

Techniques in Molecular Genetics

Protein and Enzyme Assays

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Day 4

- Protein&EnzymeAssays
- Start SDS Gels
- Immuno (Western) Blotting

Enzyme Assay for Acid Phosphatase

- **6.6 Enzyme assay for acid phosphatase activity**
(Modified from (Dassa et al, 1982) : <http://www.jbc.org/cgi/reprint/257/12/6669>)
- **Protocols**
- **Prepare sample extract**
 - Inoculate overnight cultures of WT and *rpoS* mutant in 10 ml LB and growth at 37°C
 - Collect cells by centrifugation at 4,000 rpm for 10 min
 - Discard supernatant and resuspend pellet in 10 ml LB 20mM Phosphate buffer (pH7.0) (Pi buffer), ice cold.
 - Spin down the cells at 4,000 rpm for 10 min
 - Repeat steps 3 and 4 two more times
 - Discard supernatant and resuspend pellet in 1 ml Pi buffer
 - Transfer 0.5 ml to 15 ml Falcon blue cap tube, and save the other 0.5 ml on ice
 - Sonicate until cultures turn clear (less than 5 min)
 - Transfer crude extracts to 1.5 ml Eppendorf tubes and spin down at 13,000 rpm for 10 min
 - Carefully transfer the supernatant to clean 1.5 ml tubes
 - Quantify protein concentration using Bradford assay
- **Enzyme assay**
 - Prepare reaction buffer containing : 250 mM glycine/HCl buffer with 25 mM *p*-nitrophenylphosphate (PNPP)
 - Prepare 3 tubes, labelled with (1. Control 2. WT 3. *rpoS*)
 - Add 0.5 ml reaction buffer into each tube
 - Incubate all three tubes in water bath at 37°C for 2 min to equilibrate
 - Start the reaction by adding 20 µl cell extract (if the protein concentration is higher than 1 mg/ml) of WT and *rpoS* to tube 2 and 3, respectively.
Add 20 µl of Pi buffer to control tube 1
 - Incubate at 37°C for 15 min and stop the reaction by adding 1 ml of 1N NaOH
 - Transfer 1 ml reaction mixture to cuvette to measure absorbance at 410 nm
- **Calculation of activity**
 - One unit of enzyme is defined as the amount of acid phosphatase required to release 1 nmol of PNPP/min under the above defined conditions (At 25 mM, the no enzyme rate of PNPP hydrolysis was 0.49 nmol/min/ml)
 - Alternatively, one unit of enzyme was defined as 1,000 X A410 per minute per ml (Atlung et al. 1989)

Acid Phosphatase

Expression of acid phosphatase (AP), encoded by the *appA* gene, is RpoS dependent. You will test AP activity in cell extracts prepared from wild type and *rpoS* mutant strains of *Escherichia coli*.

Replication

- **Biological Replication (experiment error)**
 - Replication of the EXPERIMENTAL UNIT
 - Use at least 3 replications
 - Contributes to statistical experimental error
 - e.g. three independently prepared cell extracts
- **Technical Replication (sampling error)**
 - Repeated sampling of the same EXPERIMENTAL UNIT
 - Can use two replicates
 - Does not contribute statistical experimental error
 - e.g. two measurements of the same cell extract.
 - Technical Error (within EU error) is usually much smaller than Biological Error (between EU error)

Replication

- All assays should be replicated. All published experiments must be performed several times.
- Types of Replication
 - Biological Replication
 - Technical Replication

Enzyme Assay for Acid Phosphatase

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Quantify protein concentration using Bradford assay

(this has been done for you..)

