bionano

Ionic® Purification System for FFPE Samples



Why We Developed a New Approach to Nucleic Acid Purification

The commonly used bead- and column-based extraction technologies have followed the same fundamental workflow for over 20 years. This workflow uses ethanol, chaotropic salts, and other solutions to bind nucleic acid to a silica membrane or surface-labeled bead, which is then washed prior to the nucleic

acid being stripped off the solid support into an elution buffer. During this typically laborious process, the nucleic acid may be denatured, dehydrated, and fragmented. The eluate is also susceptible to contamination from wash buffers or beads.

Traditional Purification

Separation Principle

Column-based Silica Solubility Bead-based Carboxyl

Process and Characteristics

- Surface-based → loss and damage
- · Wash solution contamination
- Risk of bias (e.g., GC content)

Disadvantages

Incomplete binding to or removal from the solid support

Consequences

- Nucleic acid loss compromises data quality when sample input quantity is limited
- Recovery can be biased by fragment length or GC content
- For researchers, reduced biological insight
- For clinicians, less actionable information and false negatives

Contamination from wash buffers and bead coatings

• Low purity leading to false negatives and compromised data

Workflow with multiple hands-on steps

• Throughput bottleneck and potential errors; excessive use of disposable tips and labware

Isotachophoresis, a Superior Approach to Nucleic Acid Separation

Isotachophoresis (ITP) separates and concentrates charged molecules in solution solely based on their electrophoretic mobility. Biological samples are gently lysed and added to the Ionic® Fluidic Chip. An electric field is then applied to the chip and the nucleic acid is isolated in its natural, native form. The nucleic

acid is not denatured or dehydrated, and there's no binding to, or stripping from, fixed surfaces. The result is a higher yield of pure nucleic acid that is less fragmented and free from bead or wash buffer contamination.

Isotachophoresis

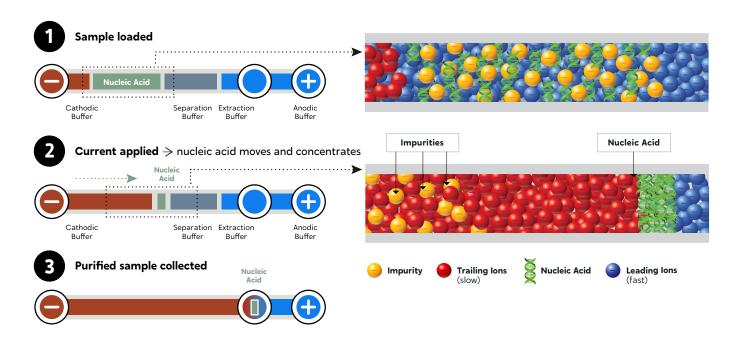
Separation Principle

Charge

Process and Characteristics

- Solution-based → high yield and integrity
- High purity
- Best representation of sample
- Result = better data

Simple, Automated Charge-based Sample Prep in Solution

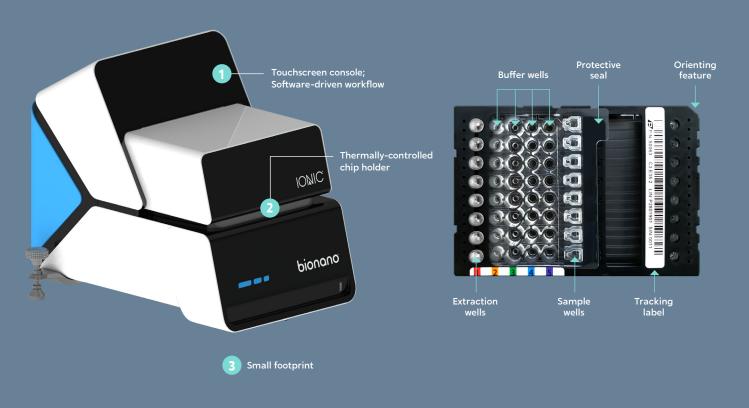


Ionic® Purification System

How it Works

The Ionic® Fluidic Chip is placed on the Ionic system and separation buffers are loaded. The chip is then primed. Next, biological samples are added into the 8 sample wells and purification using isotachophoresis begins. By applying an electric field across the length of a chip microchannel, the Ionic system separates and

concentrates nucleic acid between buffers with higher and lower mobilities. Impurities fall behind the low mobility buffer and are separated from target nucleic acids. As target nucleic acids pass through the channel, an integrated sensor stops the current once nucleic acids reach the extraction well.



