

Non-invasive Single Cell Metabolic Trajectories Using the Phasor-FLIM method

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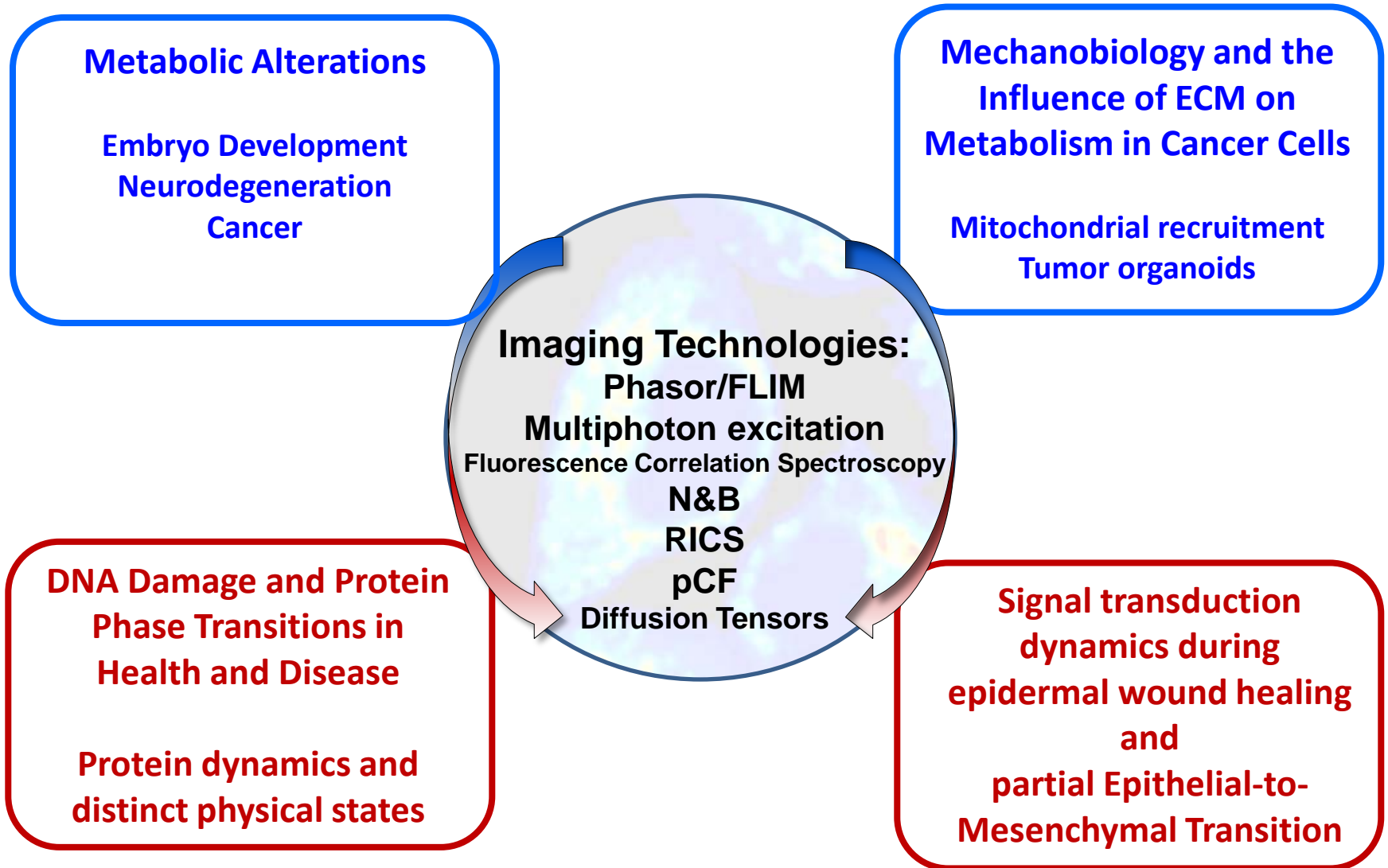


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Overview of the Lab

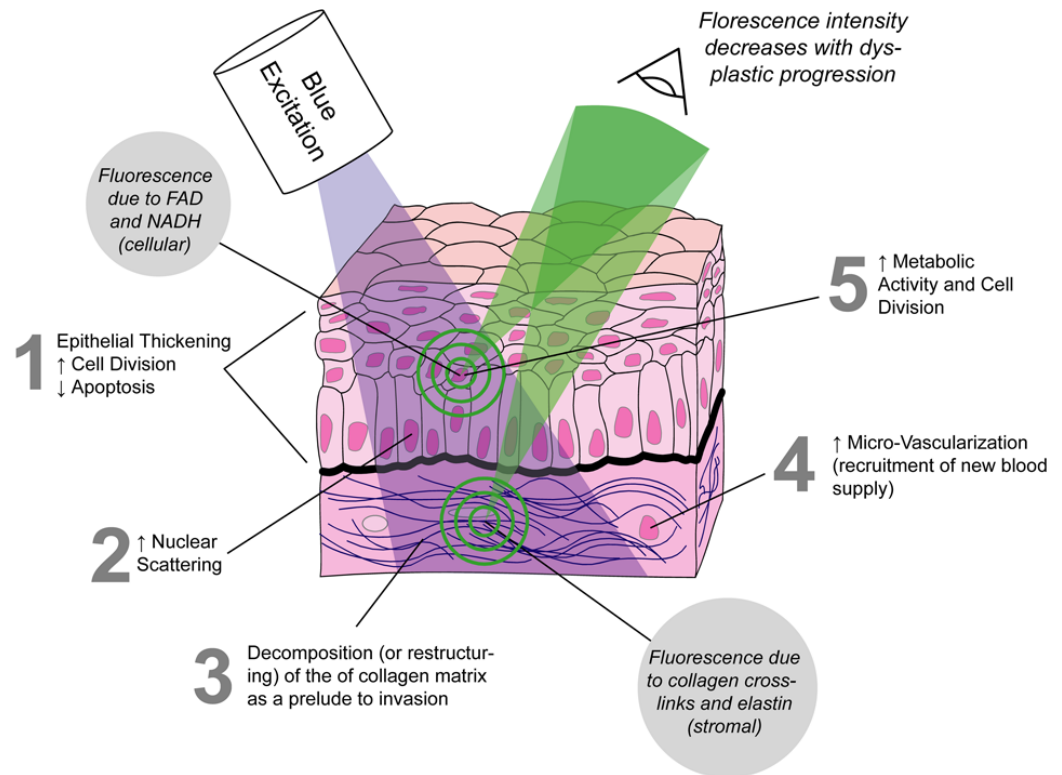


Non-invasive tissue fluorescence imaging technique:

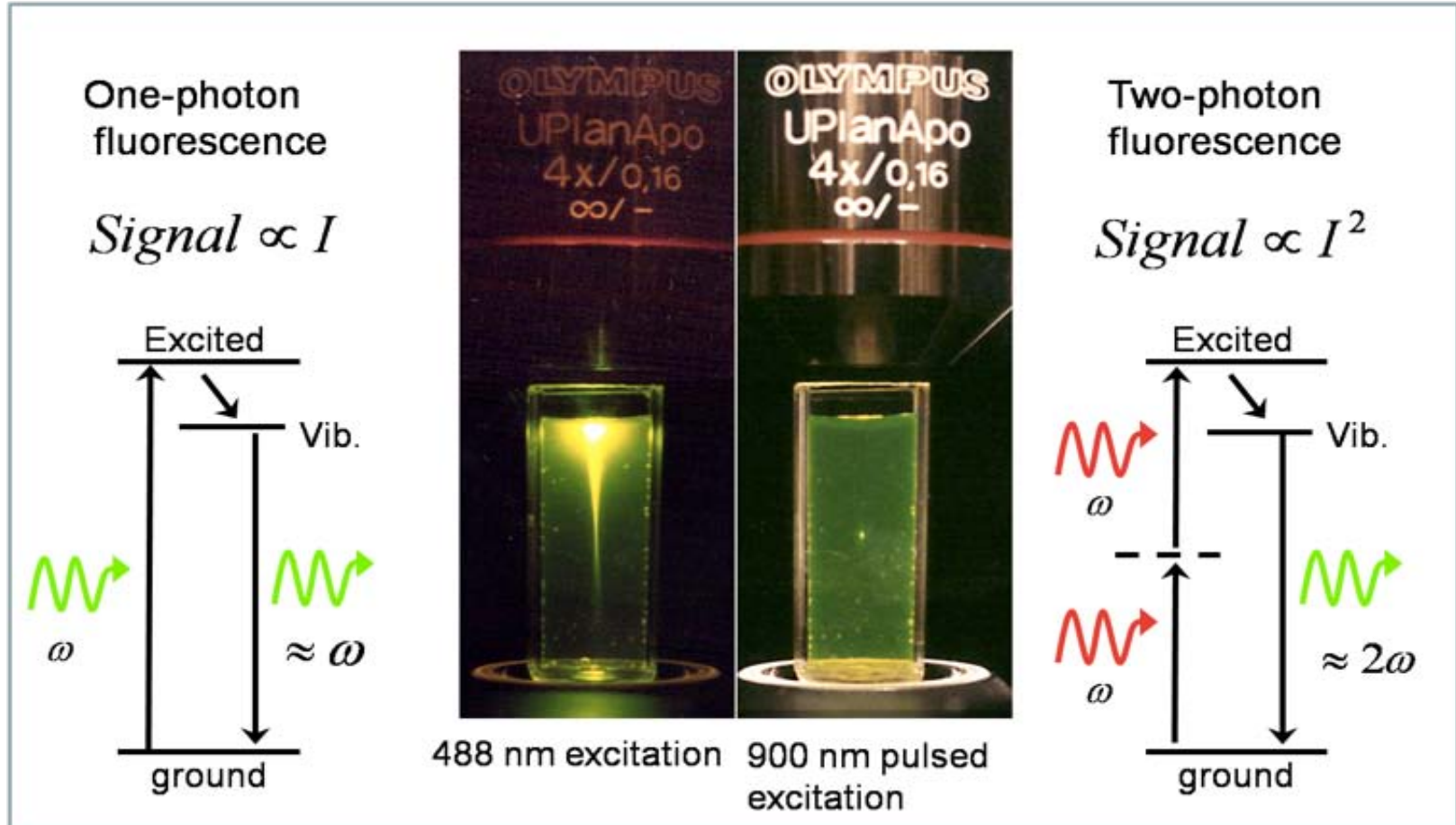
- Cellular fluorescence imaging enables researchers to see the effects of cellular, structural, and/or metabolic activity by observing the in response to light excitation
- Natural tissue fluorescence is caused by intrinsic “fluorophores”.

