



Professional Development





Comparative Proteomics Kit I: Protein Profiler Module

DNA+RNA+PROTEIN+TRAIT





Protein Profiler Kit

Instructors

Stan Hitomi

Coordinator – Math & Science San Ramon Valley Unified School District Danville, CA

Kirk Brown

Lead Instructor, Edward Teller Education Center Science Chair, Tracy High School and Delta College, Tracy, CA

Sherri Andrews, Ph.D.

Curriculum and Training Specialist Bio-Rad Laboratories

Essy Levy, M.Sc.

Curriculum and Training Specialist Bio-Rad Laboratories





Is There Something Fishy About Teaching Evolution?

Explore Biochemical Evidence for Evolution





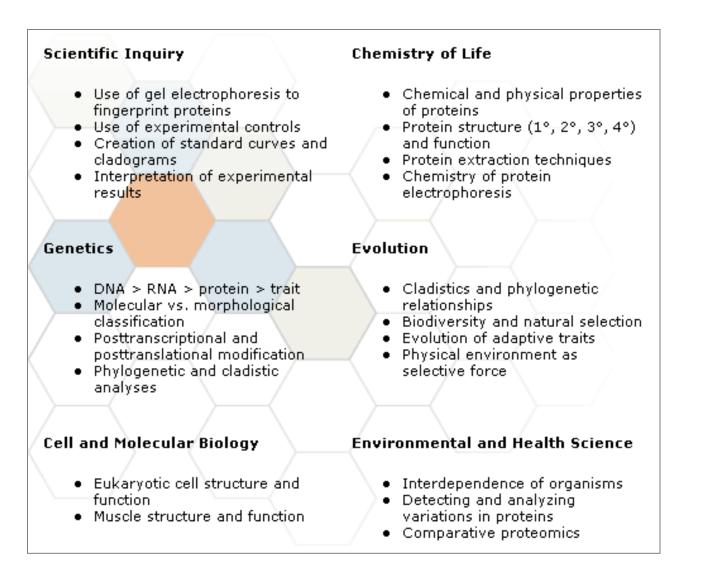
Why Teach Protein Electrophoresis?



- Powerful teaching tool
- Real-world connections
- Laboratory extensions
- Tangible results
- Link to careers and industry
- Standards-based











Comparative Proteomics I: Protein Profiler

Kit Advantages



- Analyze protein profiles from a variety of fish
- Study protein structure/function
- Use polyacrylamide electrophoresis to separate proteins by size
- Construct cladograms using data from students' gel analysis
- Compare biochemical and phylogenetic relationships. Hands-on evolution wet lab
- Sufficient materials for 8 student workstations
- Can be completed in three 45 minute lab sessions





Workshop Timeline



- Introduction
- Sample Preparation
- Load and electrophorese protein samples
- Compare protein profiles
- Construct cladograms
- Stain polyacrylamide gels
- Laboratory Extensions





Traditional Systematics and Taxonomy



- Classification
 - Kingdom
 - Phylum
 - Class
 - Order
 - Family
 - Genus
 - Species
- Traditional classification based upon traits:
 - Morphological
 - Behavioral





Can biomolecular evidence be used to determine evolutionary relationships?





Biochemical Similarities



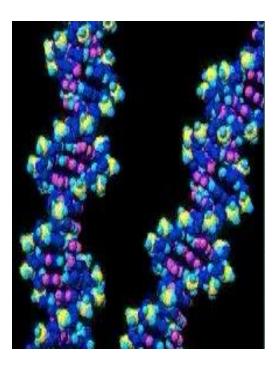
- Traits are the result of:
 - Structure
 - Function
- Proteins determine structure and function
- DNA codes for proteins that confer traits

DNA+RNA+PROTEIN+TRAIT





Biochemical Differences



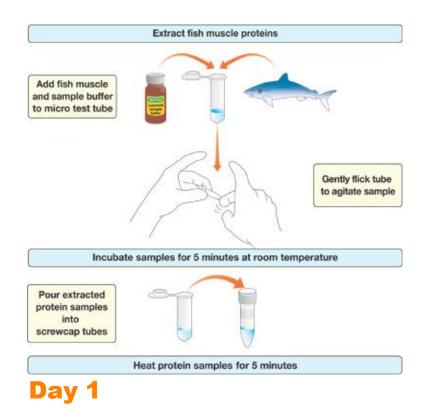
• Changes in DNA lead to proteins with:

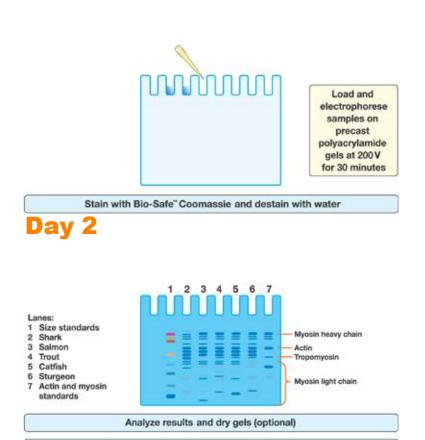
- Different functions
- Novel traits
- Positive, negative, or no effects
- Genetic diversity provides pool for natural selection = evolution





Protein Fingerprinting Procedures





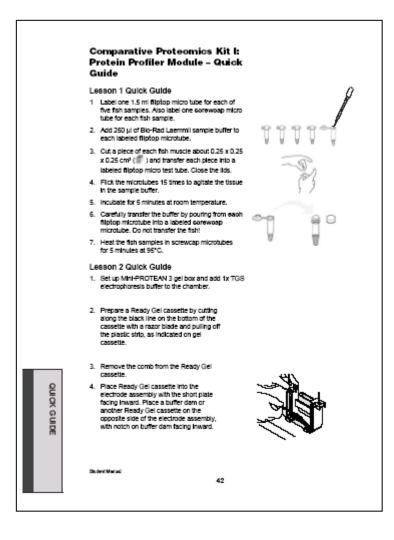
Compare molecular data to evolutionary tree

Day 3





Laboratory Quick Guide







What's in the Sample Buffer?



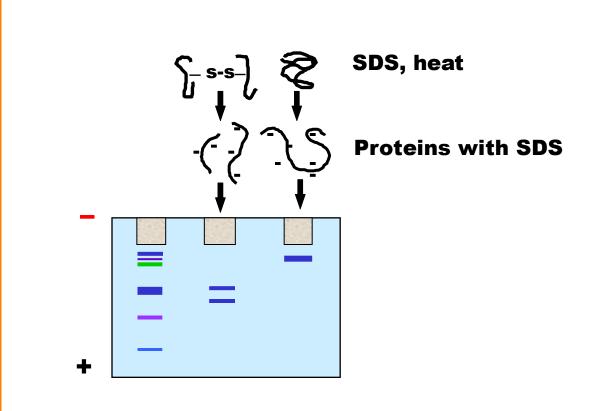
- Tris buffer to provide appropriate pH
- **SDS** (sodium dodecyl sulfate) detergent to dissolve proteins and give them a negative charge
- **Glycerol** to make samples sink into wells
- **Bromophenol Blue** dye to visualize samples





Why Heat the Samples?

• **Heating** the samples **denatures** protein complexes, allowing the separation of individual proteins by size







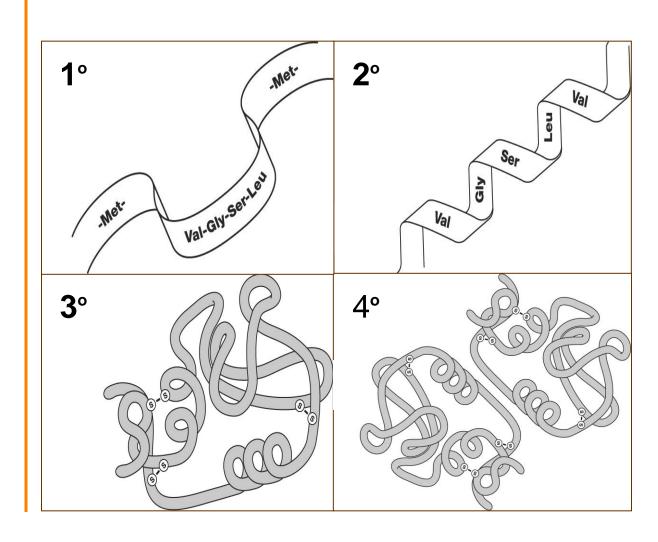
Making Proteins

DNA	TAC	GGA	TCG	AGA	TGA
mRNA	AUG	CCU	AGC	UCU	ACU
tRNA	UAC	GGA	UCG	AGA	UGA
Amino Acid	Tyr	Gly	Ser	Arg	STOP