The Next-Gen Sample Preparation System

The revolutionary Ionic Purification System requires no binding, stripping, or washing from fixed surfaces for higher yields, higher quality nucleic acids, and ultimately, better data for your research.

- · No organic solvents
- No harsh, high-salt buffers
- No system programming
- · No beads or repetitive washing
- No hands-on mixing, separation, sample transfers, or buffer exchanges
- No pumps, valves, or other moving parts

Rapid Purification of Precious Samples in Just ONE Hour

Ionic System Workflow



Load Run Buffers



Add buffers from reagent kit – use touchscreen to start priming

8 mins (manual)

2

Load Samples



Load 8 samples – use touchscreen to start the run

65 mins

3

Get Purified Nucleic Acid



Collect samples when touchscreen displays

"Run Complete"

2 mins (manual)





75 mins
Total Run-time

Simplified Nucleic Acid Preparation

The lonic system is so different, its advantages are most readily understood in contrast with conventional nucleic acid extraction and purification methods:



Higher nucleic acid yields

No sample loss associated with binding nucleic acids to, or stripping from, fixed surfaces



Simple workflows with fully automated separations

No columns or beads and no repetitive washing



Reduced nucleic acid fragmentation

No harsh high-salt buffers or organic solvents

Simplified FFPE Workflow Saves Time and Money

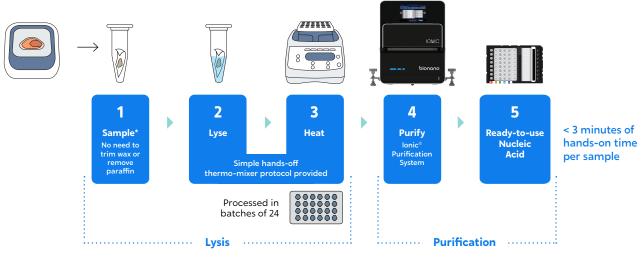
Purigen's FFPE protocols greatly simplify the processing of FFPE samples. For example, the Ionic® FFPE to Pure DNA protocol reduces the hands-on time to less than

3 minutes per sample and enables working directly from scrolls. The protocol also eliminates the need for a separate paraffin removal step.

Column-based Kit Workflow



Isotachophoresis Workflow



*Compatible with scrolls or slides

Flexibility for Working with Scrolls or Slides

The Ionic system produces more DNA and RNA from FFPE samples without requiring the use of slides (slide use and microdissection is optional). The ability to obtain comparable nucleic acid yields when using

scrolls (versus slide mounted FFPE slices) greatly simplifies the workflow when sample micro-dissection is not required. This allows projects to be completed faster and at a lower cost.

Superior Nucleic Acid Recoveries from FFPE Samples

A vast majority of clinical samples used in oncology research are stored as FFPE tissues, which often contain degraded or fragmented nucleic acid. Conventional extraction methods are labor intensive and can further damage nucleic acid during the extraction and purification process. The Ionic system simplifies and accelerates nucleic acid purification, resulting in higher yields of higher quality DNA.

Higher yields of DNA from FFPE Samples

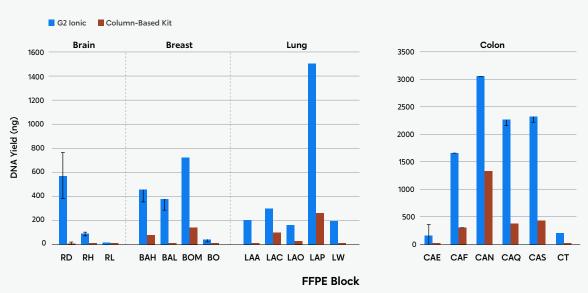


FIGURE 1: Comparison of DNA yields from consecutive sections of 18 FFPE tissue blocks purified on the lonic Purification System using the G2 FFPE DNA kit or a commercially available column-based kit. The total DNA yield for each sample was determined by multiplying the extract volume by the DNA concentration derived from the Qubit 1x dsDNA High Sensitivity assay. Error bars represent standard deviation from the average DNA yield. More DNA is recovered using the G2 FFPE DNA kit for each of the 18 FFPE blocks with expected increase in DNA yield of 3-5x on average compared to the column-based kit, with greater yield advantages for lower yielding samples.