Important Tissue Fluorophores:

- Collagen
- Elastin
- Keratin
- NADH
- FAD
- Porphyrin
- Fibrin



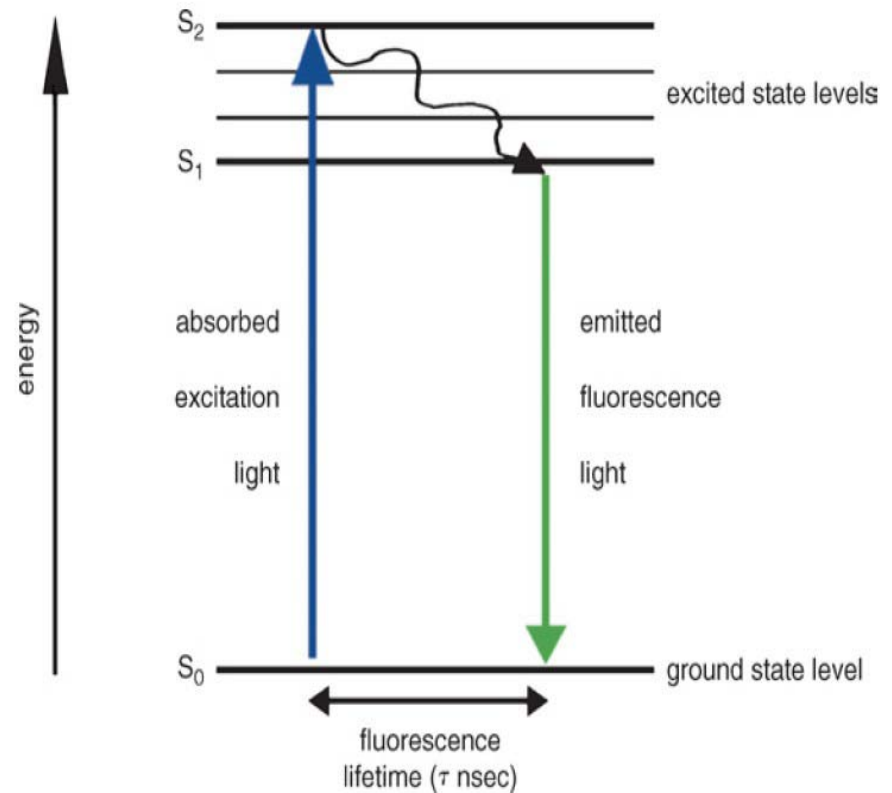
2-photon excitation offers the capability to infiltrate deeper in living tissues



How: Two Photons of half the energy (twice the wavelength), if temporally coincident, sum their energies ($2 \times 0.5 = 1 \times$) to drive fluorophore to activated state (NOTE: This requires very high powered, very expensive lasers)

Fluorescence lifetime imaging microscopy (FLIM)

- While conventional fluorescence microscopy separates fluorophores by wavelengths, FLIM separates sources by fluorescence lifetime.
- The fluorescence lifetime is the average time that a molecule spends in the excited state before emitting a photon.



Outline

- What is Fluorescence Lifetime imaging Microscopy (FLIM)?
- How is fluorescence lifetime measured in the time domain versus the frequency domain?
- How is a FLIM system set up?
- What are the advantages/disadvantages of FLIM?
- The phasor analysis for FLIM
- Metabolic trajectories of living systems

Why we do FLIM?

FLIM is used for :

- FRET (Forster Resonance Energy Transfer).
- Intracellular mapping of Ion concentration and pH imaging.
- Biochemical reactions (oxidation/reduction) processes
 - NAD and NADH. Metabolic index.
- Long lifetime imaging (phosphorescence).
 - For example O_2 concentration in the cell or in tissues

Time Resolved Fluorescence

- What's happening during the time of the fluorescence emission?
- Fluorescence Lifetime

Fluorescence Quantum Yield ϕ : important for dyes
 Ratio of the rate of fluorescence and the sum of the rates that depopulate the excited state

Quantum Yield:

Can be expressed as the lifetime

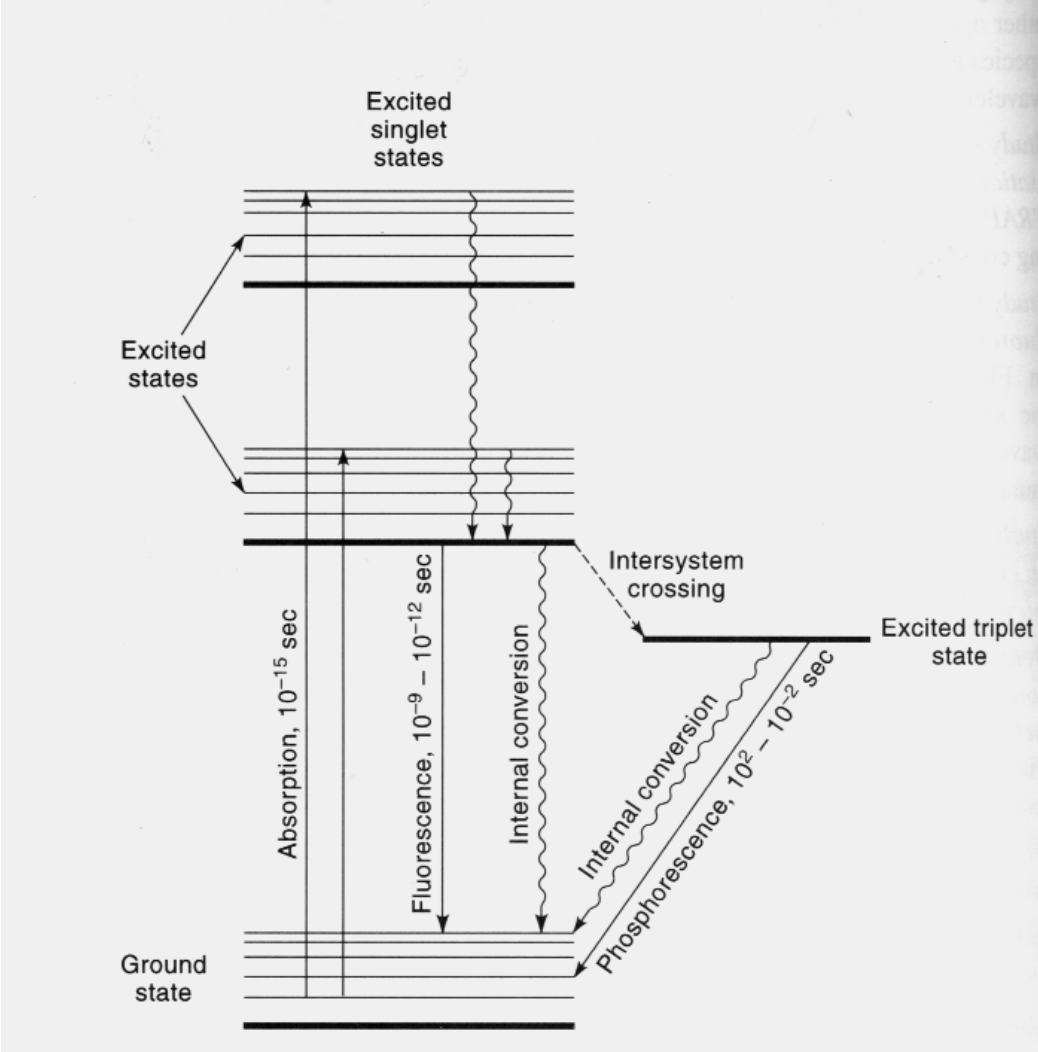
$$\phi = \frac{k_f}{k_f + k_{isc} + k_{nonrad}}$$

$$\tau_0^{-1} = k_f$$

Natural lifetime:
 Inverse of the fluorescence emission rate

$$\tau^{-1} = k_f + k_{isc}$$

Measured lifetime is sum of Rates of natural lifetime and non radiative decay paths



Fluorescence Lifetime τ

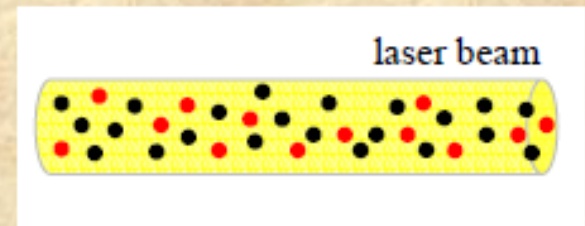
Although we often speak of the properties of fluorophores as if they are studied in isolation, such is not usually the case.

Absorption and emission processes are almost always studied on *populations* of molecules and the average properties of a molecule of the population are deduced from the macroscopic properties of the process.

The rate of the change of molecules in the excited state depends on how much you depopulate the emission plus how much you populate because of excitation

Illustration:

A laser beam passes through a solution containing fluorophores. At any given time some fluorophores will be excited, while the rest will be in its ground state.



The excited state population of fluorophores is described by a rate equation:

$$\frac{dn^*}{dt} = -n^* \Gamma + n f(t)$$

where $n + n^* = n_0$ (n_0 describes the total number of molecules and is a constant)

Meaning of the lifetimes

- Imagine that a sample is excited with an infinitely sharp pulse of light
- This results in an initial population (n_0) of fluorophores in the excited state
- The excited-state population decays with a rate of $\Gamma + k_{nr}$:

$$\frac{dn(t)}{dt} = -(\Gamma + k_{nr}) n(t)$$

$n(t)$ is the number of excited molecules at time t following excitation, Γ is the emissive rate and k_{nr} is the nonradiative rate.

Because emission is a random event, and each excited fluorophore has the same probability of emitting in a given period of time, the result is an exponential decay of the excited state population :

$$n(t) = n_0 \exp(-t/\tau)$$

Measuring fluorescence lifetime using a field programmable gate array (FPGA)

- Our approach uses serial detectors in the photon counting mode, and the digital heterodyning method to acquire data which is directly analyzed in the frequency domain.
- the sampling windows slide through the entire period of the emission response due to the slight difference in frequencies, for a total of 256 steps

