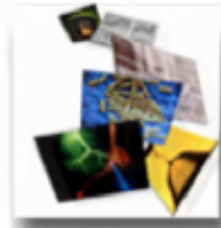




NMR  
Core facility



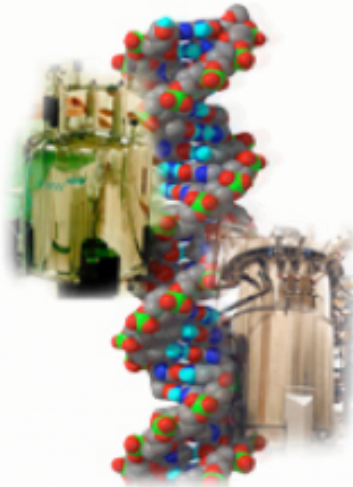
DNA : Protein  
Core facility



Molecular Interactions  
Core facility

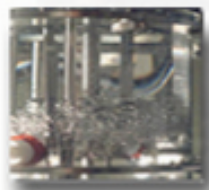


Mass Spectrometry  
Core facility



Combinatorial  
Core facility

## Advanced Biotechnology CORE Facilities



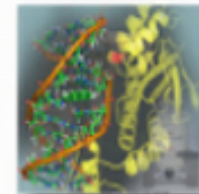
Fermentation  
Core facility



Bioimaging  
Core facility



Confocal  
Core facility



Structural  
Core facility

Welcome !!

欢迎





**Goals of the Course:** The primary objective of this course is to introduce students to the relative, sophisticated high technology that is / will be available in the ABCore Facilities at Georgia State University. To achieve this goal students will be given live instruction and directed VIDEO experience in performing “wet lab” experiments, purification of RNA and DNA as well as how to analyze and utilize these and other cellular attributes using a variety of instruments. Informational lectures will be provided, where needed, to cover the scientific background behind the techniques as well as other more advanced techniques. Students will be expected to maintain a legible and well organized notebook of the activities carried out throughout the course.

<b>ATTENDANCE</b>	<b>(70%)</b>
<b>ON-LINE QUESTIONS / PPT</b>	<b>(30%)</b>

**Attendance:** Students will be expected to attend ALL classes at the scheduled time. Owing to the nature of the course some web based activities may extend beyond the assigned time. On these occasions, students who need to leave early should email/tx the instructor **in advance** . Failure to attend two or more class days without this agreement will forfeit ALL points given for attendance.

**Ethical considerations:** Unless otherwise stated, all examination answers that are submitted for evaluation are considered to be the sole property and ideas of each individual student. Any student(s) caught copying or cheating will automatically receive **zero credit** for that particular submission, and possibly be subject to further disciplinary action.

Be aware of **Plagiarism**. All students should be aware of the academic honesty policy (<http://deanofstudents.gsu.edu/faculty-staff-resources/academic-honesty/>).

**BIOL4905: Summer Semester '22 -ON LINE: -CRN 55906**

#	Date	Time	Lecture Description
	July 5th	8:30 am -10:00 pm	Welcome Reception
	July 6th	----	----
1	July 7th	8:30 am - 11:30 am	Introduction -ABCORE -Training
2	July 8th	8:30 am - 11:30 am	Plasmid DNA Preparation
	July 9th	--	WEEKEND
	July 10th	--	WEEKEND
3	July 11th	8:30 am - 11:30 am	Proteomics I
4	July 12th	8:30 am - 11:30 am	Proteomics II
5	July 13th	8:30 am - 11:30 am	Proteomics III
6	July 14th	8:30 am - 11:30 am	RNA Preparation
7	July 15th	8:30 am - 11:30 am	---
	July 16th	--	WEEKEND
	July 17th	--	WEEKEND
8	July 18th	8:30 am - 11:30 am	qPCR & Robot
9	July 19th	8:30 am - 11:30 am	DNA Sequence Analysis
	July 20th	-	MINI BREAK
10	July 21st	8:30 am - 11:30 am	Automated Microscopy / AFM
11	July 22nd	8:30 am - 11:30 am	Next Gen. DNA Sequencing
	July 23rd	--	WEEKEND
	July 24th	--	WEEKEND
12	July 25th	8:30 am - 11:30 am	Microarray I
13	July 26th	8:30 am - 11:30 am	Microarray II
14	July 27th	8:30 am - 11:30 am	Nanostring
15	July 28th	8:30 am - 11:30 am	Flow Cytometry
-	July 29th	8:30 am - 11:30 am	- FINAL-
	July 30th - Aug 3rd	--	RECREATION



## SUMMER INSTITUTE - ONLINE MODALITY CALENDAR 2022

SUN	MON	TUE	WED	THU	FRI	SAT
June 26	27	28	29	30	31	July 02
	9:00-10:00am Virtual Program Orientation for Summer Institute Online Modality					
July 03	04	05	06	07	08	09
	Holiday (Independence Day)	8:30-10:00am -Welcome Reception and Buddy Meet & Greet Event	Free Day	Classes begin! 8:30-11am: BIOL4905 INTRODUCTION 8-10:20pm: Afternoon course	8:30-11am: BIOL4905 DNA PREPARATION 8-10:20pm: Afternoon course	
10	11	12	13	14	15	16
	8:30-11am:BIOL4905 PROTEOMICS I 8-10:20pm: Afternoon course	8:30-11am:BIOL4905 PROTEOMICS II 8-10:20pm: Afternoon course	8:30-11am:BIOL4905 PROTEOMICS III 8-10:20pm: Afternoon course	8:30-11am: BIOL4905 RNA PREPARATION 8-10:20pm: Afternoon course	Virtual Independence Day Activity	
17	18	19	20	21	22	23
	8:30-11am:BIOL4905 qPCR / ROBOTS 8-10:20pm: Afternoon course	8:30-11am:BIOL4905 DNA Sequence Analysis 8-10:20pm: Afternoon course	Midterm Break	8:30-11am:BIOL4905 Next Gen. Sequencing 8-10:20pm: Afternoon course	8:30-11am:BIOL4905 Automated Microscopy /AFM	
24	25	26	27	28	29	30
	8:30-11am:BIOL4905 Microarray I 8-10:20pm: Afternoon course	8:30-11am:BIOL4905 Microarray II 8-10:20pm: Afternoon course	8:30-11am:BIOL4905 Nanostring 8-10:20pm: Afternoon course	8:30-11am:BIOL4905 Flow Cytometry 8-10:20pm: Afternoon course	FINALS	
31	August 01	02	03			
	9:00-10:00am: Closing Reception		Grades available in PAWS			

Legend:  
Orange: Courses      Blue: Activities



NMR  
Core facility



DNA : Protein  
Core facility



Molecular Interactions  
Core facility



Mass Spectrometry  
Core facility



Combinatorial  
Core facility

## Advanced Biotechnology CORE Facilities

Director: Houghton

Cell, Protein & DNA  
Core facility



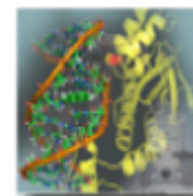
Fermentation  
Core facility



Bioimaging  
Core facility



Confocal  
Core facility



Structural  
Core facility



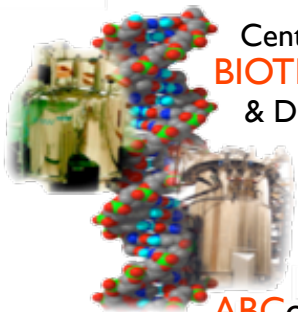


**Petit Life Science Center**



**Natural Science Center**





Center for  
**BIOTECHNOLOGY**  
& DRUG DESIGN

ABC Core Facilities

R  
P

## Genomics Core Facility



**WORKFLOW**  
MICROARRAY CBY/BCIP  
EXPRESSION  
TARGET PREPARATION  
TARGET HYBRIDIZATION  
FLUIDICS  
PROBE ARRAY WASH  
AND STAIN  
PROBE ARRAY SCAN  
DATA ANALYSIS

3. Fluidics station 450: wash & stain chip



4. Affymetrix GeneChip Scanner 7G



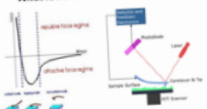
Report

Gene	Expression
Gene 1	1.2
Gene 2	0.8
Gene 3	1.5
Gene 4	0.9
Gene 5	1.1
Gene 6	0.7
Gene 7	1.3
Gene 8	0.6
Gene 9	1.4
Gene 10	0.5

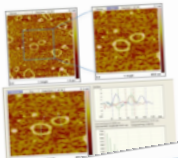
## Atomic Force Microscopy Core Facility



What is AFM and how does it work?



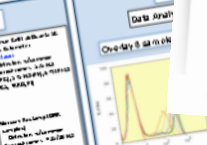
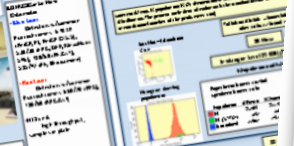
**Applications in biology**  
AFM can be used in analyzing protein structure after protein is bound to a flat surface. The plain thin, amorphous polyethylene glycol (PEG) mica, glow discharged mica, or highly Oriented Pyrolytic Graphite (HOPG). The scanning can either in air, after sample is dried, or in liquid, in the buffer or medium.



## Flow Cytometry

Debby Walthall (dwalthall@gsu.edu)

**Instrument/Services Analysis Software**  
BD FACSCanto and FACSARIA  
Diva Software



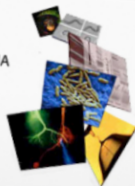
Saturday, November 28, 2009



Welcome  
Cell, Protein and DNA Core Facilities

### About

Welcome to GSU  
Cell - Protein - DNA  
Core Facilities



### Resources

Find answers  
to your  
questions  
here



### Services

Learn about  
some of the  
services that  
the core  
facility offers



### Instrumentation

Find out what equipment  
and facilities are  
available for  
your use



### Contacts

**John E. Houghton**  
Director of Core Facilities  
Tel: 404-413-5390 | Email: [jhoughton@gsu.edu](mailto:jhoughton@gsu.edu)

**Debby Walthall**  
Research Scientist  
Tel: 404-413-5363 | Email: [dwalthall@gsu.edu](mailto:dwalthall@gsu.edu)

### What's New?

Find out what's new in  
the Cell - Protein -  
DNA Core Facilities



<http://biotech.gsu.edu/core>

http://biotech.gsu.edu/core



SNP 0 Position

C / A

TAGCCATCGGTA N GTA C TCAATGATCAGCT

ATCGGTAGCCAT G CAT G AGTTACTA PM Allele A  
 ATCGGTAGCCAT C CAT G AGTTACTA MM Allele A  
 ATCGGTAGCCAT T CAT G AGTTACTA PM Allele A  
 ATCGGTAGCCAT A CAT G AGTTACTA MM Allele A

*Why not stop by for a look?*

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### Services

Learn about  
some of the  
services that  
the core  
facility  
offers



### Instrumentation

Find out what equipment  
and facilities are  
available for  
your use



### What's New?

Find out what's new in  
the Cell - Protein -  
DNA Core Facilities



## **(Level I) General Access Instrumentation**

Effectively low maintenance, “relatively” generic, but technically proficient instrumentation that is normally operated by graduate researchers and other research personnel on a daily, routine basis .

## **(Level II) Specialized Instrumentation**

High maintenance, sophisticated instrumentation, available for operation by general graduate researchers and technical support staff **after** they have received specific and thorough training in the appropriate use of the technology.

## **(Level III) Centralized Instrumentation / Service**

- (a) DNA Purification & Sequencing (mapping)
- (b) Genomics / Proteomics
- (c) Atomic Force Microscopy
- (d) Mass Spectrometry/NMR
- (e) Cell Analysis / Sorting

High maintenance, highly sophisticated instrumentation, operated by qualified, technical personnel.



# **Level I**

**General Access Equipment  
(relatively low maintenance)**





from permeablizing cell membranes -  
**Gene Pulse Electroporation System (BioRad).....**

..to Cell breaking -**Sonifier 450**  
(Branson -top right)  
and  
French Press (Aminco)





**Lyophilizer (LabConco)**



**Liquid scintillation Counter LS6500 (Beckman)**

**Thermocyclers (Eppendorff)**



**Automated Protein Separation**



**AKTA Explorers / Purifiers (GE)**





Victor X3 plate-reader  
(Perkin Elmer)



Spectramax iD5



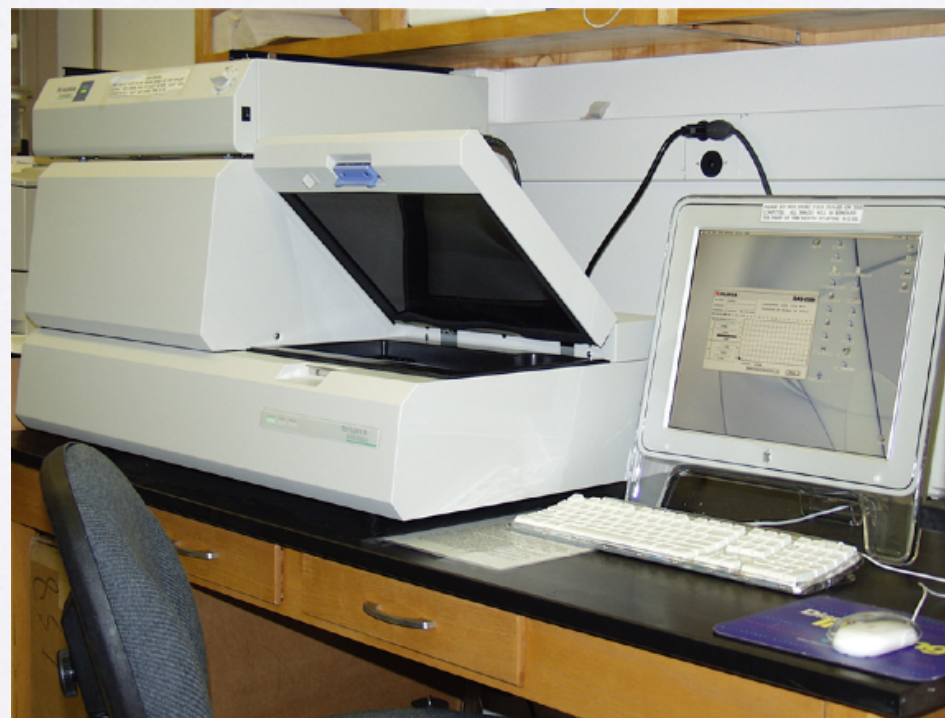
EnSpire Multimode Plate Reader

## Assorted Micro-Plate Readers





**Omega UV-Imaging System**  
(Ultra-Lum)



**Phosphoimager**  
(Fuji) Model **Bas 2500**



**UVP GelDoc**  
Gel documentation imaging system



## Fluorescent Bioimaging System

**Typhoon 7000R Ge (Cytiva)**

## Chemi-Luminescence Image Analyzer (Fuji) Model LAS-4000-mini







**Optima MAX-XP**

**Table-top Optima model-**

**Assorted Centrifuges  
and  
Ultracentrifuges-**



**to the Analytical XLA-X1**





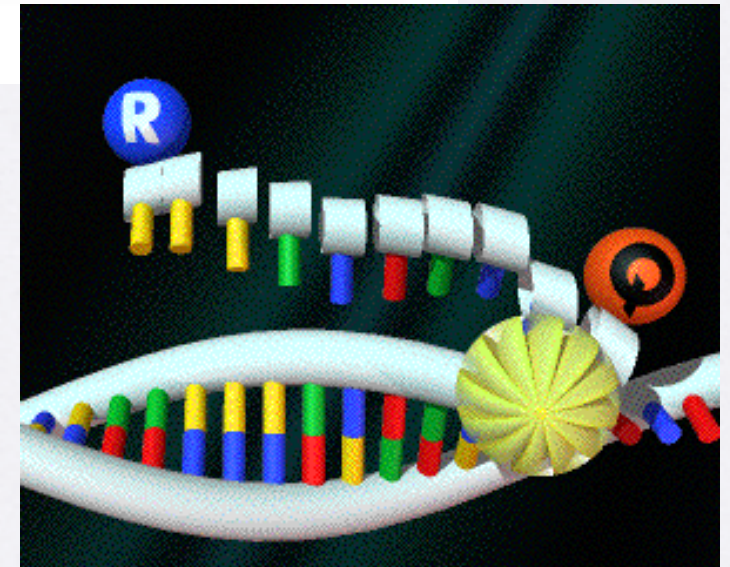
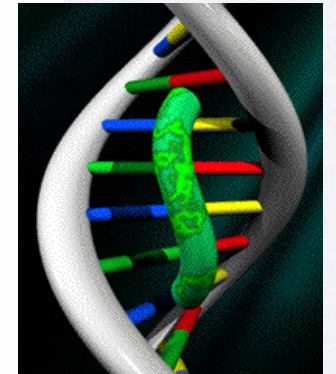
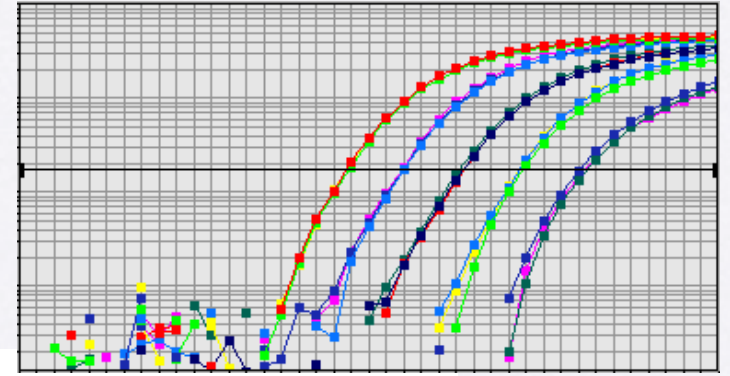
## **Level II**

