Sedimentation equilibrium



Thermodynamic information

- Mass
- Stoichiometry
- Equilibrium constants
- Nonideality



Preliminaries

What do you want to know?

Sample handling

Sample type

Optical system



What do you want to know?

Monomer molecular weight

Stoichiometry of a complexAssociation constants

Nonideality



Preliminaries

What do you want to know?

Sample handling

Sample type

Optical system



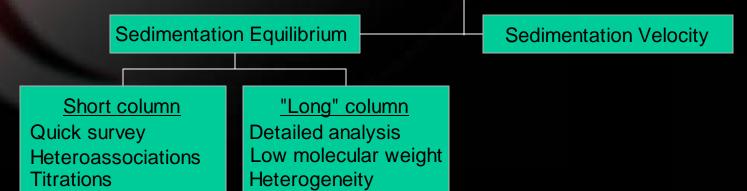
Sample Handling

Sample Arrives

If sample has not been gel filtered during purification, do so before analysis

General Sample Handling

Estimate concentration and volume Bring sample to dialysis equilibrium with buffer Choose centerpiece material





Preliminaries

What do you want to know?

Sample handling

Sample type

Optical system



		Sample type	2
	<u>Protein</u>	Polysaccharide	Nucleic Acid
	Choice of optics	Interference optics	Absorbance optics
	1 A _{230 or 280}	C > 1 mg/ml	1 A ₂₆₀
	1 mg/ml	Nonideality	Nonideality



Preliminaries

What do you want to know?

Sample handling

Sample type

Optical system



Choosing optical system

Use absorbance if:

- Need selectivity
- Added sensitivity
- Cannot dialyze sample

Use both:

- Determine extinction coefficient
- Test for sample purity
- Extend concentration range

Use interference if:

- Buffer absorbs
- Sample does not absorb
- Precision required
- g(s)
- Extinction coefficient varies
- Short columns



What do you want to know?

Monomer molecular weight

Stoichiometry of a complex

Association constant

Nonideality



Monomer molecular weight

Use denaturing conditions 6 M guanidinium chloride 8 M urea Determine partial specific volume Measurement Calculation Density Viscosity (for sedimentation velocity)



What do you want to know?

Monomer molecular weight
Stoichiometry of a complex

- Association constant
- Nonideality
- Sedimentation equilibrium
 - Shape of a complex
 - K_s the effect of C on s
 - Molecular weight (discrete mixtures)
 - Association constant (favorable conditions)
- Sedimentation velocity



Stoichiometry of a complex

- Examine each component individually Isolate components for heterooligomer M_d in denaturing buffer M_n in nondenaturing buffer Isolated components (if stable alone) Holocomplex Stoichiometry • Homooligomer $\overline{N} = M_n / M_d$
 - Heterooligomer N can be harder to determine
 - Uncertainty rises as N increases



What do you want to know?

Monomer molecular weight Stoichiometry of a complex

- Association constant
- Nonideality
- Sedimentation equilibrium
 - Shape of a complex
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Sedimentation velocity

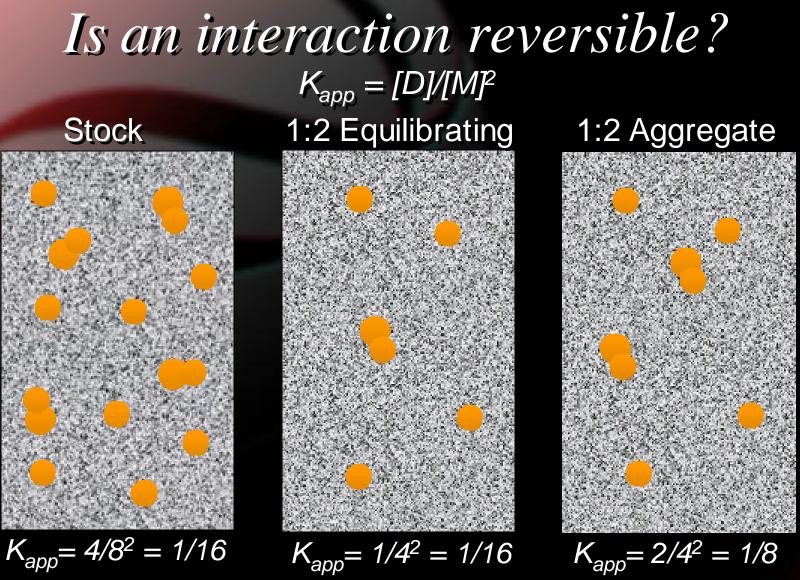


Association Constant

Depends on type of association Self association

- Must have correct assembly model
- Must cover correct concentration
- Hetero association
 - Test using different ratios of subunits
 - Model as self association for 1:1 assemblies
 - **D**s < 3.5







Nonideality

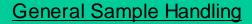
Always present Two major sources Excluded volume Constant and small for spheres Worse for rods and coils Charge-charge repulsion • Dominates for proteins of any shape at low G/2• Minimize by adding salt G/2 > 0.1 or more Minimize by extrapolation to C=0