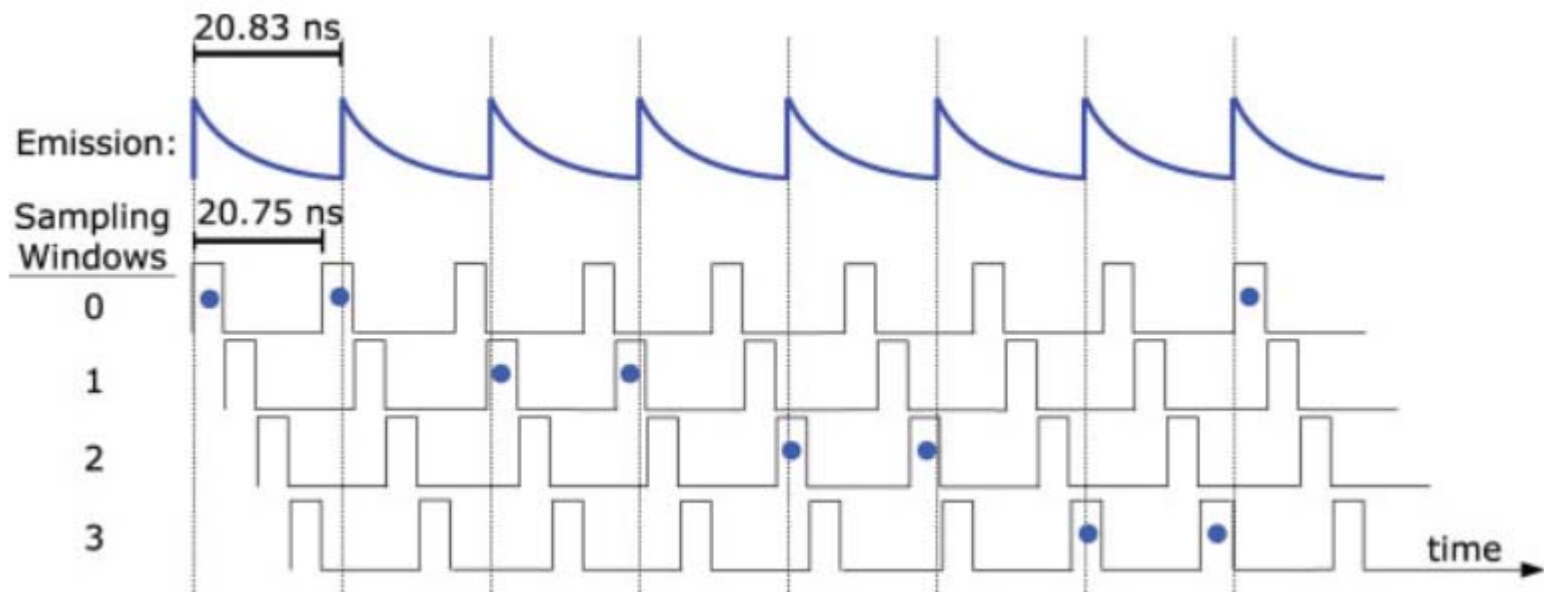
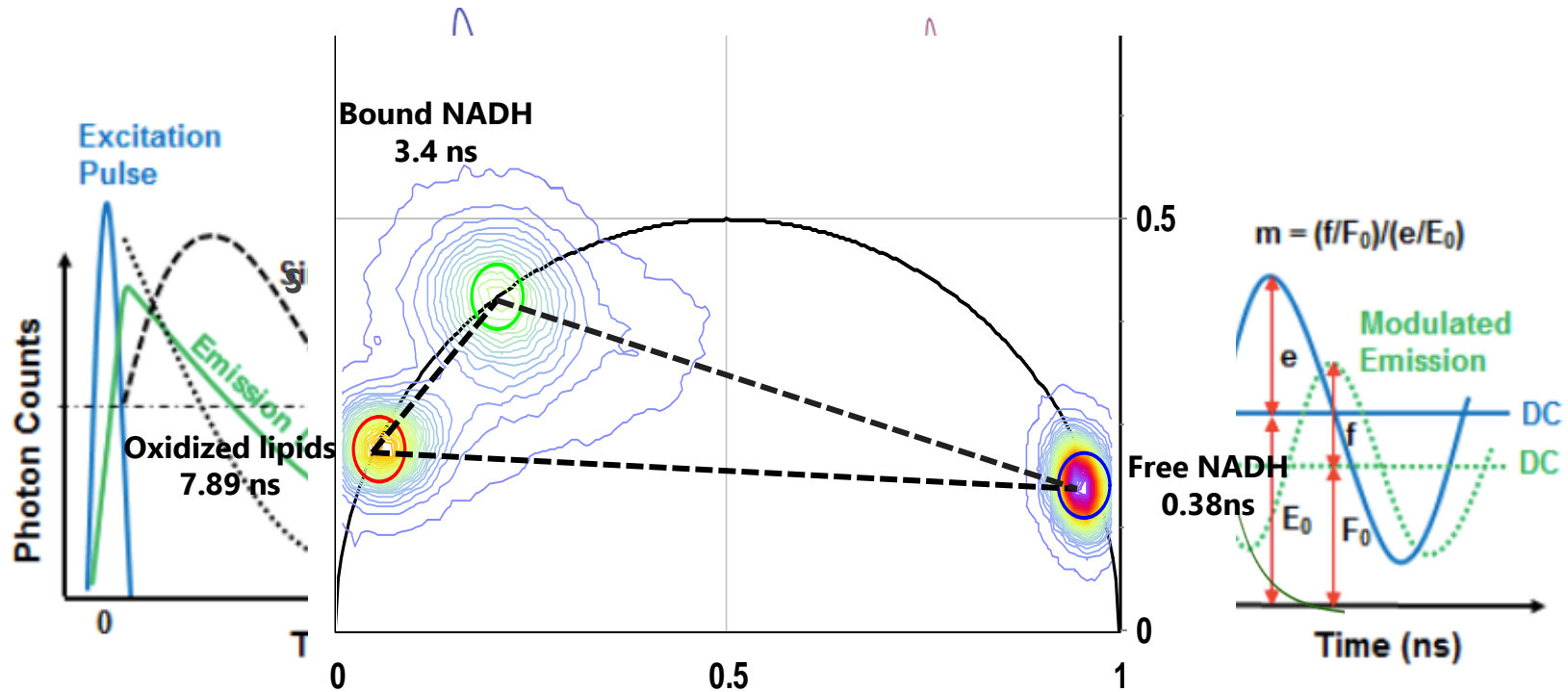


illustration of the digital heterodyning principle
Colyer et al.

Phasor - A Graphic Representation of the Raw FLIM Data



$$g_i(\omega) = \int_0^{\infty} I(t) \cos(\omega t) dt / \int_0^{\infty} I(t) dt$$

$$s_i(\omega) = \int_0^{\infty} I(t) \sin(\omega t) dt / \int_0^{\infty} I(t) dt$$

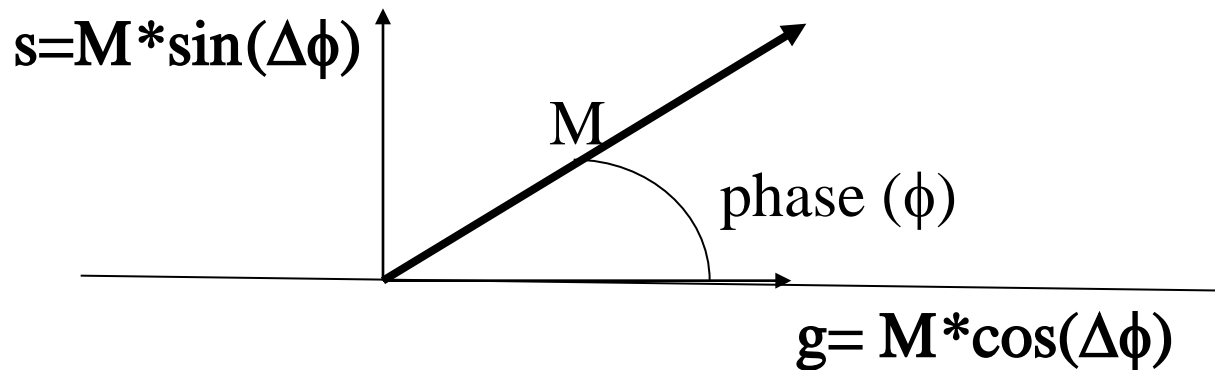
$$g(\omega) = m \cos(\varphi) \quad s(\omega) = m \sin(\varphi)$$

Using Phasors: what is a phasor??

$$g_i(\omega) = \int_0^{\infty} I(t) \cos(\omega t) dt / \int_0^{\infty} I(t) dt$$

$$s_i(\omega) = \int_0^{\infty} I(t) \sin(\omega t) dt / \int_0^{\infty} I(t) dt$$

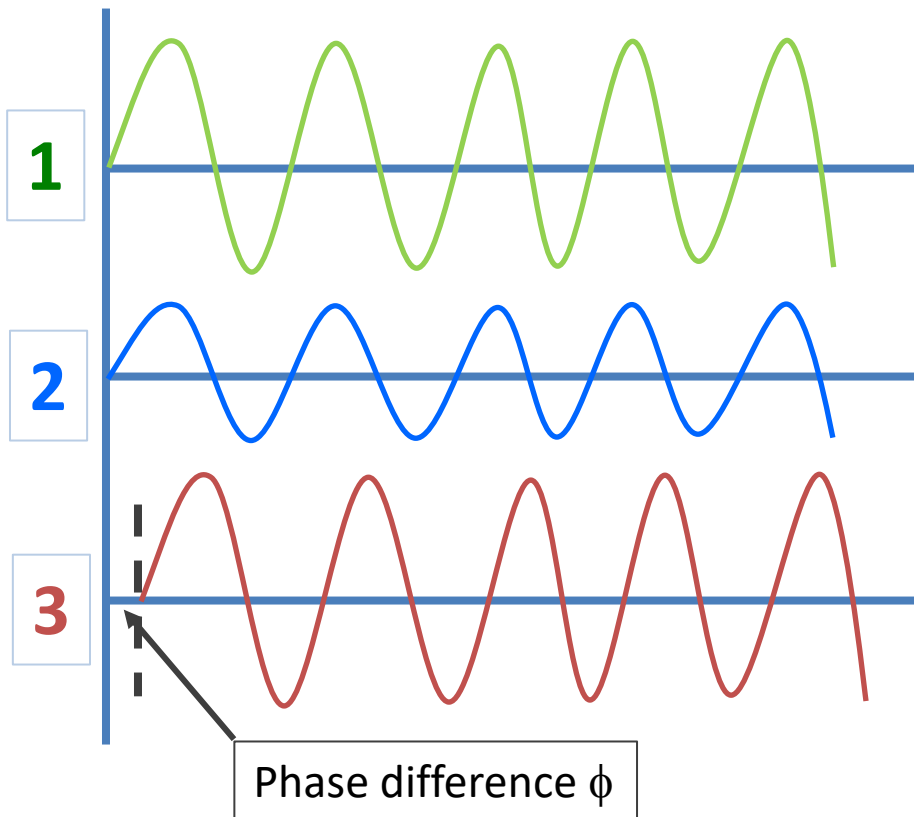
- A phasor is a quantity like a vector. Phasors is the addition of amplitudes using vectors. They can be added. You need to calculate the vector components and then add the components to obtain the vector sum



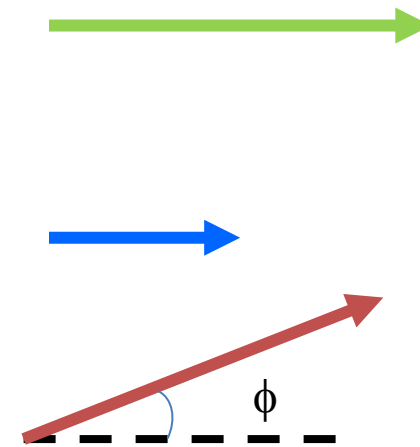
NOTE: s and g are obtained automatically from the Fast Fourier Transform (FFT) of the time domain

