Abstract

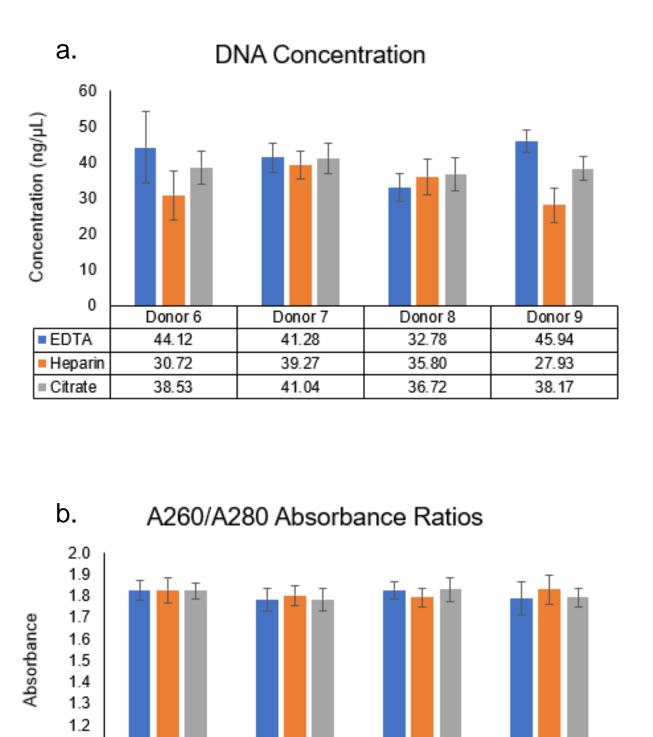
The isolation of high quality genomic DNA (gDNA) is the precursor to many molecular biology assays. The new GenFind V3 Blood and Serum DNA Isolation Kit from Beckman Coulter uses the patented SPRI (Solid Phase Reverse Immobilization) paramagnetic bead technology to isolate genomic DNA from fresh or frozen whole blood and serum containing Citrate, EDTA, or Heparin anticoagulants, as well as from cultured cells. GenFind V3 uses an improved cell lysis buffer and Proteinase K treatment to rupture cell membranes and digest proteins to give consistently high quality gDNA with improved yields and purities. The GenFind V3 kit supports up to 400 µL sample input volumes from blood or serum or two million cultured cells and can be performed in either a 96-well plate or tube-based format. Additionally, the need for large volume whole blood extractions continues to grow along with the ability to easily automate the process to alleviate user errors. To address this need, a protocol that supports >400 μ L and up to 2 mL of whole blood is available that can also be performed in an automated fashion using a 24-well plate. Volumes greater than 2 mL can be performed in a tube-based format. Currently, we are also in the process of integrating a Thermo Scientific KingFisher[™] Presto on the i-Series to allow for higher volume processing and faster throughput.

Here, we demonstrate a walk-away automated solution for GenFind V3 kit to purify up to 400 µL of whole blood using the Beckman Coulter Biomek i7 Hybrid Genomics Workstation. The Biomek i-Series method is a high yielding and robust nucleic acid purification process and can process up to 96 samples in a 96-well format in less than 3.5 hours with minimal user interaction and no off-deck centrifugation or vacuum filtration. The method is also compatible with the Biomek i-Series NGS workstation, facilitating the installation process for existing Biomek i-Series users.

Materials and Methods

Whole blood was collected from four individual donors in blood collection tubes coated with EDTA, Citrate, and Heparin anticoagulants. Four technical replicates for each donor were used and a total of 48 samples of 400 µL were aliquoted in a 2.2 mL Abgene deep well plate (16 EDTA, 16 Citrate, 16 Heparin). Genomic DNA was then extracted using the GenFind V3 Blood and Serum DNA Isolation automated method implemented on the Biomek i7 Hybrid platform. All extracted genomic DNA samples were analyzed using a NanoDrop One (Thermo Scientific). 15 samples (five EDTA, five Citrate, and five Heparin) were randomly selected and analyzed using 2200 TapeStation using the Genomic DNA ScreenTape kit (Agilent Technologies). Those same 15 genomic DNA samples were then converted to NGS sequencing libraries using the Kapa HyperPlus Kit implemented on the Biomek i-Series i5 NGS Workstation. NGS library QC was performed using the 2100 Bioanalyzer with the High Sensitivity DNA kit (Agilent Technologies).