Nucleic acid yields from scrolls using the Ionic Purification System are on average 3.5x higher when compared to yields from a column-based kit.

Higher Quality DNA vs. Column-based Kits

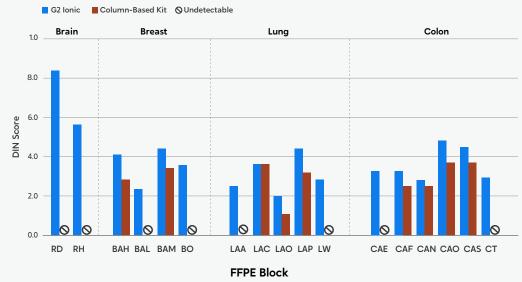


FIGURE 2: Comparison of DNA from consecutive sections of 18 FFPE tissue blocks purified on the lonic Purification System using the G2 FFPE DNA kit or a commercially available column-based kit. The DNA Intergity Number (DIN) was calculated for each extract using a Genomic DNA ScreenTape on the Agilent TapeStation. The lonic G2 DNA quality is typically higher than the column-based competitor, and nearly half of the column-based eluates had a concentration that was below that required for DIN score calculation.

Improved NGS Data Quality from FFPE Samples

In addition to increased yields and a greatly simplified workflow, data quality is also improved. Data below shows purified DNA from FFPE samples analyzed using the Agilent SureMASTR Tumor Hotspot sequencing panel which includes 252 amplicons ranging in size from 128–245 bps. To highlight coverage differences related to the sample purification technology, all data was normalized to a reference sample to

remove the effect of coverage differences introduced by differences in target amplicon amplification efficiencies. The reference data set was generated using the average results obtained from the purification of a high-quality sample using both lonic system and column-based techniques (as such the reference sample is not biased to either technology).

More Uniform Coverage vs. Column-based Kits

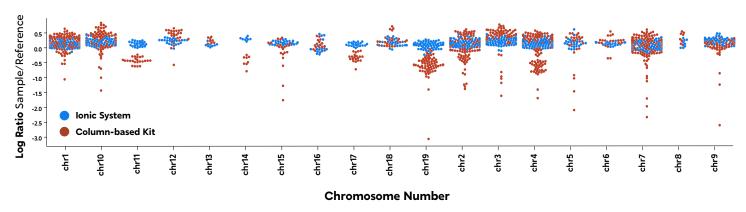


FIGURE 3: Results shown on a chromosome level. Ionic system eluates show tighter clustering more centered around the zero line. This is indicative of more uniform sequencing coverage.

Less Coverage Bias vs. Column-based Kits

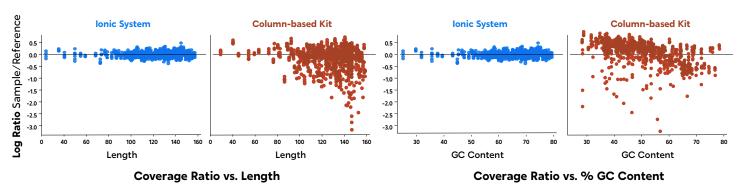


FIGURE 4: Results shown relative to amplicon length and amplicon GC content. Ionic system samples show superior uniformity for both amplicon length and GC content, with very little deviation from the expected coverage. Column-based purification shows a bias towards shorter amplicon lengths and lower GC content amplicons.

- · Ionic system purification shows no bias towards amplicon length.
- Ionic system purification shows no bias towards GC content.

A Better Solution for RNA from FFPE Samples

The Ionic Purification System provides for the automated purification of RNA from FFPE tissue samples with less hands-on time than conventional bead and column-based methods. To help scientists overcome the sample preparation bottleneck

commonly associated with FFPE samples, the Ionic system provides a simple workflow that co-purifies both mRNA and miRNA with higher yields versus column-based extraction kits.

