Utilizing 2DE gels for carcinogen screening

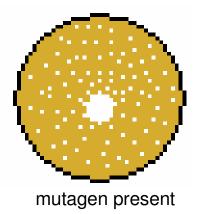
MaryEd Pratts

Overview

- Current method used for mutagen screening – The Ames Salmonella test
- 1. How it works (S9)
- 2. Why a compliment would be helpful
- Our Approach 2DE gels
- 1. How and why we use them
- 2. Current work and accomplishments
- 3. Future work

The Ames Salmonella assay





- •The Ames test consists in the detection of mutations by a bacteria that is histidine dependent: Salmonella typhimurium.
- •Several strains of this bacteria, all carriers of a mutation in the his operon, can revert spontaneously to *His*+, and thus grow in a histidine-free medium.
- •This very weak spontaneous reversion can be increased by mutagens, which allows the qualification of the mutagenic potential of these substances.

S9 Rat Liver Enzymes

- Bacteria lack the oxidative enzyme systems for metabolizing some compounds to electrophilic metabolites capable of reacting with DNA.
- The metabolic activation system usually consists of a 9000xg supernatant fraction of rat liver homogenate in the presence of NADP and NADPH cofactors.



Limitations of Ames

- False positives
- False negatives
- Only compliments are extensive and usually require live animals
- Only cellular growth level, not any protein expression change

Use of 2DE gels

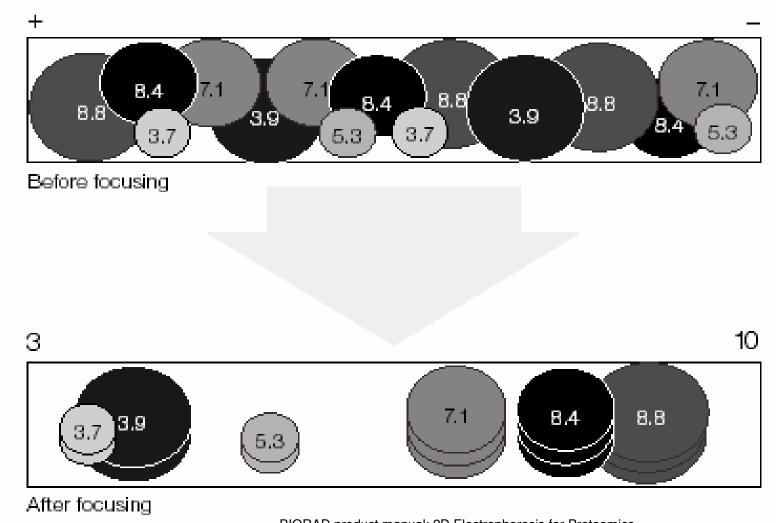
- Most commonly used technique in proteomics
- Proteins are separated by two different chemical properties – pH dependent isoelectric point (pl) and molecular weight
- Can easily be coupled with mass spec to identify the isolated proteins

How 2DE gels work

The first dimension:

- Isoelectric Focusing (IEF)
- Protiens are applied to strip containing an immobilized pH gradient (IPG strip)
- Migrate to their pI point net charge is zero

IEF



BIORAD product manual: 2D Electrophoresis for Proteomics

Slide taken from Kelly Fowlkes

How 2DE gels work

The second dimension:

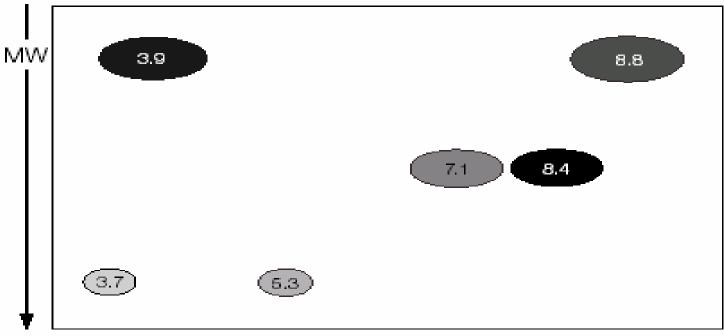
- IEF-focused proteins are equilibrated in SDS and reducing agents
- SDS coats the proteins in proportion to their mass
- Electric current is applied through the polyacrylamide gel and the proteins will migrate.