Results: Automated gDNA Extraction



Donor 7

1.78

1.80

1.78

Donor 6

1.83

1.83

1.83

EDTA

Heparin

■ Citrate

Whole blood samples were used for genomic DNA extraction using the Biomek i7 Dual Hybrid Genomic Genomic DNA was Workstation. using NanoDrop One quantified and size distribution (Figure 1) DNA Integrity (Figure 2) and Numbers (DIN) was calculated with Agilent TapeStation using the Genomic ScreenTape (Table 1).

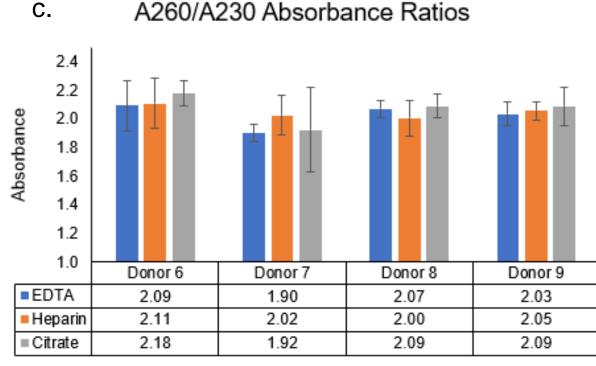


Figure 1: gDNA Concentration and purities determined using NanoDrop

Donor 8

1.83

1.79

1.83

Donor 9

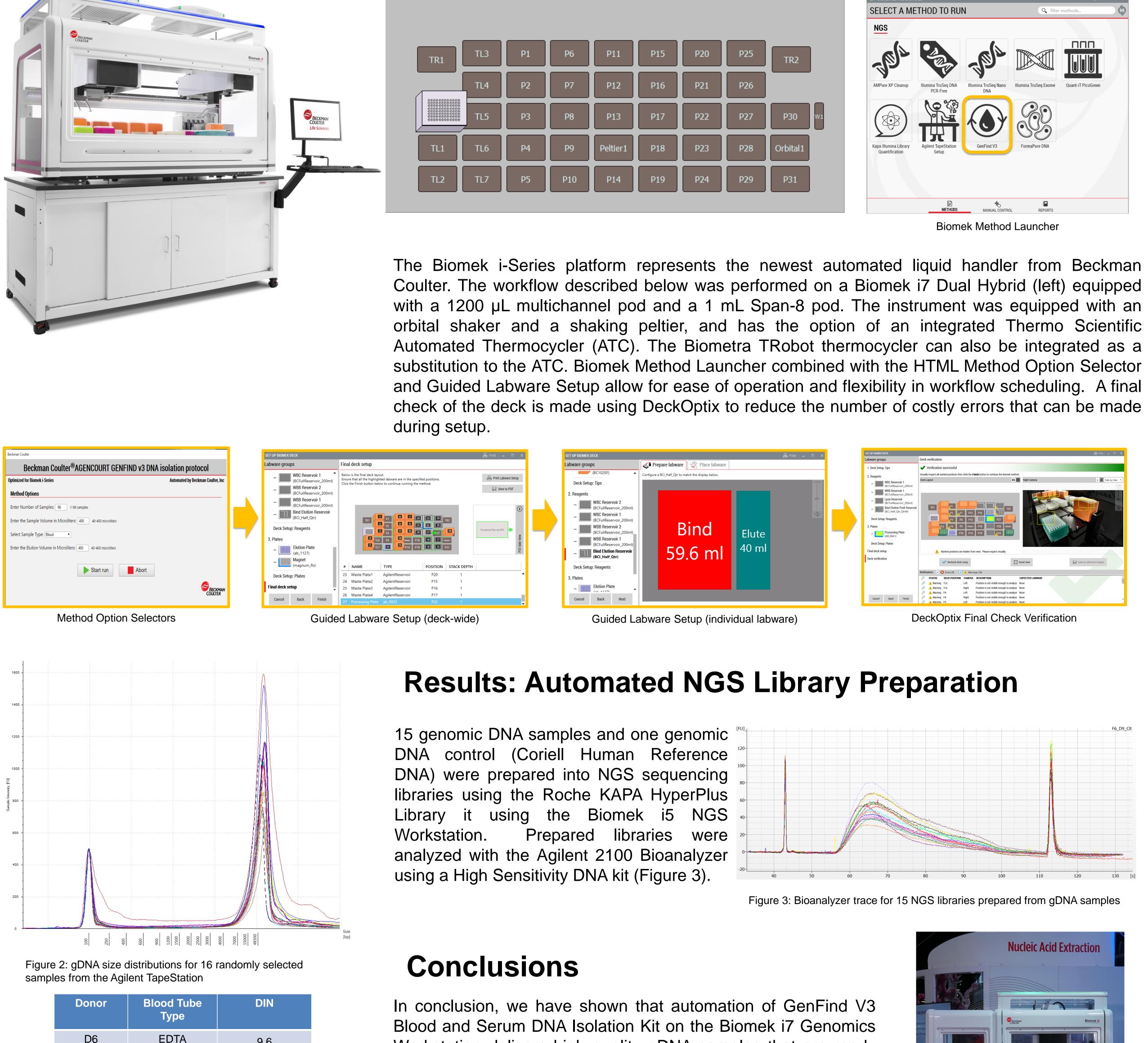
1.79

1.83

1.79

Automating Genomic DNA Extraction from Whole Blood and Serum with GenFind V3 on the Biomek i7 Hybrid Genomic Workstation Randy Pares, M.S., Li Liu, Ph.D., Han Wei, Ph.D., Mary Blair

