

Nucleotide preparations (100mM)

By following the procedure below you should end up with a large stock of free nucleotides that can be used for sequencing, PCR, random hexamer/oligo nucleotide labeling, etc. And it is real cheap! For additional info see also Sambrook Appendix B10.

Resuspend the entire vial into the indicated volume of sterile ddH₂O (or 1X T₁₀E_{0.1}).

<u>Nucleotide</u>	<u>Quantity in Sigma vial</u>	<u>Volume ddH₂O</u>
dATP (Sigma D 6500)	25mg	400μl
dGTP (Sigma D 4010)	10mg	160μl
dCTP (Sigma D 4635)	25mg	400μl
dTTP (Sigma T 0251)	25mg	400μl

Bring the pH of each nucleotide to ~7. To do this Sambrook recommends adding 1M Tris. I have found that adding 10μl of 1M Tris is insufficient to raise the pH to 7. Therefore I add 10μl of 1M Tris and then 1μl aliquots of 10N NaOH until the pH reaches about 7. It takes 2 to 4μl of 10N NaOH to do this. To determine the pH, simply spot 10μl aliquots onto pH paper.

Determine the exact concentration of each nucleotide by first diluting 1.0μl of the dNTP into 2000μl TE (0.5μl into 1000μl) and measuring the OD at the appropriate wavelength (λ). You then need to divide λ by the extinction coefficient (ϵ). For a cuvette with a 1cm path length (which we assume), absorbance (A) at the $\lambda = \epsilon M$ where M is the molarity.

<u>Nucleotide</u>	<u>λ</u>	<u>Extinction coefficient (ϵ)</u> (M ⁻¹ · cm ⁻¹)
dATP	259	1.54 X 10 ⁴
dGTP	253	1.37 X 10 ⁴
dCTP	271	9.10 X 10 ³
dTTP	260	7.40 X 10 ³

For example: Lets say that you diluted 0.5μl of dATP into 1000μl. The OD₂₆₀ is then determined to be 0.6922.

$$\begin{aligned}M &= \lambda/\epsilon \\M &= (0.6922 \times 2000)/(1.54 \times 10^4) \\M &= 0.090 = \mathbf{90mM}\end{aligned}$$

Store at -80°C. Repeated freezing and thawing is not good for nucleotides so Sambrook recommends aliquoting the dNTPs into small volumes.

Materials Needed:

Nucleotides

1M Tris

10N NaOH

pH paper

Spectrophotometer