SDS-PAGE



SDS-charged proteins in IPG strip

Apply to SDS-PAGE get



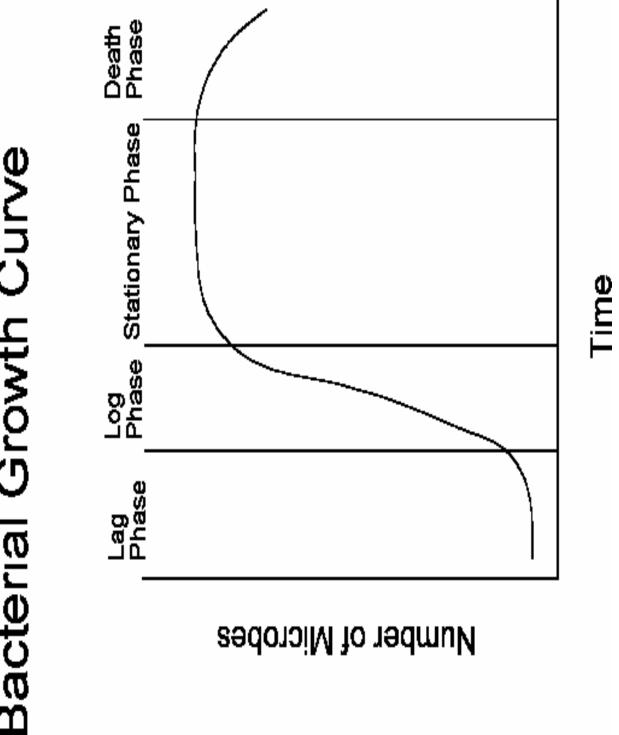
SDS-charged proteins resolved according to size in SDS-PAGE gel

How we use 2DE

- Grow the bacteria Pseudomonas putida on standard carbon sources in the presence of different carcinogens
- 2. Examine the proteins by 2DE
- 3. Identify the expressed proteins
- 4. Eventually replace the Ames Test

Sample Prep

- A bio-safe soil bacteria: Pseudomonas putida strains P.p.F1and P.p.KT2440 are used
- To determine when to extract the protein from cells (mid-log), a growth curve for a specific set of conditions is conducted



Sample Prep

- The preparation to extract the protein from the cells consists of: Centrifugation, Sonication, Microfugation, RNAase treatment, protein assay, and rehydration
- 2DE gels are then run on the protein sample, and the gels are then stained
- The gels are then scanned and dried
- Progenesis software is then used to analyze and compare gels

Gel Results

- We have the complete proteomic signature for that set of conditions
- Once the gels have been analyzed, we can identify biomarkers under specific growth conditions
- Mass spectrometry or other methods can then be used to find the identity of the biomarker proteins.

2DE gels - why

- In determining the biomarkers, we can also determine the differences in protein expression when a carcinogen is present in the growth conditions
- This method can be used as a test for carcinogens if common proteins (such as DNA repair) are expressed in the presence of all carcinogenic compounds
- This would be a more specific test than Ames which only tests on the cellular growth level

Group Accomplishments

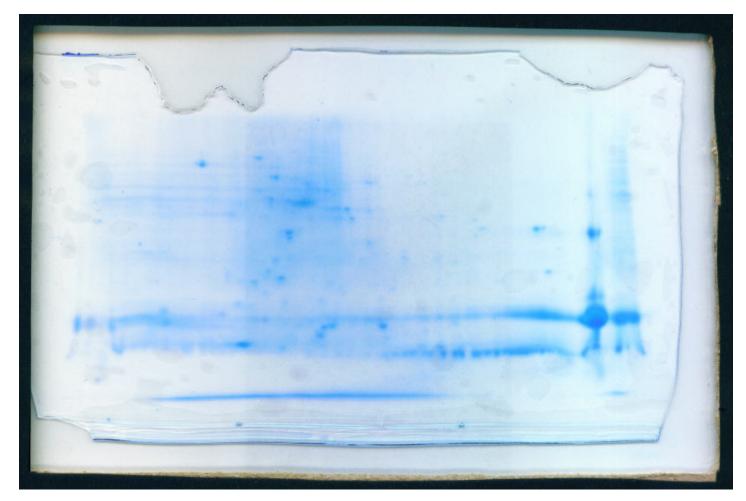
- Growth curves for P.p.F1 have been conducted using: Succinic Acid, Phenylethylamine, Benzoic acid, and Succinic Acid in the presence of 9-Aminoacridine.
- Growth curves for P.p.KT2440 have been conducted using: Succinic Acid, Phenylethylamine.

Group Accomplishments

- Gels have been run under all of these growth conditions
- Some gels have been analyzed and compared using the Progenisis software at the U of R proteomics lab
- Similarities and differences have been found in some of the gel comparisons

