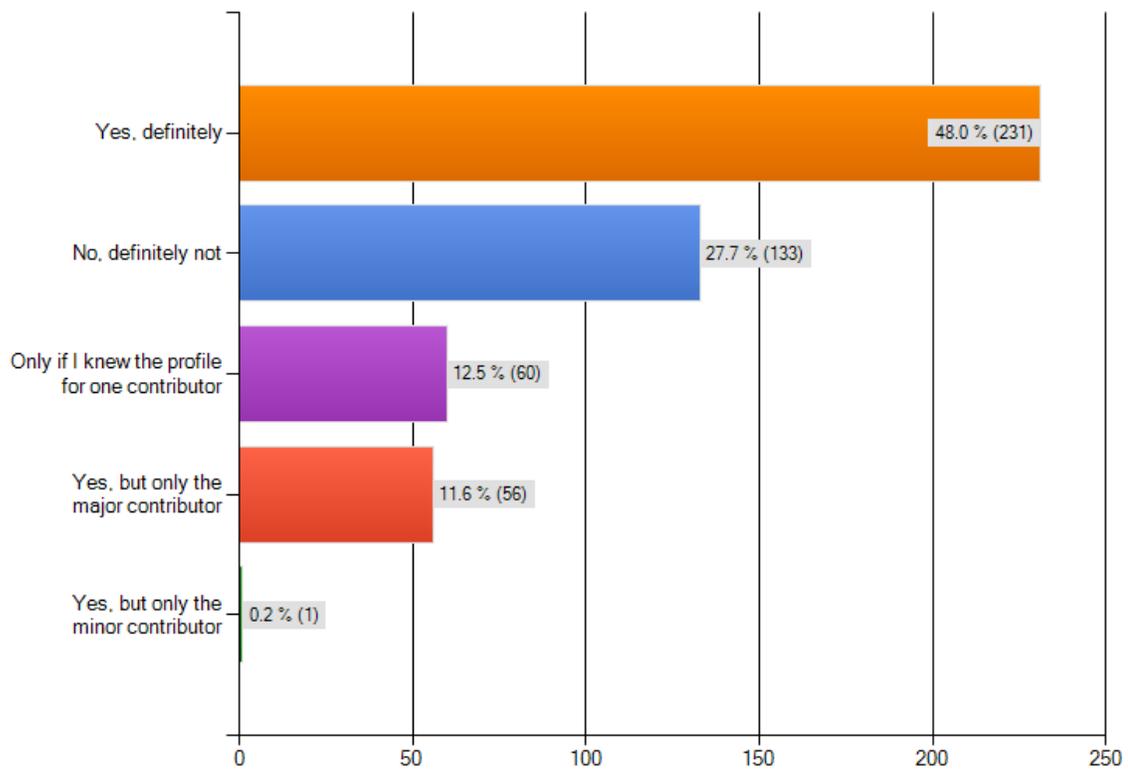
Poll Question 10: If a sample has more than one possible conclusion using different assumptions, then...



