| STANDARD OPERATING PROCEDURE | | | | | | |
|---|--|-----------------------|--------|-------------|--|--|
| A Ragon Institute of MGH, MIT and Harvard | DNA Measurement using NanoDrop ND-1000 | | SOP | PL-SOP-006 | | |
| | instrument | | # | | | |
| | Originated by: | Ildiko Toth | Date: | 27 April 17 | | |
| | Laboratory: | Processing Laboratory | Pages: | 1 of 2 | | |
| | Approved by: | Alicja Trocha | | | | |
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I. PURPOSE:

The purpose of this procedure is DNA quantification using NanoDrop ND-1000 instrument.

II. SAFETY:

This protocol needs to be carried out in a BSL1 Lab. Personnel are required to wear gloves and a lab coat before initiating any work.

III. PROCEDURE:

Wear Gloves.

- 1. Open program on computer.
- 2. Initialize spectrophotometer:
 - a. Load 2ul distilled water on measurement pedestal.
 - b. Lower sampling arm gently.
 - c. Click "OK".

3. Measure Blank:

- a. Load 2ul blank buffer (the buffer, solvent, or carrier used in your sample).
- b. Lower sampling arm gently.
- c. Type BLANK into the Sample ID window.
- d. Click "BLANK" (F3).

4. Ensure Quality of Blank:

- a. Load 2ul of same blank buffer.
- b. Lower sampling arm.
- c. Click "MEASURE" (F1).
- d. Check that the value in the concentration window is within -0.9 and +0.9.
- e. If the value exceeds these limits, repeat the blanking process. If not, proceed to measure your DNA.
- f. Repeat blanking process after every 10 samples.
- 5. Briefly vortex or mix DNA samples before loading on the NanoDrop.
- 6. Measure DNA concentration and purity:
 - a. Type in the Sample ID.
 - b. Load 2ul of DNA sample.
 - c. Lower sampling arm.
 - d. Click "MEASURE" (F1).
 - 260/280 window: a value of ~1.8 indicates pure DNA. Lower values indicate contaminants such as protein.
 - 260/230 window: a value of 1.8-2.2 indicates pure DNA. Lower values indicate contaminants

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- ng/ul window: displays DNA concentration.
- 7. Between each measurement, clean upper and lower measurement pedestals with de-ionized (miliQ) water on a kim-wipe.
- 8. When you are finished measuring your samples clean the pedestals again and leave a dry, folded kim-wipe between the sampling arm and the lower measurement pedestal.
- 9. Your measurements are automatically saved and can be found in: My Computer → Local Disk C → NanoDrop Data → Default → Nucleic Acids
 - a. Look for the date on which the measurements were taken.
- 10. Remember to clean the area after your work is done

IV. REFERENCES/ ADDITIONS/ NOTES:

N/A