The addition of amplitudes using vectors

Wave



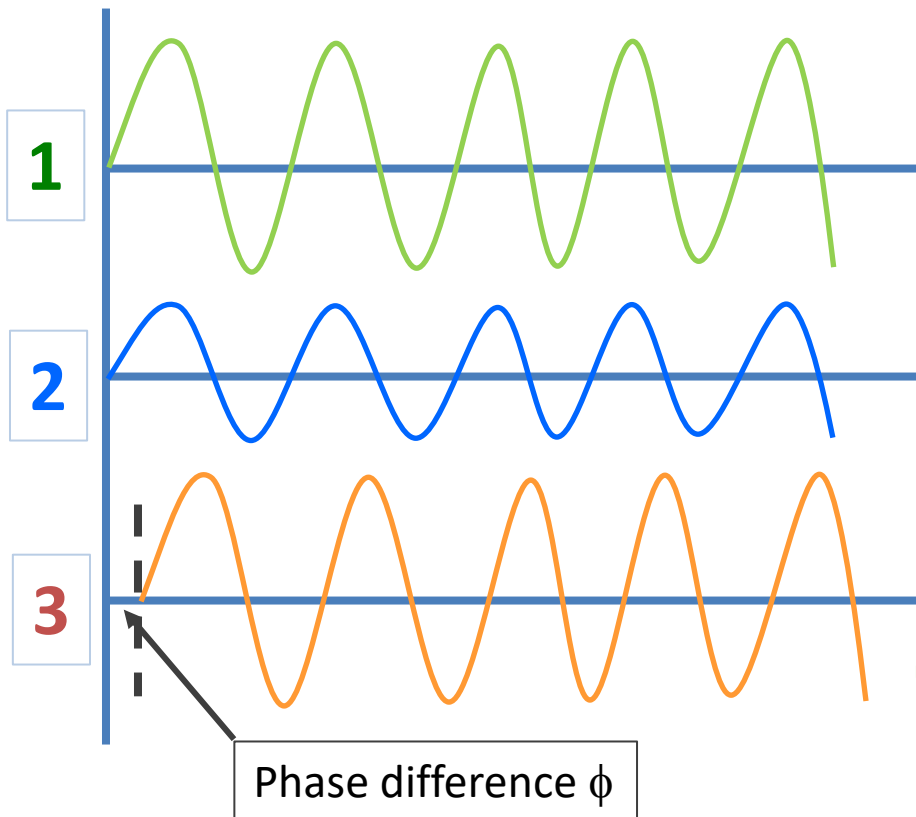
Vector representation



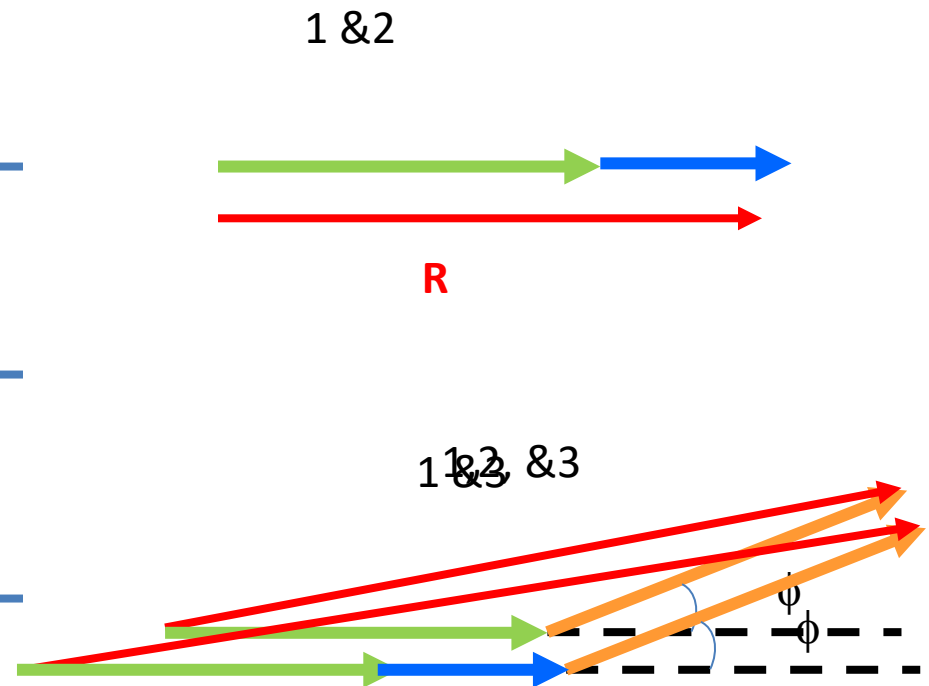
Amplitude of the wave \sim Magnitude or length of the vector
Phase of the wave \sim Direction of the vector

The addition of amplitudes using vectors

Wave



Adding vectors



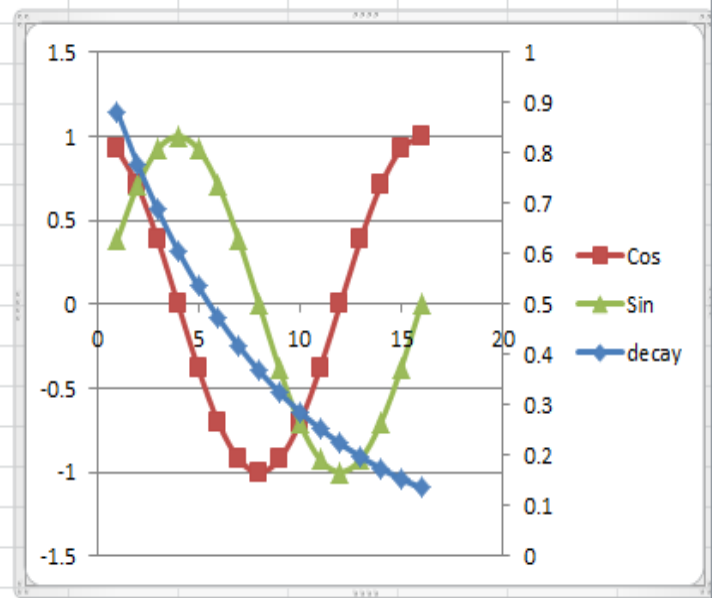
Amplitude of the wave \sim Magnitude or length of the vector
Phase of the wave \sim Direction of the vector

The resultant Vector (**R**) is the
resultant Amplitude

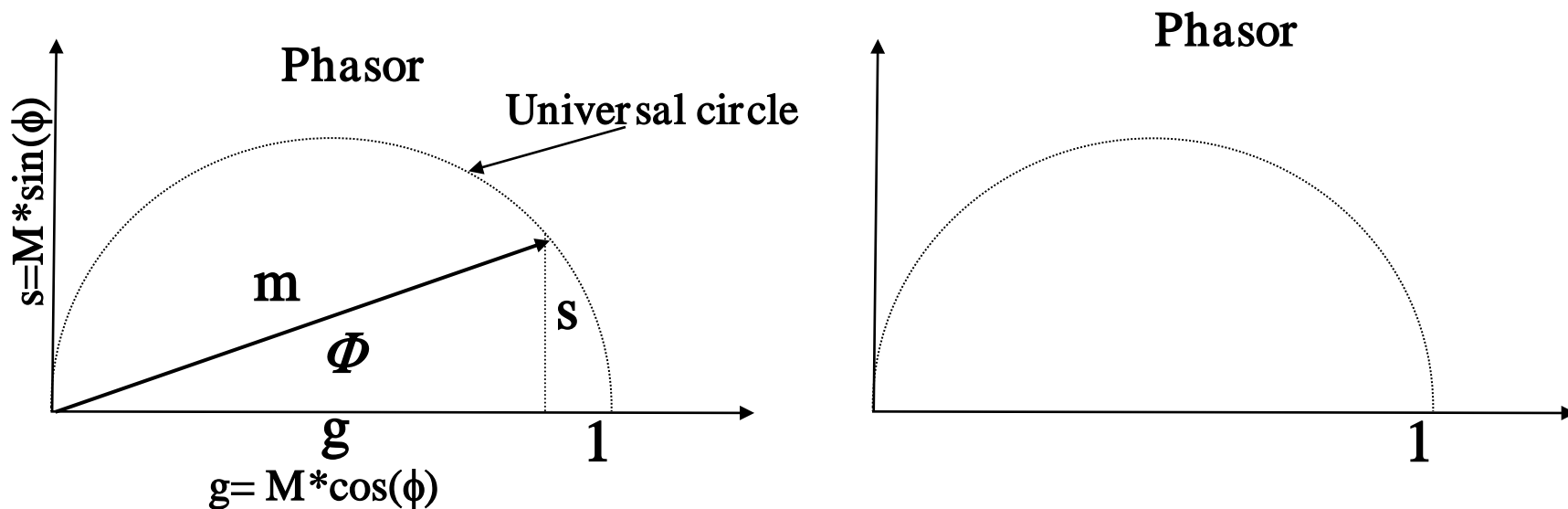
Change Chart Type Save As Template Switch Row/Column Select Data Chart Layouts Chart Styles Move Chart Location

Chart 1

	A	B	C	D	E	F	G	H	I	J	K	L	M	N	O
1															
2		t	i(t)	cos(2pi*f)	sin(2pi*f)		i(t)Cos(2pi*f)	i(t)*sin(2*pi*f)							
3		1	0.882497	0.923879659	0.382683126		0.815320938	0.337716673							
4		2	0.778801	0.70710725	0.707106312		0.55069568	0.55069495							
5		3	0.687289	0.382684352	0.923879152		0.263014852	0.634972236							
6		4	0.606531	1.32679E-06	1		8.04742E-07	0.60653066							
7		5	0.535261	-0.3826819	0.923880167		-0.204834861	0.494517418							
8		6	0.472367	-0.707105374	0.707108188		-0.334012928	0.334014257							
9		7	0.416862	-0.923878644	0.382685578		-0.385129917	0.159527083							
10		8	0.367879	-1	2.65359E-06		-0.367879441	9.76201E-07							
11		9	0.324652	-0.923880675	-0.382680674		-0.299940141	-0.124238225							
12		10	0.286505	-0.707109127	-0.707104436		-0.202590157	-0.202588813							
13		11	0.25284	-0.382686803	-0.923878136		-0.096758377	-0.233592975							
14		12	0.22313	-3.98038E-06	-1		-8.88144E-07	-0.22313016							
15		13	0.196912	0.382679449	-0.923881183		0.075354051	-0.181922991							
16		14	0.173774	0.707103498	-0.707110065		0.122876163	-0.122877304							
17		15	0.153355	0.923877628	-0.382688029		0.141681223	-0.05868711							
18		16	0.135335	1	-5.30718E-06		0.135335283	-7.18249E-07							
19															
20		Sum	6.49399				0.213132286	1.970935956							
21						g and s	0.032819929	0.303501541							



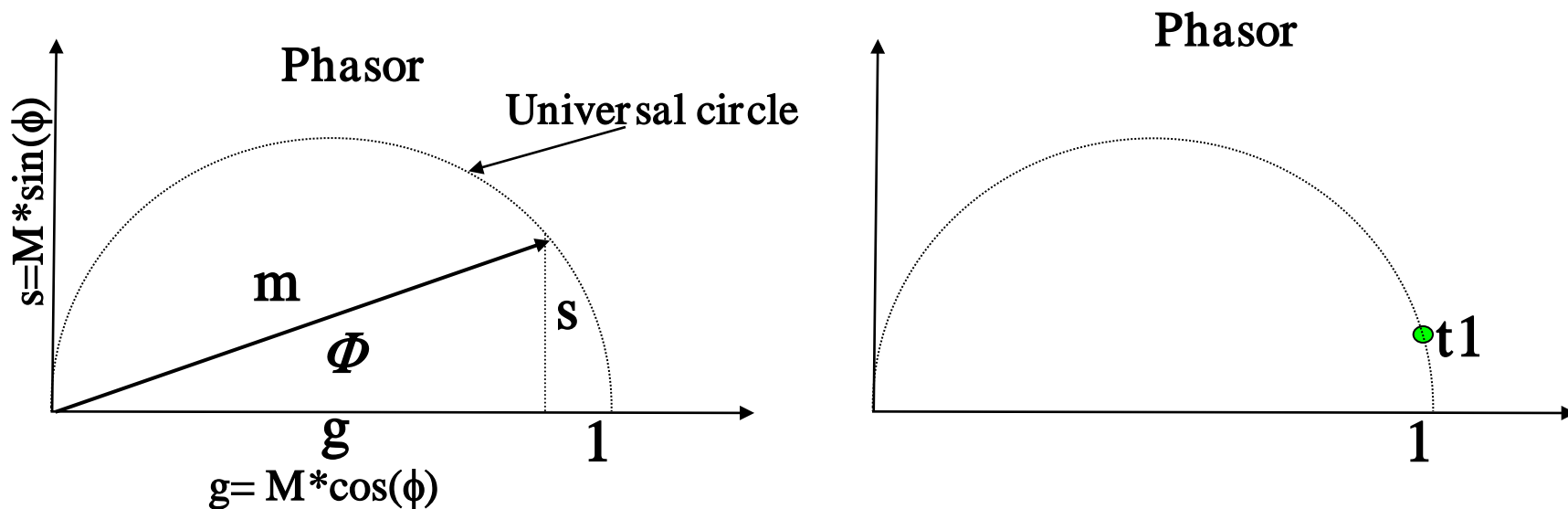
The algebra of phasors: A fit free method



Simple rules to the Phasor plot:

- 1) All single exponential lifetimes lie on the “universal circle”
- 2) Multi-exponential lifetimes are a linear combination of their components
- 3) The ratio of the linear combination determines the fraction of the components

