

NanoDrop Spectrophotometer Checklist

Revised May 2008

Introduction

This document presents the guidelines for using the Nanodrop ND-1000 spectrophotometer at the UHN Microarray Centre (UHNMAC).

The Nanodrop ND-1000 spectrophotometer is able to perform full spectrum UV-Vis absorbance analyses (220-750nm) and enables researchers to quantify DNA, RNA, dyes, proteins, and other samples. Only 1 μ L is required by the Nanodrop to quantify nucleic acid samples, however, we ask that 2 μ L be sent. 2 μ L of sample is required by the instument for proteins (please send 4 μ L). Since the Nanodrop ND-1000 has a large dynamic range (2-3700 ng/ μ L for dsDNA), dilutions are not necessary for most samples. The accuracy of each measurement is typically within 2%. Using the microarray application, the concentration of nucleic acid and up to 2 dyes can be calculated and displayed at the same time.

Unlike traditional spectrophotometers, the Nanodrop does not require cuvettes or capillaries. Instead, the sample is pipetted directly onto the measurement pedestal. When the apparatus is closed, the sample arm compresses the droplet and a sample column is drawn (to a controlled path length of 1mm) and surface tension holds the sample in place. After the 10 second measurement, the sample is simply wiped off and the instrument is ready for the next sample. The sample cannot be collected following its measurement.

Another advantage of the Nanodrop ND-1000 is that there are few solvent restrictions. Although, one solvent that is restricted is hydrofluoric acid. Even samples dissolved in high vapour pressure solvents such as acetone can be measured, but only using the short path length to reduce evaporation.

The Nanodrop ND-1000 spectrophotometer service is available to all customers for a fee. Samples can received Monday-Friday, 9am-4pm, excluding holidays. Please contact geneservice@microarrays.ca_____ or call 416-581-7463. The cost of Nanodrop is \$5 per sample.

The UHNMAC is located at: MaRS Centre, Toronto Medical Discovery Tower 101 College St, Room 9-601 Toronto, ON, Canada						
M5G 1L7						
Please complete the following:						
Your name:						
P.I.:						
Billing Address:						
Phone Number:						
e-mail address:						
Preferred method of contact: Phone \Box	e-mail □					
Number of samples (\$5 per sample):						

<u>Checklist</u>

 \Box We request that you submit/bring enough sample to allow for 2 measurements, if possible. (For nucleic acids, 1µL of sample is required for each measurement but we request 2µL; and for proteins, 2µL of sample is required for each measurement, but we request 4µL.)

 \square Please bring a tube of your buffer/solvent *in case* your sample needs to be diluted. The concentration of the samples should be between 2 and 3700 ng/µL (for dsDNA).

□ Please complete the following table (add another page if necessary).

Sample ID	sample	of interest (ie. 260, 280, 550,	Approximate (estimated) concentration	Purification/isolation method	B u f f e r / solvent