Levels of Protein Organization







Protein Size Comparison

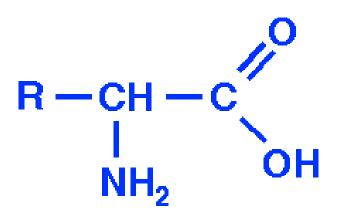
- Break protein complexes into individual proteins
- Denature proteins using detergent and heat
- Separate proteins based on size





Protein Size

- Size measured in kilodaltons (kD)
- Dalton = approximately the mass of one hydrogen atom or 1.66 x 10⁻²⁴ gram
- Average amino acid = 110 daltons







Muscle Contains Proteins of Many Sizes

Protein	kD	Function
Titin	3000	Center myosin in sarcomere
Dystrophin	400	Anchoring to plasma membrane
Filamin	270	Cross-link filaments
Myosin heavy chain	210	Slide filaments
Spectrin	265	Attach filaments to plasma membrane
Nebulin	107	Regulate actin assembly
α -actinin	100	Bundle filaments
Gelosin	90	Fragment filaments
Fimbrin	68	Bundle filaments
Actin	42	Form filaments
Tropomysin	35	Strengthen filaments
Myosin light chain	15-25	Slide filaments
Troponin (T.I.C.)	30, 19, 17	Mediate contraction
Thymosin	5	Sequester actin monomers





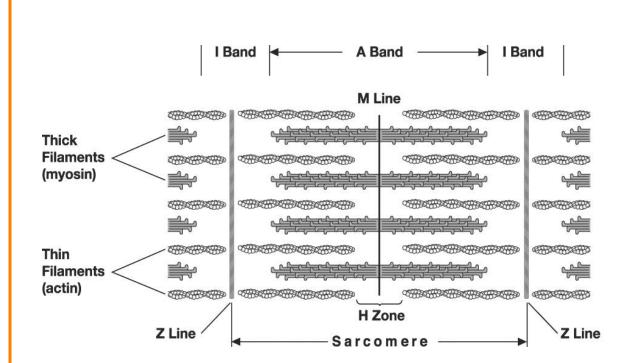
Actin and Myosin

Actin

- -5% of total protein
- -20% of vertebrate muscle mass
- 375 amino acids
 = 42 kD
- Forms filaments

• Myosin

- Tetramer
- two heavy subunits (220 kD)
- two light subunits (15-25 kD)
- Breaks down ATP for muscle contraction

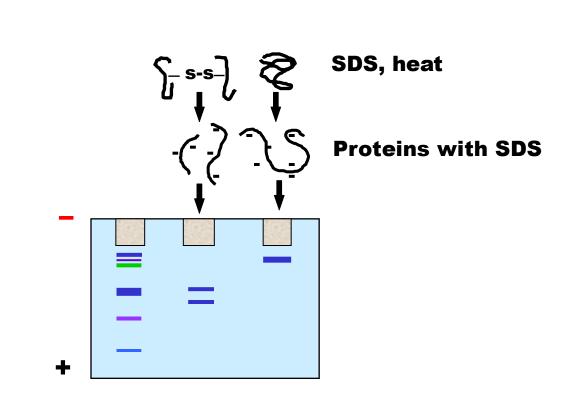






How Does an SDS-PAGE Gel Work?

- Negatively charged proteins move to positive electrode
- Smaller proteins move faster
- **Proteins** separate by size







SDS-Polyacrylamide Gel **Electrophoresis** (SDS-PAGE)

• **SDS detergent** (sodium dodecyl sulfate)

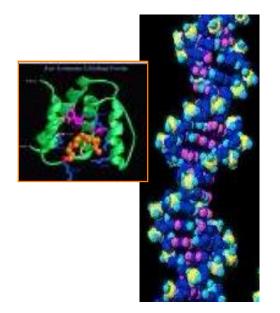
- Solubilizes and denatures proteins
- Adds negative charge to proteins
- Heat denatures proteins

$$\begin{array}{c}
CH_{3} \\
CH_{2} \\
CH_{$$





Why Use Polyacrylamide Gels to Separate Proteins?