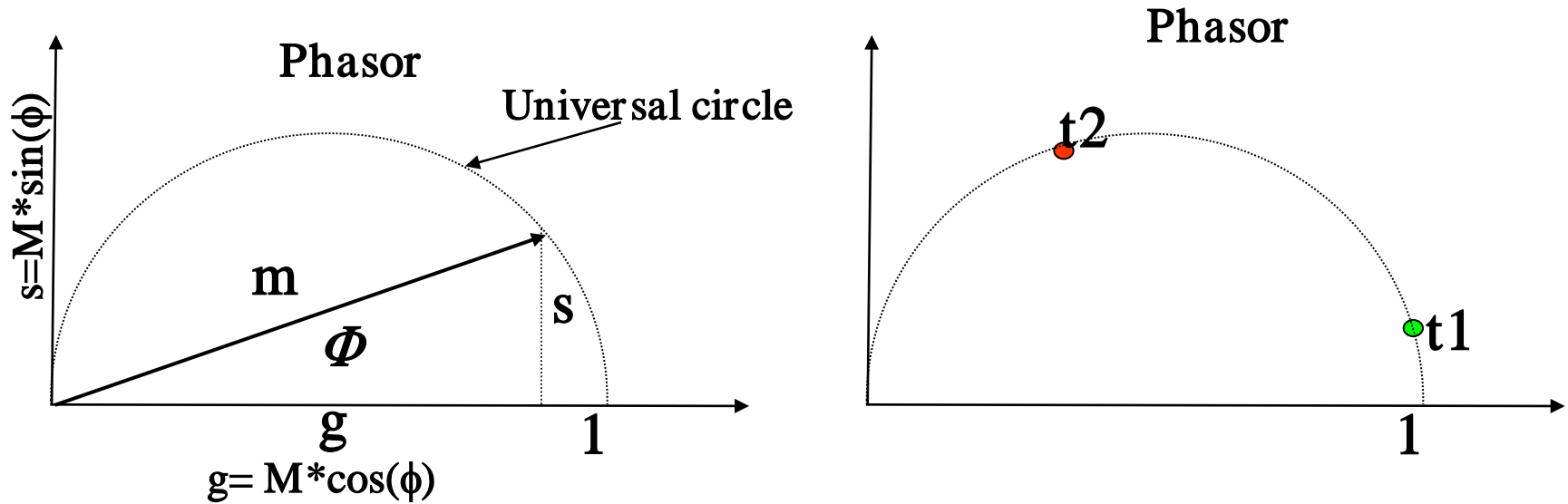
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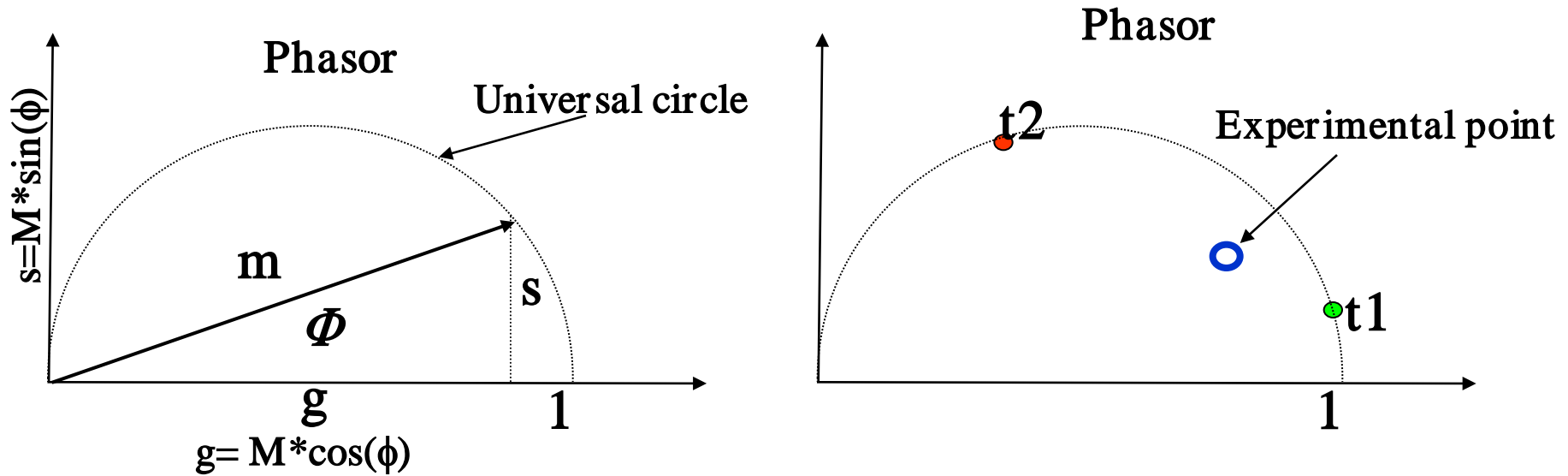
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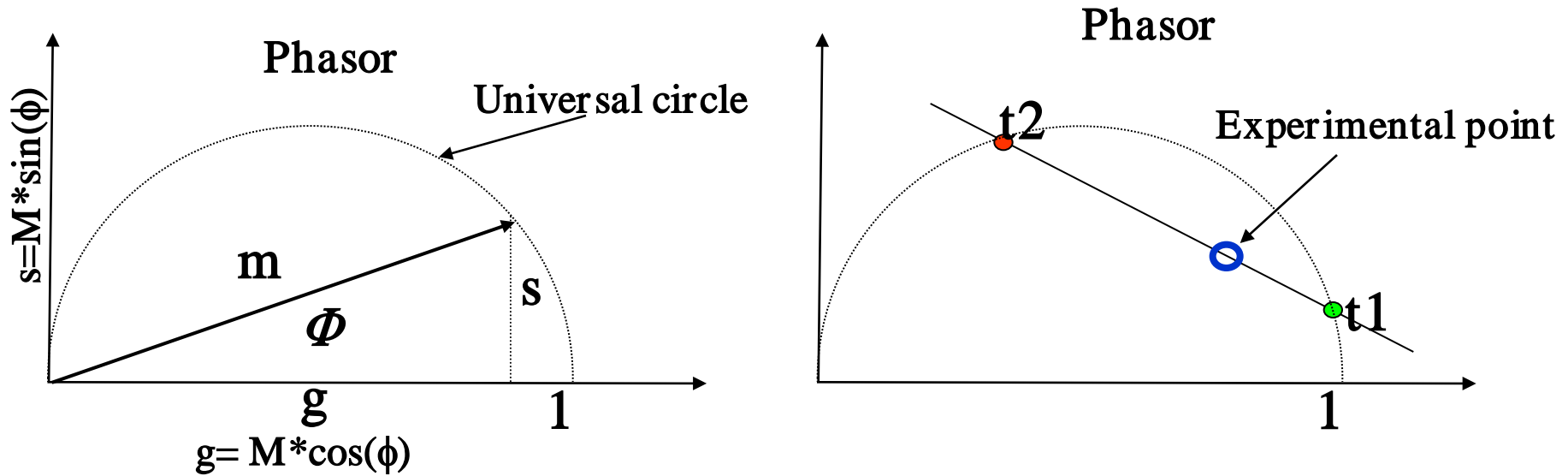
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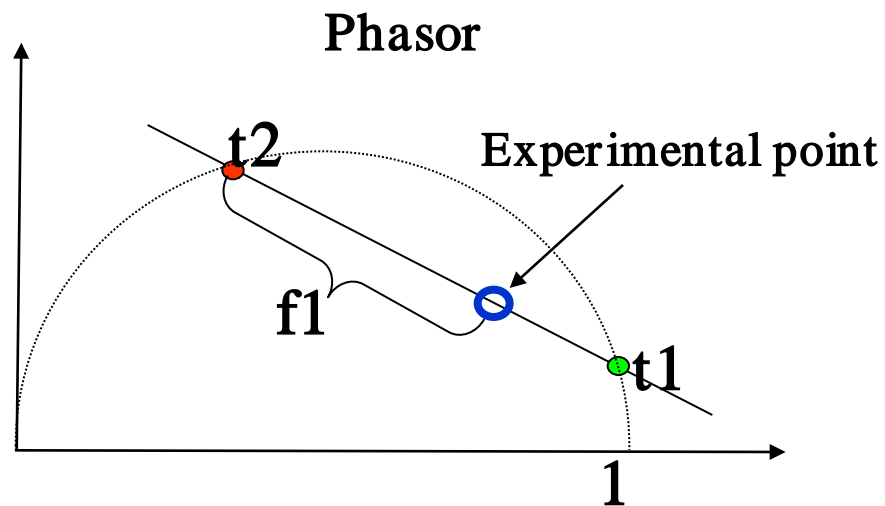
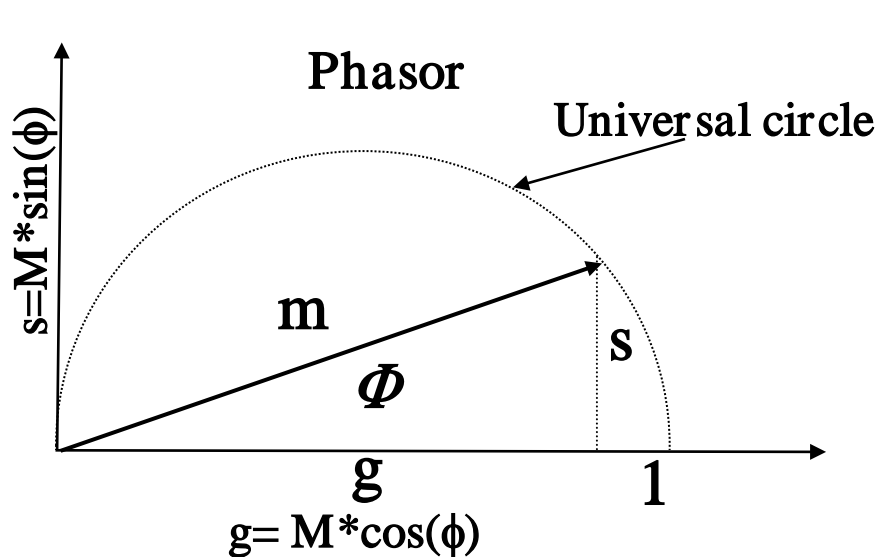
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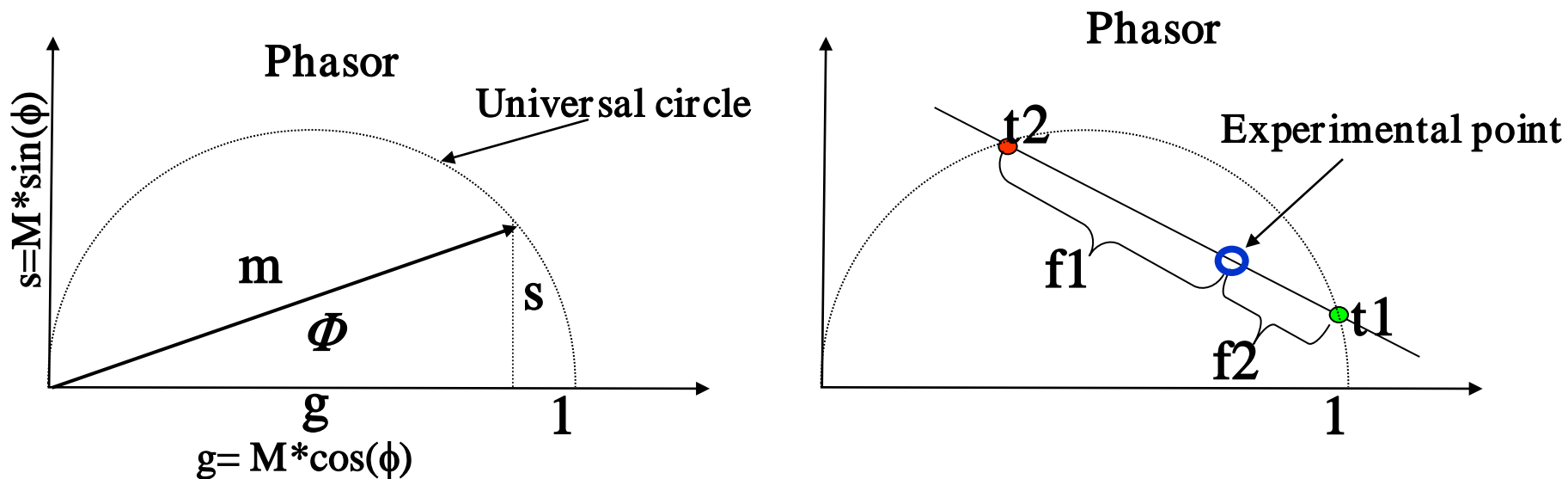
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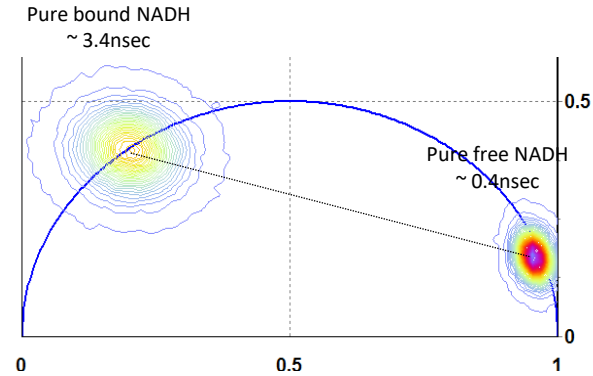
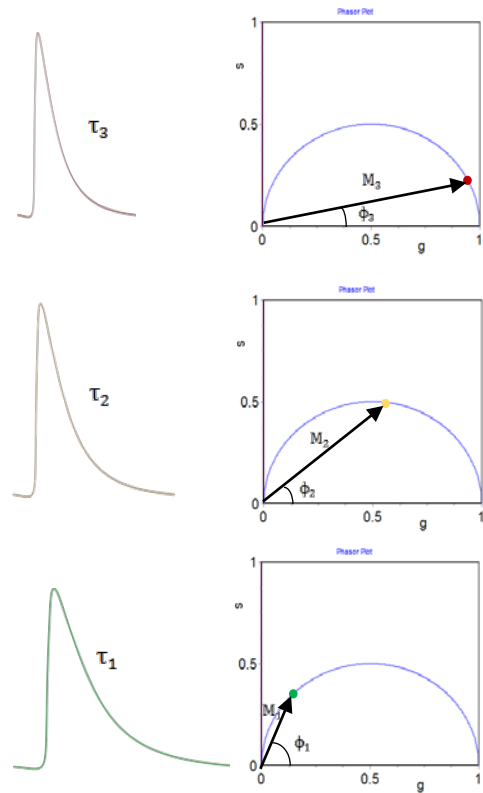
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Phasor approach quickly reveals %Bound NADH

