Measuring gene expression and protein production

Module 3, Lecture 5

20.109 Spring 2013

Lessons from Lecture 4

- Always look for the recovery file
- The myth of multitasking
 - task-switching taps processing power
 - can't stay in tune with your audience
- Sunk costs and irrational decisions
- So... concrete examples of potential standards:
 - characterizing TE constructs (for the same application) by the same methods across labs, to facilitate efficacy comparisons: e.g., cartilage TE proteoglycans all measured by DMMB
 - for a given method, have base set of analysis points to clarify:
 e.g., threshold for positive signal as n std dev above average
 - having a base set of characteristics to test, even if not by the same exact assay: e.g., for all TE scaffolds test cell viability, compressive strength, standard gene expression, etc.

Existing ASTM standards for TE

Designation: F2212 - 11

Standard Guide for Characterization of Type I Collagen as Starting Material for Surgical Implants and Substrates for Tissue Engineered Medical Products (TEMPs)¹

5.7 Impurities Profile—The term impurity relates to the presence of extraneous substances and materials in the collagen. These impurities can be detected by Western blots, ELISAs, GC-MS, and other types of assays. The user is also directed to Guide E1298 for additional information. If there is

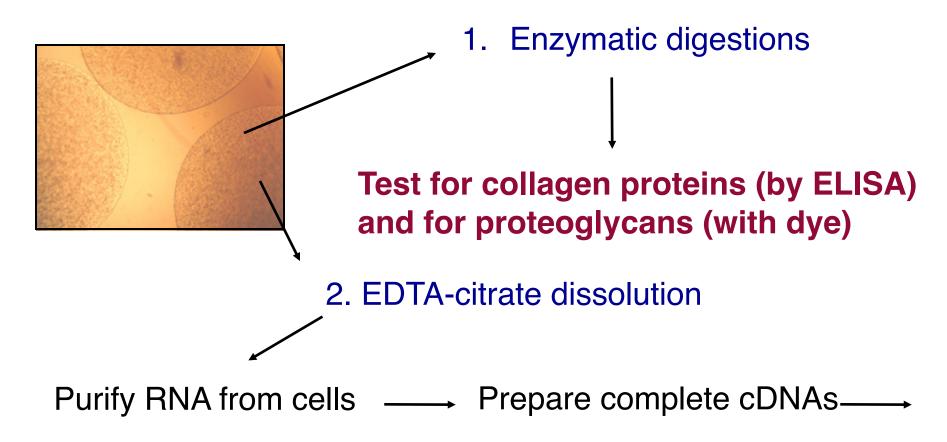
Topics for Lecture 5

- Measuring protein levels
- Measuring transcript levels
- Imaging assays

Sounds boring! Why bother?

- In 20.109, we tell you what assays to perform
 - designs vary, but measurement paths are identical
- In real research, you must decide not only what is worth measuring but how to measure it
 - sometimes just choosing among existing technologies
 - sometimes inventing something novel or customized
- Hey... this type of thinking also happens to be relevant to the M3 proposal!

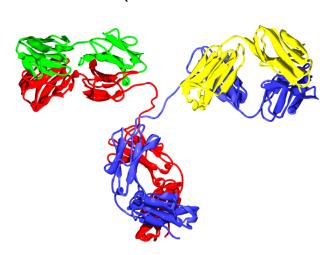
Module overview: 2nd half

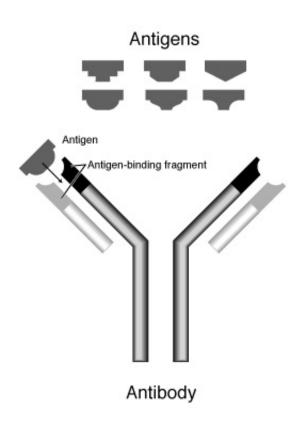


Run qPCR to measure CN II, CN I, and 18S RNA.

Antibodies are specific and diverse

- Specificity
 - variable region binding, K_D ~ nM
 - linear or conformational antigens
- Diversity
 - gene recombination
- Production
 - inject animal with antigen, collect blood
 - hybridomas (B cell + immortal cell)





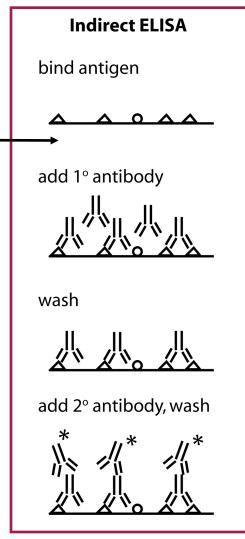
Public domain images (Wikimedia commons)