**Specialized Equipment  
(high maintenance)**

## **Level III**

**Centralized Equipment. Service  
and Molecular Analysis**



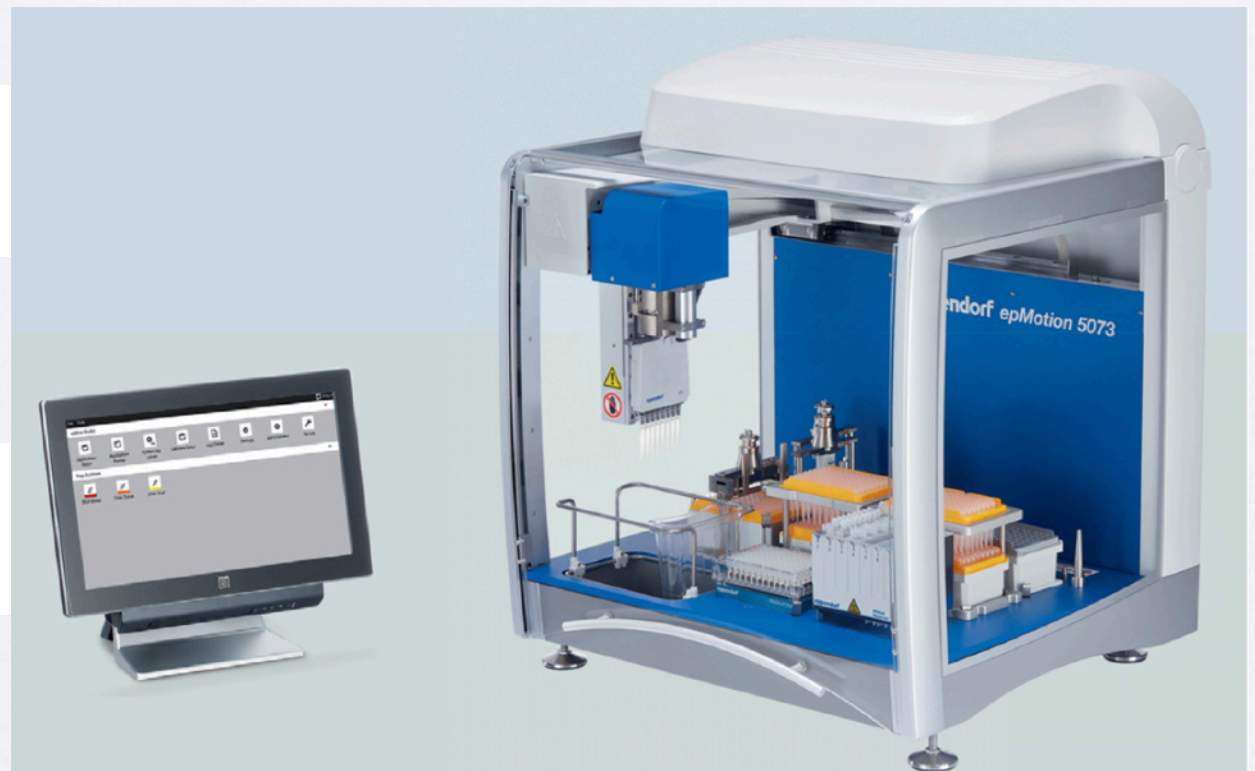


**Quantitative PCR (qPCR)**  
**Amplification of DNA/RNA**  
**Models ABI/Life 7500 FAST,**



**Robotic  
Workstation**

**Integra Assist Plus  
(Integra)**



**Epimotion 5073  
(Eppendorf)**



SNP 0 Position

C / A

TAGCCATCGGTA N GTA C TCAATGATCAGCT

ATCGGTAGCCAT	G	CAT G	AGTTACTA	PM Allele A
ATCGGTAGCCAT	C	CAT G	AGTTACTA	MM Allele A
ATCGGTAGCCAT	T	CAT G	AGTTACTA	PM Allele B
ATCGGTAGCCAT	A	CAT G	AGTTACTA	MM Allele B

*Why not stop by for a look?*

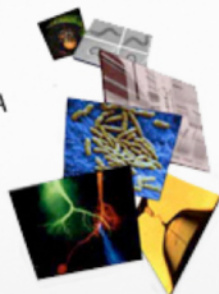
## Welcome

Cell, Protein and DNA Core Facilities



### About

Welcome to GSU  
Cell - Protein - DNA  
Core Facilities



### Resources

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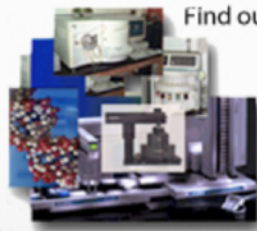
### Services

Learn about  
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services that  
the core  
facility  
offers



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available for  
your use



### What's New?

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the Cell - Protein -  
DNA Core Facilities





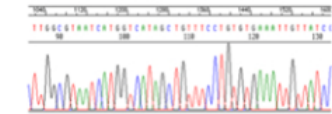
# GSU Biology Core Facility

## Supporting Life Sciences at GSU

[http://biotech.gsu.edu/core\\_facility/index.html](http://biotech.gsu.edu/core_facility/index.html)

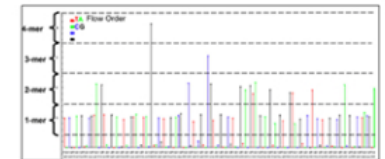


### DNA Sequence Analysis: Profiling DNA



Sanger Sequencing –  
>800 base pairs/run

High Throughput Genomic Sequencing –  
100,000 base pairs/run



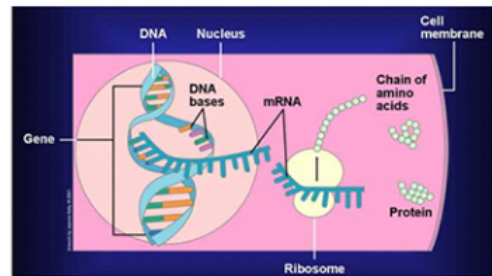
### RNA Expression

### Microarray: Analysis Profiling mRNA

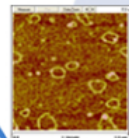


Colour of pin-point dots demonstrates the presence / absence of gene sequences

### DNA Replication



### Atomic Force Microscopy Imaging at the Ångström level



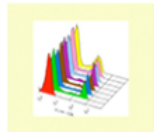
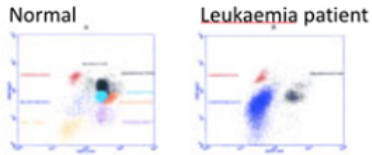
Protein structure analysis

### Protein Expression

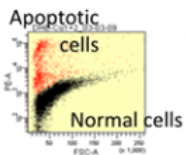
### Proteomics Profiling Proteins

2D Protein gel  
Protein separation using Electric charge and molecular weight

### Flow Cytometry Profiling Cells



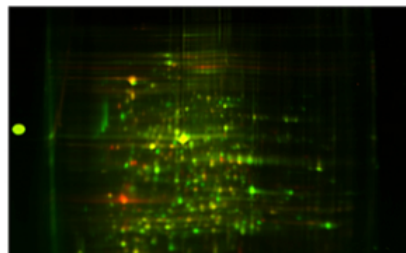
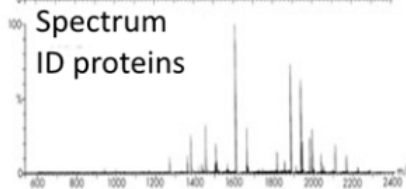
See effects of different drugs on Cell cycle



Apoptosis -programmed cell death

### Cellular Functions

### Mass Spectrometry







# GSU Biology Core Facility

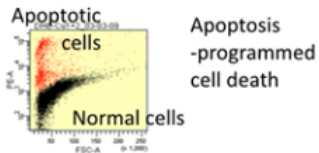
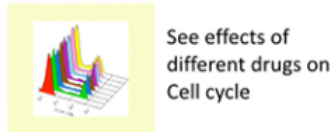
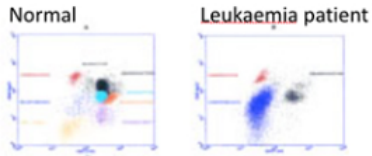
## Supporting Life Sciences at GSU

[http://biotech.gsu.edu/core\\_facility/index.html](http://biotech.gsu.edu/core_facility/index.html)



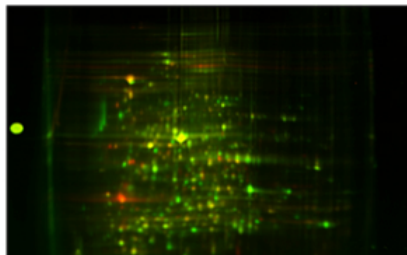
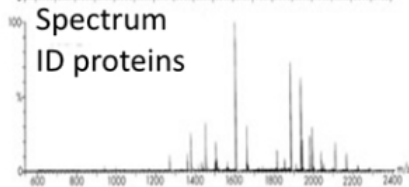
### Flow Cytometry

#### Profiling Cells



### Cellular Functions

### Mass Spectrometry



### Proteomics

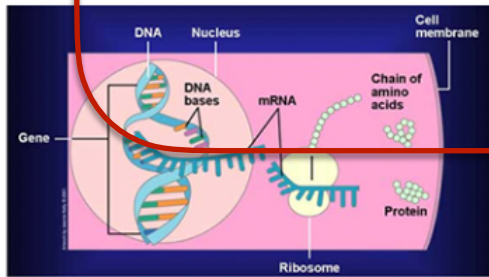
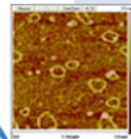
#### Profiling Proteins

2D Protein gel  
Protein separation using Electric charge and molecular weight

### Atomic Force Microscopy

#### Imaging at the Ångström level

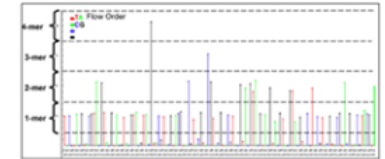
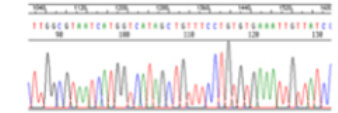
Protein structure analysis



Sanger Sequencing –  
>800 base pairs/run

High Throughput Genomic Sequencing –  
100,000 base pairs/run

### DNA Sequence Analysis: Profiling DNA



### RNA Expression

### Microarray: Analysis Profiling mRNA



Colour of pin-point dots demonstrates the presence / absence of gene sequences



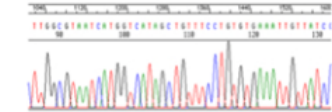
# GSU Biology Core Facility

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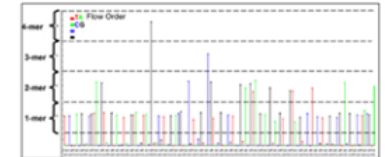


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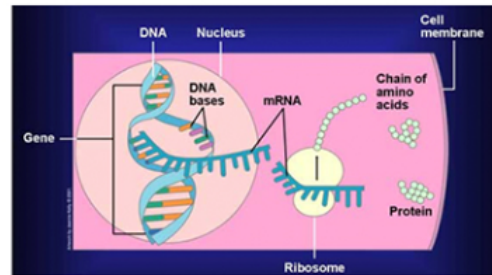
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### Microarray: Analysis Profiling mRNA

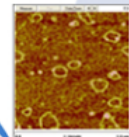


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### DNA Replication



### Atomic Force Microscopy Imaging at the Ångström level



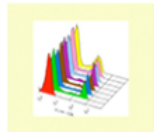
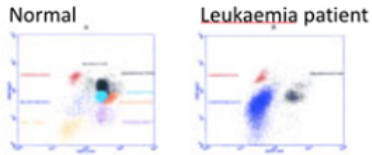
Protein structure analysis

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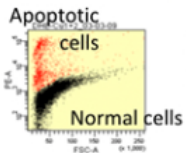
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2D Protein gel  
Protein separation using Electric charge and molecular weight

### Flow Cytometry Profiling Cells



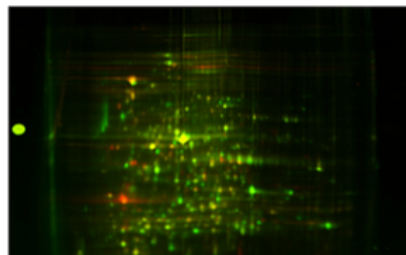
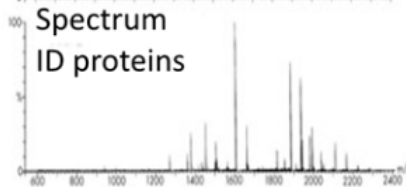
See effects of different drugs on Cell cycle



Apoptosis - programmed cell death

### Cellular Functions

### Mass Spectrometry







# GSU Biology Core Facility

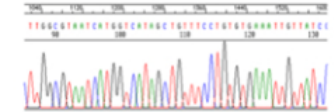
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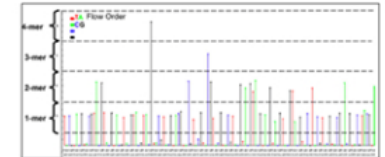


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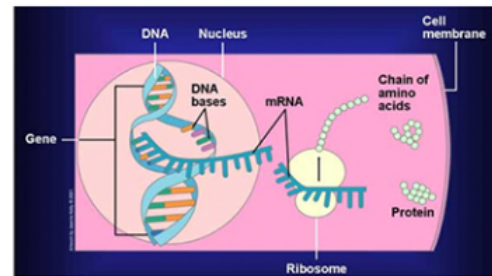


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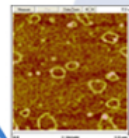


Colour of pin-point dots demonstrates the presence / absence of gene sequences

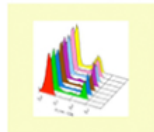
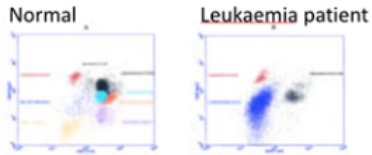


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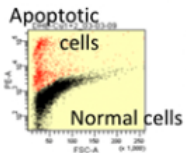
Protein structure analysis



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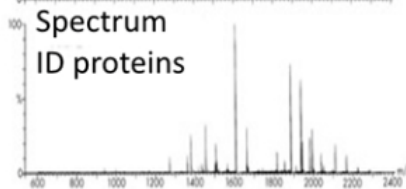
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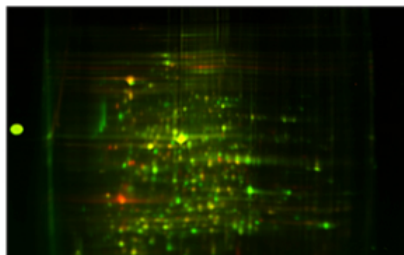


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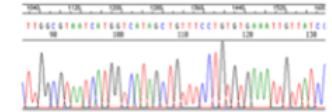
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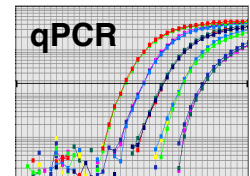
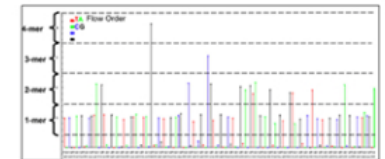


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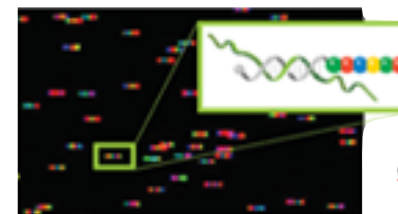
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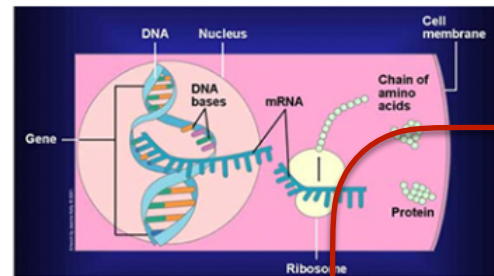
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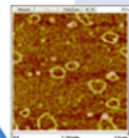
### Nanosttring

### RNA Expression

### DNA Replication



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Protein structure analysis

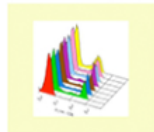
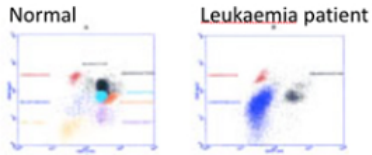
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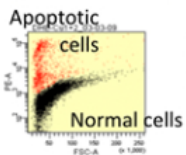
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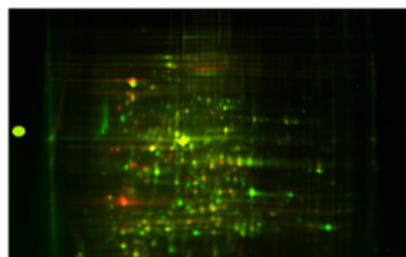
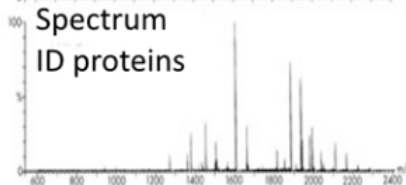
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Apoptosis - programmed cell death

### Cellular Functions

### Mass Spectrometry







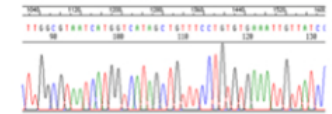
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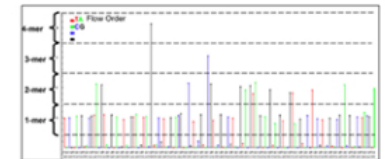


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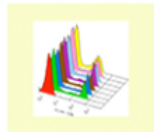
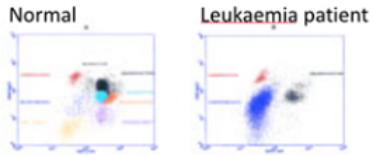


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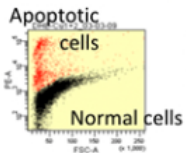
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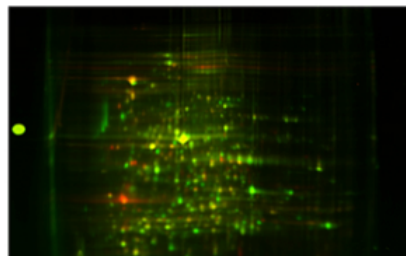
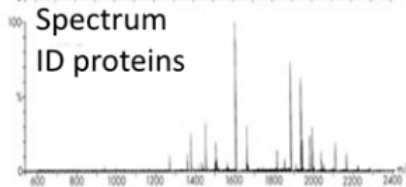
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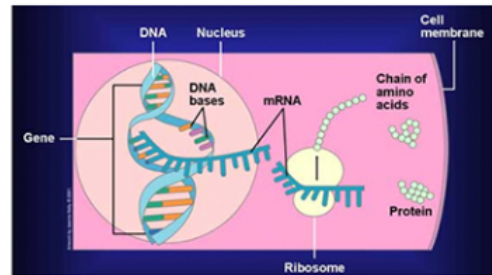
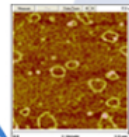
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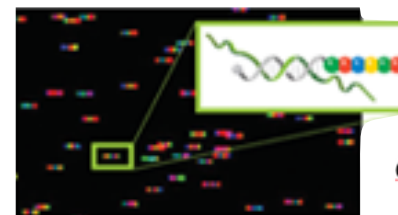
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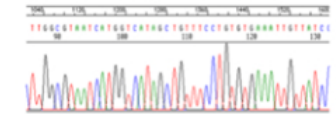
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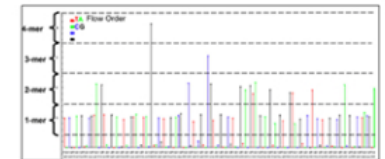


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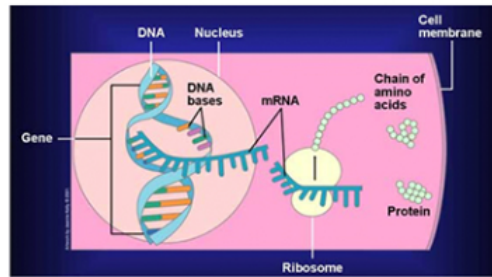
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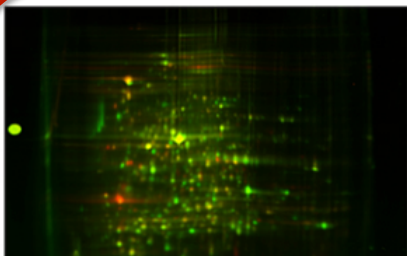
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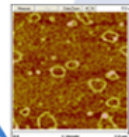
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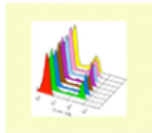
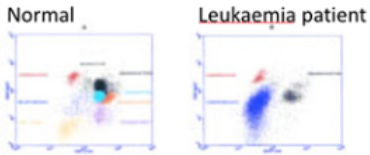


