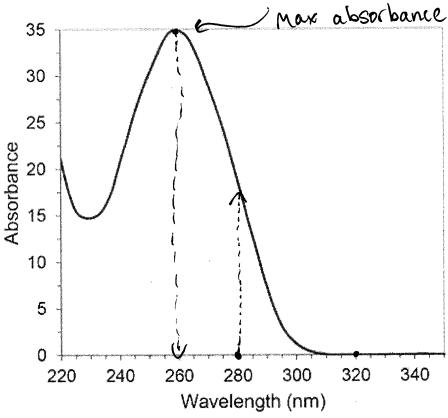
Answer Key

CHEM1501 Midterm Practice Problems

Written by Theo Nelson

Question 1: The following is a spectrum graph for purified nucleic acids, e.g. DNA and RNA.



a.) Please identify the λ_{max} for DNA/RNA:

4 mas = 260 nm

b.) For these kinds of purified nucleic acids the absorbance at 280 nm often represents the amount of protein content, while the absorbance at 320 nm can represent general background noise in the solution. Calculate the approximate ratio between protein content and nucleic acid content:

Itein content, while the absorbance localized the approximate ratio between protein content and nucleon 320 nM = 0 Abs. $\frac{260}{280} = \frac{35}{7.7} \approx 2.0 \text{ to } 2.2$

You May also Calculate the 280/200 (atio for this of Assume for the remaining questions that the calculated ratio indicates an adequate nucleic acid purification

Question 2: There is a mysterious tube in the lab. You measure the absorbance on a spectrophotometer, with the machine providing the "raw" values in the table below.

sample	260 Raw	280 Raw	320 Raw	260	280	260/280	ng/µl
blank	.072	.069	.060	n/a	n/a	n/a	n/a
tube	.084	.075	.061	?	?	?	?

Recall that the 320 nm signal is general background noise - which allows you to ensure that samples have the same amount of background signal. The amount of nucleic acid/protein material should be calculated in relation to the amount absorbed by a tube with the liquid which the nucleic acid is dissolved in (blank).

a.) Calculate the amount of the 260 and 280 nm signal attributed to the nucleic acids in the tube?

b.) Is this an "adequate nucleic acid purification," when comparing the 260/280 ratio to the optimal ratio as calculated in 1-b?

$$\frac{260}{280} = \frac{.012}{.006} \approx 20 \text{ Nucleic and puission.}$$

Question 3: The Beer-Lambert Law is a relationship between absorbance and concentration: $A = \epsilon lc$. If A = absorbance of nucleic acids, $\epsilon_{DNA} = 0.020 \, (\mu g/mL)^{-1} cm^{-1}$ and I = path length. You have been provided a diagram of the equipment you utilized to make these measurements.

Question 4: After completing your experiment on the spectrophotometer, you have 25 μ l left in the mystery tube. The label states: "LINC00173 // DNA // 3093 bases." You assume that this means the tube contains identical particles of DNA 3092 bases long called LINC00173.

Each base is on average 650 g/mol. You wish to phosphorylate the 5' ends of the DNA strands; each DNA strand has one 5' end. You will design a reaction totaling 50 μl to accomplish this task. You have been provided the following instructions:

• 10 units (or 1 µl) of T4 PNK-enzyme can phosphorylate 300 pmol of 5' termini

a.) Calculate all the necessary volumes to perform a successful phosphorylation reaction:

- You should have a final concentration of 1X T4 PNK Buffer
- You should have a final concentration of 1 mM ATP
- Nuclease-free water is not involved in the reaction; it is simply to ensure that the final reaction volume is 50 μl.

You have access to the reagents described in the following table:

T4 PNK	201pl
10X T4 PNK Buffer	? 5 pl
10 mM ATP	? 5 pl
DNA (20 ma)	? 25 pl ? 14-15 pl
Nuclease-free Water	? 14-15pl

10 x (x ml) = 1x (50 ml) Amount of 10x TY HOK X = 5 ml Buffer required 10 nM (x ml) = 1 nM (50 ml) Amont of 10 nM ATP x=5 ml required DNA = 25 ml => given in the mystem the How much enzyme. 25 ml DNA. (2 ng) = 300 ng DNA 30972 bases DNA. (650a) = 2009800 g 109 ng) (109 ng)

= 2.0098×105/10073

300 ng DNA. (1 mol LINCO0173) (6.022 ×10²³ nolealor LINCO0173) =8.989×1010 moleuler LINCOOFT3 (8,989×10° nolander UNXXXII) (15' tennini) 1940 1844 198 Mills = 8.981×1006 f fermin. 8.969×10°5' tenimi. (1 nol 5' tennini) = 1.49×10⁻¹³ nol 5' tennini 6.022×10²³ tennini 1.49×10⁻¹³ nol 5' temini (1012 pmol) = . 149 pmol 5' temin (149 pmol 5' Jenni) (1 mlty PNK) = . 000498 pl Ty PNK Note: you need very little enough to perform this reaction. All annen up to I pl anepted if you addrew the fearibility of pipetting out. 000498 pl. Nucleare-free Water: 50 pl-25 pl-5 pl-60-1 pl) = 14-15 pl