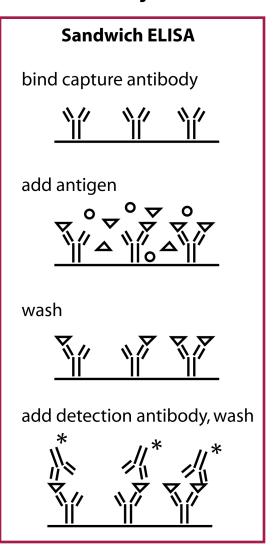
Day 5-7: protein analysis by ELISA

- ELISA: enzyme-linked immunosorbent assay
 - specific
 - sensitive
 - multiple kinds

"blocking" step also needed

= protein
of interest





Common protein-level assays

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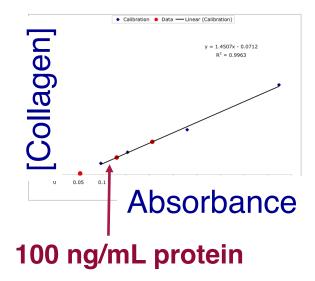
- simple and low cost
- Coomassie detection limit ~ 0.3-1 ug/band (2-5 ng/band for silver staining)
- cannot distinguish two proteins of same MW

Western blot

- identifies specific protein
- detection limit ~1 pg (chemiluminescent)
- only simple for denatured proteins

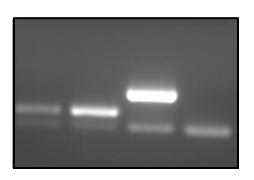
ELISA

- detects native state proteins
- quantitative
- high throughput

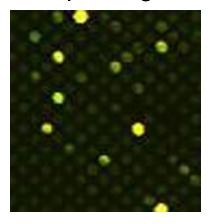


Common transcript-level assays

- RT-PCR (end-point)
 - simple, low cost
 - can be semi-quantitative
- Microarrays (end-point)
 - spotted c(cDNA)
 - high cost, need specialty equipment
 - complicated and fraught analysis
 - high throughput
- q-PCR (real-time)
 - some special equipment, medium cost
 - highly quantitative
 - multiplexing potential
 - requires optimization (primers)



Sample 1: red Sample 2: grn



End-point RT-PCR

- Co-amplification in one tube
 - collagen + GAPDH
- Optimize primers
 - no cross-hybridization
 - similar signals (vary [primer])
 - similar efficiency
- Reliability issues
 - must be in exponential phase
 - sensitive to change in [RNA]
- Visualize on a gel
 - measure band intensity/area
 - low dynamic range

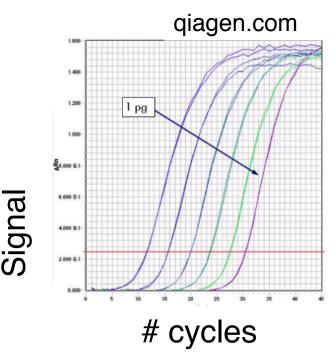


Collagen (upper band)
GAPDH (lower band)

Which sample is from chondrocytes, and which from stem cells?

Introduction to qPCR

- Real-time tracking of [DNA]
- Uses probes that fluoresce
 - when bind to any DNA
 - when bind to specific DNA (FRET)
- How and why does [DNA] change during PCR?
 - first plateau
 - exponential phase
 - linear phase
 - second plateau
 - 1: detection limit
 - 2: competition, reagent limits, inhibition
- Starting point for analysis: threshold cycle C_T



Current Protocols in Cell Biology, Molecular Biology

Interlude: grad school – to go or not?

February 25, 2013

For Graduate Science Programs, It's Time to Get Real

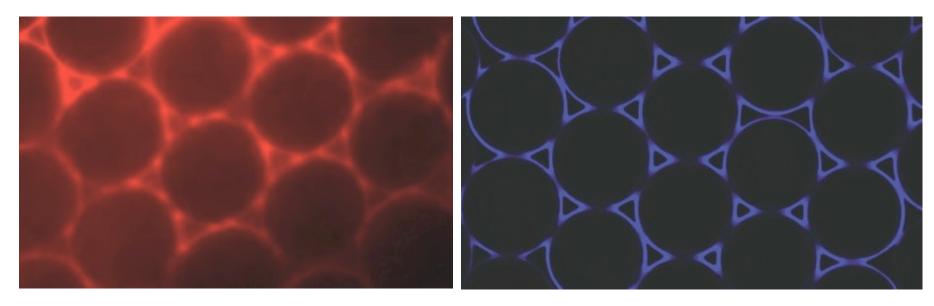
By Amanda A. Shea

The relationship between the number of Ph.D.'s in the sciences and the academic jobs available to them is, to put it scientifically, inversely related. Money for science has stagnated over the last decade and will most likely continue to do so, leading to too many Ph.D.'s competing for too few teaching and research jobs in academe.