- Polyacrylamide gel has a tight matrix
- Ideal for protein separation
- Smaller pore size than agarose

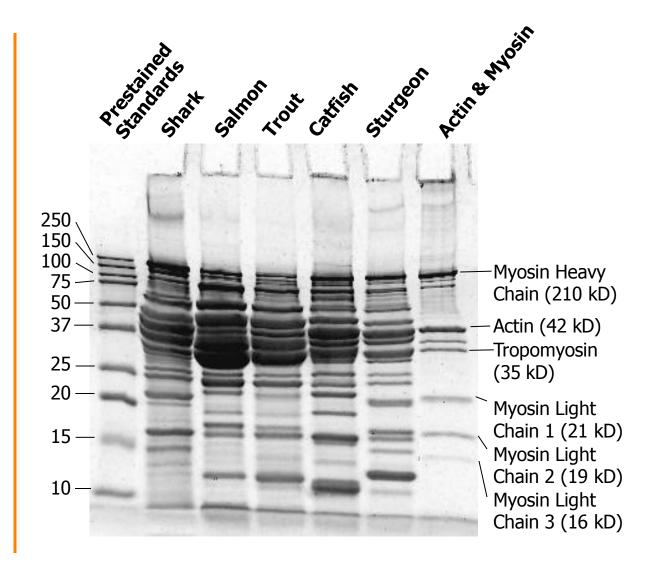
Proteins much smaller than DNA

- Average amino acid = 110 daltons
- Average nucleotide pair = 649 daltons
- -1 kilobase of DNA = 650 kD
- 1 kilobase of DNA encodes 333 amino acids = 36 kD





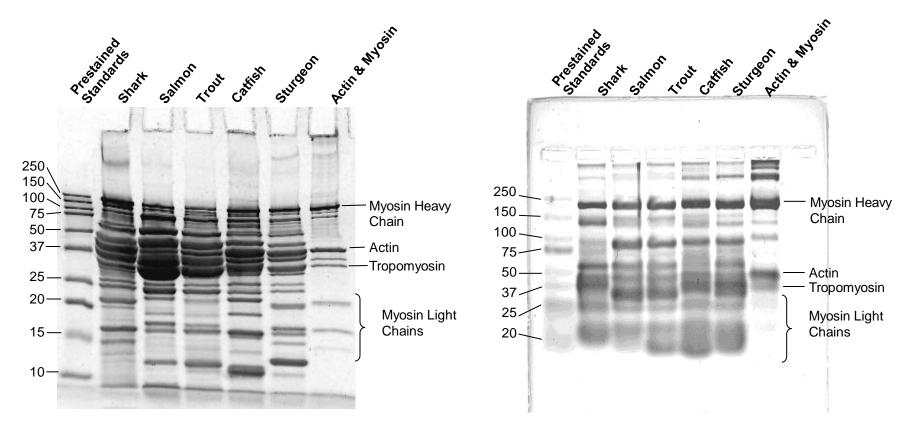
Polyacrylamide Gel Analysis







Can Proteins be Separated on Agarose Gels?



