



The Benefit of Using the QIAGEN MinElute® PCR Purification Kit for Post PCR Cleanup on Low Level DNA Samples

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Abstract:
This study was designed to evaluate the similarities and differences in capillary electrophoresis signal detection when using the QIAGEN MinElute® PCR Purification Kit on amplified DNA obtained from commonly used short tandem repeat (STR) commercial amplification kits.

The following commercially available STR amplification kits used in this study were:

- Applied Biosystems' Ampf/STR®:** Promega's:
- Profiler Plus® ID kit
 - Cofiler® kit
 - Identifiler® kit
 - MiniFiler™ kit
 - Yfiler® kit
 - PowerPlex® 16 system
 - PowerPlex® Y system
 - PowerPlex® S5 system

Serial dilutions prepared from DNA extracts ranging from 1.0 ng to 0.0078 ng were evaluated on each of the amplification kits. Concentrations at which data fell below 75 RFUs were run through the clean up columns.

Introduction:
This study was conducted to demonstrate the benefits of including the QIAGEN MinElute® PCR Purification Kit into the analysis scheme on forensic samples containing low quantities of DNA.

The QIAGEN MinElute® PCR Purification Kit uses a silica membrane to bind DNA fragments ranging in size from 70 bp to 4kb. While the DNA is bound to the membrane, impurities such as unwanted primers, salts, enzymes, unincorporated nucleotides, dyes, oils, and detergents flow through the column. This causes an increase in the signal from the amplified DNA which in turn increases the peak height of the resulting data allowing previously low level samples to reach to a callable level. Removal of these impurities ensures that more DNA is injected during the electrokinetic injection on the instrumentation, thus increasing the fluorescent signal intensity.

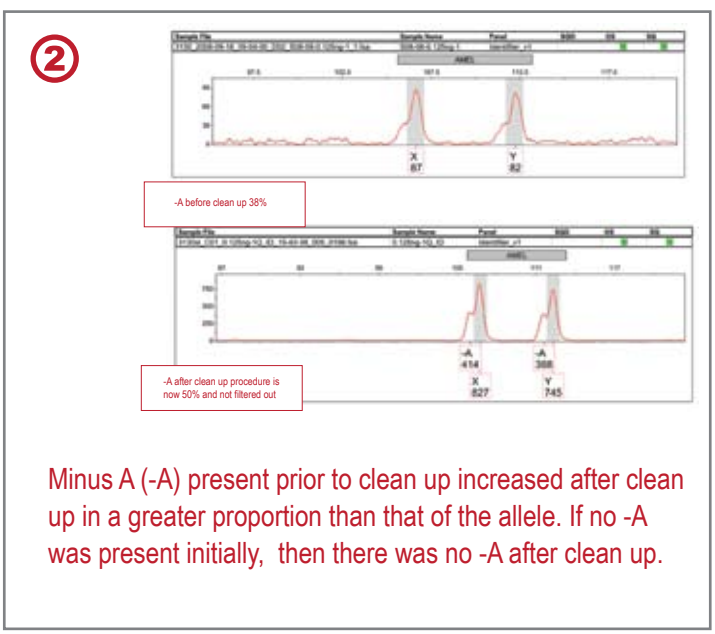
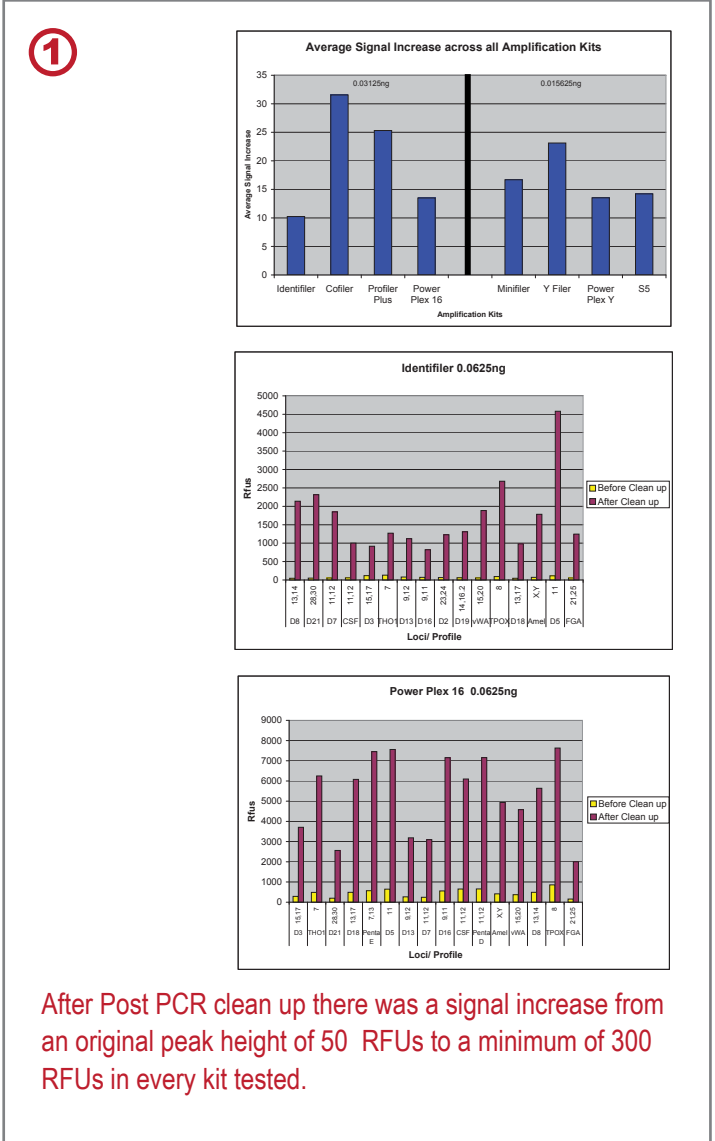
- Materials:**
- Standards from 2 male donors
 - Phenol:chloroform:Isoamyl alcohol (25:24:1)
 - TE Buffer, DTT and Proteinase K (10ng/ul)
 - Applied Biosystems Human DNA Quantifiler® Kit
 - Applied Biosystems' Ampf/STR® Profiler Plus® kit, Cofiler® kit, Identifiler® kit, MiniFiler™ kit, and the Yfiler®
 - Promega's PowerPlex® 16 system, PowerPlex® Y system,