Total Responses: 513

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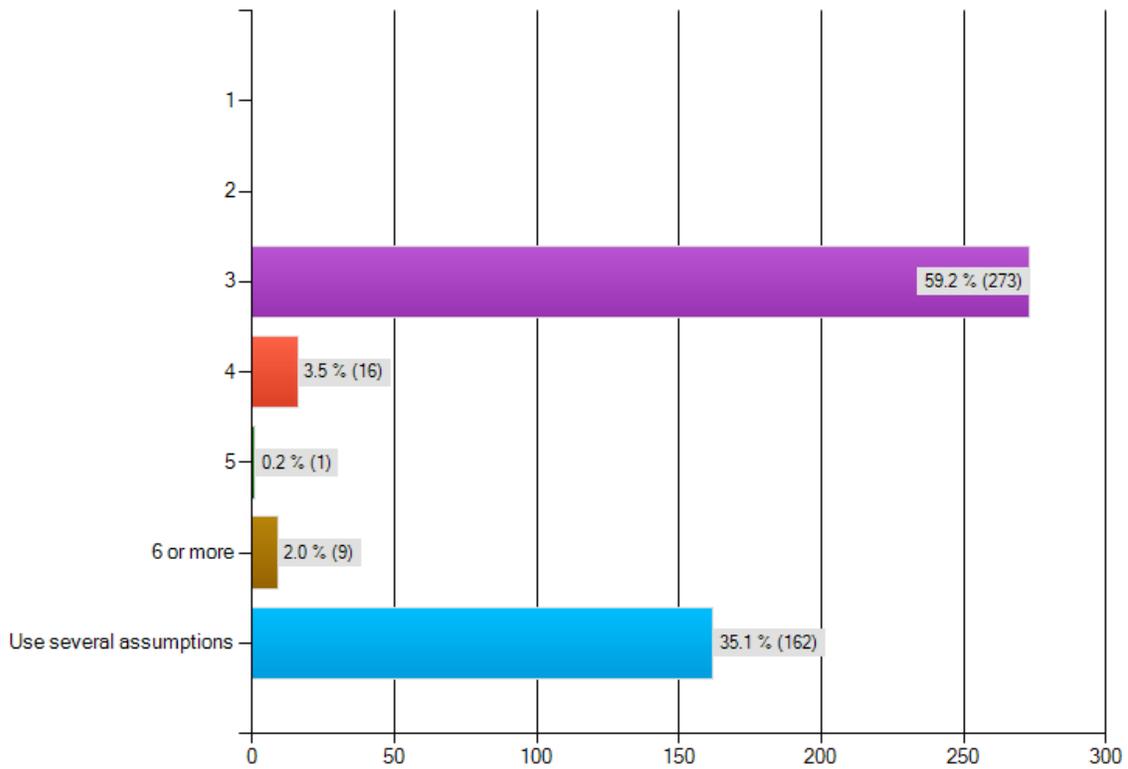
Poll Question 11: Would you attempt to interpret the mixture in Profile 9 (see Example Profiles)?



Total Responses: 481

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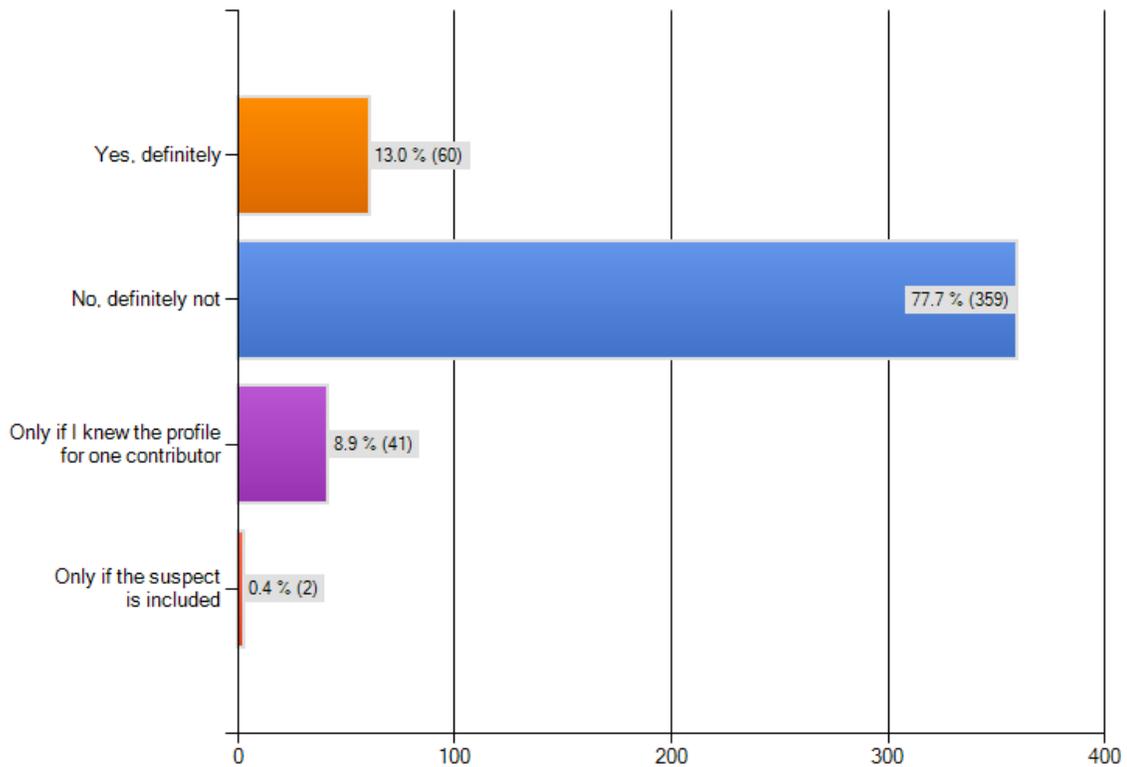
Poll Question 12: How many contributors should be assumed for interpretation of Profile 9 (see Example Profiles)?