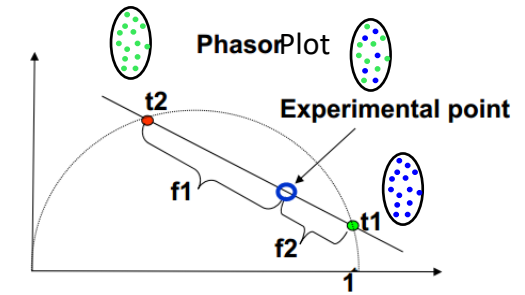
Phasors allow us to visualize data easily and efficiently, in our case the free to bound ratios of NADH.

- * Each phasor represents the lifetime data for one pixel from the observation.
- * Lifetimes are represented as points, called phasors, on the diagram.

Each pixel on the acquired image is converted through Fourier Transform to create coordinates in phasor space.

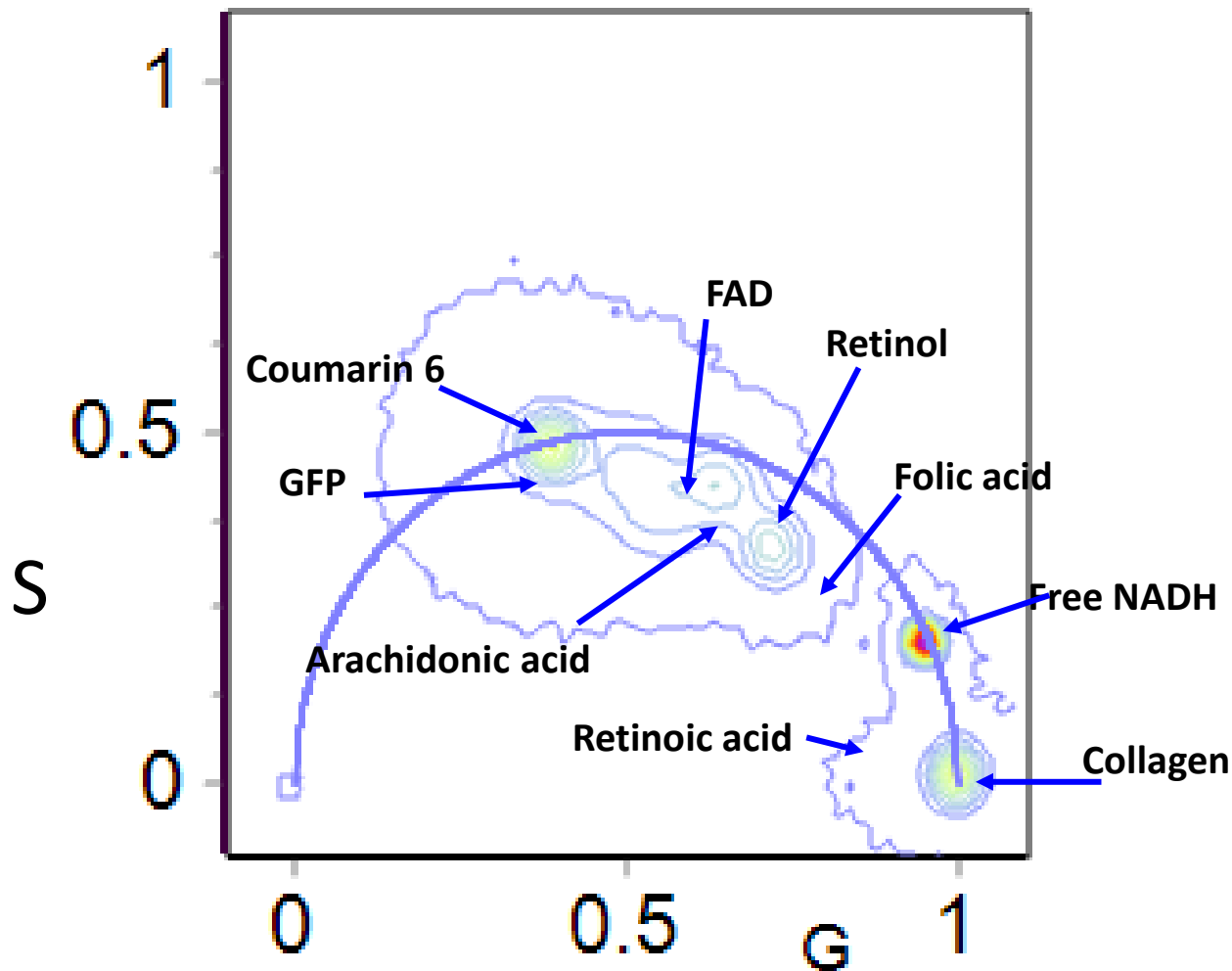


Law of linear addition of phasors:



$$\text{Fraction of Bound NADH in Experimental Point} = \frac{f_2}{f_1 + f_2}$$

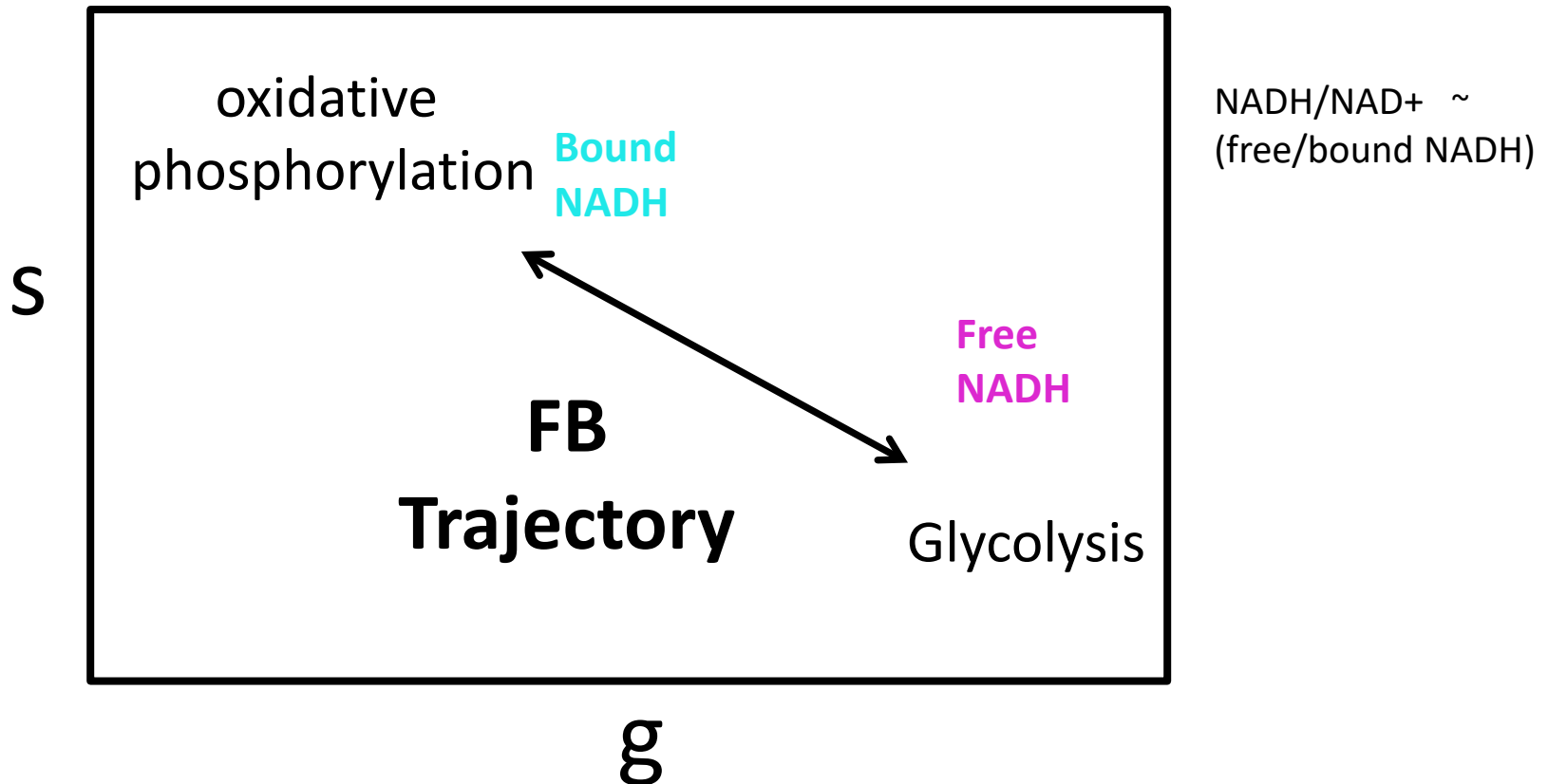
Phasor Fingerprint of pure chemical species....