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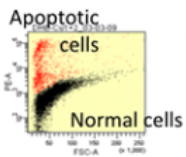


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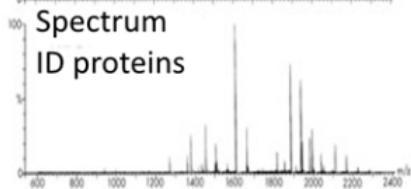
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# GSU Biology Core Facility

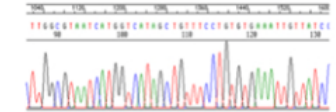
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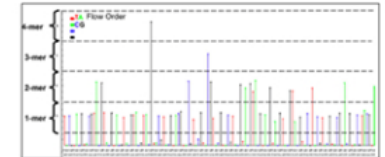


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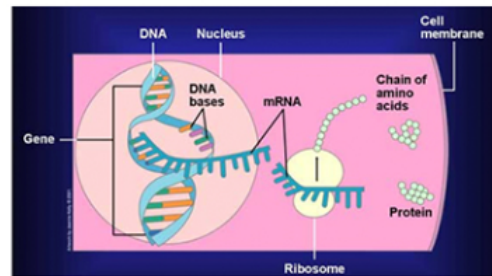


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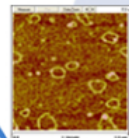


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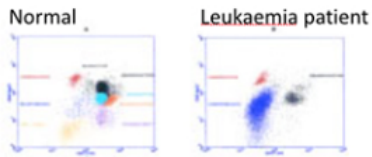
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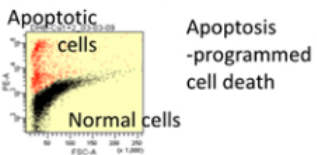


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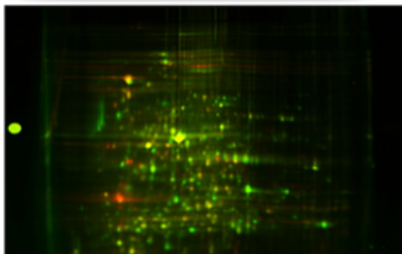
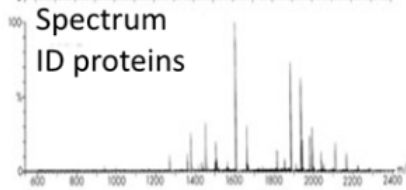
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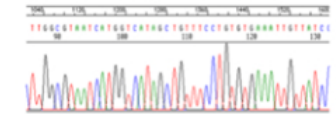
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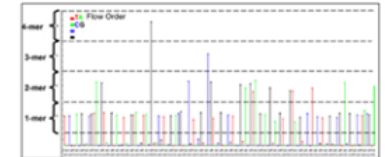


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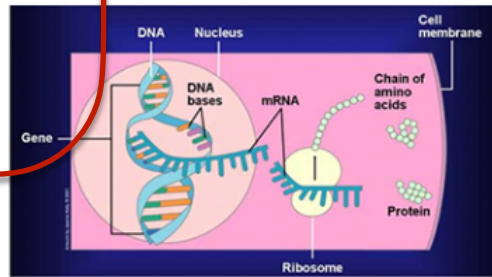
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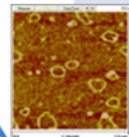
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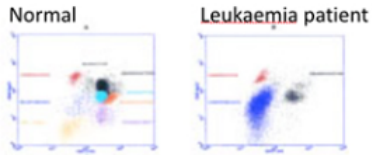


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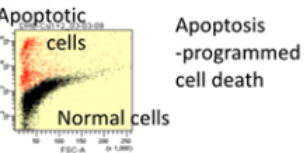
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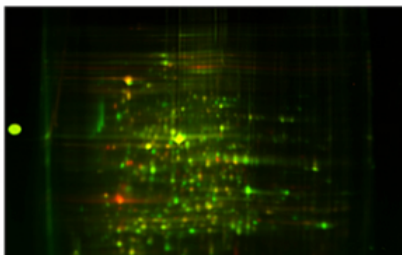
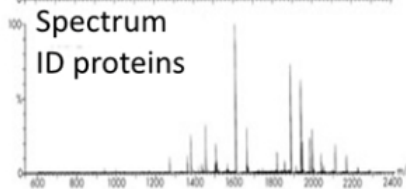


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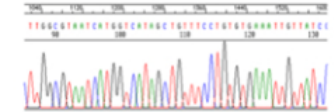
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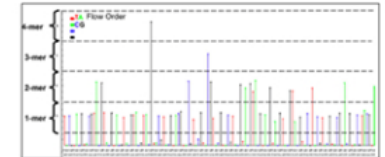


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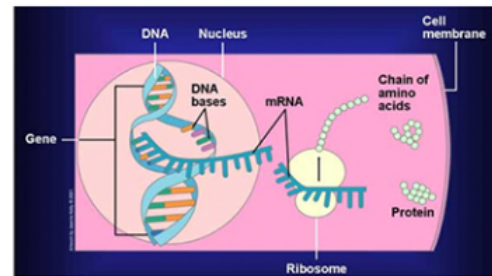


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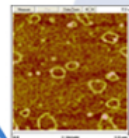


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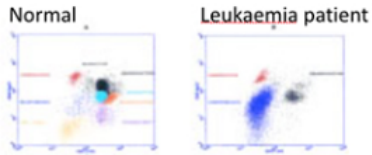


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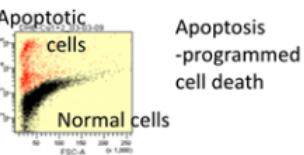
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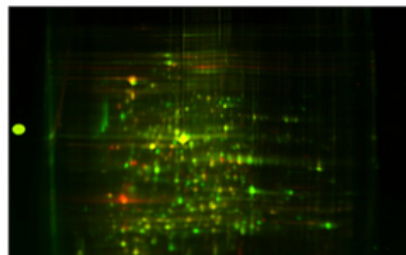
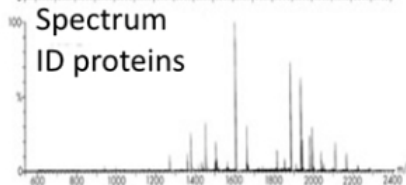


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## SUMMER INSTITUTE - ONLINE MODALITY CALENDAR 2022

SUN	MON	TUE	WED	THU	FRI	SAT
June 26	27	28	29	30	31	July 02
	9:00-10:00am Virtual Program Orientation for Summer Institute Online Modality					
July 03	04	05	06	07	08	09
	Holiday (Independence Day)	8:30-10:00am -Welcome Reception and Buddy Meet & Greet Event	Free Day	Classes begin! 8:30-11am: BIOL4905 INTRODUCTION 8-10:20pm: Afternoon course	8:30-11am: BIOL4905 DNA PREPARATION 8-10:20pm: Afternoon course	
10	11	12	13	14	15	16
	8:30-11am:BIOL4905 PROTEOMICS I 8-10:20pm: Afternoon course	8:30-11am:BIOL4905 PROTEOMICS II 8-10:20pm: Afternoon course	8:30-11am:BIOL4905 PROTEOMICS III 8-10:20pm: Afternoon course	8:30-11am: BIOL4905 RNA PREPARATION 8-10:20pm: Afternoon course	Virtual Independence Day Activity	
17	18	19	20	21	22	23
	8:30-11am:BIOL4905 qPCR / ROBOTS 8-10:20pm: Afternoon course	8:30-11am:BIOL4905 DNA Sequence Analysis 8-10:20pm: Afternoon course	Midterm Break	8:30-11am:BIOL4905 Next Gen. Sequencing 8-10:20pm: Afternoon course	8:30-11am:BIOL4905 Automated Microscopy /AFM	
24	25	26	27	28	29	30
	8:30-11am:BIOL4905 Microarray I 8-10:20pm: Afternoon course	8:30-11am:BIOL4905 Microarray II 8-10:20pm: Afternoon course	8:30-11am:BIOL4905 Nanostring 8-10:20pm: Afternoon course	8:30-11am:BIOL4905 Flow Cytometry 8-10:20pm: Afternoon course	FINALS	
31	August 01	02	03			
	9:00-10:00am: Closing Reception		Grades available in PAWS			

Legend:  
Orange: Courses      Blue: Activities



ABC Core  
**F**acilities



Jesse

Chip

Sandy

Kyu

Sonja

## Personnel

Cell, Protein and DNA Core Facilities



**John E. Houghton, PhD**  
Director of ABCore Facilities  
Location: 520 PSC  
Tel: 404-413-5390 (O)  
Email: [jhoughton@gsu.edu](mailto:jhoughton@gsu.edu)



**"Sandy" Ying Sin Hsieh, PhD**  
Research Scientist II  
Rm: 519 / 535 / 659B PSC  
Tel: 404-413-5363 (O)  
Email: [yhsieh3@gsu.edu](mailto:yhsieh3@gsu.edu)  
Instrument Training, Flow Cytometry,  
Genomics, qPCR, plate readers



**Sonja R. Young**  
Laboratory Supervisor  
Rm: 519 / 661 PSC  
Tel: 404-413-5363 (O)  
Email: [sstovall@gsu.edu](mailto:sstovall@gsu.edu)  
Genomics, Instrument Maintenance,  
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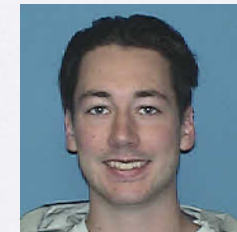
**Hyuk-Kyu Seoh, PhD**  
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Rm: 521 / 537 PSC  
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Email: [hseoh@gsu.edu](mailto:hseoh@gsu.edu)  
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purification/separation Sequence  
Analysis



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DNA/RNA purification/troubleshooting  
AFM Service



**"Chip" Foster**  
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Autoclaves, Instrument  
Maintenance



**Shawn Canavan**  
Teaching Asst.





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"How to" GUIDES



Other, Related  
ABCore Facilities

## CoreLab Technology Training & Access

Cell, Protein and DNA Core Facilities

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Technology Training & Access **COVID-19 CHANGES for SPRING '21'**

### Registration

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## ABCORE FACILITIES

Training Videos -some with, some without audio

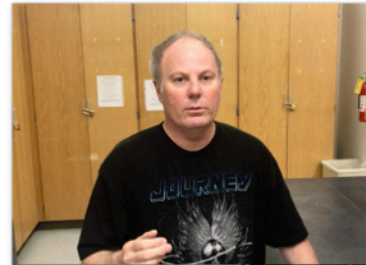
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- High Speed, floor standing Centrifuge

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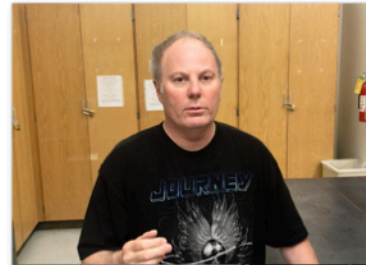
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SNP 0 Position

C / A

TAGCCATCGGTA N GTA C TCAATGATCAGCT

ATCGGTAGCCAT G CAT G AGTTACTA PM Allele A  
 ATCGGTAGCCAT C CAT G AGTTACTA MM Allele A  
 ATCGGTAGCCAT T CAT G AGTTACTA PM Allele A  
 ATCGGTAGCCAT A CAT G AGTTACTA MM Allele A

*Why not stop by for a look?*

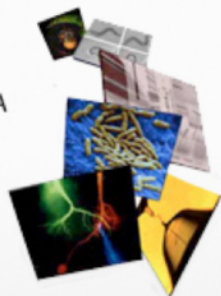
## Welcome

Cell, Protein and DNA Core Facilities



### About

Welcome to GSU  
Cell - Protein - DNA  
Core Facilities



### Resources

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### Contacts

**John E. Houghton**

Director of Core Facilities

Tel: 404-413-5390 | Email: [jhoughton@gsu.edu](mailto:jhoughton@gsu.edu)

**Debby Walthall**

Research Scientist

Tel: 404-413-5363 | Email: [dwalthall@gsu.edu](mailto:dwalthall@gsu.edu)

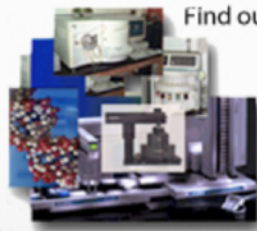
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some of the  
services that  
the core  
facility  
offers



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Find out what equipment  
and facilities are  
available for  
your use



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Legend:

Orange: Courses

Blue: Activities

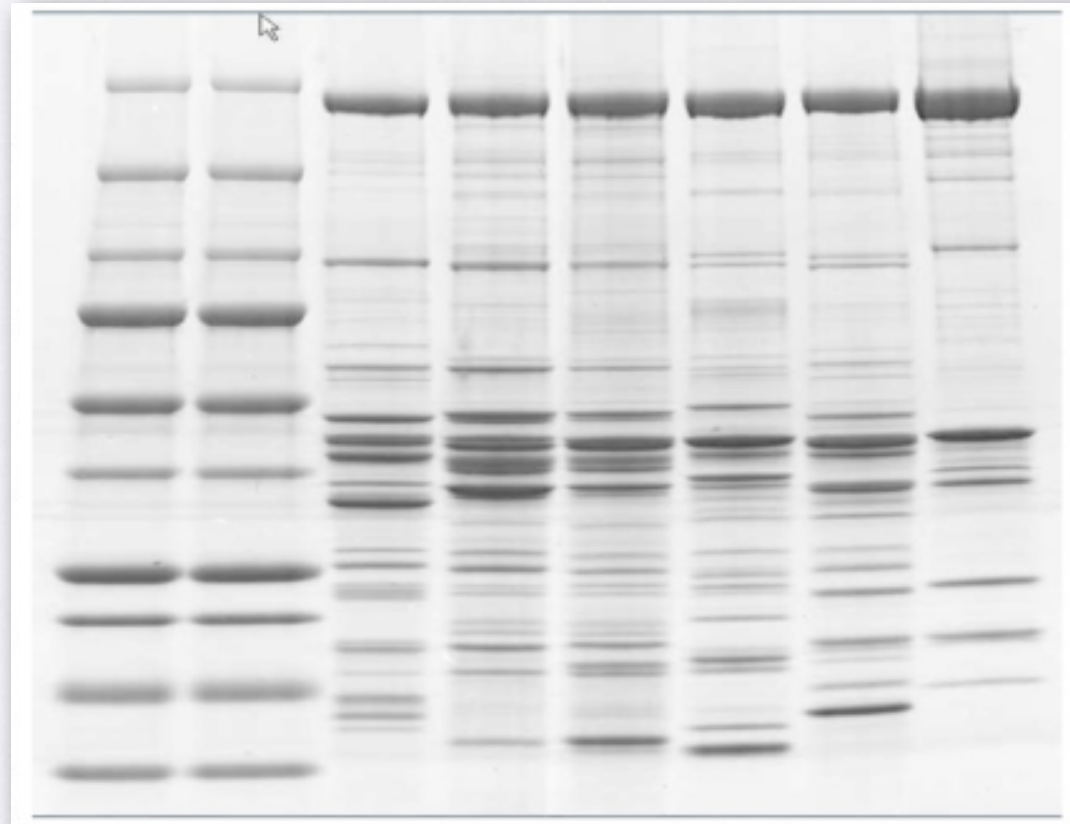


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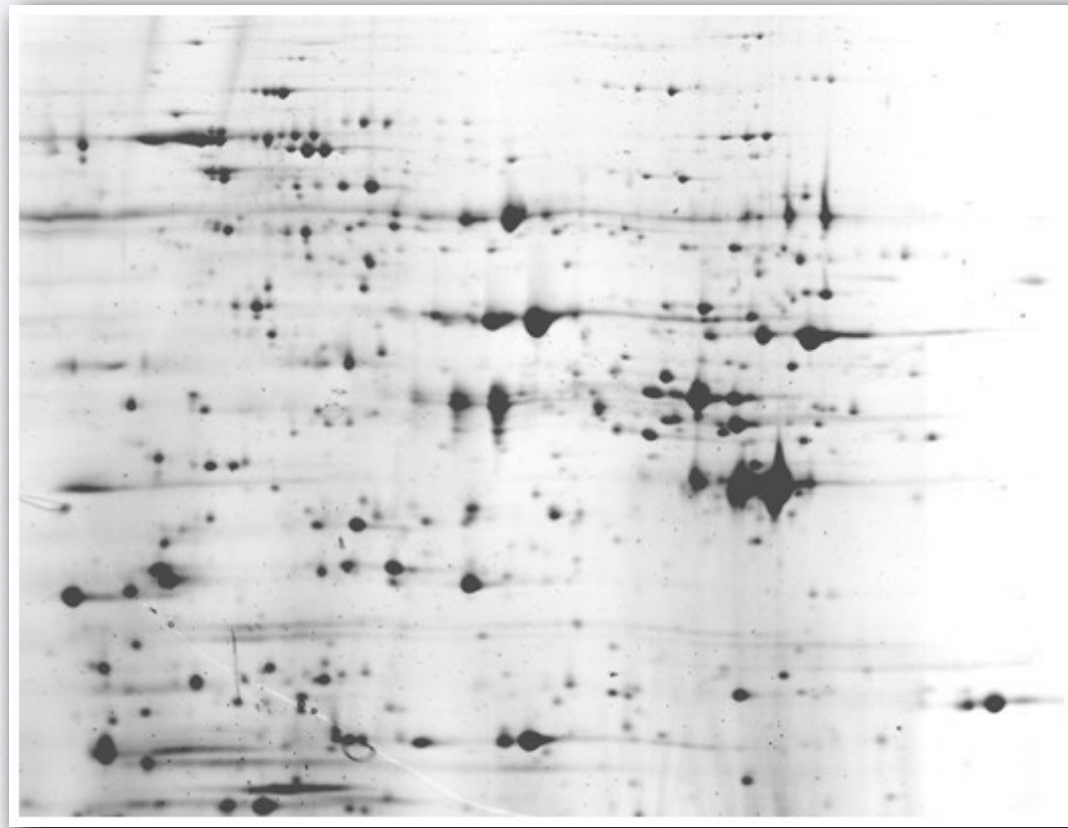
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# Proteomics:





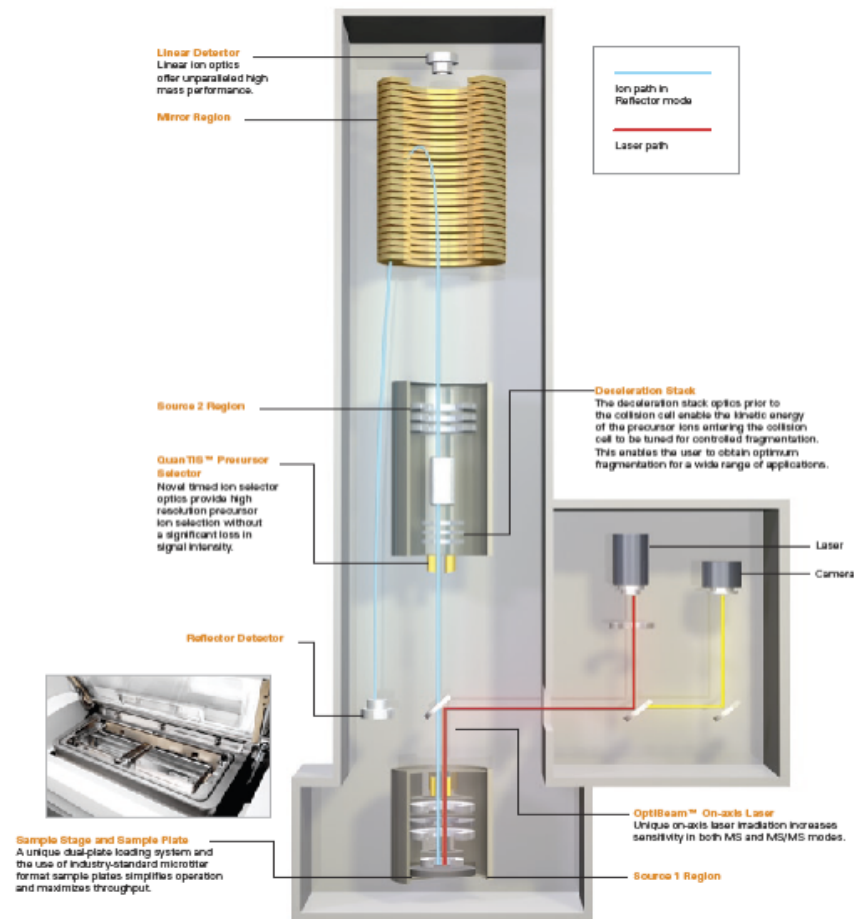
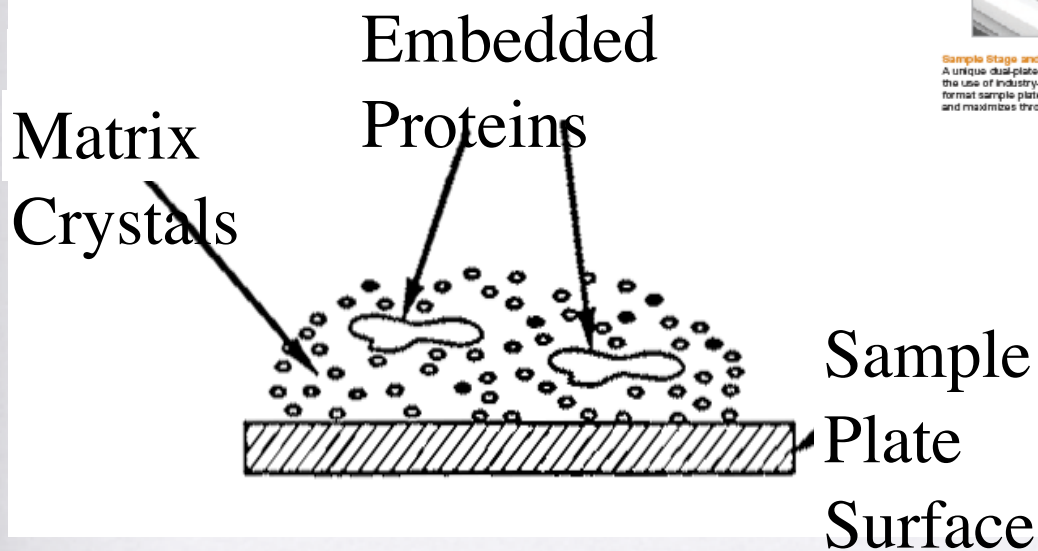
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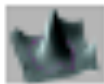


**Ettan II 2D gel Proteomics System complemented by a MALDI TOF/TOF(ABI) Model 4800+**



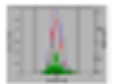


# Proteomics:



## Fluorescence 2-D differential in-gel electrophoresis platform

Amer sham Pharma cia Biotech UK Limited, Amer sham Place, Little Chalfont, Buckinghamshire, HP7 0NA, England



2-D differential in-gel electrophoresis (DIGE) is a powerful tool for identifying and quantifying protein spots, especially in complex samples (Figure 1).

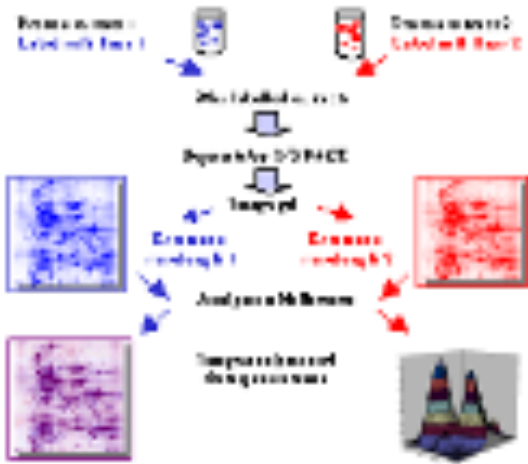


Figure 1: Overview of the 2-D DIGE technology (fluorescence dye and dye with Cy5 or Cy3 labels) (Cy5/Cy3/DAPI).

The major limitation of the DIGE technology is the low resolution of the protein spots. However, the development of a novel 2-D DIGE approach allows protein spots to be analyzed with an accuracy of protein identification that is not possible with the DIGE platform. The DIGE platform allows protein spots to be analyzed with an accuracy of protein identification that is not possible with the DIGE platform.

- The DIGE platform offers several advantages over traditional 2-D PAGE:
- Increased sensitivity and resolution
  - Ability to analyze complex samples
  - High resolution and accuracy
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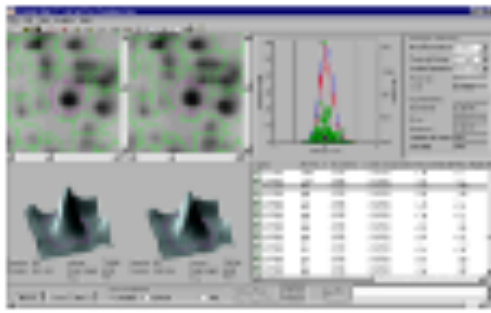


Figure 2: DIGE software interface showing protein spots identified in a sample.

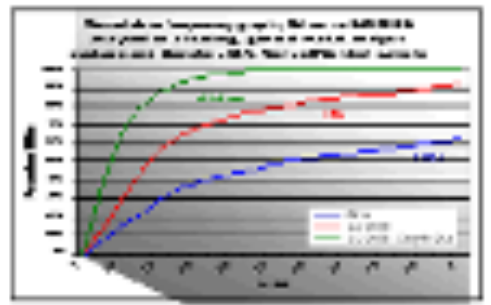


Figure 3: Comparison of protein spot identification rates for different DIGE methods.

To further improve the 2-D DIGE platform, we have developed a novel 2-D DIGE platform that allows protein spots to be analyzed with an accuracy of protein identification that is not possible with the DIGE platform.

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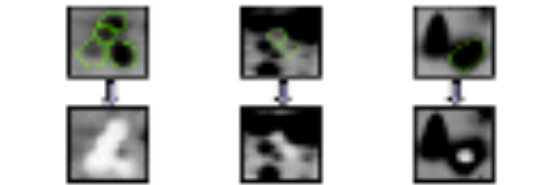


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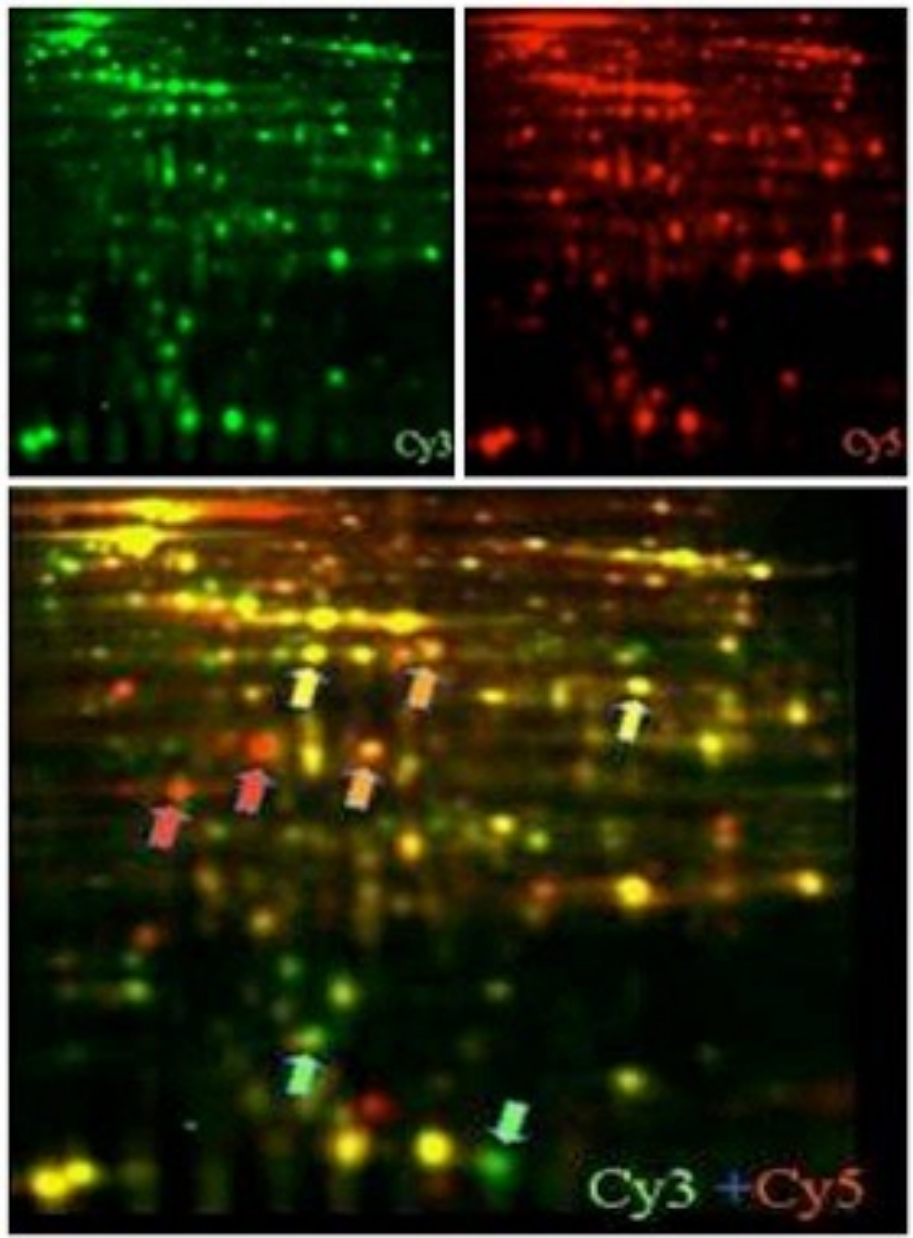
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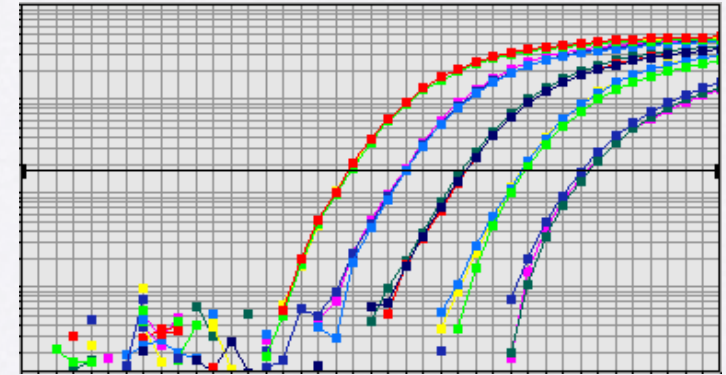
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Legend:  
Orange: Courses     Blue: Activities





**Quantitative PCR (qPCR)**  
**Amplification of DNA/RNA**  
**Models ABI/Life 7500 FAST,**

# SARS-CoV-2 Pandemic

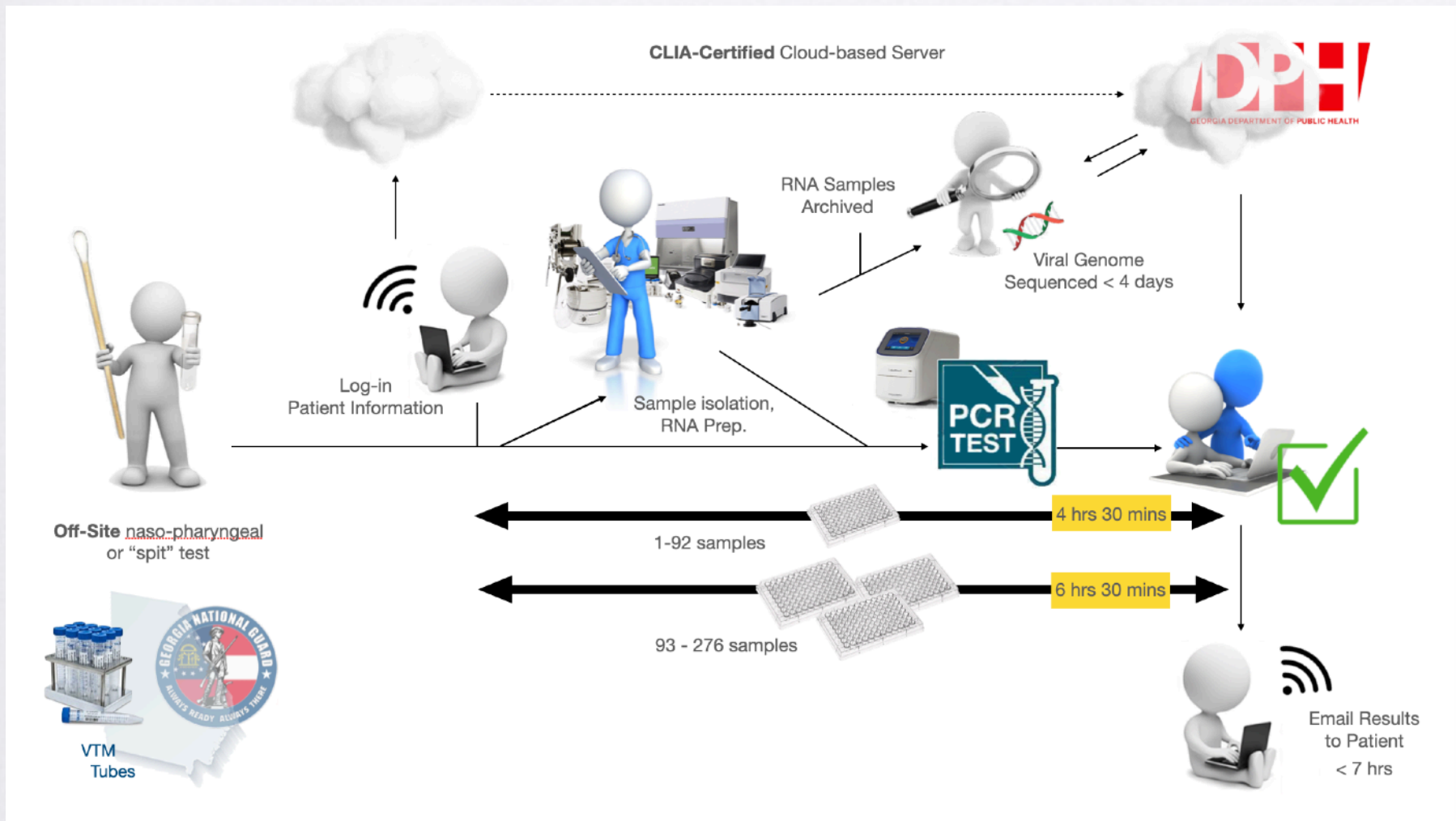
## CLIA-certified COVID-19 PCR testing lab

# 2020-22



**F**acilities

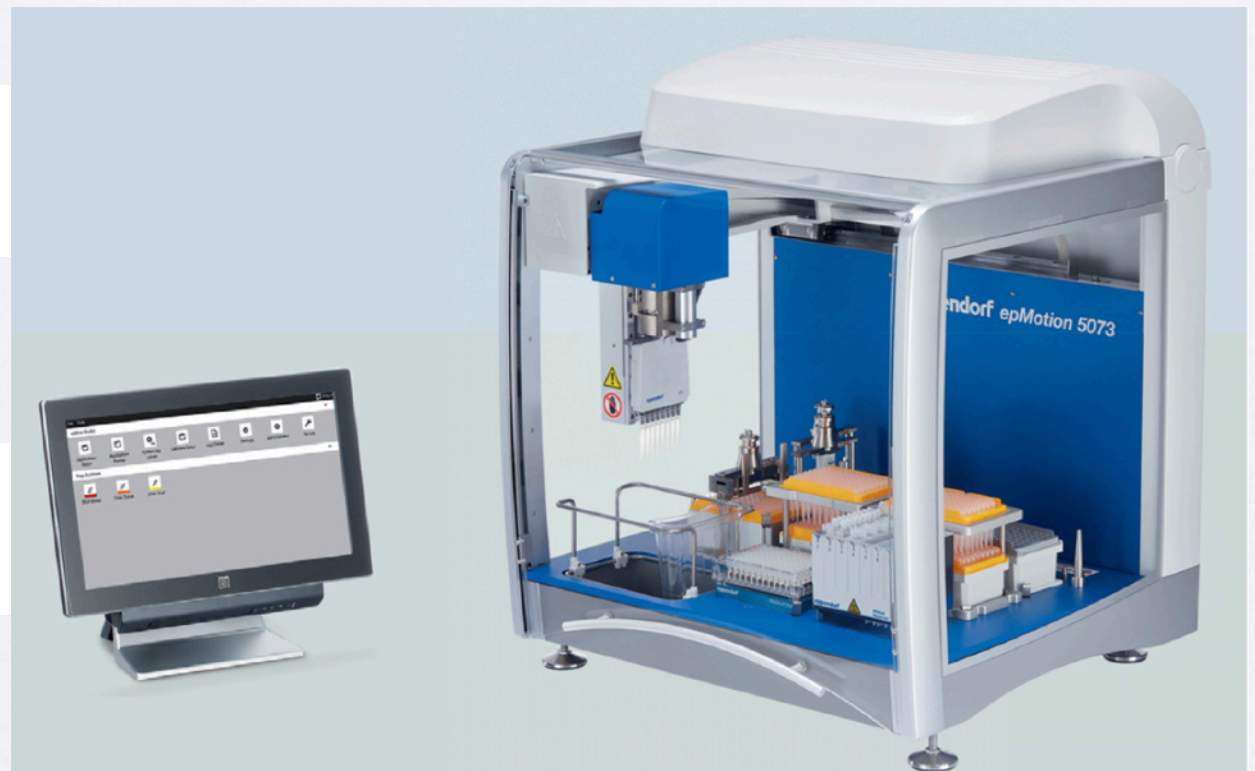






**Robotic  
Workstation**

**Integra Assist Plus  
(Integra)**



**Epimotion 5073  
(Eppendorf)**



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**Capillary DNA Sequencers  
(ABI/Life Technologies) Model 3500xl**