Greater RNA yields from FFPE Samples

(Qubit yield by purification method - 10 µm sections)

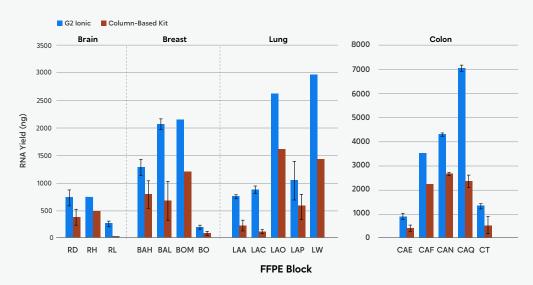


FIGURE 5: Comparison of RNA yields from consecutive sections of 17 FFPE tissue blocks purified on the lonic Purification System using the G2 FFPE RNA kit or a commercially available column-based kit. The total RNA yield for each sample was determined by multiplying the extract volume by the RNA concentration derived from the Qubit HS RNA assay. Error bars represent standard deviation from the average RNA yield. More RNA is recovered using the G2 FFPE RNA kit for each of the 17 FFPE blocks with expected increase in RNA yield of 2-3x on average compared to the column-based kit

RNA Quality Assessment from FFPE Blocks

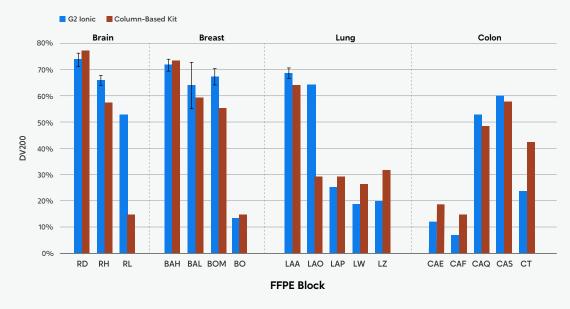


FIGURE 6: Comparison of RNA quality from consecutive sections of 17 FFPE tissue blocks purified on the lonic Purification System using the G2 FFPE RNA kit or a commercially available column-based kit. All RNA extracts were normalized to 1.5 ng/ul input concentration and the DV200 (percentage of RNA fragments >200 bp) was calculated using a High Sensitivity RNA ScreenTape on the Agilent Tapestation. For higher quality FFPE blocks (DV200 >30%) the G2 lonic RNA quality is consistent with or higher than the column-based kit. Since the lonic purification process co-extracts short RNAs ,such as miRNA, the G2 lonic DV200 values may appear lower than the column-based kit for especially poor quality FFPE blocks (DV200 <30%).

Get Higher miRNA Yields from FFPE Samples

For researchers studying either the relationship between gene expression and microRNA expression or focusing purely on microRNA expression in FFPE tissue samples, the lonic system provides more miRNA than the market-leading column-based miRNA kit. More impressive, is that no additional steps are required. The Ionic FFPE to Pure RNA kit produces both mRNA and more miRNA from FFPE samples in a single, simple workflow.

Total RNA Purification with Higher Yields of miRNA

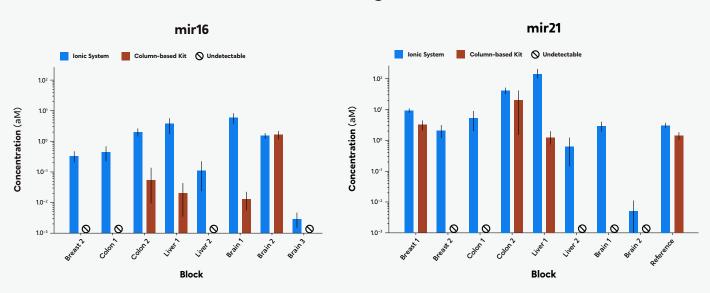


FIGURE 7: RNA from replicate sections of 8 FFPE sample blocks were purified using either the lonic FFPE to Pure RNA Kit or a column-based miRNA extraction kit. The extracted and purified samples from each kit were analyzed by qPCR and the Applied Biosystems TaqMan Advanced miRNA Assays for miR-16 and miR-21. The concentration of the target miRNA represented in each sample was extrapolated and plotted against the tissue type of the source FFPE sample block. The lonic system produced samples with a higher concentration of miRNA in all but one of the samples tested. For several samples the column-based miRNA extraction kit did not yield a detectable amount of miRNA.