- and the PowerPlex® S5 system
- Running Buffer, 10X
 - 16 capillary array, 36cm
 - POP-4™ polymer for 3130x/
 - Matrix standards
 - Internal Lane Size Standards
 - Hi-Di™ Formamide
 - 96-Well GeneAmp® PCR System 9700
 - 7500 Real-Time PCR System
 - 3130x/ Genetic Analyzer
 - QIAGEN MinElute® PCR Purification Kit.

- Method:**
- Two separate known human male DNA standards were prepared utilizing a standard organic extraction method in conjunction with the Millipore Microcon® 100 centrifugal filter device.
 - Serial dilutions were prepared from DNA extracts at the following concentrations: 1.0, 0.5, 0.25, 0.125, 0.0625, 0.03125, 0.015625, and 0.0078 ng.
 - The samples were quantitated using the Applied Biosystems Quantifiler® Human Quantification Kit on an Applied Biosystems 7500 Real-Time PCR System. The results were normalized with NIST quantitation standards.
 - Samples were amplified on an Applied Biosystems GeneAmp® PCR 9700 thermal cycler following manufacturer's specifications.
 - The samples were then separated and detected using an Applied Biosystems 3130x/ Genetic Analyzer using manufacturer's recommendations:

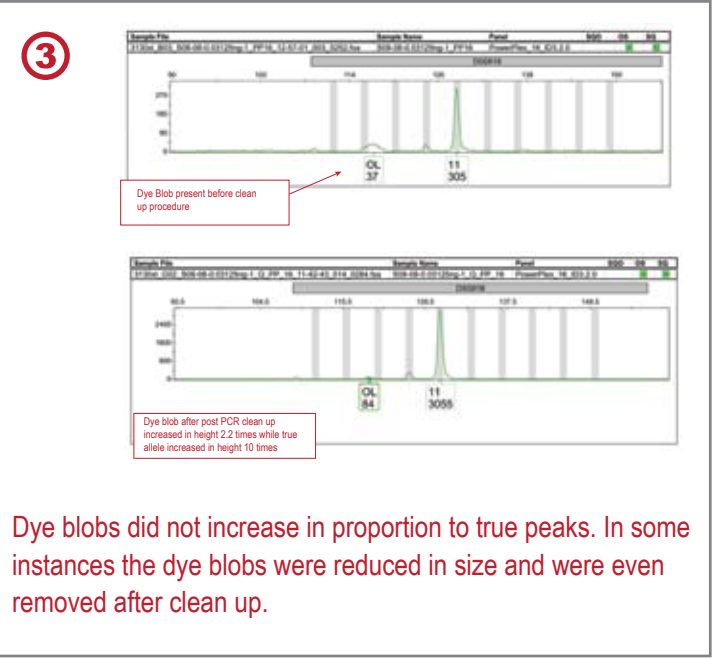
- Applied Biosystems kits: 3kv, 10 sec injections, 8.7 µl Formamide, 0.3 µl GS 500, 1 µl sample.
- Promega kits: 3kv, 10 sec injections, 1200 sec run time, 9.5 µl Formamide, 0.5 µl ILS 600, 1 µl sample
- Data were analyzed using GeneMapper® ID Software v3.2 using a threshold of 75 RFUs.
- Amplified samples from these dilutions which had data falling below a 75 RFU threshold were purified using the QIAGEN MinElute® PCR Purification Kit. A total of 4 washes were performed manually on each sample.
- After the post PCR clean up procedure the samples were then run on the Applied Biosystems 3130x/ Genetic Analyzer under the same conditions. All of the purified amplification product, which ranged between 9 and 10 µl, was consumed