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## Automated Fluorescent Microscopes

**Axiomager II /  
Observer  
(inverted)  
(Zeiss)**

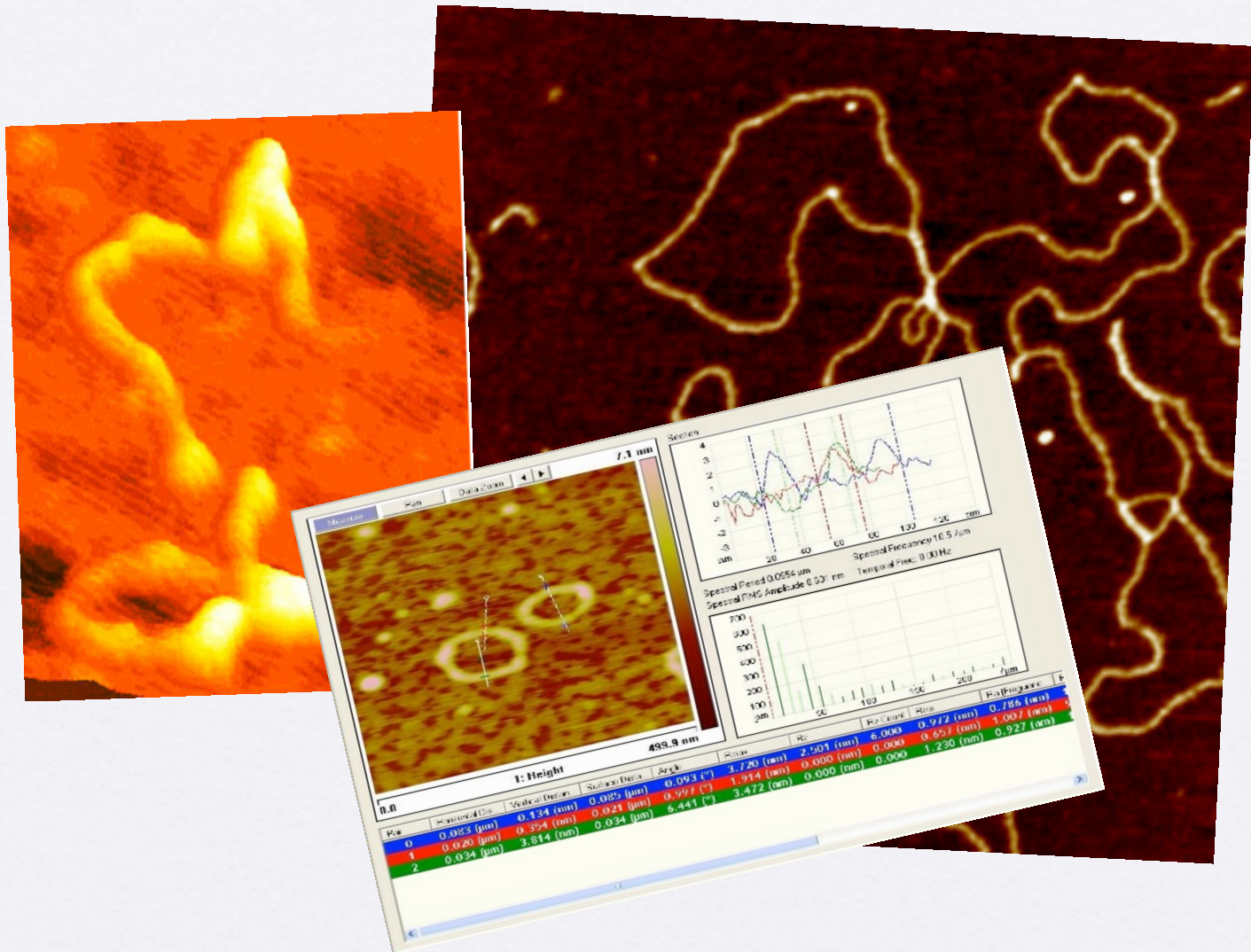


## Automated Fluorescent Digital Microscope

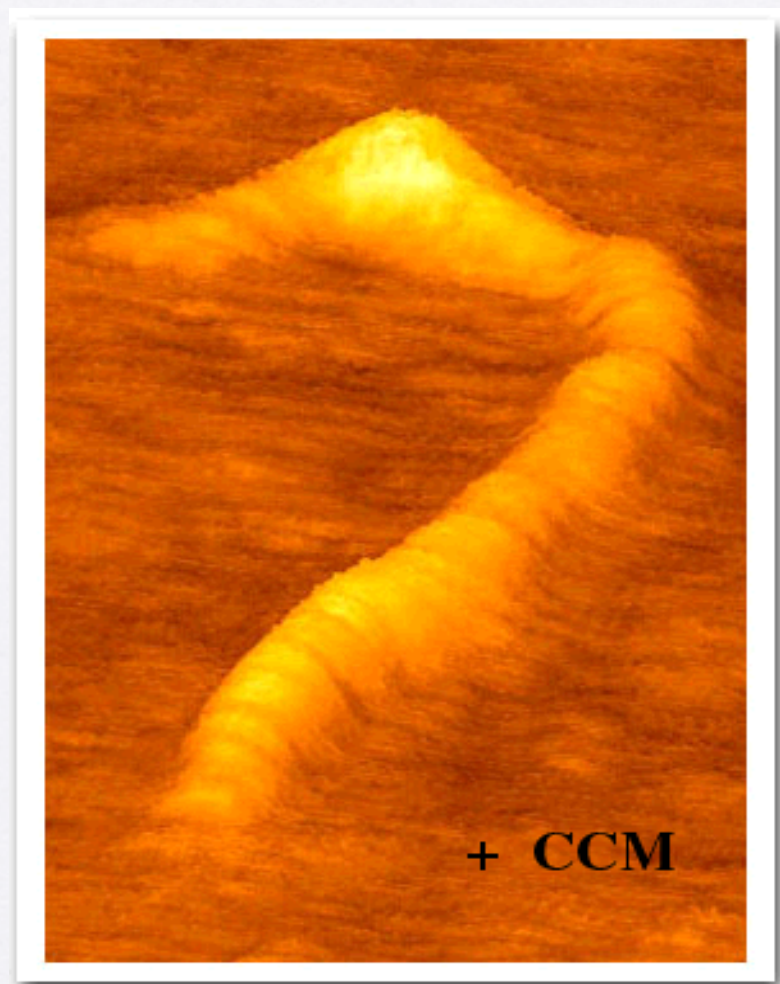
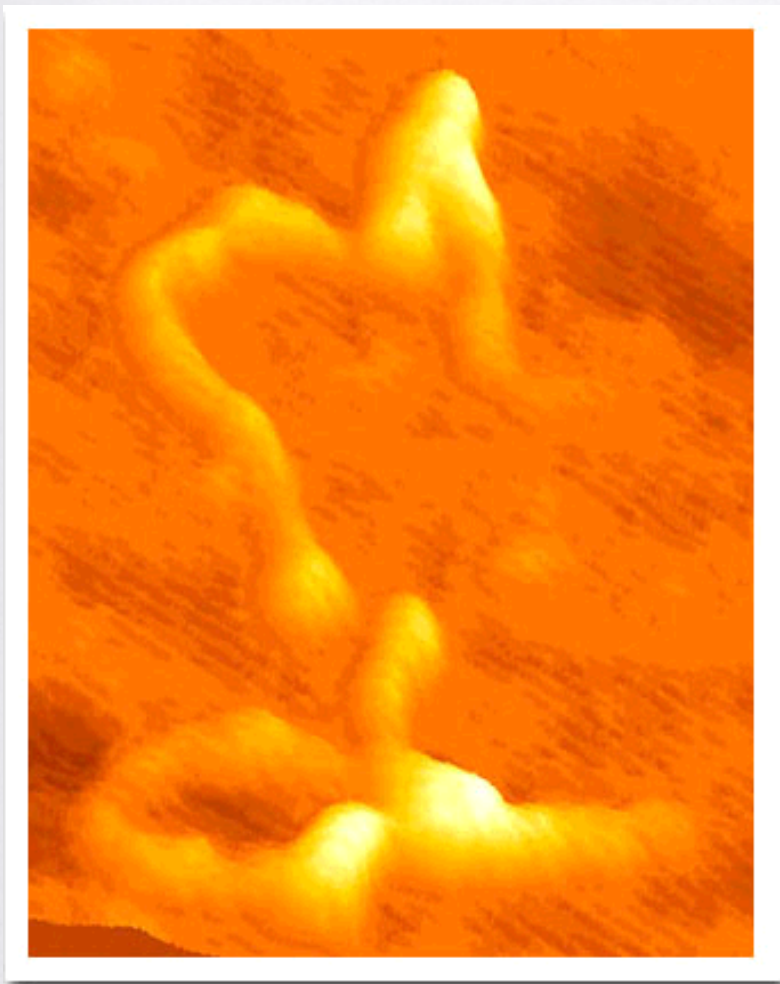
**(Keyence) BZ-X700**

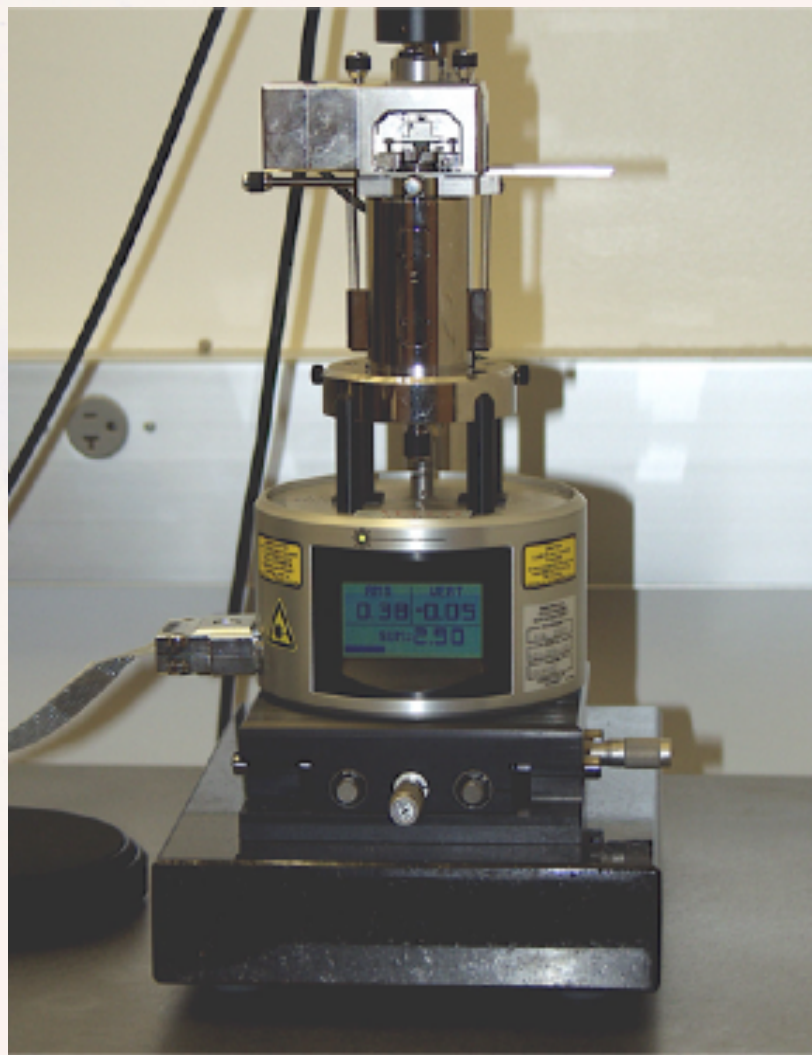


# Atomic Force Microscopy:

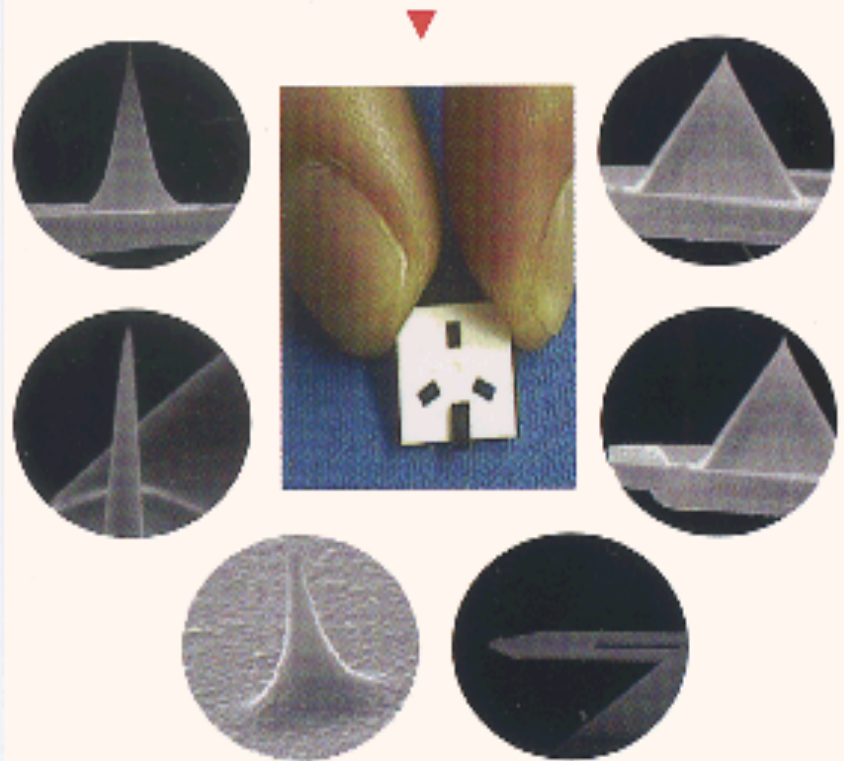




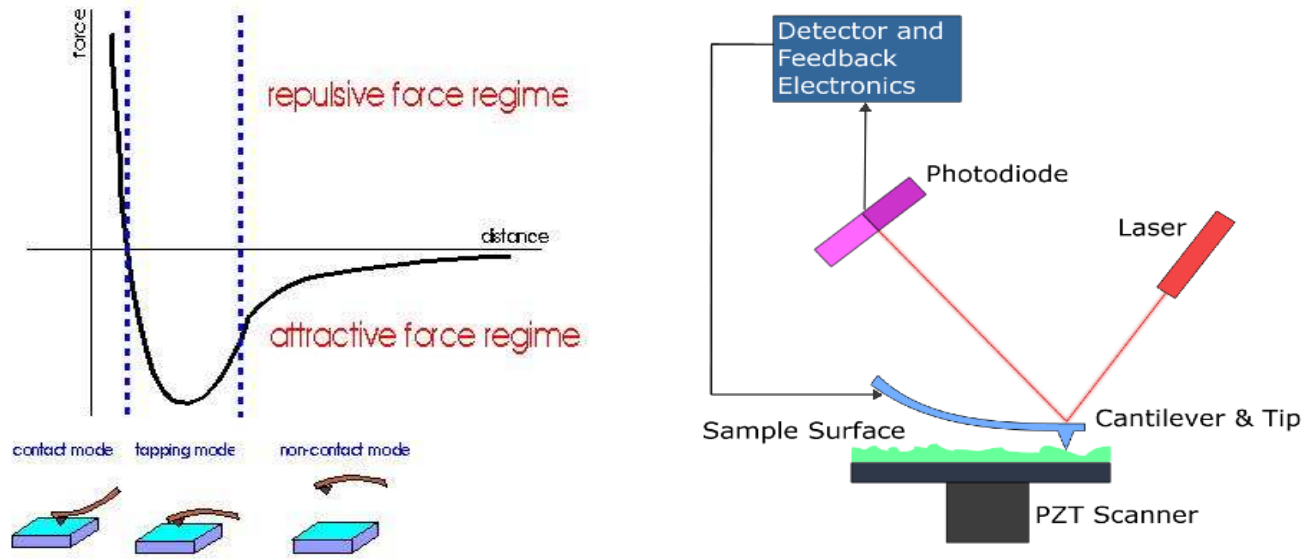




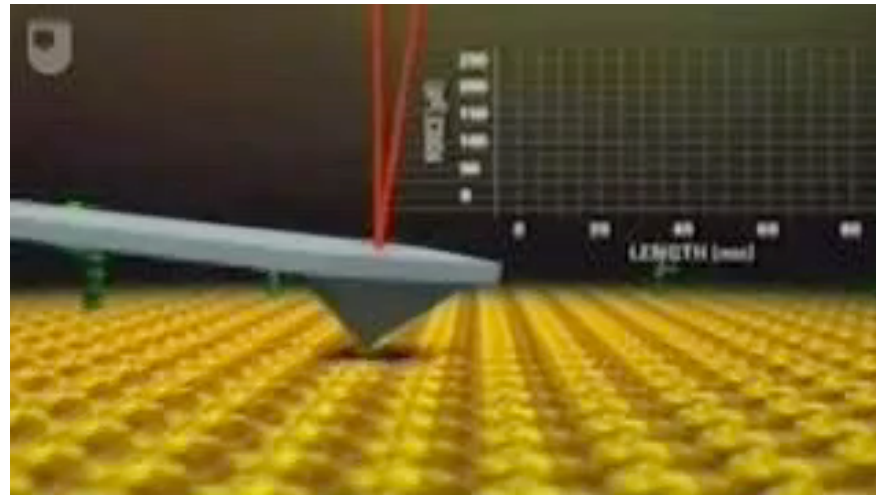
## Cantilevers







Atomic Force Microscope (AFM) operates by measuring attractive or repulsive forces between a probe or "tip" and the sample. The tip is located at the end of a leaf spring or "cantilever". A laser beam is reflected off the cantilever. Any angular deflection of the cantilever caused by the change of the force between tip and sample is represented by the angular deflection of the laser beam. Images are taken by scanning the sample relative to the tip and measuring the deflection of the cantilever as a function of lateral position. Different from traditional microscope, image from AFM is three dimensional.