Reproducible miRNA Expression Profiles

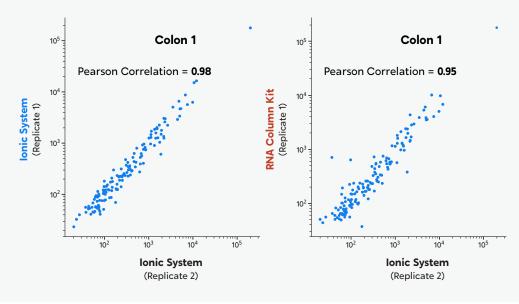


FIGURE 8: Samples from "Colon 1" of FIGURE 9 were analyzed for miRNA expression using the NanoString nCounter Human miRNA panel. The level of miRNA expression between replicate samples purified using the lonic system has a Pearson correlation of 0.98. The level of miRNA expression between replicate samples purified using the lonic system and the column-based miRNA kit has a Pearson correlation of 0.95. This analysis indicates a high reproducibility of miRNA expression across replicate samples purified using the lonic system that is comparable to that of the column-based miRNA extraction kit.

Simultaneous Extraction of RNA and DNA from FFPE

The Ionic® FFPE Complete Purification Kit is used with the Ionic system to enable the automated purification of DNA and RNA, including microRNA from FFPE tissue samples. The kit provides a protocol, Ionic® Fluidic Chips and reagents to enable the Ionic system to automate DNA and RNA purification using an innovative isotachophoresis technology. Samples are prepared for purification on the lonic system using a simple lysis procedure that can be automated using a programmable thermomixer without any need for micro-dissection or de-paraffinization using harsh chemicals.

Comparison of Total Hands-on Time vs. Manual Methods

	IONIC	Manual Bead-based	Manual Column-based
Lysis time	1.5 hrs	Overnight	1 hr
RNA isolation	2 hrs	2 hrs	2.5 hrs
Lysis time	1.7 hrs	3 hrs	3.5 hrs
Total time	5.2 hrs	13 hrs	6.5 hrs
Total hands-on time	1.5 hrs	6 hrs	7 hrs

TABLE 1: In a study conducted by a third-party genomic services lab, this table shows 3 extraction methods that were used to compare the hands-on time and total time to extract and purify RNA and DNA from 8 samples. Replicate 10 µm sections of FFPE samples were extracted and purified using either the lonic system, a market-leading manual bead-based kit, or a manual column-based kit.

Six adjacent sections of a 10 μ m thickness were harvested from 6 FFPE tissue blocks containing brain, breast, colon, or lung tissue. DNA and RNA were extracted and purified from 4 of the 6 sections using the published workflow for the lonic FFPE Complete Purification Kit. DNA and RNA were extracted and purified from the remaining sections using the published workflow for either a market-leading manual column-based kit or a market-leading manual bead-based kit.

The average estimated time to process 8 samples through the Ionic FFPE Complete Purification kit was 5 hours and 12 minutes with a hands-on time of 1 hour and 30 minutes (TABLE 1). This results in 11.25 minutes of hands-on time per sample to extract both DNA and RNA. The estimated time to process 8 samples through the column-based kit was 7 hours with most of that time being hands-on. This results in a hands-on time of 52.5 minutes per sample. Using a similar calculation, the hands-on time for the manual bead-based approach was 45 minutes per sample.

Improvement to DNA Yield with Comparable RNA Yield

The simplified workflow of the Ionic FFPE Complete Purification Kit provides simultaneous extraction and purification of FFPE samples without compromising yield.