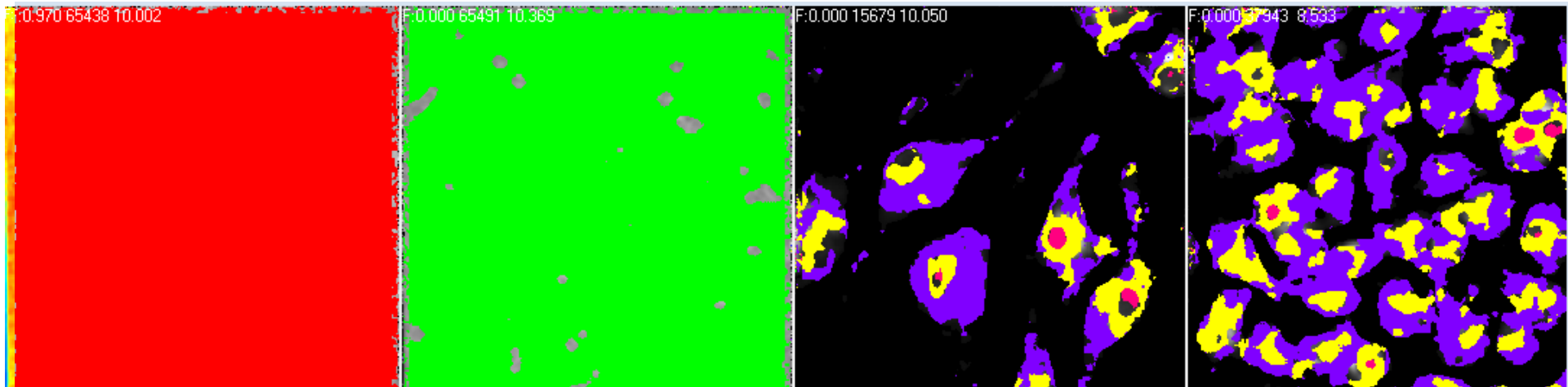
Metabolic Trajectory in the Phasor Plot

Free/bound NADH gradients associated with:

- Glycolysis/oxidative phosphorylation
- Oxidative stress
- cell proliferation
- differentiation
- Cancer



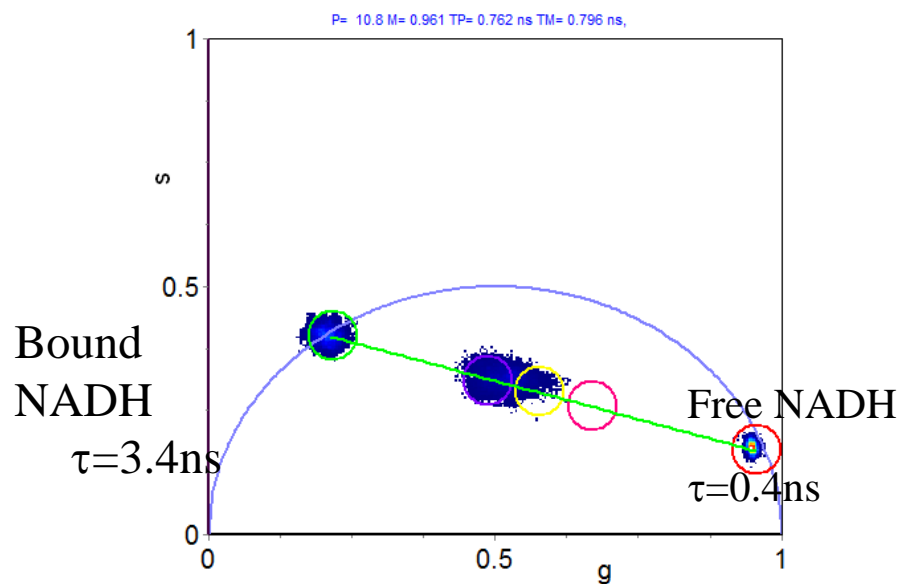
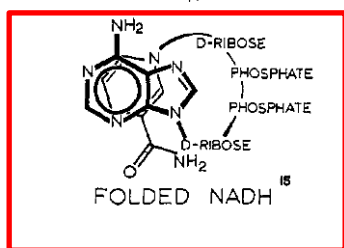
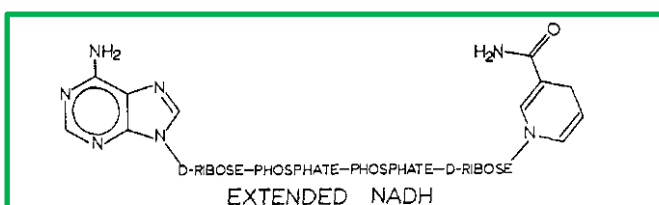
Linear combination of free and bound NADH