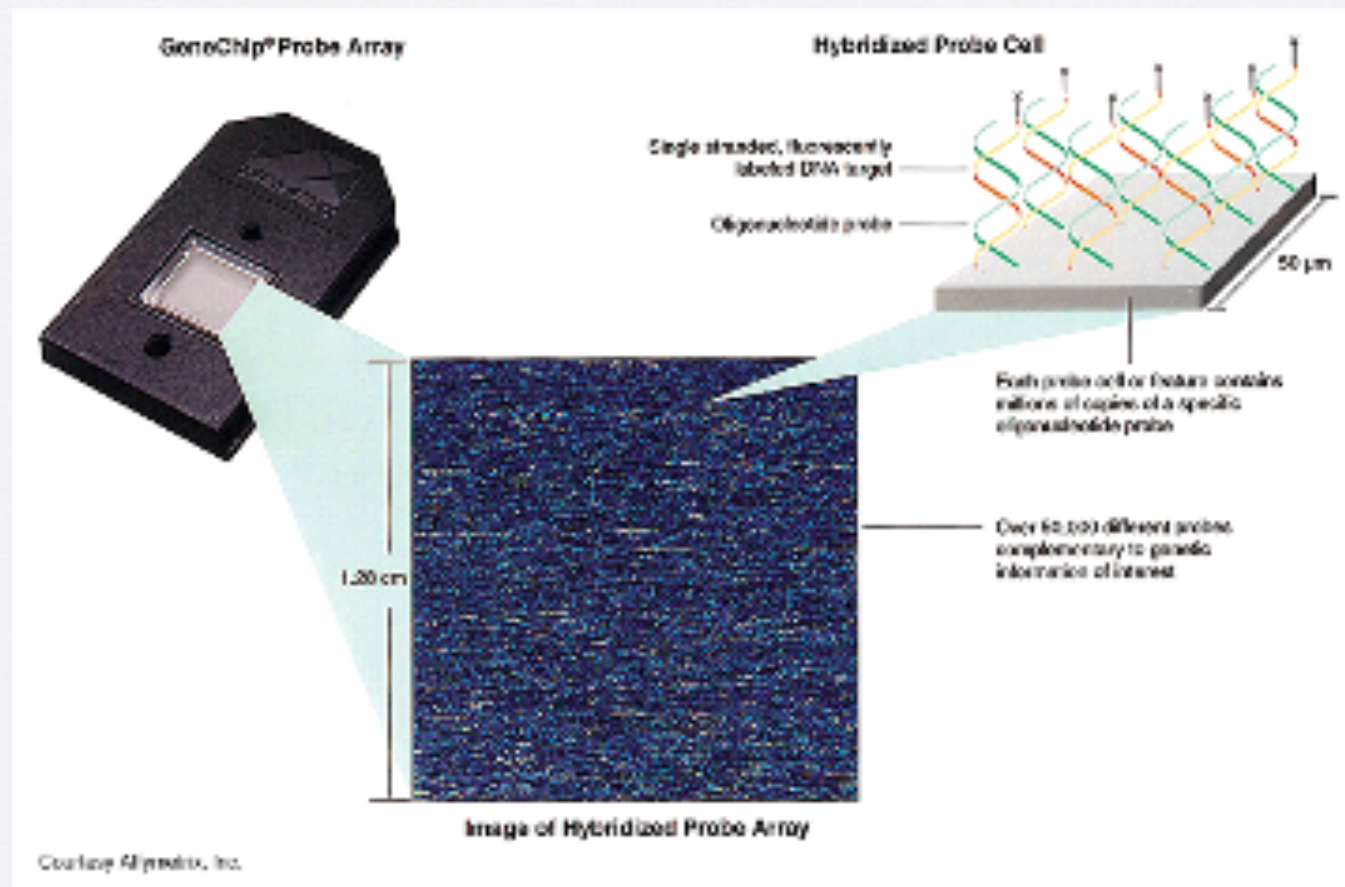
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# Genomics:



**Microarray Technology:** enables us to monitor variable mRNA expression -at the level of the genomic mRNA expression.

# Transcriptomic / Genomic Analysis

Design  
Experiment

Prepare  
Sample

Hybridize

Wash  
&  
Stain

Scan

Analysis



Probe  
Array



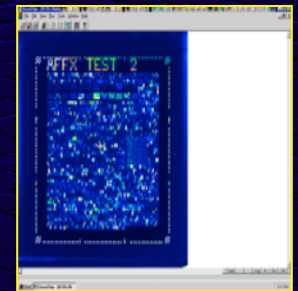
Hybridization  
Oven



Fluidics Station



Scanner



Software

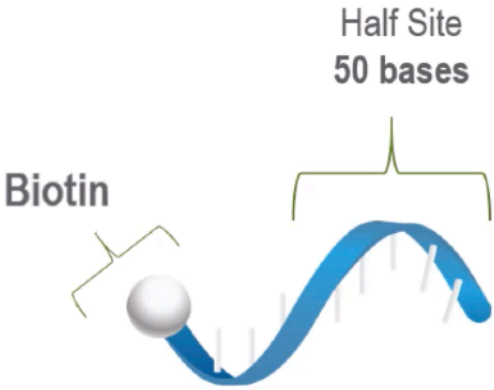


## SUMMER INSTITUTE - ONLINE MODALITY CALENDAR 2022

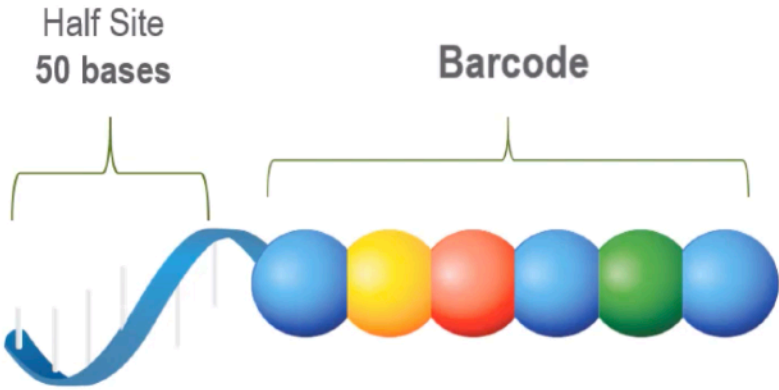
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# Digital Counting Using Barcoded Probes



Target-specific **Capture** Probe



Target-specific **Reporter** Probe

Barcode	Identity
	XLSA
	FOX5
	PDCD1

FOR RESEARCH USE ONLY. Not for use in diagnostic procedures.



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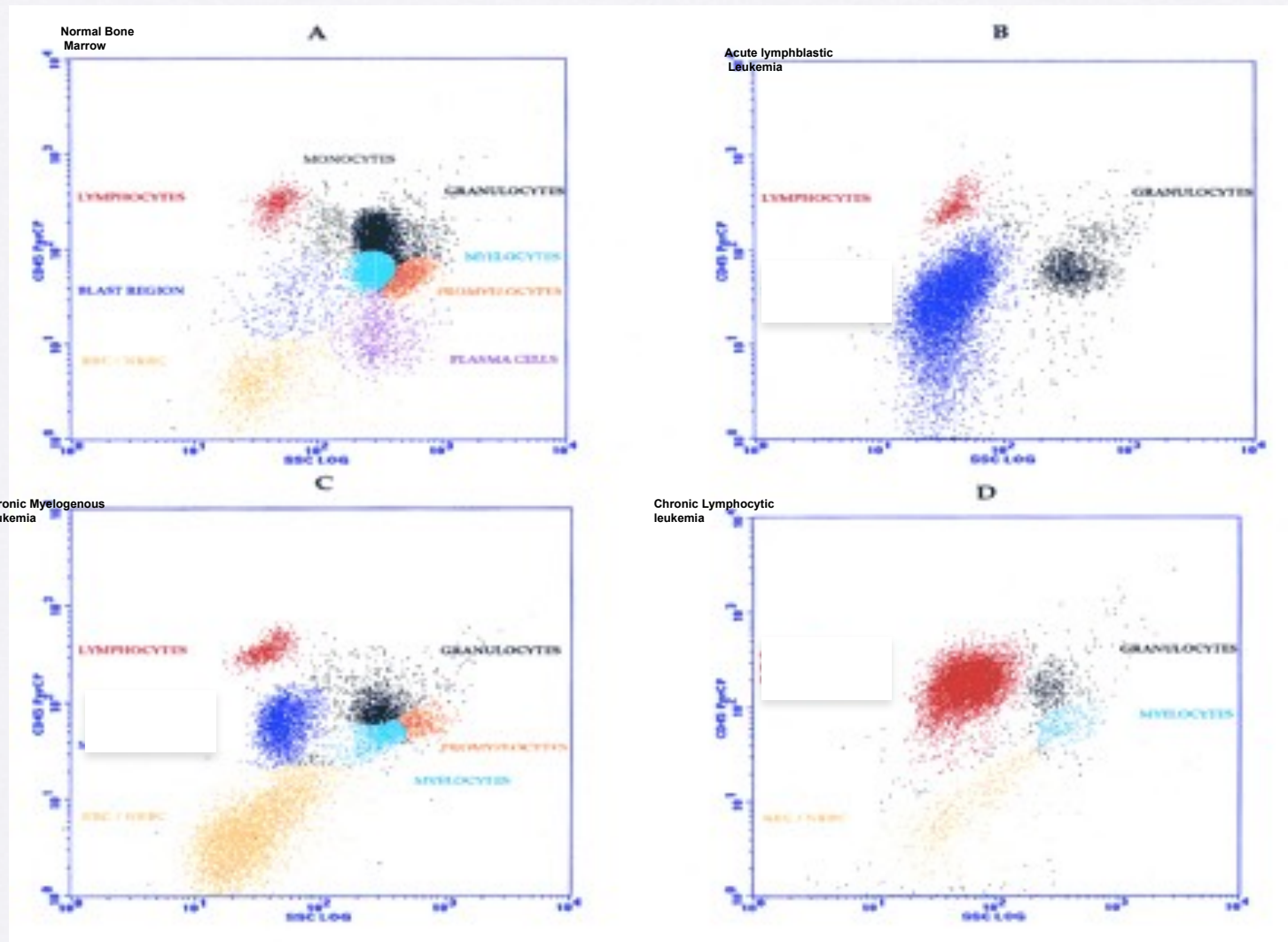


**Automated Flow Cytometry**  
**LSR Fortessa (Becton Dickinson)**

**Automated Flow Cytometry**  
**LSR Fortessa (Becton Dickinson)**







Use of up to six fluorescent markers to various specific types of cancer enables precise diagnosis.

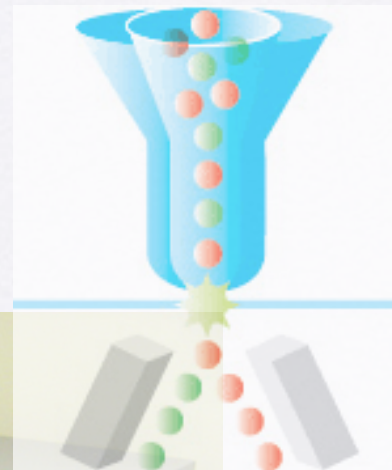
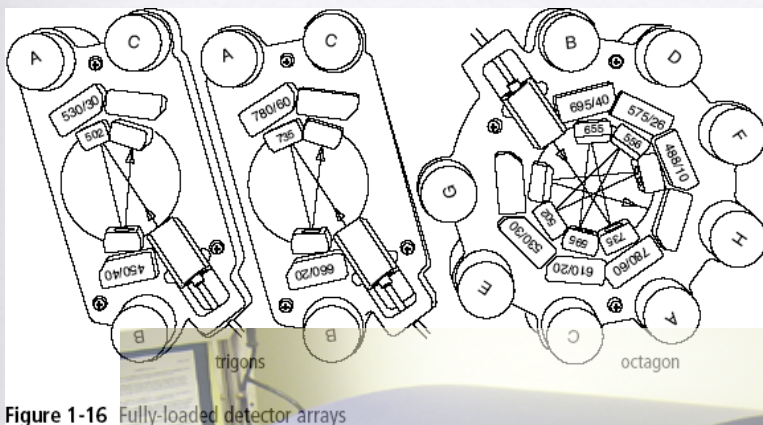
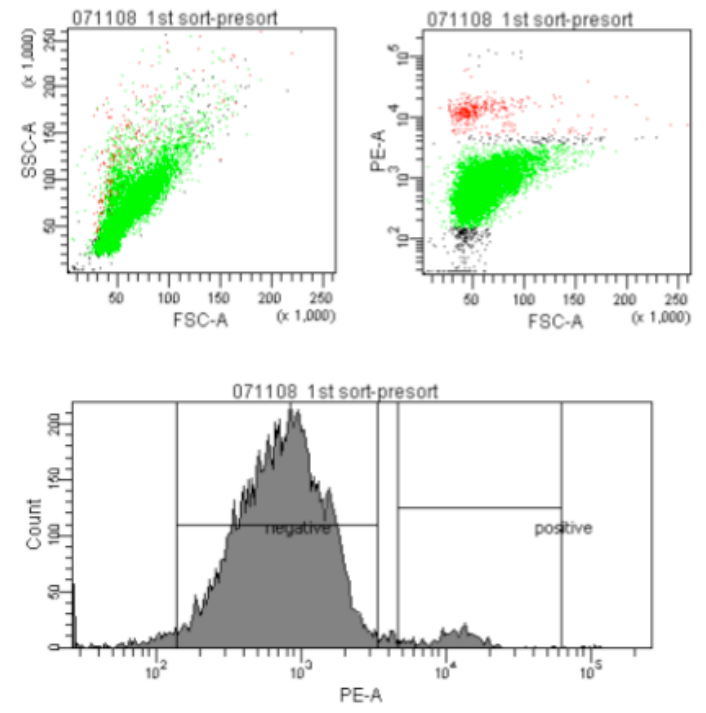


Figure 1-16 Fully-loaded detector arrays



FACSDiva Version 6.1





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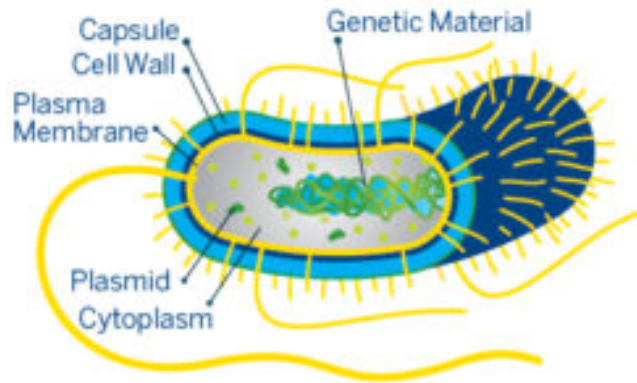
Legend:  
Orange: Courses    Blue: Activities



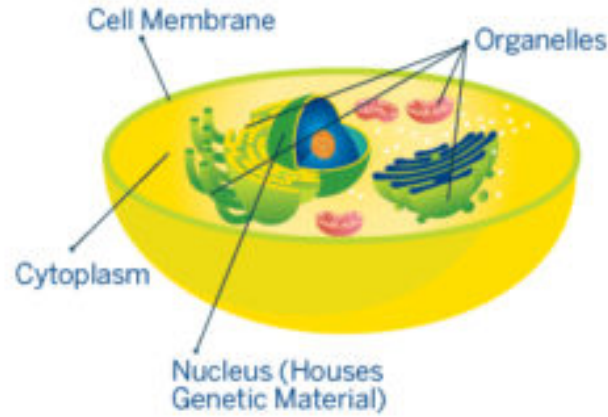




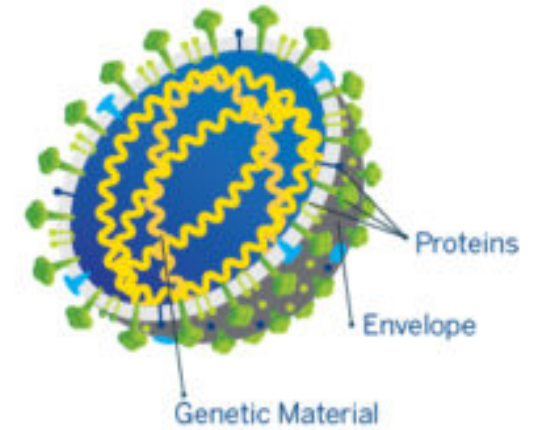
## PROKARYOTE



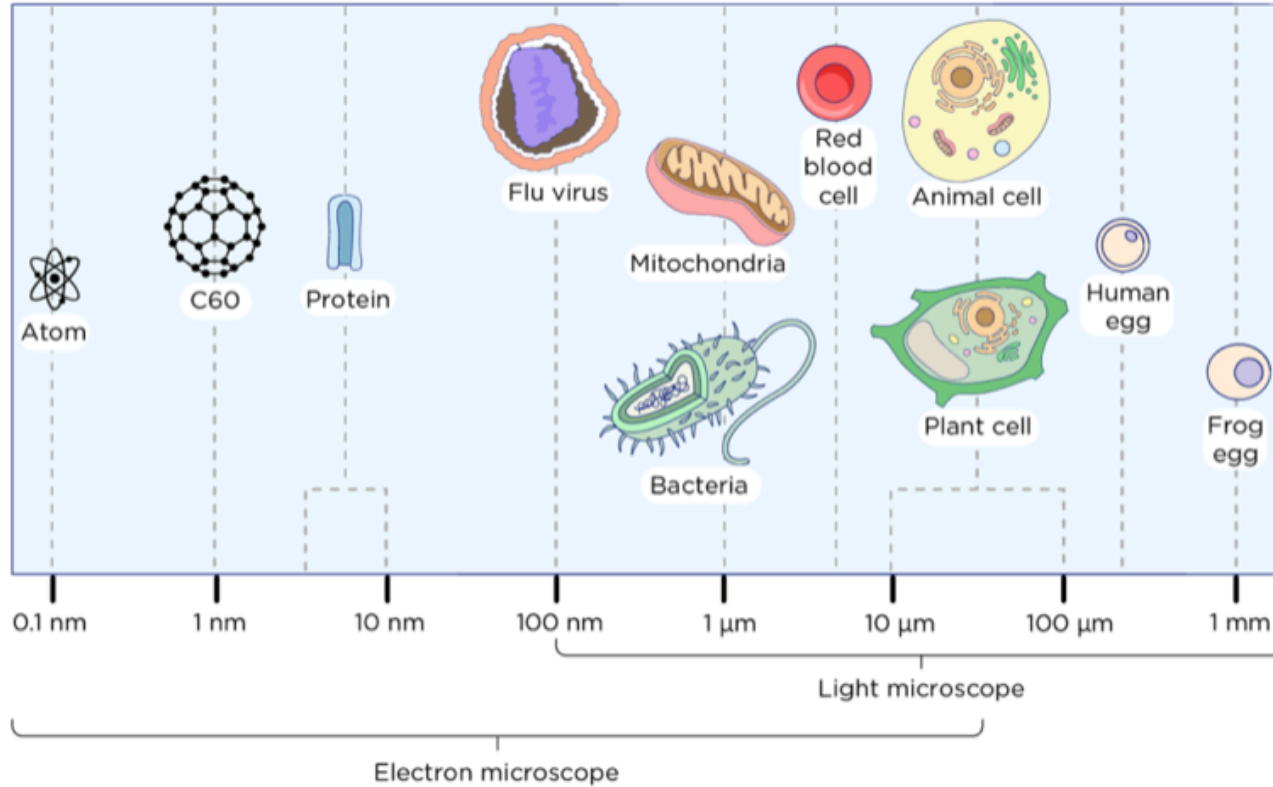
## EUKARYOTE

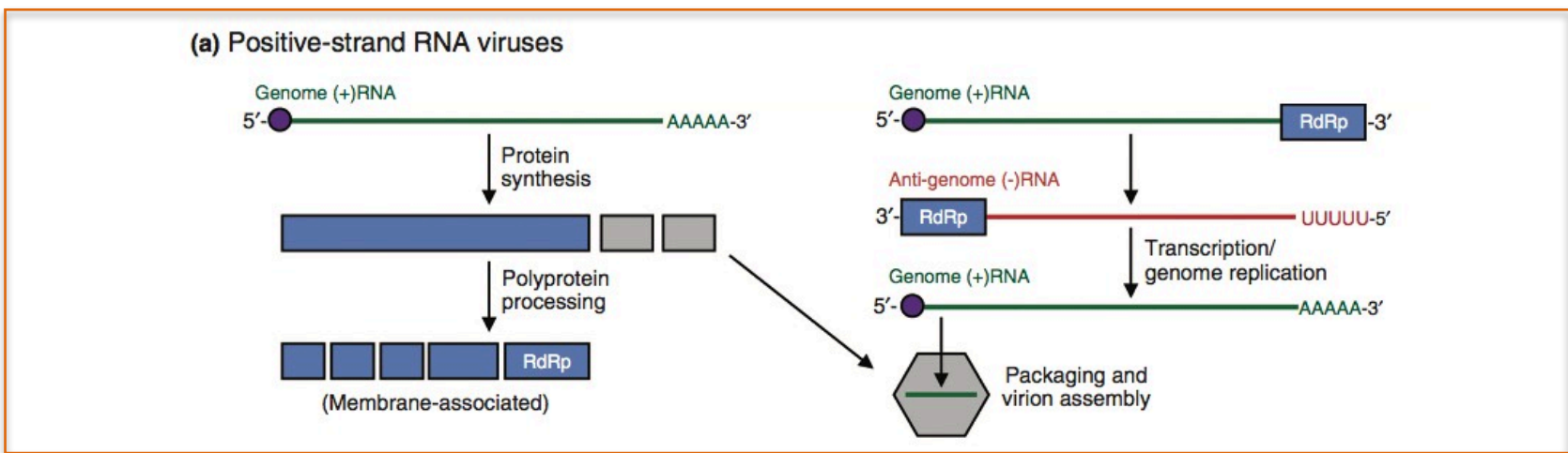


## VIRUS



Relative size of prokaryotes and viruses





Overview of transcription and replication strategies for various types of (a) **Positive-strand RNA [(+)RNA] viruses.**

The genomes of (+)RNA viruses are message-sense (green), and they often contain a 5' m<sup>7</sup>G cap (purple circle) and 3' poly-A tail (AAAAA).