Enzyme Assay for Acid Phosphatase

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Enzyme assay

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Prepare 3 tubes in , labelled with (1. Control 2. WT 3. rpoS)

Add 0.5 ml reaction buffer into each tube

Incubate all three tubes in water bath at 37°C for 2 min to equilibrate

Start the reaction by adding 20 μ l cell extract (if the protein concentration is higher than 1 mg/ml) of WT and rpoS to tube 2 and 3, respectively. Add 20 μ l of Pi buffer to control tube 1

Incubate at 37°C for 15 min and stop the reaction by adding 1 ml of 1N NaOH

Transfer 1 ml reaction mixture to cuvette to measure absorbance at 410 nm

Calculation of activity

One unit of enzyme is defined as the amount of acid phosphatase required to release 1 nmol of PNPP/min under the above defined conditions (At 25 mM, the no enzyme rate of PNPP hydrolysis was 0.49 nmol/min/ml)

- Alternatively, one unit of enzyme is defined as $1,000 \times A_{410}$ per minute per ml (Atlung et al. 1989)
- Molar extinction coefficient for *p*-nitrophenol = $17.700 \times 10^3 \text{ M}^{-1} \text{ cm}^{-1}$ (this has been incorporated into the general spreadsheet I will give out)

Calculations

- Prepare a table beforehand
- Label Tubes before doing experiment
- Use a spreadsheet table to record results.
- Put spreadsheet in labbook (raw and calculated data)

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Calculations-Spreadsheet

- The general spreadsheet I will post will be slightly modified from the following sheets given in class.
- Features
 - Formats numbers and decimals properly
 - Calculates data
 - Organizes information including experimental details
 - Provides some quality control indicators
 - Graphs protein standard curve

Protein..

Bradford Protein Assay (MICROASSAY) Date 5/8/2013 20:14

- add water to standard tubes in triplicate User
- add protein standard volume to each tube, use a fesh tip for each tube (total volume)
- add Protein dye reagent (200ul) to each tube: allow 15s between additions
- mix thoroughly with a vortex mixer
- incubate at least 5 min (preferably 30min) and read at 595 in the same order as protein reagent addition

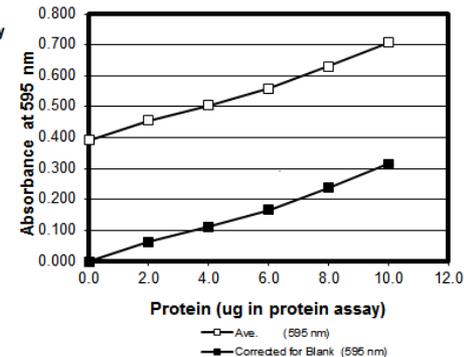
- Notes: (1) the "0" protein control as it is used for both the standard curve and for background subtraction (make sure calculated and measured values are in good agreement)
- (2) make sure that all unknown values fall in the linear portion of the curve--if not, dilute and reassy
- (3) if possible, use Corrected A595 (Average-blank) values in the range 0.01 to 0.400
- (4) if the calculated Intercept differs from the Corrected for Blank (below) by more than .005, check standard curve
- (5) Use 1.0 mg/mL BSA Standard

YOU CAN ONLY TYPE INTO YELLOW CELLS -OTHER CELLS ARE LOCKED

Standard Curve

BSA Protein (ug)	Volume of Standard (ul)	Volume of Water (ul)	Total Vol.	Abs At 595 nm (triplicate)			Ave. (595 nm)	Corrected for Blank
0	0.0	800.0	800.0	0.392	0.400	0.384	0.392	0.000
2	2.0	798.0	800.0	0.450	0.455	0.460	0.455	0.063
4	4.0	796.0	800.0	0.501	0.499	0.510	0.503	0.111
6	6.0	794.0	800.0	0.570	0.555	0.550	0.558	0.166
8	8.0	792.0	800.0	0.620	0.630	0.640	0.630	0.238
10	10.0	790.0	800.0	0.700	0.720	0.700	0.707	0.315

Bradford Assay Standard Curve



Regression

Slope	0.0307619
Intercept	-0.0049206

Unknown Samples

Sample Number	SampleID	Volume of Sample (ul)	Volume of Water (ul)	Readings		Average	Average-blank	Protein (mg/mL)	Error Flags	
				Rep #1	Rep#2				2.5%	5.0%
1	First sample	2.0	798.0	0.552	0.558	0.555	0.163	2.65		
2										
3										
4										
5										
6										
7										
8										
9										
10										
11										

- User input
- Spreadsheet calculated values
- Protein value wrong--check standard curve

Enzyme..