Results and Discussion:

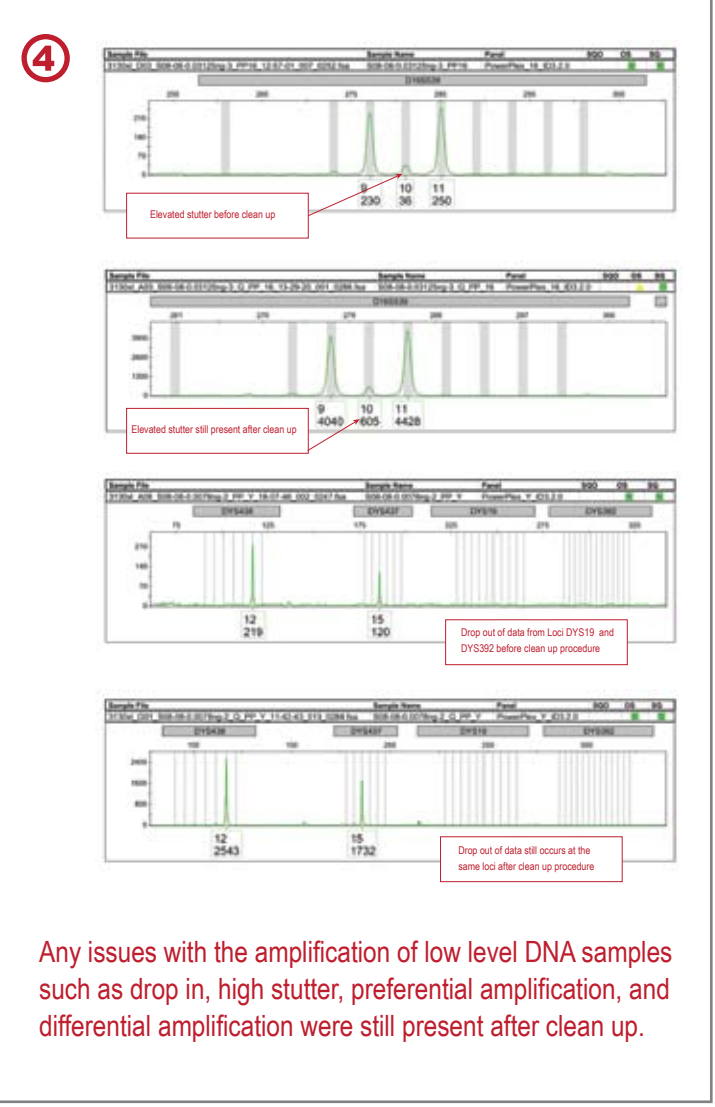


After Post PCR clean up there was a signal increase from an original peak height of 50 RFUs to a minimum of 300 RFUs in every kit tested.

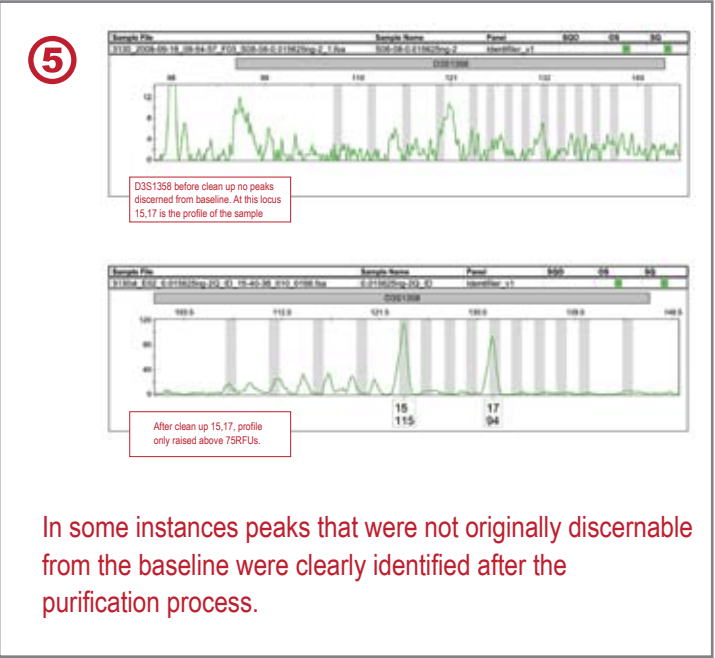
Minus A (-A) present prior to clean up increased after clean up in a greater proportion than that of the allele. If no -A was present initially, then there was no -A after clean up.