Host cell ribosomes translate the genome into one or more polyproteins, which are co-translationally and post-translationally processed by virally encoded proteases.

Some of the mature polyprotein processing precursors and products include the **RNA-dependent RNA polymerase (RdRp; light blue rectangle)** and cofactors (**light blue squares**) that mediate viral RNA synthesis in association with cellular membranes.

Other proteins made by the virus include those that will assemble into viral particles (gray squares). The **RdRp** mediates the synthesis of negative-strand RNA [(-)RNA] antigenome (red) using the genome as template.

The antigenome is then converted into new (+)RNA genome by the **RdRp** and then packaged into nascent virion particles (gray hexagon).



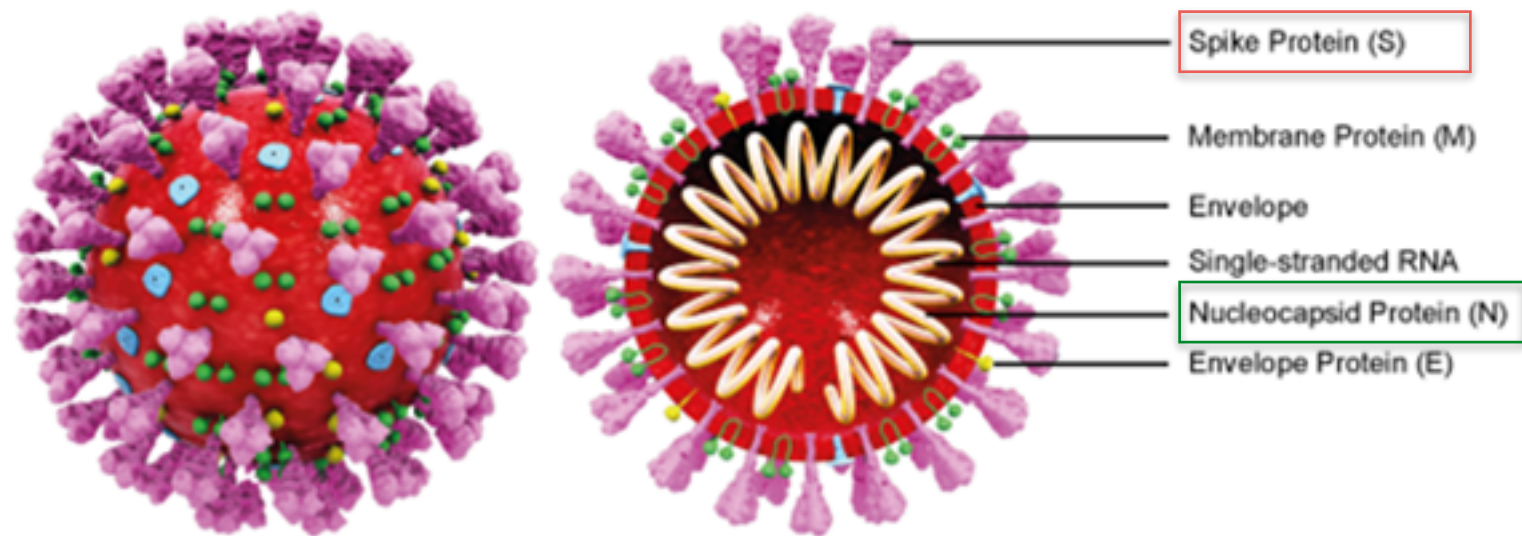


FIGURE 1: Schematic diagram of SARS-CoV-2.

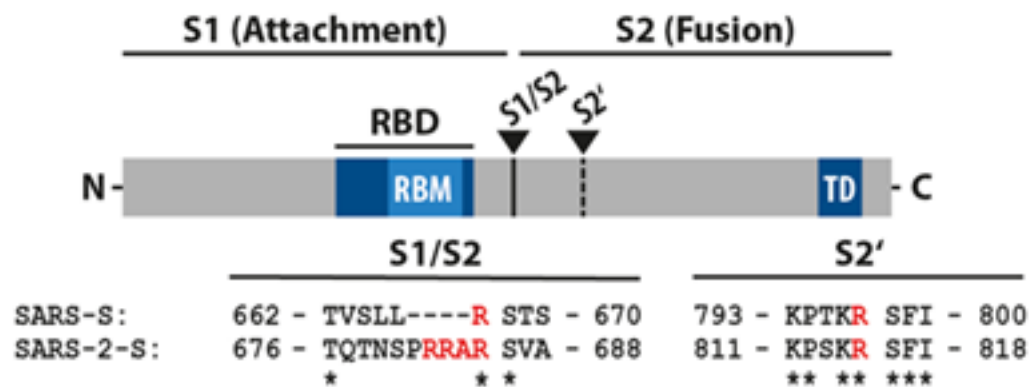
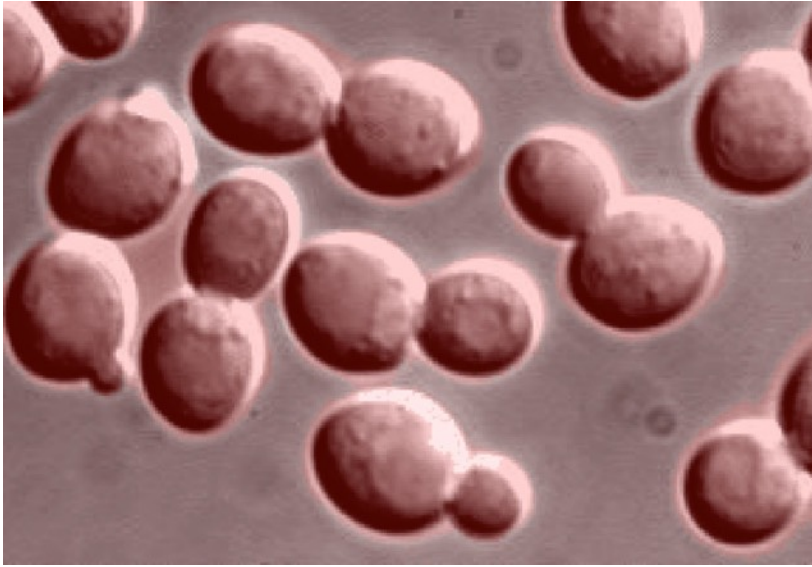


FIGURE 2: Domain comparison overview of Spike Protein S of SARS and SARS-CoV-2. Schematic illustration of SARS-S including functional domains (RBD, receptor binding domain; RBM, receptor binding motif; TD, transmembrane domain) and proteolytic cleavage sites (S1/S2, S2', see arrows). Amino acid sequences around the two protease recognition sites (red) are shown for SARS-S and SARS-2-S (conserved residues are indicated as asterisks).



## Baker's Yeast: *Saccharomyces cerevisiae*

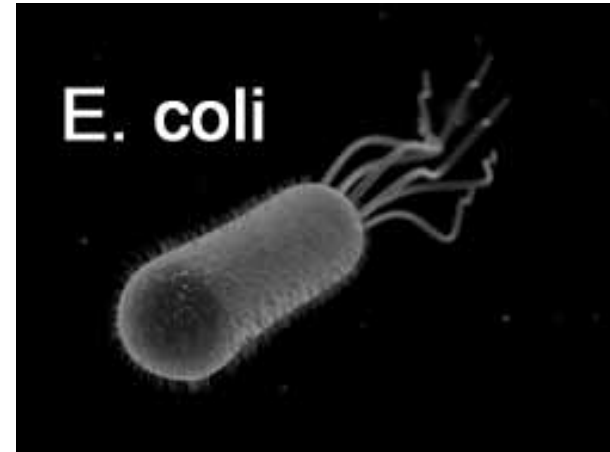
Yeasts are small, *single-celled plants*. They are members of the family *fungi* (singular, *fungus*), which also includes mushrooms. Fungi differ from other plants in that they have no chlorophyll.

~50  $\mu\text{m}$

Eukaryote

DNA is linear

Cell wall



Bacteria thrive on many different types of food. But most yeasts can live only on sugars and starches. From these, they produce carbon dioxide gas and alcohol. Thus, they have been useful to man for centuries in the production of certain foods and beverages. They are responsible for the rising of bread dough and the fermentation of wines, whiskey, brandy and beer. They also play the initial role in the production of vinegar.

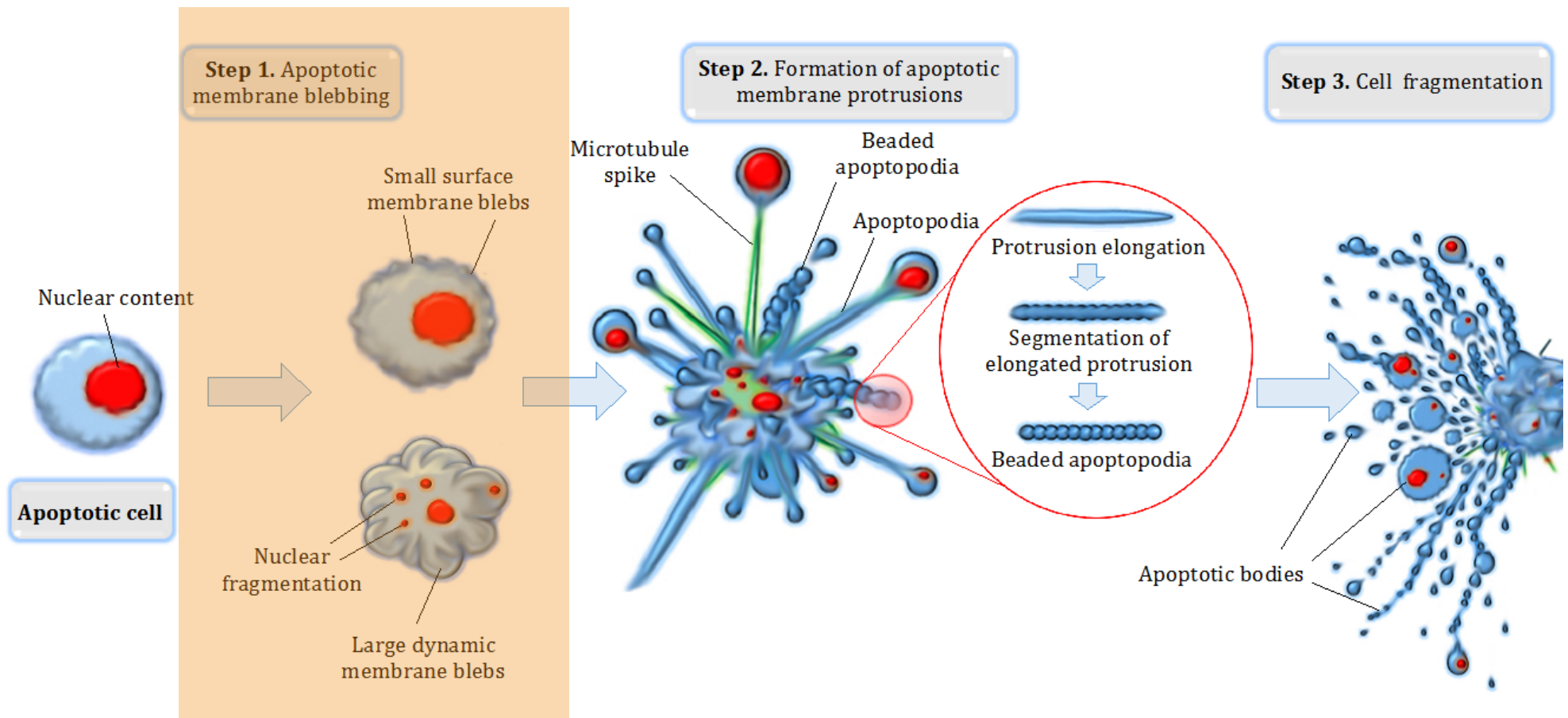
~1.5 -3  $\mu\text{m}$

Prokaryote

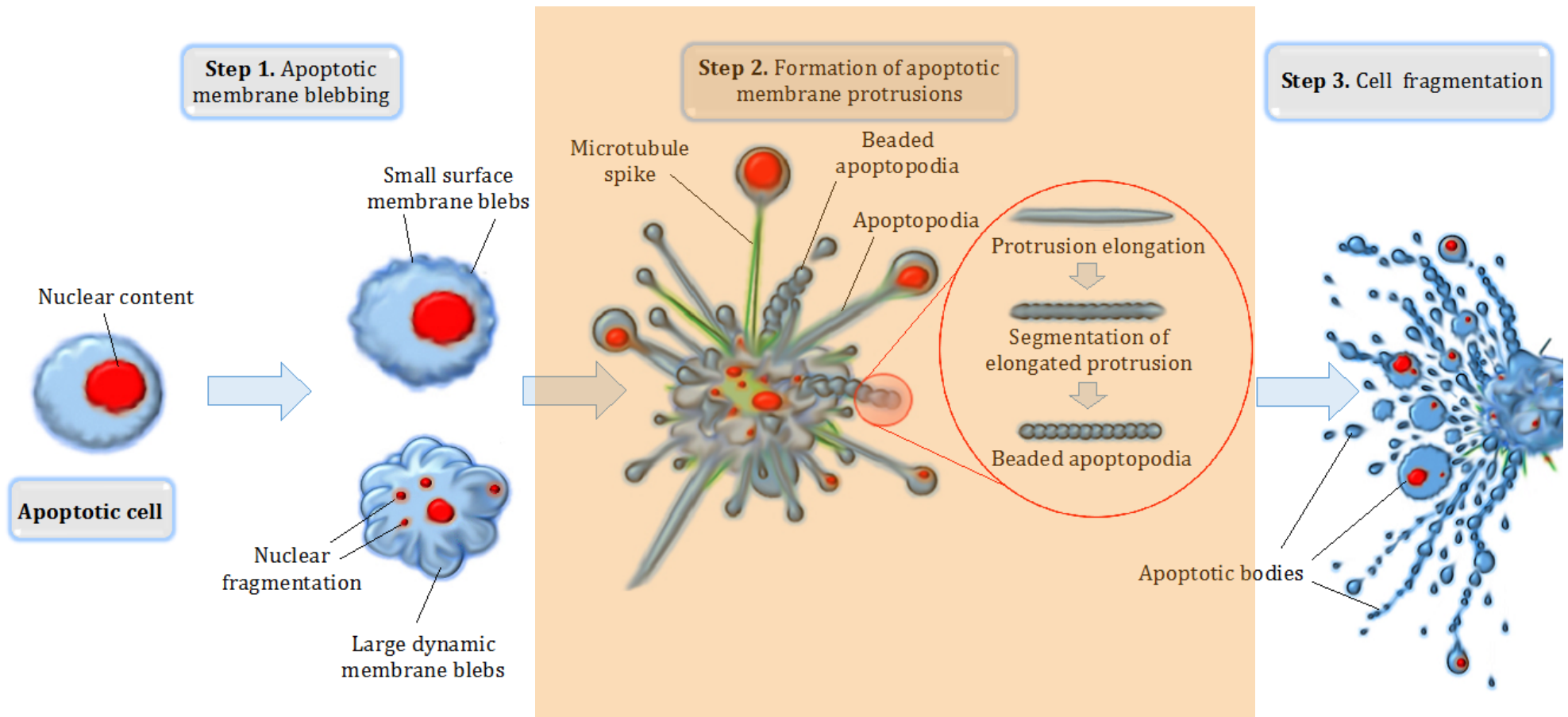
DNA is Circular

cell membrane



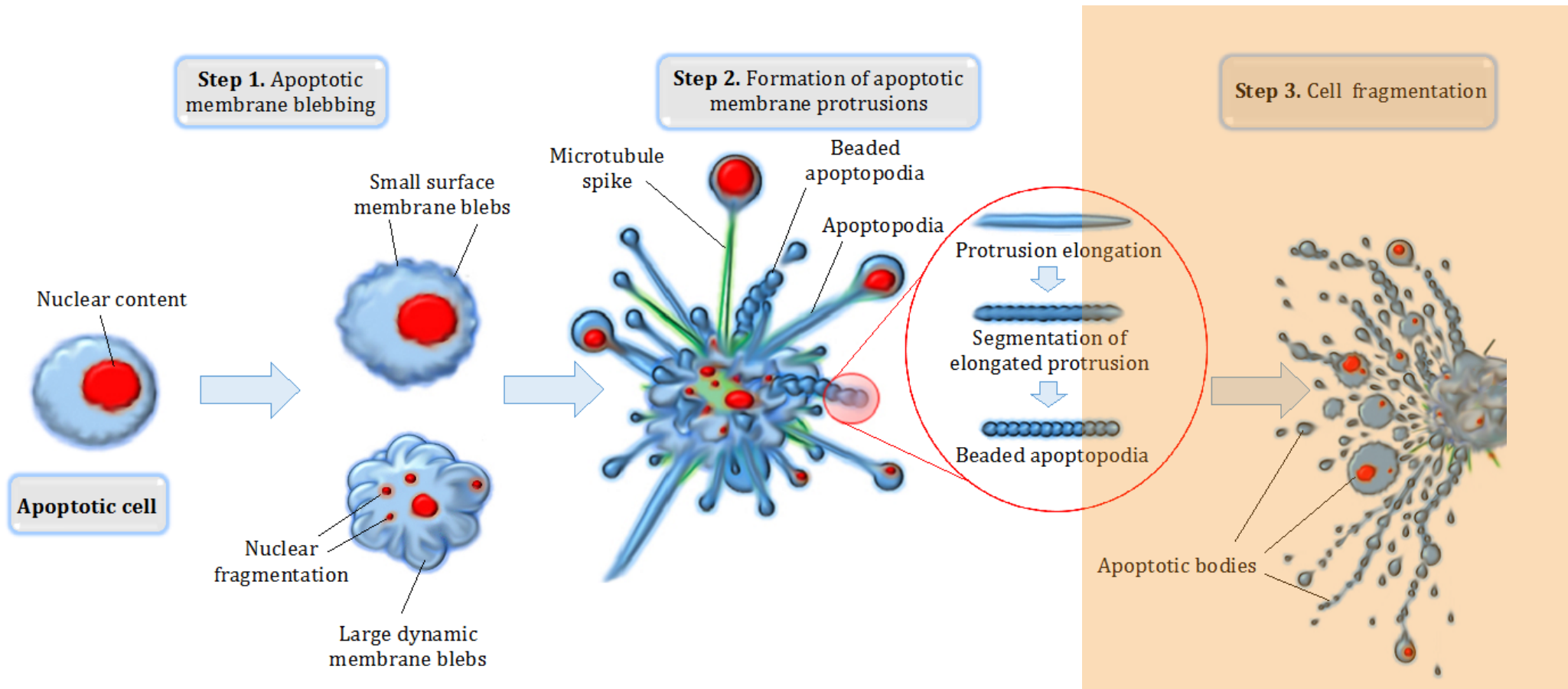


Different steps in apoptotic cell disassembly.