Total Responses: 461

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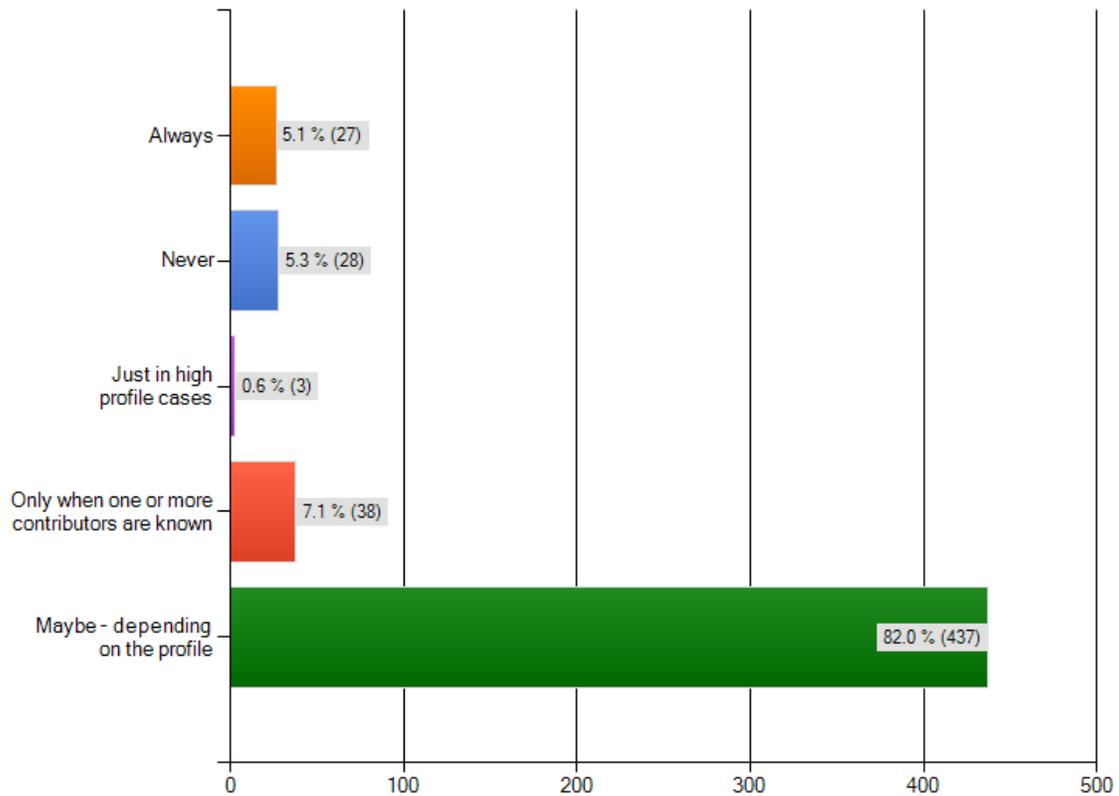
Poll Question 13: Can alleles for a major contributor be determined for Profile 9 (see Example Profiles)?



