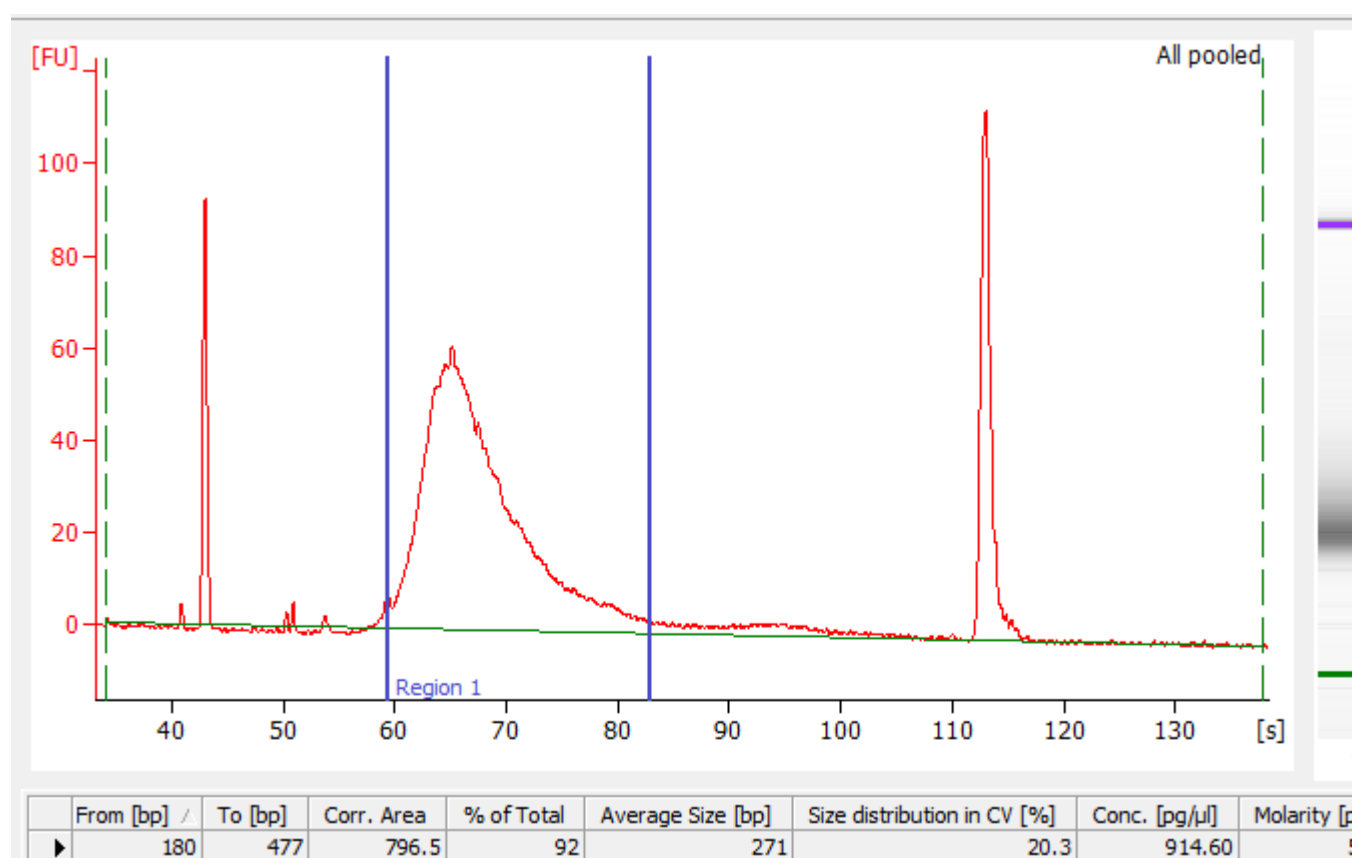


Subject: Next-Seq Submission
Date: Tuesday, September 21, 2021 at 8:57:53 AM Eastern Daylight Time
From: Caffrey, Paul
To: seqlab
Attachments: image001.png

Hi Sequencing Core!

I have an Illumina library to run. The sample is about 5.3 nM in 21 uL. Attached is the Bioanalyzer trace. Below is the information I believe you need. Please let me know if we need to provide any more information. I can drop it off later today if that works for you but there's no rush on it or anything.



Sender name: Paul Caffrey
 Lab Group: Gardner Lab
 Extension number: 6591
 Email: pcaffrey@neb.com
 Date Submitted: 092121
 Requested Date of Completion: whenever
 Project Name: Fosmid screening RNA-seq
 Sample Name: PJC_fosmid_library_screen
 Requested read lengths: Mid Output 75 Paired-End

Is your sample Low Diversity? Y, spike in 5% phi-X

Is your sample barcoded? Y

Please indicate which indexes were used: 3-6, 8-10

Thanks so much!

Paul