Different steps in apoptotic cell disassembly.





Different steps in apoptotic cell disassembly.

**Caspases** (cysteine-aspartic proteases, cysteine aspartases or cysteine-dependent aspartate-directed proteases) are a family of protease enzymes playing essential roles in programmed cell death. ... These are signalling molecules that allow recruitment of immune cells to an infected cell or tissue.

# Apoptosis in yeast

Frank Madeo <sup>1</sup>, Eva Herker, Silke Wissing, Helmut Jungwirth, Tobias Eisenberg, Kai-Uwe Fröhlich

Affiliations + expand

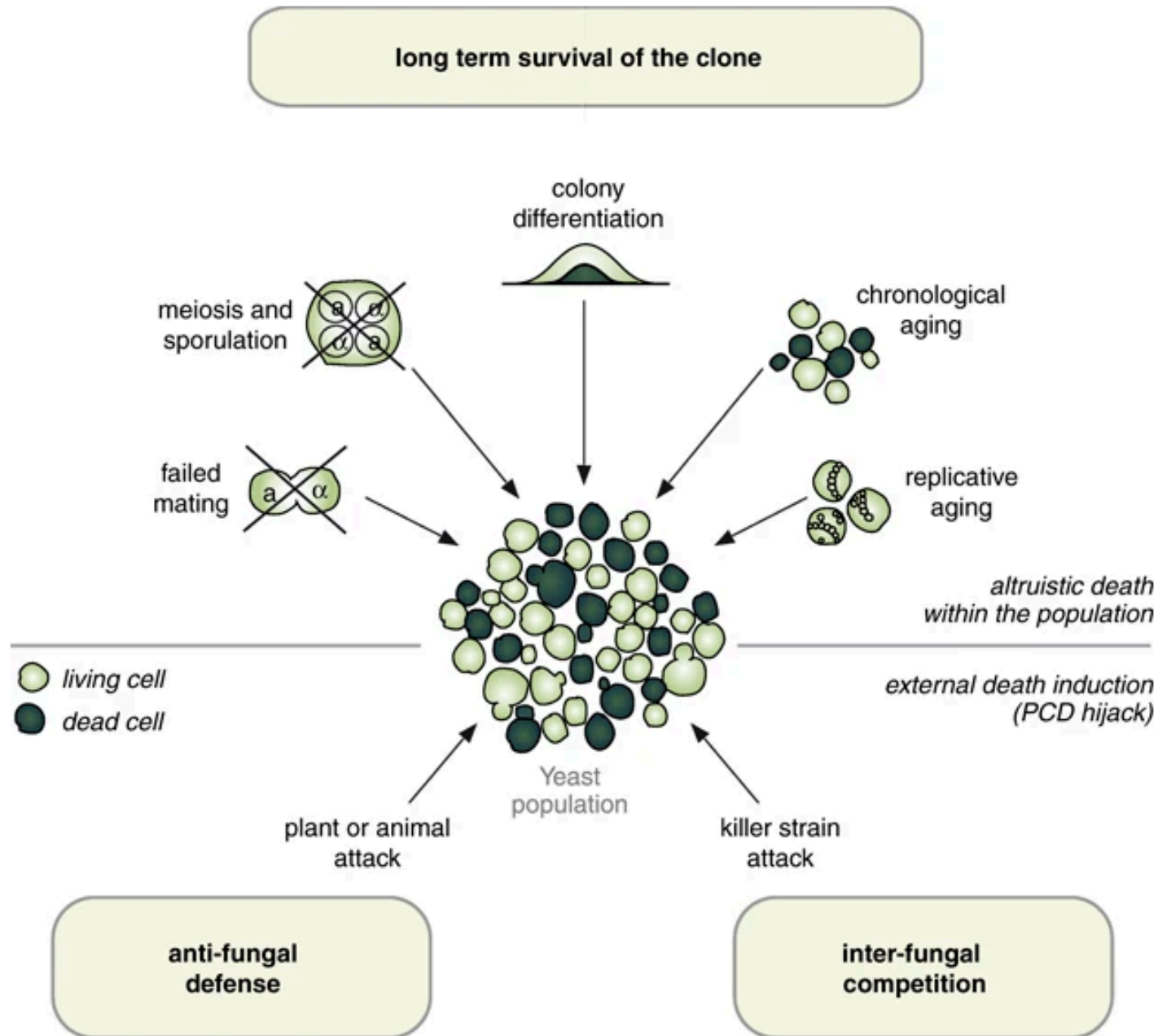
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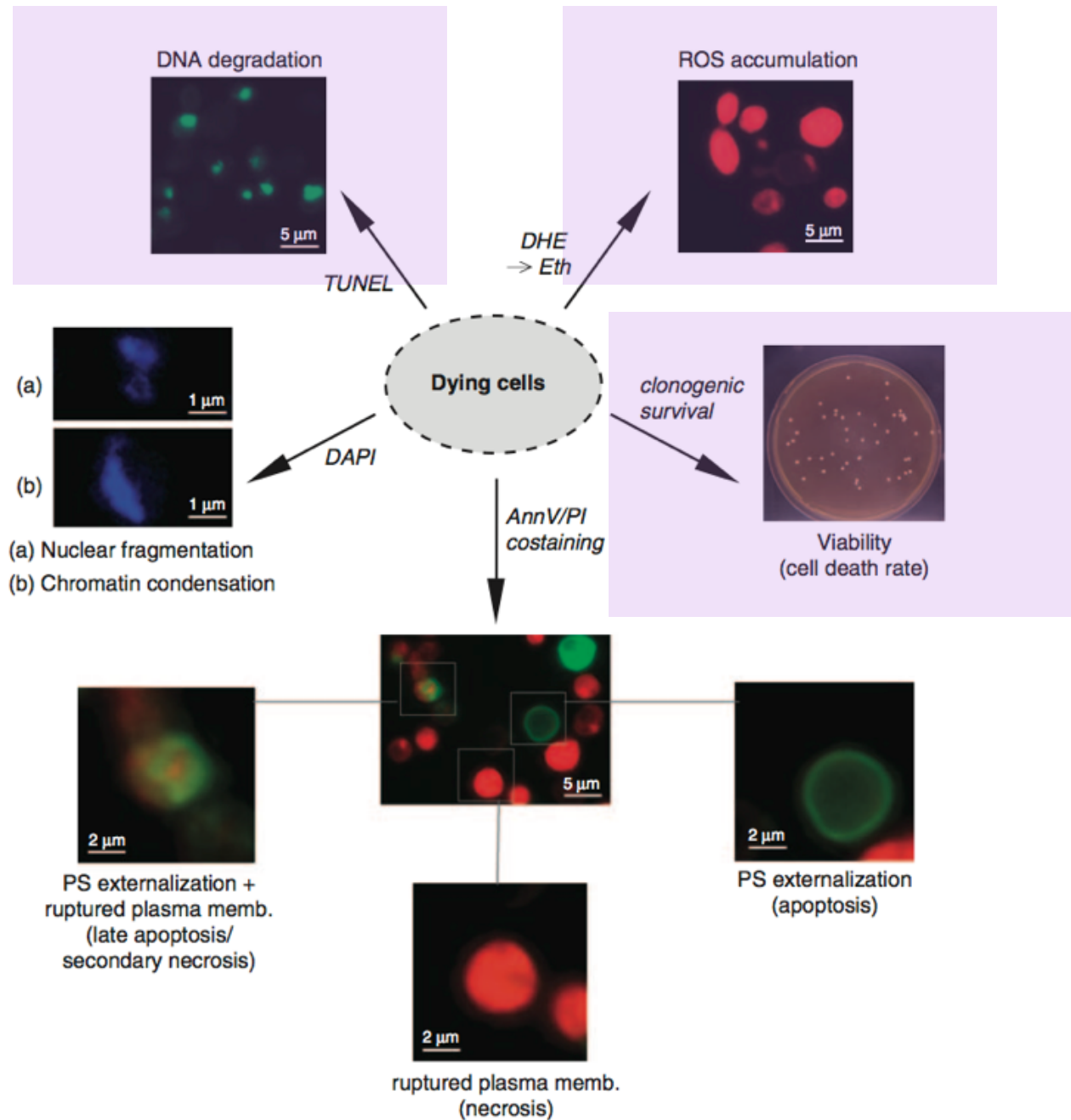
## Abstract

Apoptosis is a highly regulated cellular suicide program crucial for metazoan development. However, dysfunction of apoptosis also leads to several diseases. Yeast undergoes apoptosis after application of acetic acid, sugar- or salt-stress, plant antifungal peptides, or hydrogen peroxide. Oxygen radicals seem to be key elements of apoptotic execution, conserved during evolution. Furthermore, several yeast orthologues of central metazoan apoptotic regulators have been identified, such as a caspase and a caspase-regulating serine protease. In addition, physiological occurrence of cell death has been detected during aging and mating in yeast. The finding of apoptosis in yeast, other fungi and parasites is not only of great medical relevance but will also help to understand some of the still unknown molecular mechanisms at the core of apoptotic execution.

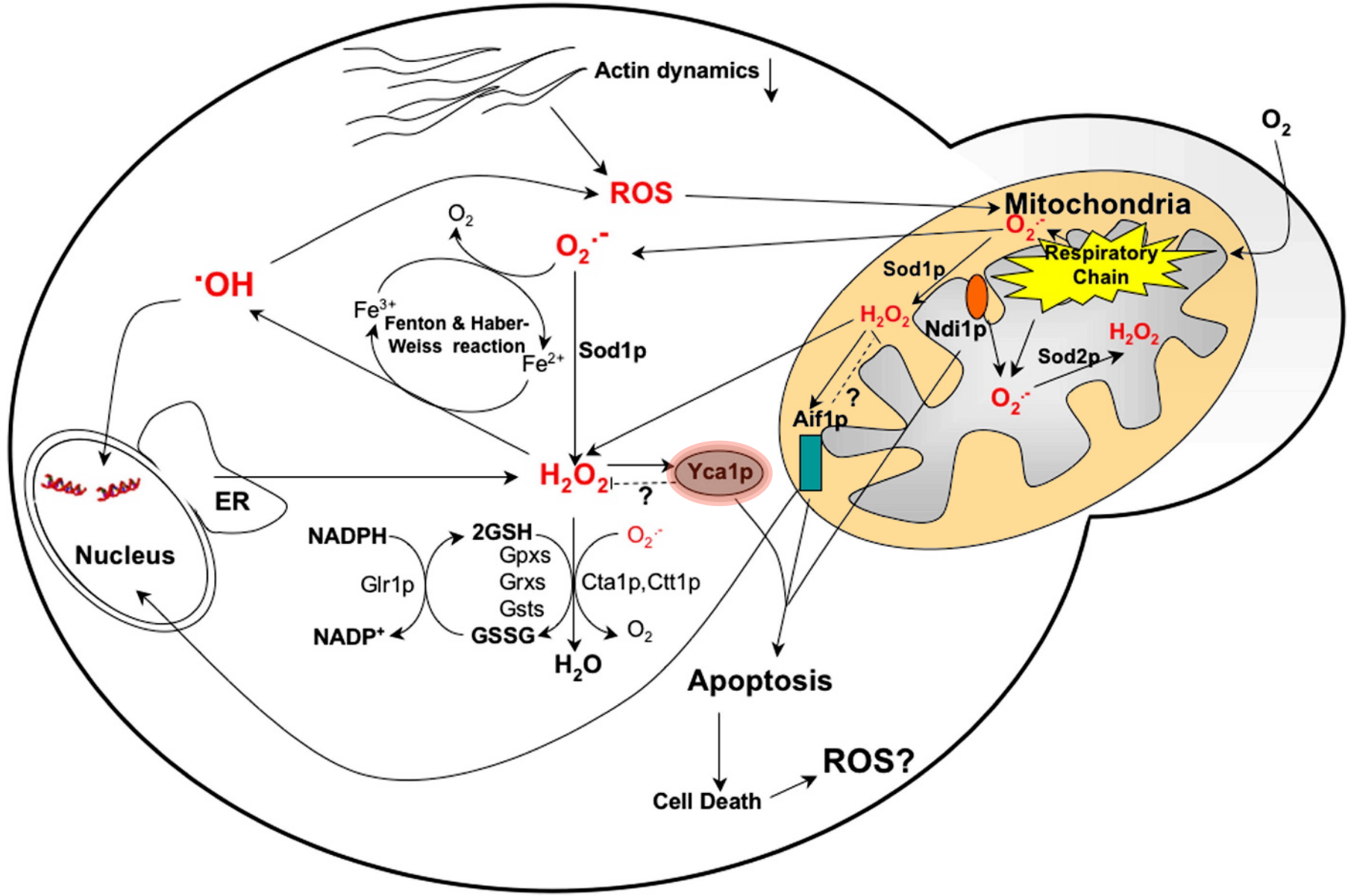


Figure 2

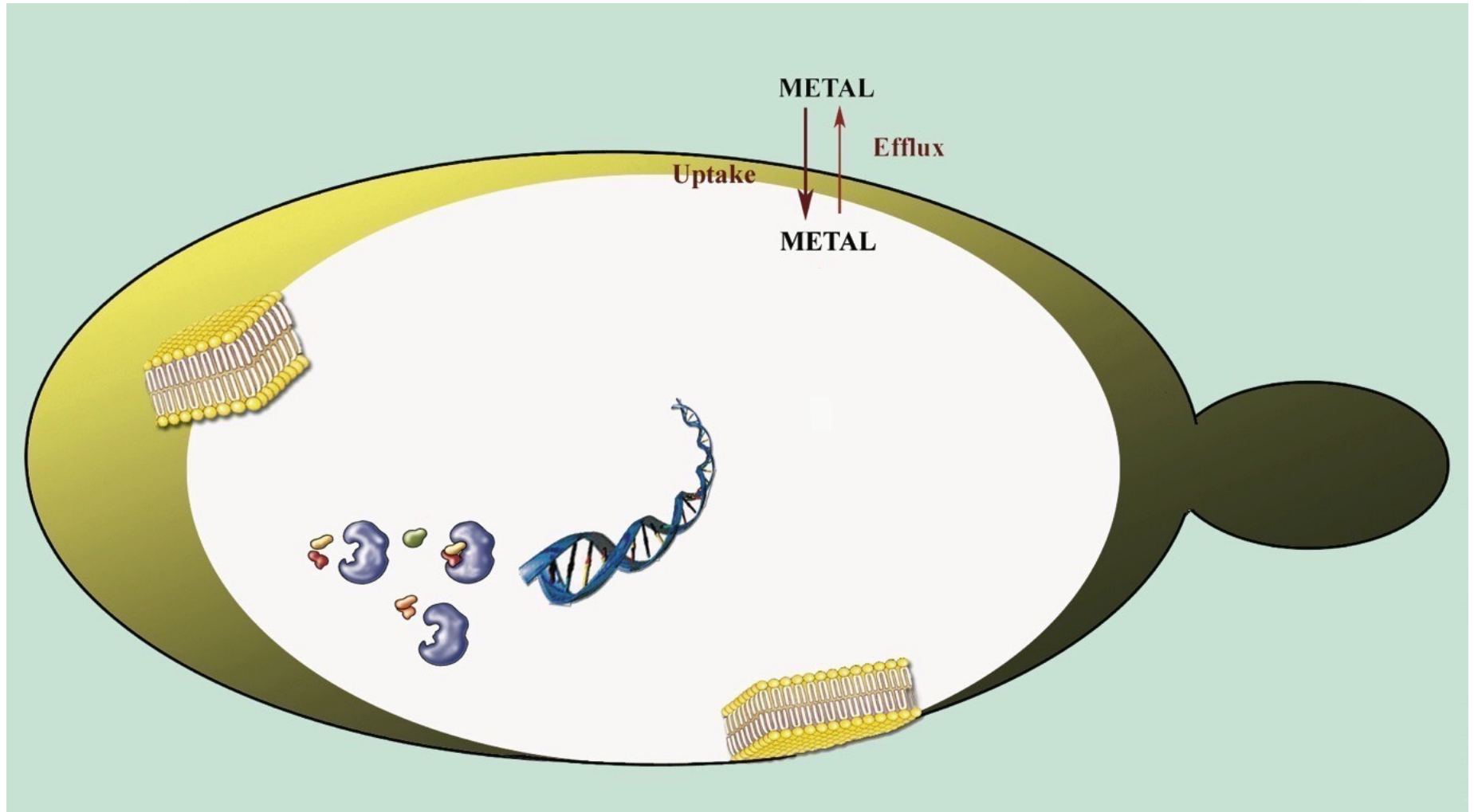








# Metals generate ROS and cause oxidative stress





# Metals generate ROS and cause oxidative stress