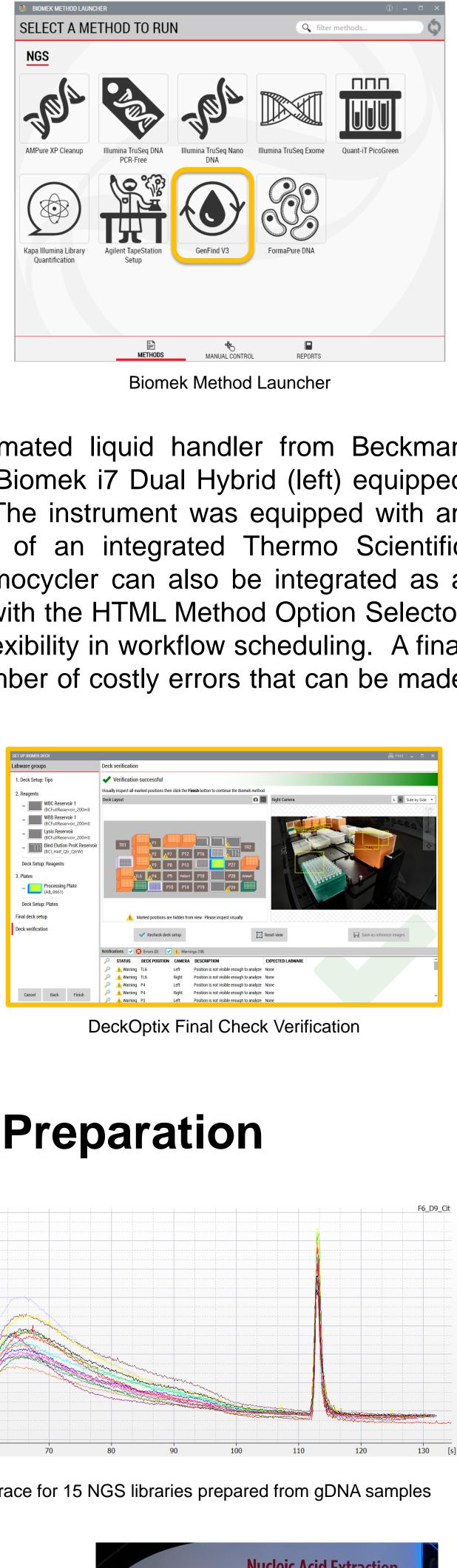
Biomek i-Series Automation

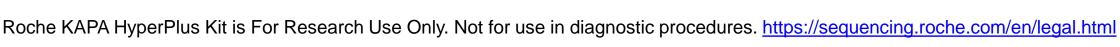


Donor	Blood Tube Type	DIN
D6	EDTA	9.6
D6	Heparin	9.5
D6	Citrate	9.6
D7	EDTA	9.2
D7	Heparin	9.4
D7	Heparin	9.6
D7	Citrate	9.5
D8	EDTA	7.8
D8	EDTA	9.1
D8	Heparin	9.6
D8	Citrate	9.5
D8	Citrate	9.2
D9	EDTA	9.5
D9	Heparin	9.6
D9	Citrate	9.2
Table 1: DIN for 16 randomly selected samples from		

Table 1. Din for to randomly selected samples from the Agilent TapeStation

Workstation delivers high quality gDNA samples that are ready for sequence ready libraries and downstream NGS applications. Future directions for automation of GenFind V3 include standardized integration of the Thermo Scientific KingFisher™ Presto on i-Series for higher sample input.





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