Total Responses: 462

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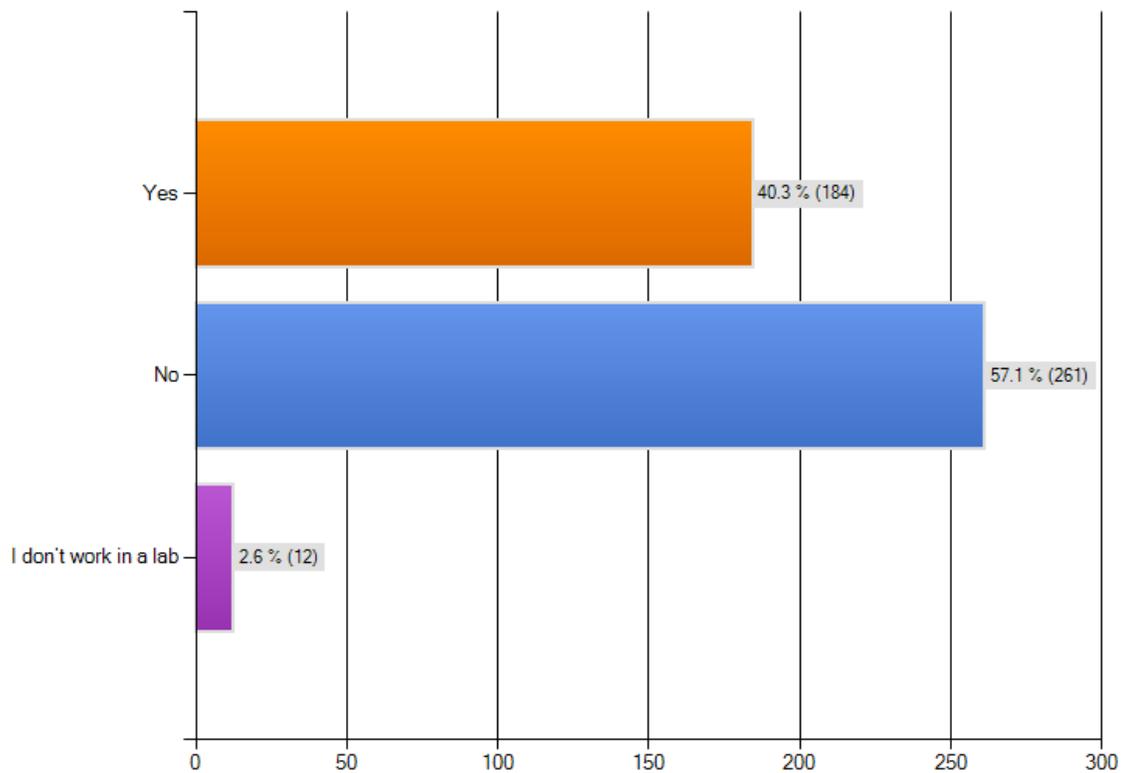
Poll Question 14: Should we be interpreting mixtures with 3 or more contributors?



Total Responses: 533

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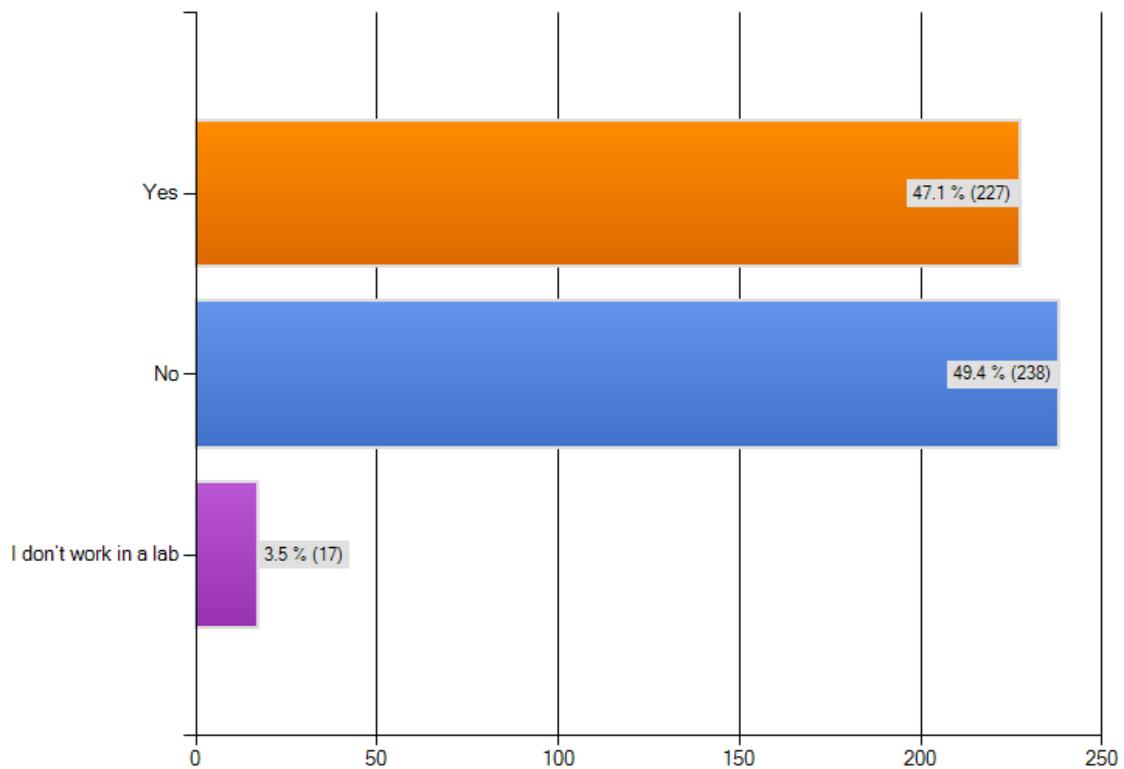
Poll Question 15: Looking at Profile 10 (see Example Profiles), would you do a CPE/CPI statistic on TPOX and D5S818 because all alleles are above the stochastic threshold?