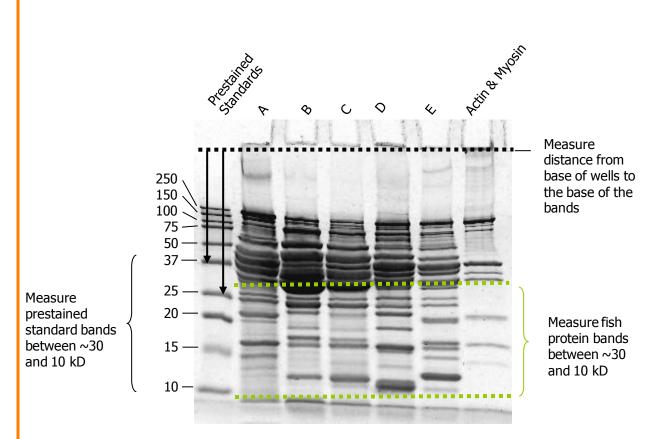
Polyacrylamide

Agarose





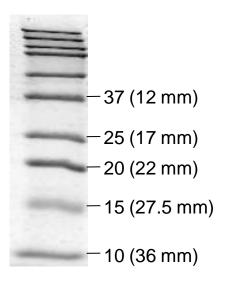
Determine Size of Fish Proteins

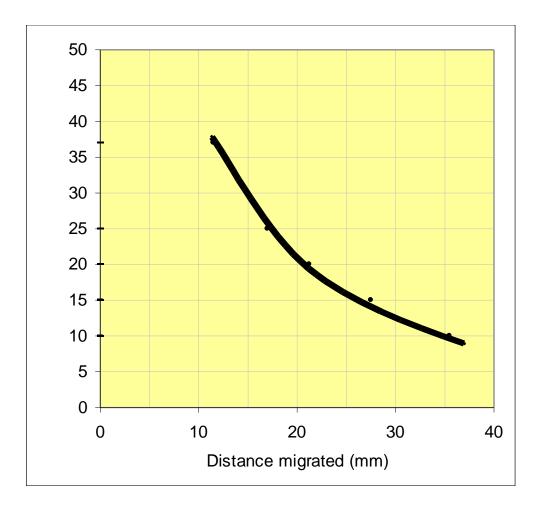






Molecular Mass Estimation

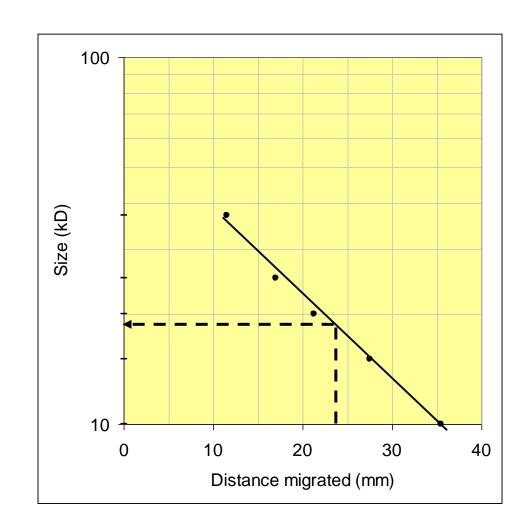








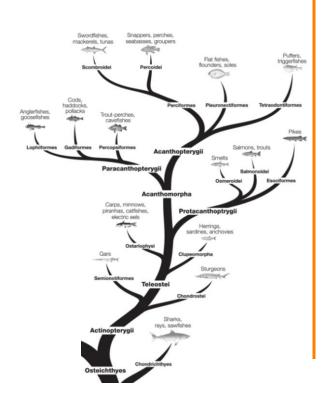
Molecular Mass Analysis With Semi-log Graph Paper

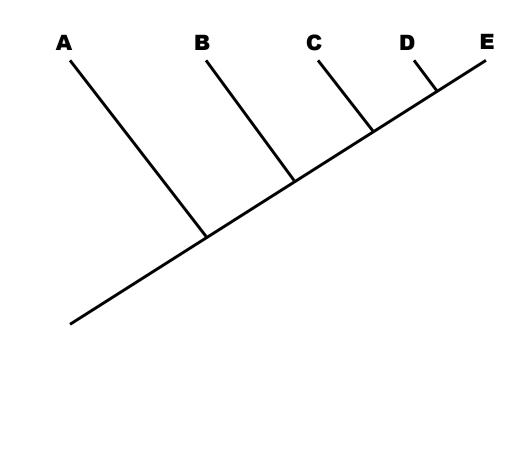






Using Gel Data to Construct a Phylogenetic Tree or Cladogram









Each Fish Has a Distinct Set of Proteins

	Shark	Salmon	Trout	Catfish	Sturgeon
Total # proteins	8	10	13	10	12
Distance proteins migrated (mm)	25, 26.5, 29, 36, 36.5, 39, 44, 52	26, 27.5, 29, 32, 34.5, 36.5, 37.5, 40.5, 42, 45	26, 27.5, 29, 29.5, 32, 34.5, 36.5, 37.5, 40.5, 42, 45, 46.5, 51.5	26, 27.5, 29, 32, 36.5, 38, 38.5, 41, 46, 47.5	26, 27.5, 30, 30.5, 33, 35.5, 37, 39, 39.5, 42, 44, 47





Some of Those Proteins Are Shared Between Fish

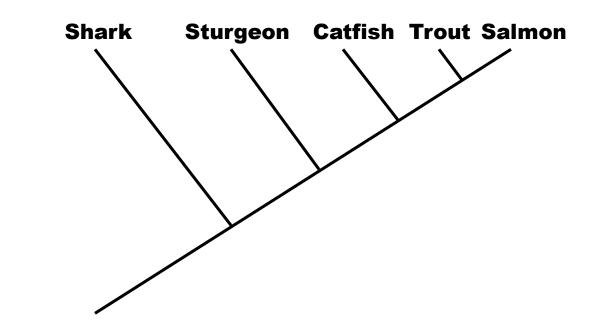
Distance (mm) 25	(KD) 32.5	Shark X	Salmon	Trout	Catfish	Sturgeon
26	31.5		X	X	X	X
26.5	31.0	X				
27.5	30.0		Х	X	X	X
28.5	29.1					
29	28.6	X	X	X	X	
30	27.6			X		X
30.5	27.1					X
32	25.6		X	X	X	
33	24.7					X
34.5	23.2		X	X		
35.5	22.2					X
36	21.7	X				
36.5	21.2	X	X	X	X	
37	20.7					X
37.5	20.2		X	X		
38	19.7				X	
38.5	19.3				X	