RNA Yield by Purification Method

10 µm sections

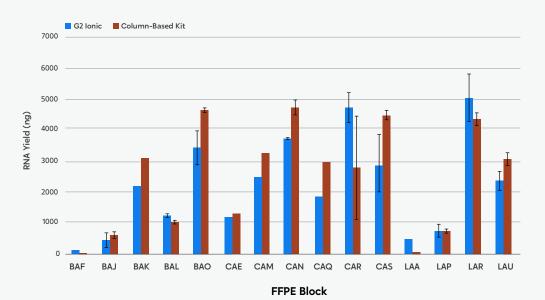


FIGURE 9: Comparison of RNA yields from consecutive sections of 15 FFPE tissue blocks purified on the Ionic Purification System using the G2 FFPE Complete kit or a commercially available column-based kit. The total RNA yield for each sample was determined by multiplying the extract volume by the RNA concentration derived from the Qubit H5 RNA assay. Error bars represent standard deviation from the average RNA yield. A similar or higher amount of RNA is recovered using the G2 FFPE Complete kit for 8 of the 15 FFPE blocks with an expected RNA yield increase of ~15% on average compared to the column-based kit

Improved DNA Yield with Optional Secondary Incubation

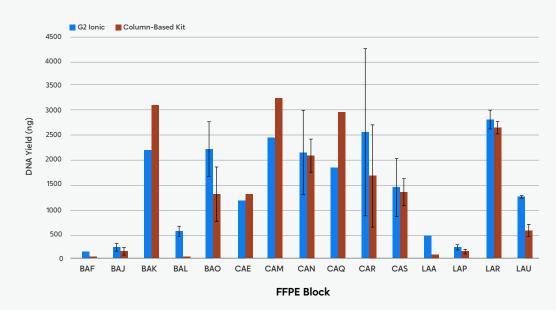


FIGURE 10: Comparison of DNA yields from consecutive sections of 15 FFPE tissue blocks purified on the lonic Purification System using the G2 FFPE Complete kit or a commercially available column-based kit. The total DNA yield for each sample was determined by multiplying the extract volume by the DNA concentration derived from the Qubit 1x dsDNA High Sensitivity assay. Error bars represent standard deviation from the average DNA yield. More DNA is recovered using the G2 Complete kit for 12 of the 15 FFPE blocks with increased DNA yield of 1.5-2x on average compared to the column-based kit and greater yield advantages for low yielding samples.

A Better Solution for DNA Purification from Tissue Samples

The Ionic[®] Tissue to Pure DNA Kit provides automated purification of DNA from fresh frozen tissue samples with less hands-on time than conventional bead and column-based methods.

Comparison of Ionic Tissue to Pure DNA Kit to 3 Column-based Methods

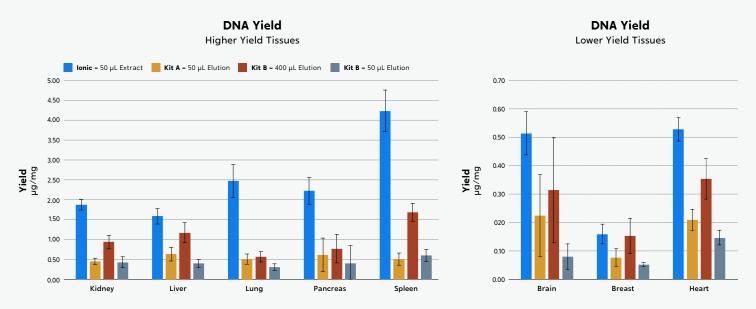


FIGURE 11: DNA extraction was performed on 8 different tissue types with four different extraction methods on a total of 632 specimens. Each tissue type/condition has an n-value ranging from 6 to 47 individual specimens. Pancreas, brain, breast, and heart specimens were derived from one donor and kidney, liver, lung, spleen were from two donors. Error Bars represent the 95% CI on the mean DNA yield (µg) per mg of tissue from replicate specimens (1-10 mg).