From: Chronicle of Higher Education

Image quality in microscopy

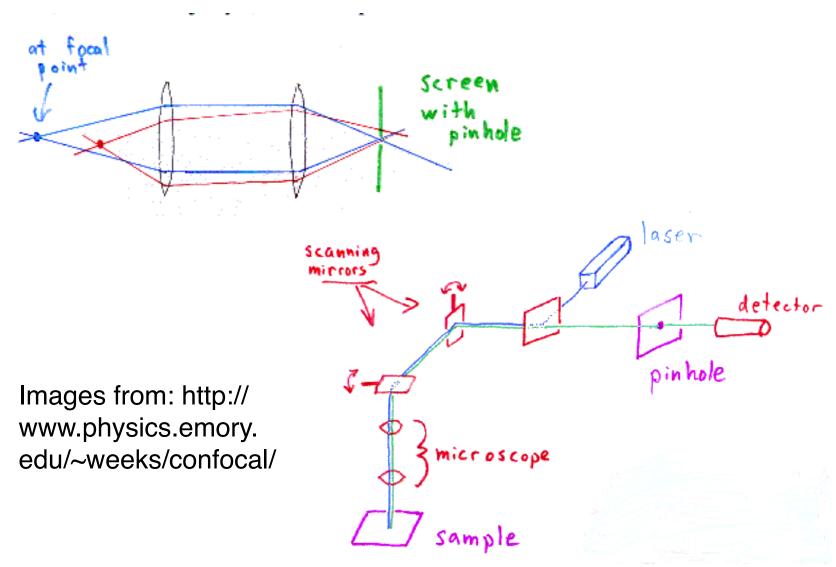
- Epifluorescence: noisy due to out-of-plane light
- · Confocal: pinhole rids out-of-plane light; scanning



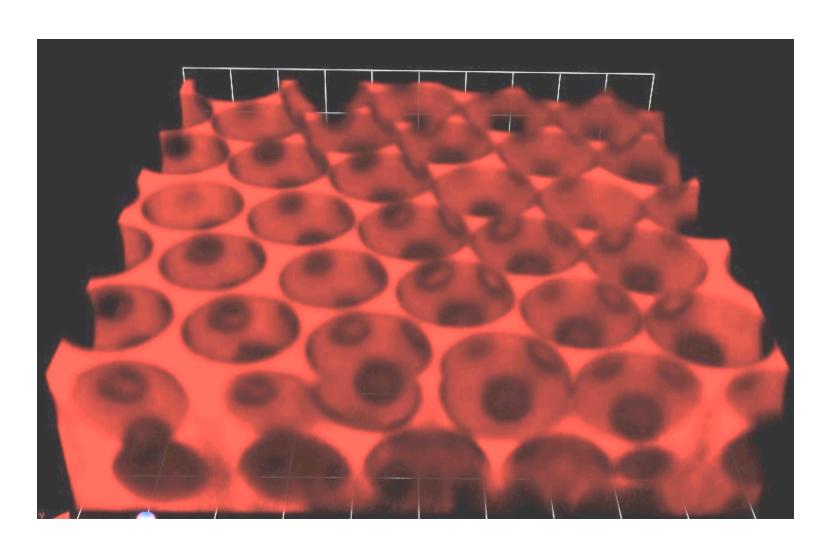
Epifluorescence

Confocal

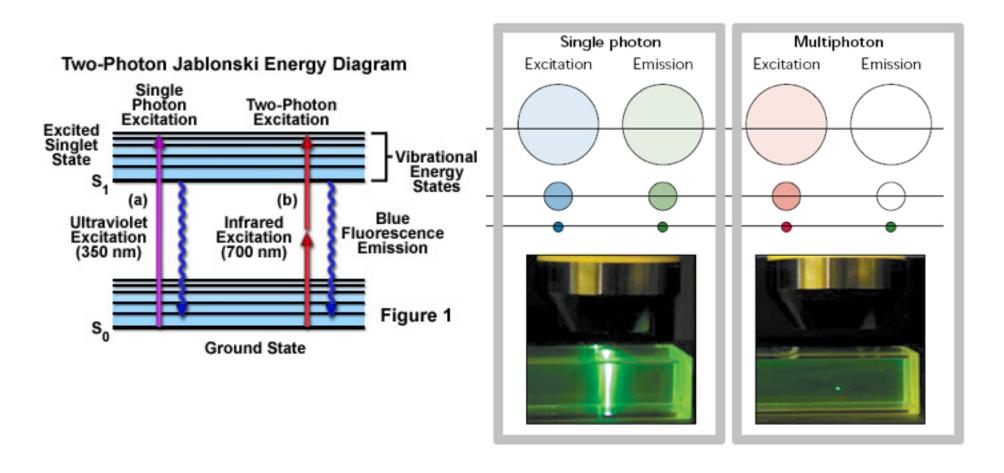
Confocal uscopy: theory



Confocal uscopy permits 3D reconstruction



2-photon microscopy: theory



Images: (1) microscopyu.com; (2) http://parkerlab.bio.uci.edu/ microscopy_construction/build_your_own_twophoton_microscope.htm ¹⁷