Character Matrix Is Generated and Cladogram Constructed

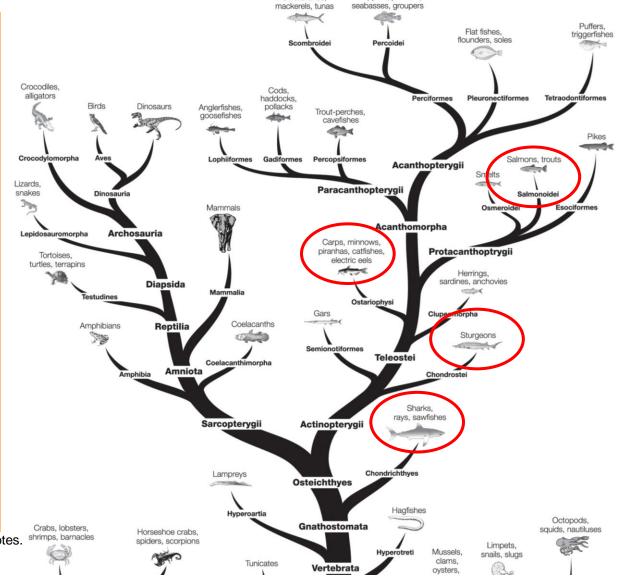
	Shark	Salmon	Trout	Catfish	Sturgeon
Shark	8	2	2	2	2
Salmon	2	10	10	5	3
Trout	2	10	13	5	4
Catfish	2	5	5	10	2
Sturgeon	2	3	4	2	12







Phylogenetic Tree



Swordfishes,

Snappers, perches,

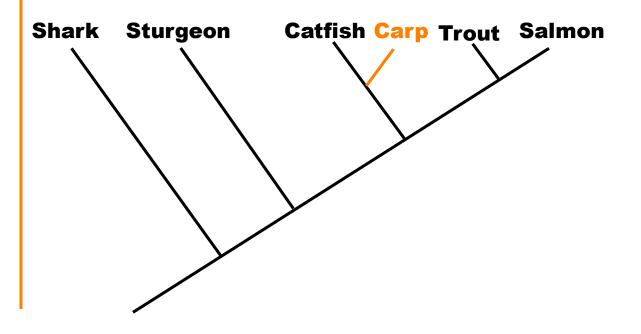
Evolutionary tree showing the relationships of eukaryotes. (Figure adapted from the tree of life web page from the University of Arizona (www.tolweb.org).)





Pairs of Fish May Have More in Common Than to the Others

	Shark	Salmon	Trout	Catfish	Sturgeon	Carp
Shark	8	2	2	2	2	2
Salmon	2	10	10	5	3	5
Trout	2	10	13	5	4	5
Catfish	2	5	5	10	2	8
Sturge on	2	3	4	2	12	2
Carp	2	5	5	8	2	11



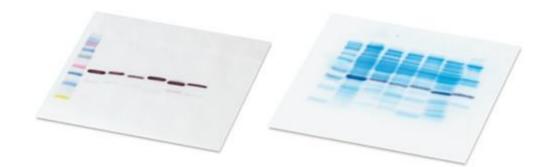




Extensions



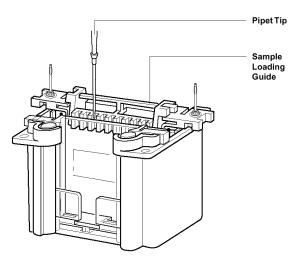
- Independent study
- Western blot analysis

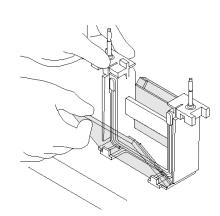




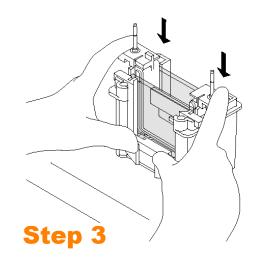


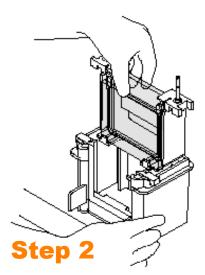
Ready Gel[®] Precast Gel Assembly

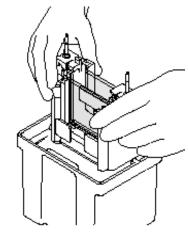




Step 1







Step 4