Linearity of Yield Across Tissue Input Amounts

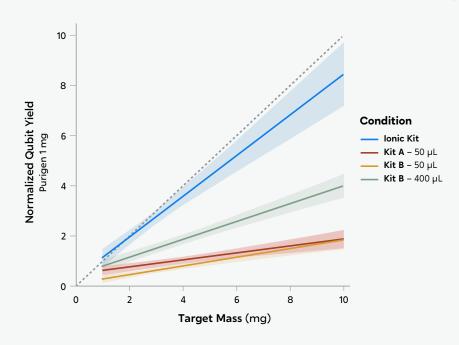


FIGURE 12: DNA yields were normalized to the lonic kit yield for 1 mg on a by-tissue basis. The dotted line represents a theoretical linear extraction efficiency up to 10 mg tissue (assuming optimal recovery by lonic kit at 1 mg specimens). The lonic kit tracked the theoretical linear extraction efficiency more closely than the column-based methods.

Maximize Yields For WBCs, PBMCs, and Cultured/Sorted Cells

The Ionic® Cells to Pure DNA Kit supports a wide range of cell types including white blood cells (WBC) or peripheral blood mononuclear cells (PBMC) isolated from blood as well as cultured or sorted

cells. The standard input range is 50k to 5 million cells. Customized protocols for as few as 10 cells are available upon request.

Higher Yields for Blood-based, Cultured, and Sorted Cells

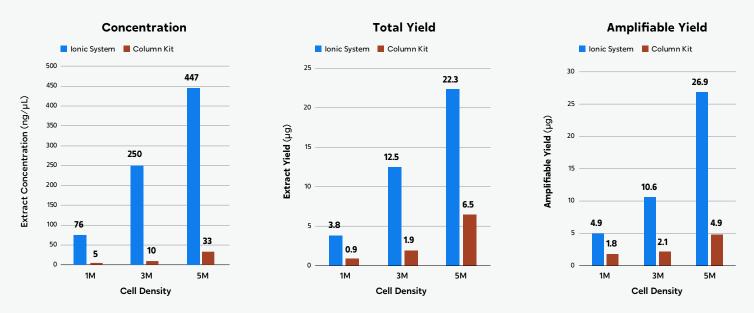


FIGURE 13: Peripheral Blood Mononuclear Cells (PBMCs) were isolated via Ficoll gradient, White Blood Cells (WBCs) were pelleted from lysed whole blood, and GM24385 cells were pelleted from culture media. Extractions were performed for each cell type at amounts ranging from 1-5 million cells then quantified via Qubit assay.

Consistent Length Profiles with Average Length Above 20k bp

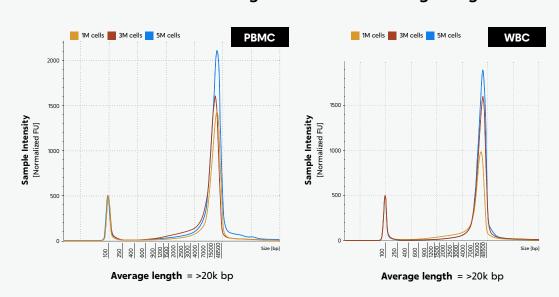
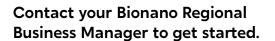


FIGURE 14: DNA Extracts from the lonic System were evaluated on a TapeStation to determine their size profile. Size profiles were consistent across cell type and quantity.

Products

Instrument			Configura	tion	Part No.
Some	Ionic® Purification Sy Includes: Power cord, in Warranty Information 12 months coverage Initial response withi On-site response with Includes parts and none parts, materials On-site labor	nstallation, warranty n 8 business hours hin 3 business days	Standard		44001
Kits			Configura	tion	Part No.
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lamona house and received and	Ionic® G2 FFPE to RNA Kit Includes: Fluidic chips, room temp reagents, -20°C reagents		6-Chip Kit	0000000	90167
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Service Contracts		Includes		Term	Part No.
1 planned maintenance visit Initial response within 8 business had on-site response within 3 business of Minimum 90-day warranty on replace Parts, materials On-site labor, per diem charges Phone and email support Remote support sessions Software upgrades		ys	12 months	44900	
Planned Maintenance-o		 1 planned maintenance visit Initial response within 12 business hou On-site response within 5 business da 90-day warranty on all replacement p 10% discount on Parts and Materials 10% discount on on-site labor, per die Phone and email support Remote support sessions Software upgrades 	ys parts	12 months	44901
Time and Materials Serv	vice	Hourly service work		N/A	44902





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