Total Responses: 457

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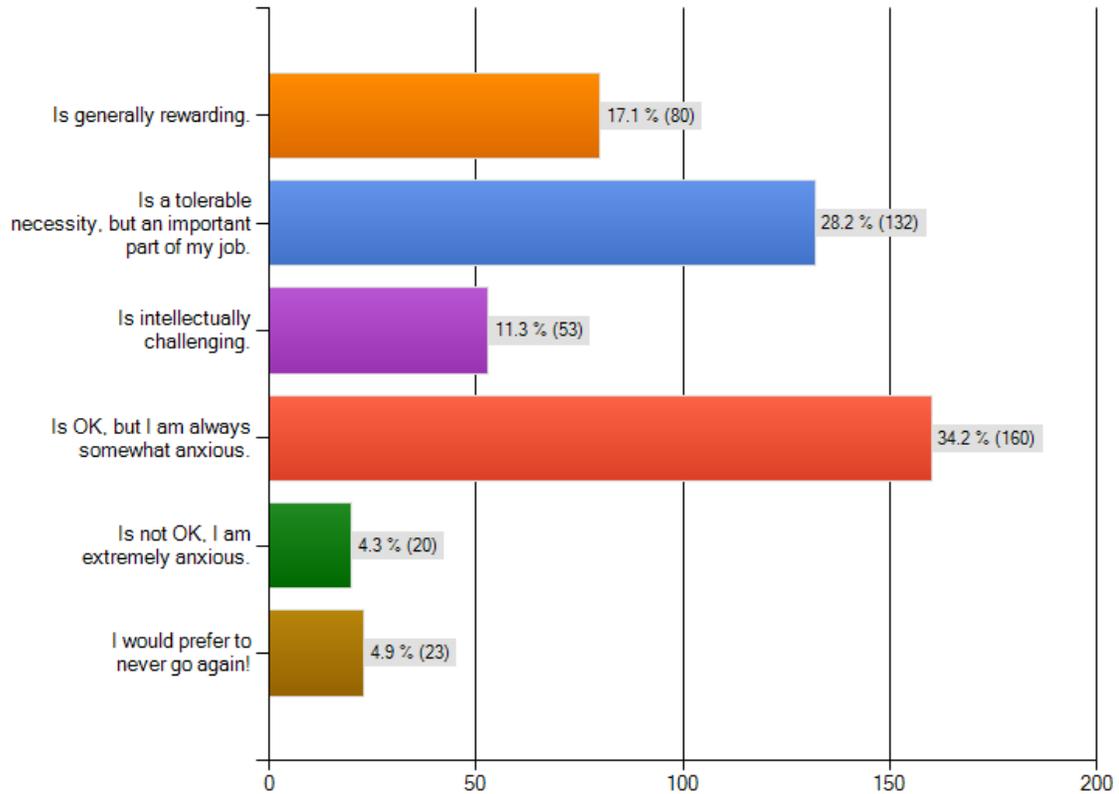
Poll Question 16: Has your laboratory implemented a “stop testing” approach with complex and/or low-level DNA mixtures?