2-photon microscopy permits deep imaging, even *in vivo*

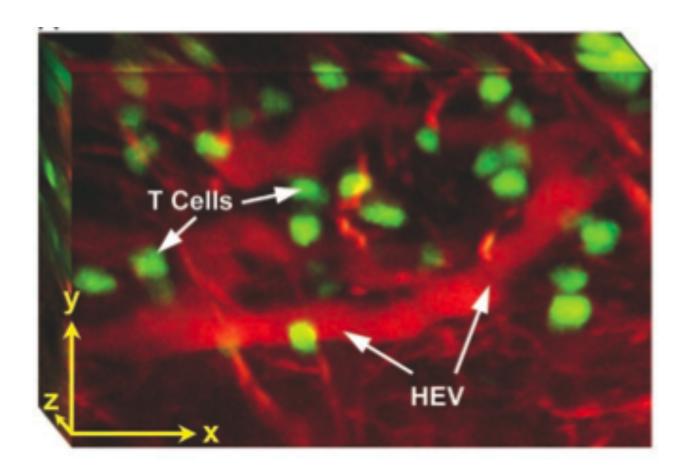


Image from: M.J. Miller, et al. *PNAS* **100**:2604 (2003)

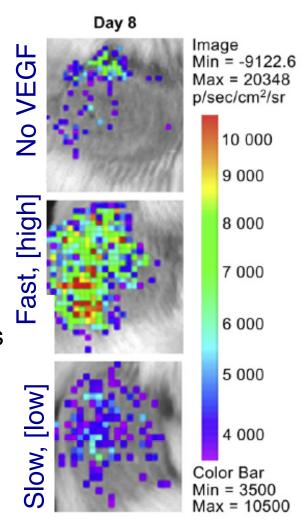
What kinds of information can imaging provide?

- Static measurements
 - overall cell state: viability, apoptosis, signaling
 - specific organelles, cytoskeleton
- Dynamic measurements
 - calcium (or other) fluxes
 - cell motility
- What is learned from single-cell (uscopy) vs. population (e.g., flow) assays? Pros/cons?
- Different modalities -> different information
 - vis-à-vis resolution, depth, coverage, signal:noise

Non-invasive imaging

- MRI, tomography, ultrasound, etc.
 - medical diagnostics
 - also measure gene expression
 - fuse gene with reporter
 - whole-body imaging with bioluminescence
- Example: monitoring angiogenesis
 - VEGF<u>R</u>2-*luc* (luciferase reporter)
 - slow- & fast-release VEGF in fibrin scaffolds
 - mice injected with luciferin (substrate) and observed for VEGF receptor upregulation
- Other uses? (think tumors)

M. Ehrbar, et al. *Biomaterials* **29**:1720 (2008) *Nature* News Feature **412**:372 (2001)

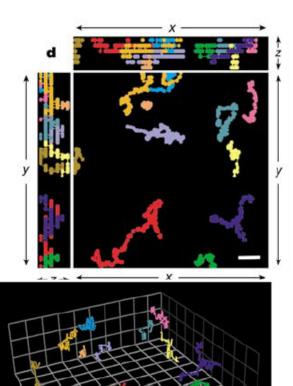


Day 5-6: image analysis

- Imaging data is often high throughput
 - 4D: time, *x-y-z*
 - requires computation, and
 - human design/interpretation
- Many available analysis packages
 - some ~ \$20-30K
 - NIH ImageJ = free
- Your analyses
 - automated cell counts
 - optional: explore other features

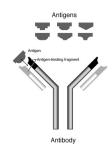
Images from: T.R. Mempel, et

al. Nature 427:154 (2004)

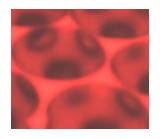


Lecture 5: conclusions

 Antibodies to diverse targets (e.g., proteins) can be made and used for detection/measurement.



- Trade-offs exist (e.g., between simplicity and accuracy) for different transcript-level assays.
- Fluorescence imaging is a powerful tool for studying cells and materials.



Next time: cartilage TE, from *in vitro* and *in vivo* models to the clinic; qPCR analysis.