General Enzyme Calculation Template

Date 5/8/2013 20:16

- determine protein concentration of samples using ProteinMicroassay or ProteinMicrotiterPlateassay sheet
- determine activity of samples using standard curve calibration assay below to obtain Total Units per mL
- divide Total Units per mL by Protein (mg/ml) to obtain specific activity

A. Enzyme assay

1. Prepare reaction buffer containing : 250 mM glycine/HCl buffer with 25 mM p-nitrophenylphosphate (PNPP)
2. Prepare 3 tubes, labelled with (1. Control 2. WT 3. rpoS)
3. Add 0.5 ml reaction buffer into each tube
4. Incubate all three tubes in water bath at 37oC for 2 min to equilibrate
5. Start the reaction by adding 20 µl cell extract (if the protein concentration is higher than 1 mg/ml) of WT and rpoS to tube 2 and 3, respectively. Add 20 µl of Pi buffer to control tube 1
6. Incubate at 37oC for 15 min and stop the reaction by adding 1 ml of 1N NaOH (TOTAL Volume = 1.52 mL)
7. Transfer 1 ml reaction mixture to cuvette to measure absorbance at 410 nm

Assay Notes : (1) make sure that you perform linearity checks
 (2) use only determine unknown values with in the standard curve (do not extrapolate..dilute and re-assay instead)
 (3) biological replicates usually yields standard errors of less than 10% (5% is quite good). If yours is higher check your technique.

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ACID PHOSPHATASE ASSAY DATA

Sample	Sample ID	Protein concentration (mg/mL)				Volume of Sample (ul)	Time of Incubation (min)	OD410 READINGS						Total Activity (nmol/min/mL)				SPECIFIC ACTIVITY	Standard Error	Error Flags	
		Rep1	Rep2	Rep3	Average			Rep1	Rep2	Rep3	Rep1	Rep2	Rep3	Average	10%	20%					
		Sample1	Sample2	Sample1	Sample2			Sample1	Sample2	Rep 1	Rep 2	Rep 3	Average								
1	Your first sample	1.30	1.50	1.45	1.417	10.0	20.0	0.600	0.600	0.610	0.605	0.650	0.650	2576.3	2597.7	2608.5	2594.2	1831.2	75.6		
2																	#DIV/0!	#DIV/0!	#DIV/0!		
3																	#DIV/0!	#DIV/0!	#DIV/0!		
4																	#DIV/0!	#DIV/0!	#DIV/0!		
5																	#DIV/0!	#DIV/0!	#DIV/0!		
6																	#DIV/0!	#DIV/0!	#DIV/0!		
7																	#DIV/0!	#DIV/0!	#DIV/0!		
8																	#DIV/0!	#DIV/0!	#DIV/0!		
9																	#DIV/0!	#DIV/0!	#DIV/0!		
10																	#DIV/0!	#DIV/0!	#DIV/0!		
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19																	#DIV/0!	#DIV/0!	#DIV/0!		
20																	#DIV/0!	#DIV/0!	#DIV/0!		

CALCULATION NOTES

Protein Concentration (from Protein Assay Table)

Rep - Biological Replicate of Experimental unit

Sample - Technical Replicate

TOTAL ACTIVITY = ((AverageOD410/1770 L mol⁻¹cm⁻¹) x (10E9 nmol/mol) x (1.0 l/1000mL) x 1.52mL (total assay volume) x ((1/incubation time) (min)) / sample volume (mL)

SPECIFIC ACTIVITY = TOTAL ACTIVITY/PROTEIN CONC. (e.g. S25/G25)

STANDARD ERROR = STANDARD DEVIATION of Biological Replicate Specific Activities (e.g. Standard Deviation (P25/D25, Q25/E25, R25/F25) / SQRT(3))

ERROR FLAGS - these will indicate if variation is high. If high, check to make sure numbers are typed correctly--it is easy to mistype a decimal

-this is not operational

MOLAR EXTINCTION COEFFICIENT (ε) OF pPNP = 17,700 M⁻¹ cm⁻¹

	=user inputted data
	= spreadsheet calculated data
	= negative number (this occur if blank values are higher than measured values)
	error flag - excess variation
	error flag - very high variation that will likely affect ability to employ statistical tests of significance