Total Responses: 482

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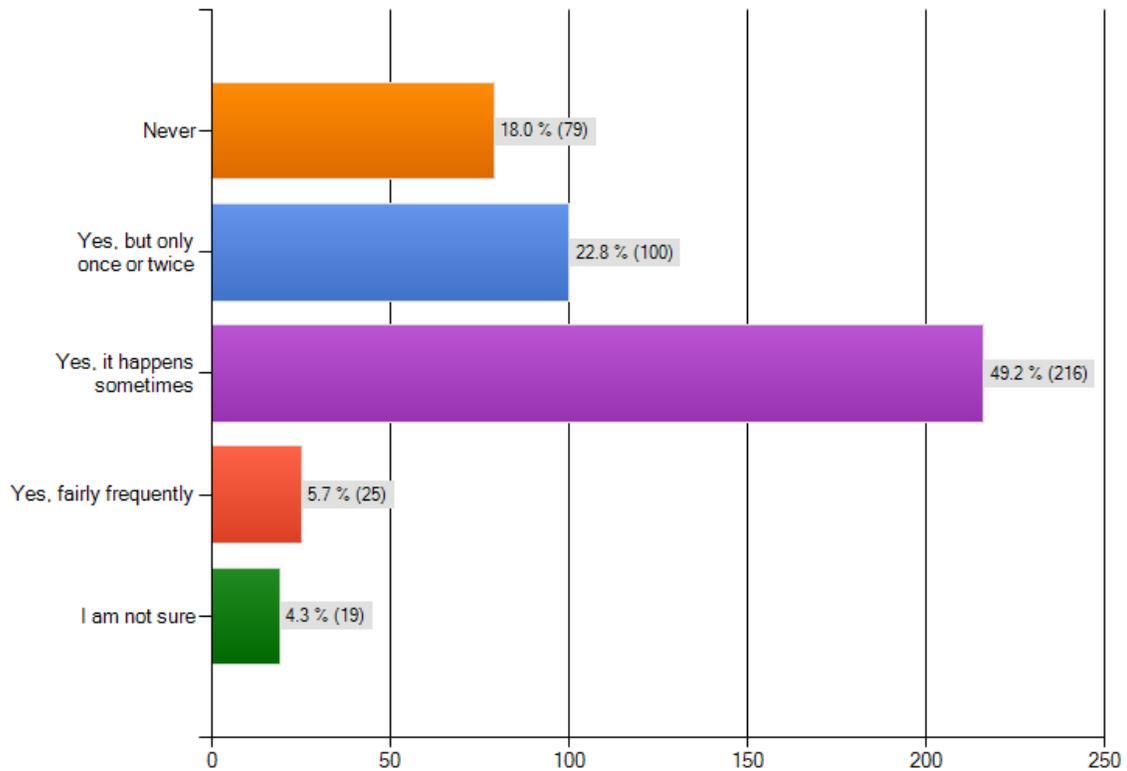
Poll Question 17: For me, testifying in court...



Total Responses: 468

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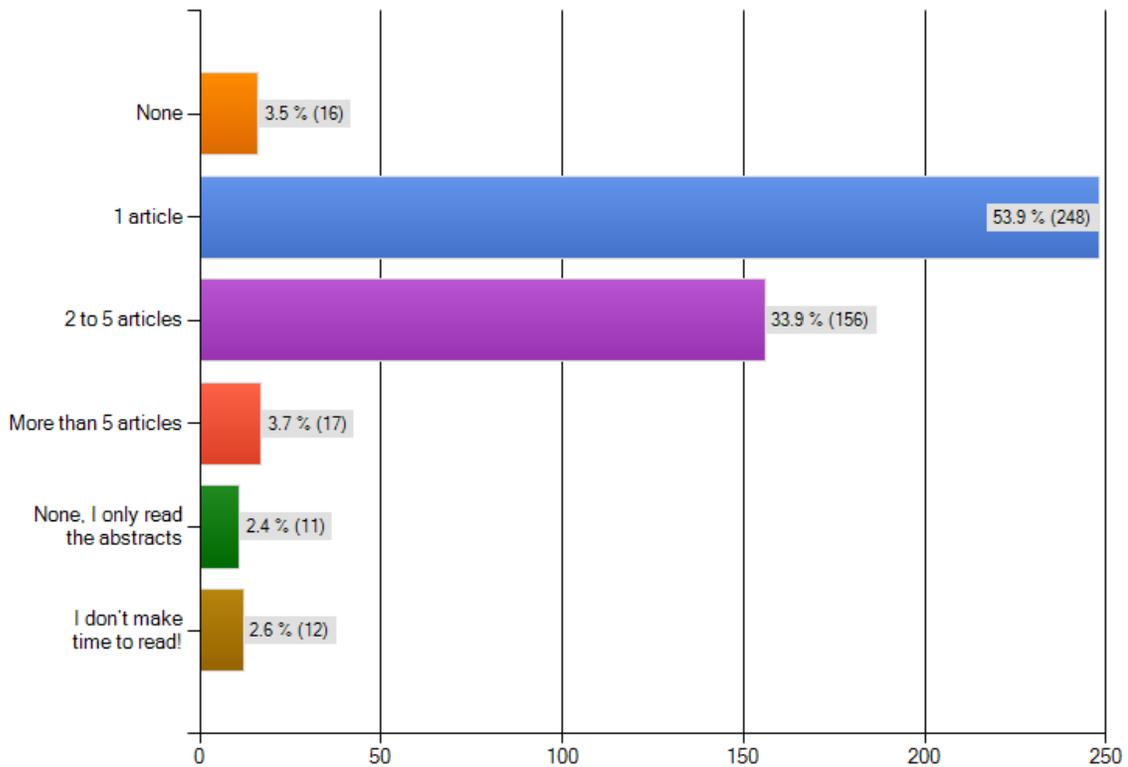
Poll Question 18: Have there been any occasions when you have felt pressure from an attorney to make stronger statements than you would otherwise make?