Two types of equilibrium expts

Sample Arrives

If sample has not been gel filtered during purification, do so before analysis



Estimate concentration and volume Bring sample to dialysis equilibrium with buffer Choose centerpiece material

Sedimentation Equilibrium

Sedimentation Velocity

Short column Quick survey Heteroassociations Titrations <u>"Long" column</u> Detailed analysis Low molecular weight Heterogeneity



Short Column Equilibrium

Solution Column < 1 mm

Select Conditions Rotor speed Temperature Use interference optics

General requirements

>30 uL sample at >1 mg/ml Dialyzed against solvent Use 15 uL in channel A Make 3 serial 1:2 dilutions for channels B->D Run duplicate sample treated with 6 M GuHCI Cannot use centrifugal gel filtration w/ 6 M GuHCI



Two types of equilibrium expts

Sample Arrives

If sample has not been gel filtered during purification, do so before analysis

General Sample Handling

Estimate concentration and volume Bring sample to dialysis equilibrium with buffer Choose centerpiece material

Sedimentation Equilibrium

Short column Quick survey Heteroassociations Titrations <u>"Long" column</u> Detailed analysis Low molecular weight Heterogeneity Sedimentation Velocity



"Long" Column Equilibrium

Solution Column > 2 mm

Select Conditions:

Rotor speed Temperature Optical system

General requirements

250 uL @ 1 unit (e.g. 1 OD or 1 mg/ml) Use 110 uL in channel A Make 2 1:3 serial dilutions for B and C



Notes on rotor speeds

Use 4 rotor speeds
Go from lowest to highest
Lowest should have monomer s ~ 2

starting rpm ~ 5x10⁶ (1/M)^{1/2}
start at 3000 rpm if working with complete unknown

Space rotor speeds to cover 2 to 3 fold range

• *s* ~ 2 - 15



General considerations

Correcting for buoyancy

Determining density

Partial specific volume

Correcting for viscosity



Useful references

Books:

Analytical Ultracentrifugation in Biochemistry and Polymer Science. (1992) S.E. Harding, A.J. Rowe, and J.C. Horton, eds. Royal Society of Chemistry, Cambridge.

Modern Analytical Ultracentrifugation. (1995) T.M. Schuster and T.M. Laue, eds. Birkhauser, Boston. *Two fairly recent books devoted entirely to this field*

K.E. van Holde, <u>Physical Biochemistry</u>. (1985) Prentice Hall, Englewood Cliffs, New Jersey. Good introductory text for general theory of sedimentation, frictional coefficients, diffusion, and other hydrodynamic analysis

Freifelder, D. (1982). Physical Biochemistry: Applications to biochemistry and molecular biology. W.H. Freeman, New York.

Regarded as a good introductory text that is strong on centrifugation methods

van Holde, K.E., W.C. Johnson, Jr., and P.S. Ho. .(1998). Principles of physical biochemistry. Prentice-Hall, Upper Saddle River.

Cantor, C.R. and Schimmel, P.R. (1980). Biophysical chemistry. Part II: Techniques for the study of biological structure and function. W.H. Freeman, San Francisco. *These two are more advanced texts with good coverage of centrifugation methods*

Special Journal Issue:

Chemtracts Biochemistry and Molecular Biology, vol. 11 no. 13 (pp. 933-1004), December 1998 (Jeffrey C. Hansen, Guest Editor) *Several review articles and condensation commentaries on current research*



Useful references

Review Articles:

Stafford, W.F. III. (1997). Sedimentation velocity spins a new weave for an old fabric. *Curr. Opin. Biotechnol.* 8, 14-24.

Laue, T.M. (1995). Sedimentation equilibrium as thermodynamic tool. Methods Enzymol. 259, 427-452.

Laue, T.M. Stafford, W.F., III (1999). Modern Applications of Analytical Ultracentrifugation. *Annu. Rev. Biophys. Biomol. Struct.* 28, 75-100.

Articles:

Laue, T.M., Shah, B.D., Ridgeway, T.M., and Pelletier, S.L. (1992). Computer-aided interpretation of analytical sedimentation data for proteins. In: Analytical ultracentrifugation in biochemistry and polymer science. S.E. Harding, A.J. Rowe, and J.C. Horton, eds. Royal Society of Chemistry, Cambridge, pp. 90-125. *Procedures for calculating partial specific volume, density, sedimentation coefficient (corrected for water @ 20°C and extrapolated to zero concentration) hydration, frictional rations, ellipsoidal shapes, etc; basis for SEDNTERF software (but note that the formulae and tables contain a number of typographical errors that were corrected in SEDNTERP Help file for corrected formulas)*

Stafford, W.F., III. (1992). Boundary analysis in sedimentation transport experiments: A procedure for obtaining sedimentation coefficient distributions using the time derivative of the concentration profile. *Anal. Biochem.* 203, 295-301. *Initial publication describing the dc/dt method*

Johnson, M.L. and Frasier, S.G. (1985). Nonlinear least-squares analysis. *Methods Enzymol*. 117:301-342. *Good overview of the fitting of experimental data*

