Lecture 14: **Quantifying Cell Function**

c. Cell Proliferation Assays

Importance:

- wound healing
- immune host defense
- cancer therapy
- tissue engineering
- bioprocessing/synthesis of theraputic glycoproteins

> Cell number

- 1. count via microscopic observation or Coulter counter
- 2. compute specific *cell proliferation rate const*:

$$k_g = \frac{1}{N} \frac{dN}{dt} = \frac{d(\ln N)}{dt} \quad \text{(units: } t^{-1}\text{)}$$

N = # of cells at time t

> Cell phase populations

In some cases, we want to know the population of cells in each phase of the cell cycle (eukaryotes):



- M = Mitosis (~ 1 h)
- G_1 = gap between cell division & DNA synthesis (~18-72 h)
- S = DNA synthesis (~6-8 h)
- G_2 = gap between DNA synthesis & mitosis ($\sim 2-3$ h)

 G_0 = quiescent cells

tritiated-thymidine uptake

1. cells exposed to [³H]thymidine pulse which labels S-phase cells only

2. % S obtained from *autoradiography* (Ag precipitates in an overlying emulsion film reveal S-phase cells, similar to a photograph emulsion*)



3. % M and t_M obtained from optical microscopy (visually distinct)



4. by following % labeled mitoses after [³H]thymidine pulse through 2 cell divisions, t_{G2} , t_S and t_{cycle} can be determined $\rightarrow t_{G1}$



flow cytometry*

- 1. cells' DNA labeled with DNA-specific fluorescent dye (ex. acriflavine, ethidium bromide)
- 2. count cell # vs. fluorescence using laser excitation
- 3. fluorescence intensity ~ DNA present



G₁:S:G₂:M DNA (& fluorescence) ratio is 1:1-2:2:2

- no radiolabeling

- faster than autoradiography

- M & G₂ populations lumped

*flow cytometry is also used to determine expression of specific cell receptors by labeling with fluorescent antibodies

d. Cell Differentiation Assays

Importance:

- change in phenotype renders specialization of function
- characterized by changes in protein synthesis, genes expressed & secretions

2D Gel Electrophoresis (SDS-PAGE): cell proteins separated on basis of mass and charge—providing a "signature" of the cell

Method

1. cell is lysed (contents extracted)

2. protein solution is separated by pI (isoelectric point = pH where net charge = 0) using *isoelectric focusing*



3. gel strip is next placed crosswise on a second gel incorporating sodium dodecyl sulfate (SDS)

4. SDS binds & denatures proteins ⇒ native charge becomes negligible ~1 SDS per 2 amino acids

charge/mass ≈ constant

5. gel is placed under E-field, separating protein chains via molecular weight

low molecular weight species move faster

$$\mu = \mu(c_{gel}, E, MW_{protein})$$

SDS-PAGE: Sodium dodecyl sulfate polyacrylamide gel electrophoresis



6. compare "before" and "after" signature to discern differentiation

DNA microarrays: test for gene expression (first commercialized by Affymetrix)

array of "probes" comprising surface-bound gene fragments (~20 bases/fragment)



Method

- 1. cells lysed to retrieve mRNA
- 2. mRNA transcribed to cDNA (complement), which is then transcribed to cRNA
- 3. cRNA is cut into 35-200 base fragments & labeled with biotin (vitamin H)—referred to as "biotinylation"





4. cRNA "cocktail" is added to array, hybridizes with DNA fragments

5. surface-bound cRNA-biotin is labeled with fluorophore-labeled avidin



6. fluorescence spatial readout determines genes being expressed

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DNA Array Fabrication

Lithographic masks used to build up A-C-T-G sequences base-by-base



S.P.A. Fodor et al., Science 251, 767 (1991).

e. Secretion

Enzyme-linked immunosorbent assay (ELISA): tests for a particular protein's presence in cell secretions

1. cell secretions aspirated from culture medium



- 4. Color intensity is read in spectrophotometer
- 5. Nonspecific adsorption is determined and subtracted



Alternately, proteins directly adsorbed on a surface may be probed by ELISA in which the second (enzyme-linked) antibody is an antibody to the first antibody.