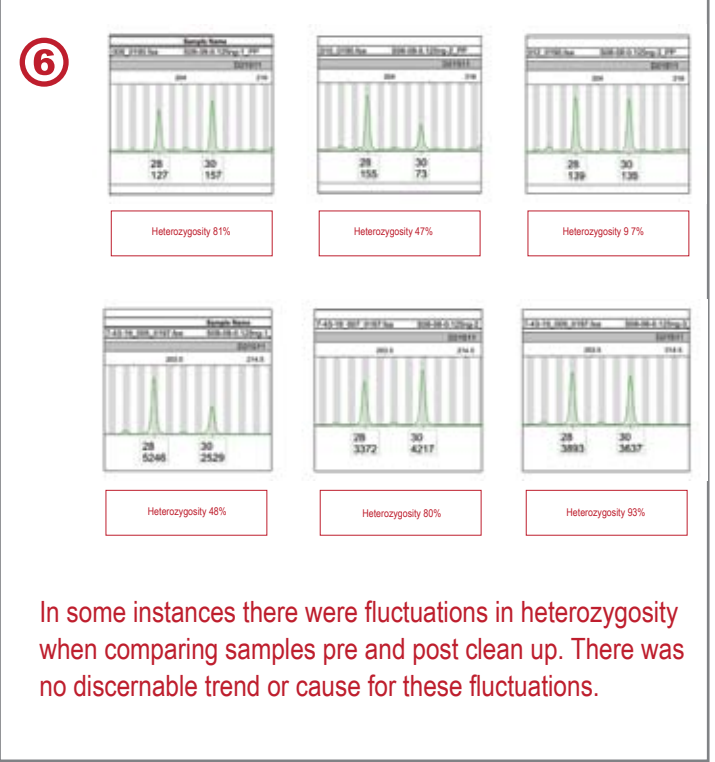
Dye blobs did not increase in proportion to true peaks. In some instances the dye blobs were reduced in size and were even removed after clean up.



Any issues with the amplification of low level DNA samples such as drop in, high stutter, preferential amplification, and differential amplification were still present after clean up.



In some instances peaks that were not originally discernible from the baseline were clearly identified after the purification process.



In some instances there were fluctuations in heterozygosity when comparing samples pre and post clean up. There was no discernable trend or cause for these fluctuations.

Conclusions:
This method of post PCR clean up consistently increased the signal of samples which were previously at levels too low to be characterized. Laboratories should perform appropriate validation studies to establish interpretation guidelines to account for the issues which occur with amplification of low level DNA samples.

Benefit to the Forensic Community:
Crime laboratories have seen an increase in the submission of requests for analysis on evidentiary items with low quantities of DNA. This study demonstrated that the QIAGEN MinElute® PCR Purification Kit consistently increased the fluorescent signal with the eight evaluated commercial STR amplification kits. This purification kit can be integrated into the laboratory process with little effort for method validation and at minimal cost. Integration of the QIAGEN MinElute® PCR Purification Kit into the DNA analysis procedure is a simple, cost effective method that can be easily implemented by a crime laboratory to increase the overall sensitivity of their DNA analysis methods.

References:

Smith, P.J, Ballantyne J. "Simplified Low-Copy-Number DNA Analysis by Post-PCR Purification." Journal of Forensic Science, 52(4). (2007):820-829

MinElute Handbook [Internet]. [updated 2008 Mar]. QIAGEN. [cited 2008 August 14]. Available from: <http://www1.QIAGEN.com/Products/DNACleanup/GelPcrSiCleanupSystems/MinElutePCR PurificationKit.aspx#Tabs=12>

The most current user and technical manuals for each kit were referenced in this study. Manuals procured from www.appliedbiosystems.com (published 1998-2006) and www.promega.com/tbs/ (published 2006.)

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