

The samples :

- The analyzed macromolecules must be soluble and stable (especially for sedimentation equilibrium experiments)

- The concentration depends on what you want to see and of what you have of available:

In Absorbance (sedimentation velocity), the comfort is $DO=1$ (we can modulate: optical path = 1.2, 0.3 or 0.15 cm, λ between 230 and 600 nm).

Using interference optics, there is no limit considering the large concentrations, but a number of reasonable fringes is needed to have informative data (the series 1.2/0.8/0.3 mg/ml for proteins is generally suitable using absorbance and interference). The minimum concentration depends on the homogeneity of the sample.

Usually, we make two dilutions - 2/3 and 1/3 - from a sample of $DO=1.2$. When we expect (or we observe) effects of the concentration, we are often brought to use various systems of detection and/or optical routes.

For polysaccharides: in our study on heparin fragments, we tested typically our samples at 3, 6 and 10 mg / ml (sometimes 1.5, 3 and 5 mg / ml).

- The usual volume in sedimentation velocity experiments is 450 μ l for cells with an optical path of 12mm (it can be 150 or 80 μ l for cells of optical path of 3 or 1.5 mm). If you have no constraints, prepare 1 mL (we cannot exclude a leak, thus a sample to be redone).

For sedimentation equilibrium experiments, the volumes are lower, typically 180 μ L in 12 mm, as the concentrations (typically $DO=0.6$), but we often duplicate samples and we make series of dilution.

The solvent :

- It is necessary to know its exact composition, especially for sedimentation equilibrium experiments (dialysis is wished)
- It has to contain some salt (for example 100 mM), so that the transport is not perturbed by electrostatic interactions (the macromolecule migrating faster than ions).
- For acquisitions using absorbance: the solvent does not have to absorb (or little) at the considered wavelength.
- For acquisitions using interference: the composition of the solvent must be strictly controlled and identical to that of the reference solvent (except for membrane proteins: the reference solvent does not have to contain detergent)
- The volume of solvent: the comfort is 1ml reference solvent by sample. If the volume of solvent does not raise problem, plan 5-10 ml, for possible measurements of density and viscosity.
- In the case of membrane proteins, it is necessary to have 2 solvents: a solvent with the detergent (for possible dilutions and measurements of density and viscosity) and a second one without detergent (for the reference).