Gels





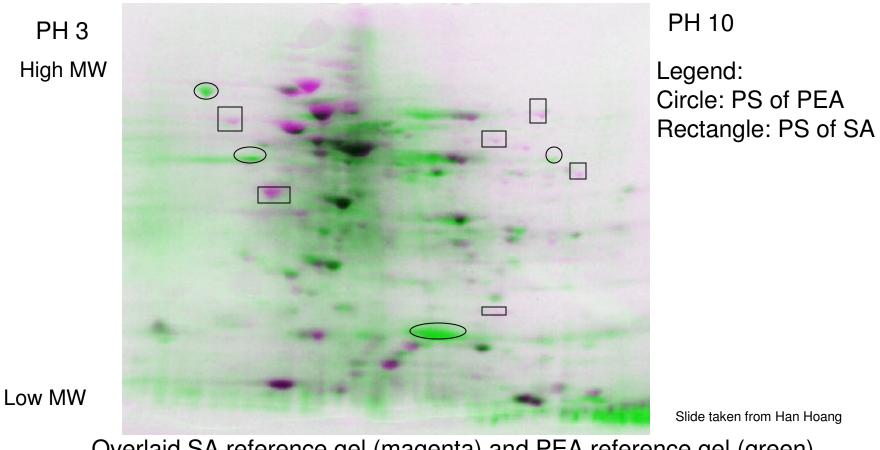
Low MW

Carbon Source: Succinic Acid Date:07

Date:07-29-2004 Mary Pratts

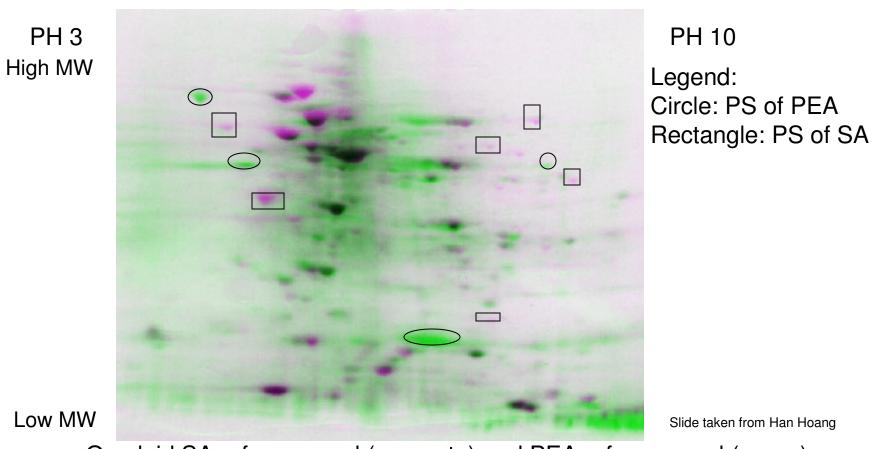
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2DE Gels – Compare – SA vs. PEA



Overlaid SA reference gel (magenta) and PEA reference gel (green)

2DE Gels – Compare – SA vs. PEA



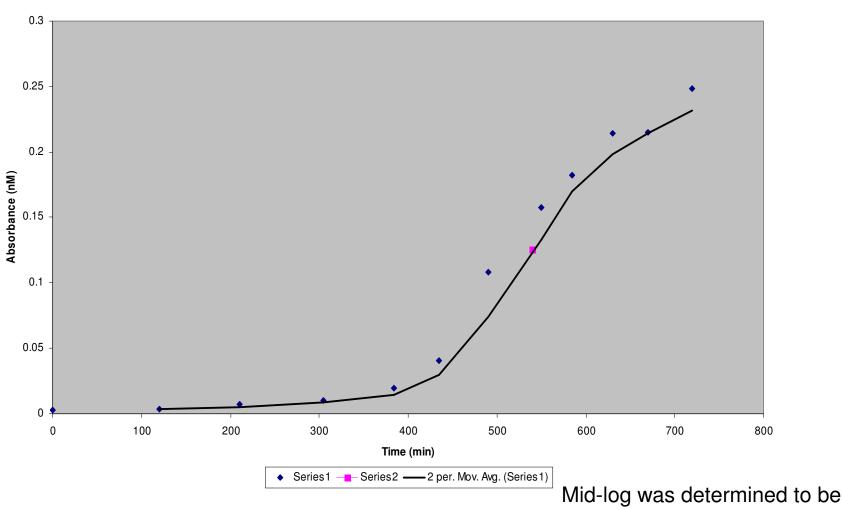
Overlaid SA reference gel (magenta) and PEA reference gel (green)

Summer Research

- Learn the SOP specific to our research lab and study goals
- Practice these techniques and become proficient in running gels
- Conduct growth curve for P.p.F1on a Succinic Acid carbon source in the Presence of the mutagen 9-Aminoacridine
- Run initial gels for this set of conditions

Growth Curve: SA and 9-AA

#2 SA and 9- AA



Mid-log was determined to be approx. 9hrs.

Future Plans

- Run and analyze gels with Succinic acid and Benzoic acid carbon sources in the presence of: 9-Aminoacridine, Sodium azide, and Benzo(a)pyrene
- Consider the addition of S9 for metabolic activation when using Benzo(a)pyrene

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- U of R Proteomics lab

References

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