Total Responses: 439

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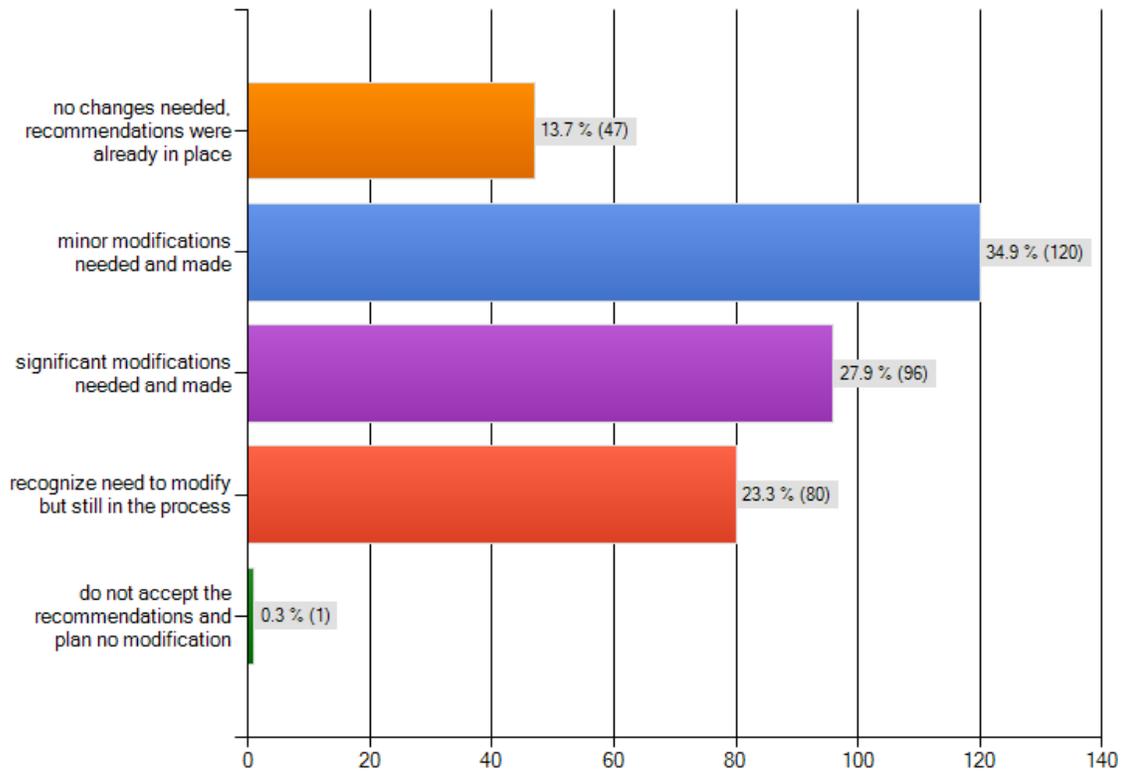
Poll Question 19: How many DNA-related articles would you estimate that you read in a typical month?



Total Responses: 460

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Poll Question 20: Our laboratory SOPs have been reviewed and the following is occurring in response to the 2010 SWGDAM Interpretation Guidelines:



Total Responses: 344

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