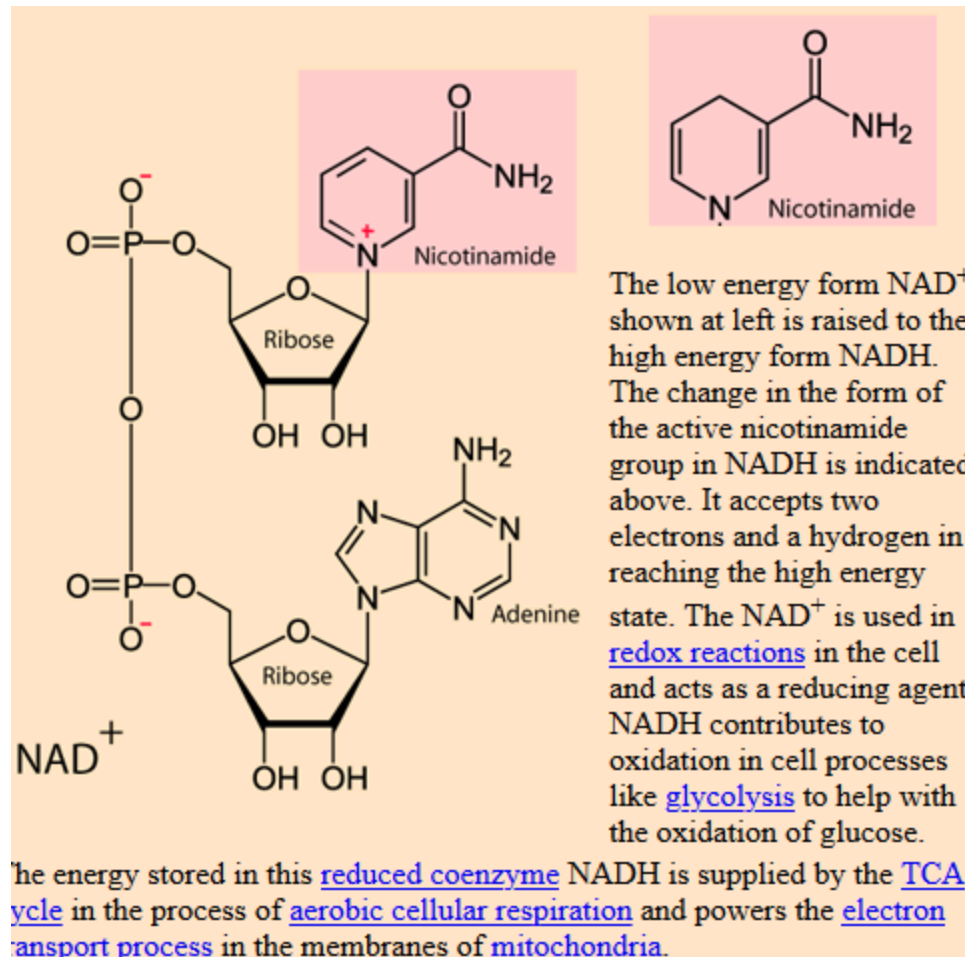
Free NADH

Bound NADH to LDH

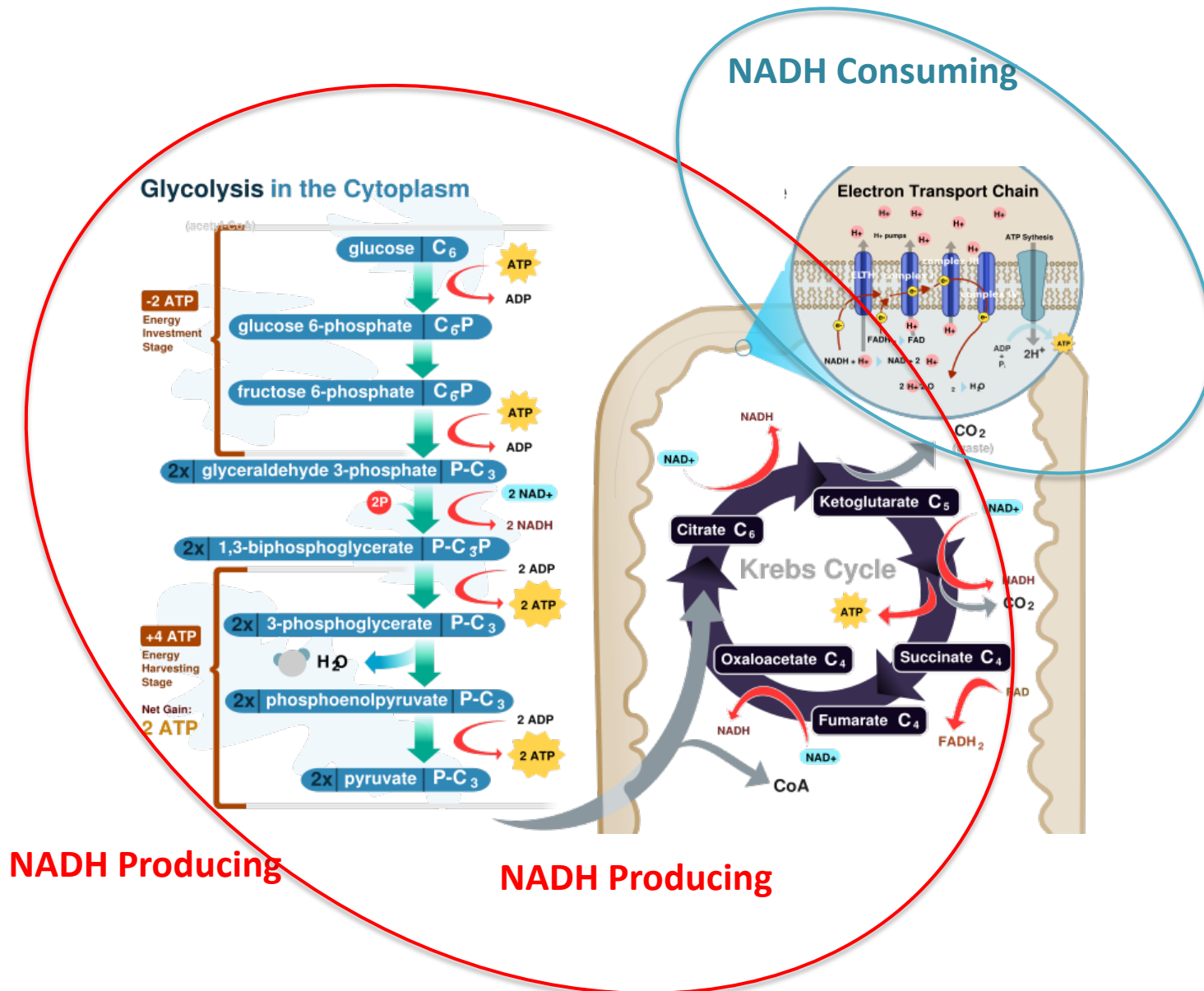
M0 macrophages on glass



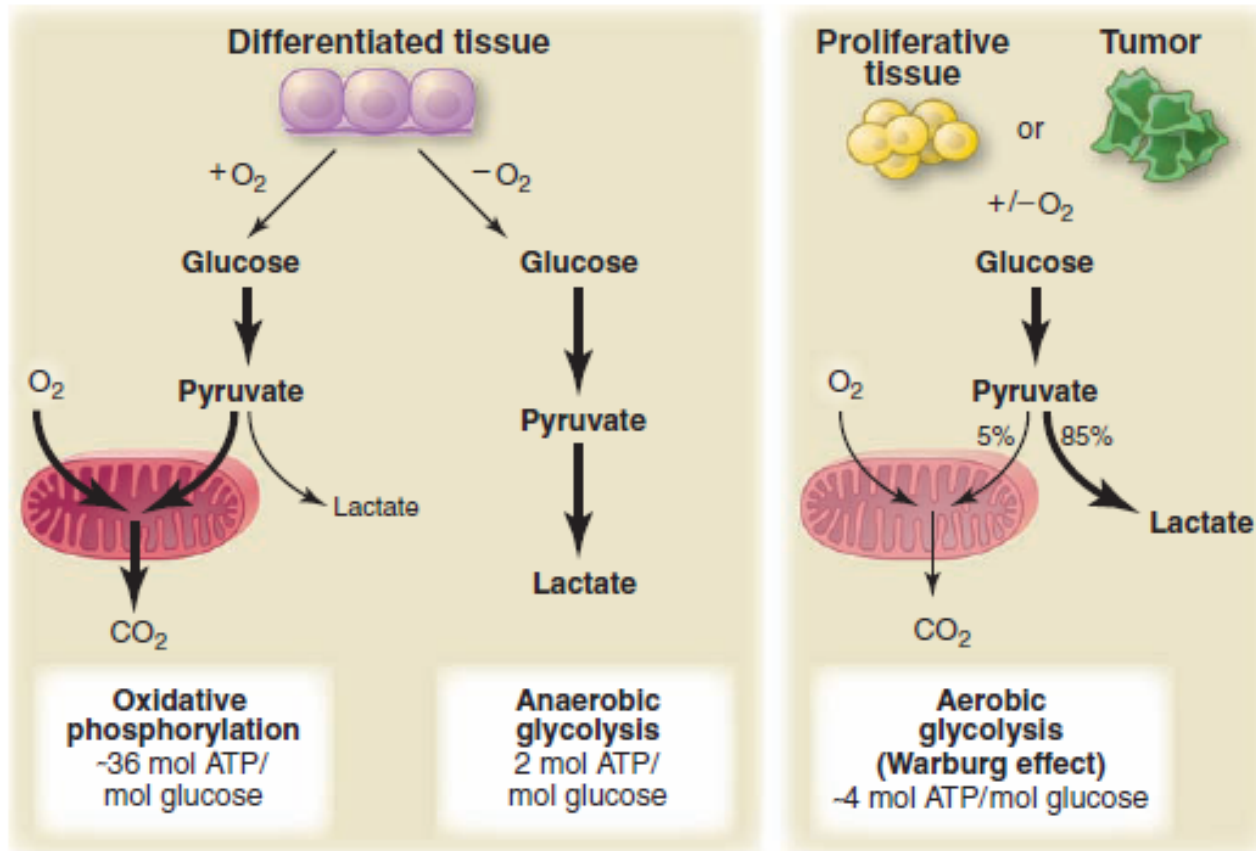
Why is NADH fluorescent while NAD⁺ is not?



Phasor FLIM measurement of Free/Bound NADH in tissue to assess metabolism



Metabolism in tissues



LOW NADH/NAD⁺
(LOW free/bound NADH)

HIGH NADH/NAD⁺
(**HIGH** free/bound NADH)

Cairns et al. Nature reviews Cancer 2011

Heiden, et al. Science 2009

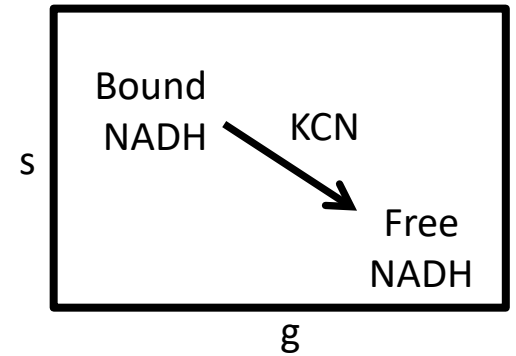
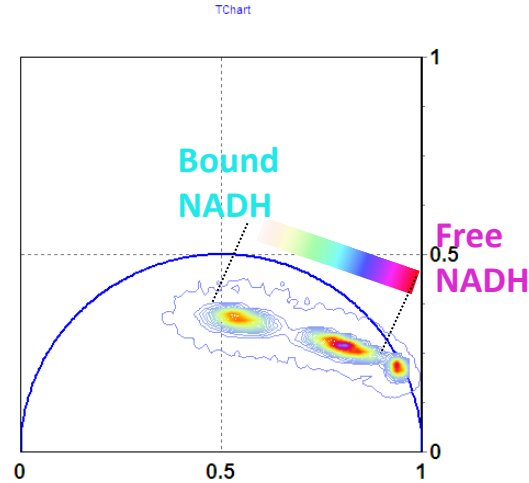
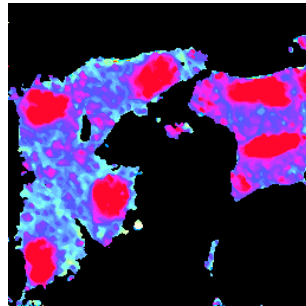
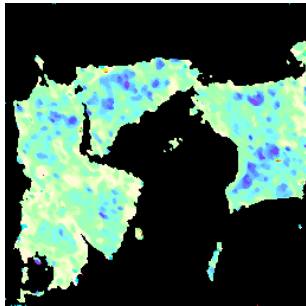
Metabolic Trajectory

Oxidized NAD⁺/ Reduced NADH --> Free/bound NADH

Electron chain
inhibition

Before

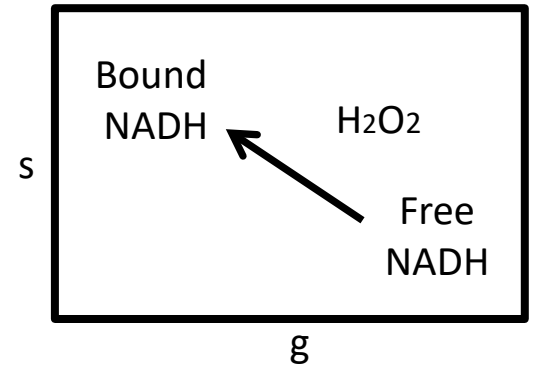
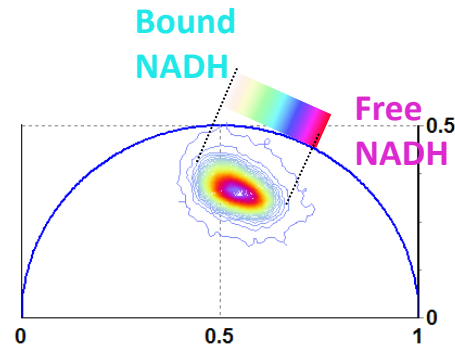
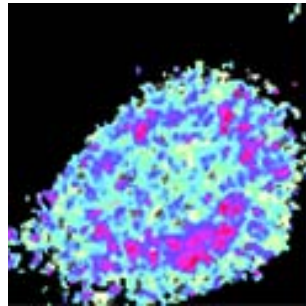
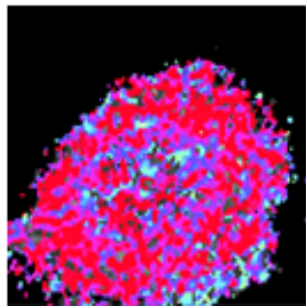
KCN



Oxidative
stress

Before

H₂O₂

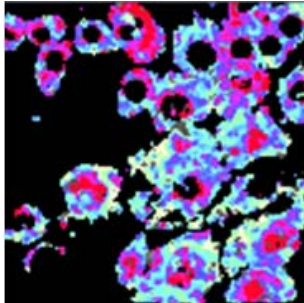


Metabolic Trajectory

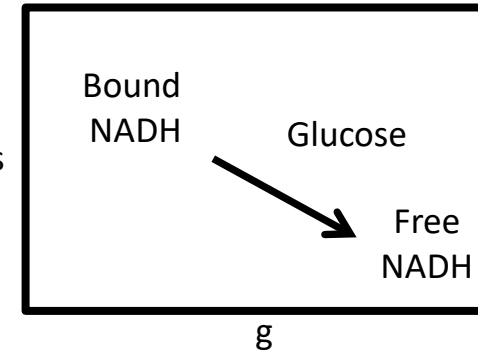
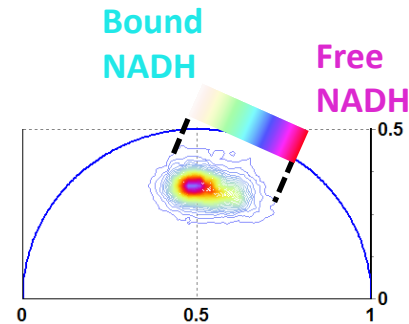
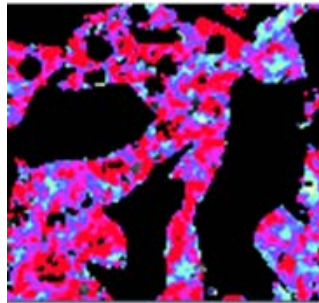
glycolysis/ oxidative phosphorylation --> Free/bound NADH

Glucose uptake

4.5 mM glucose

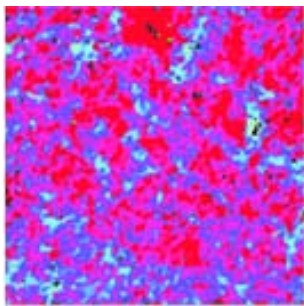


22 mM glucose

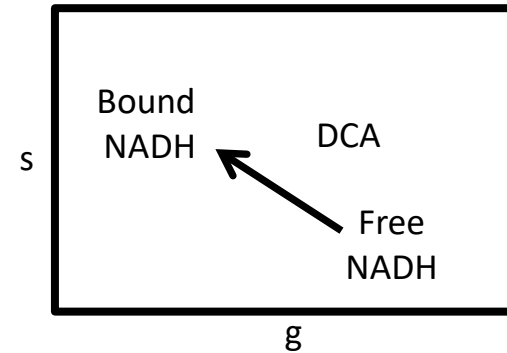
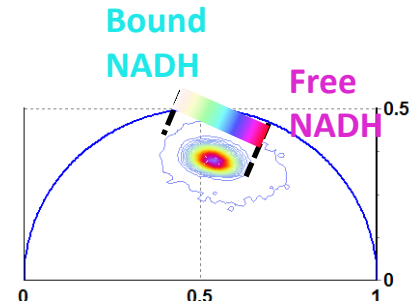
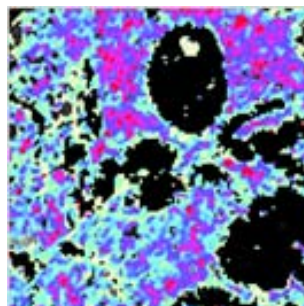


Glycolysis inhibition
Dichloroacetate (DCA)
metabolic-targeting
cancer drug

control



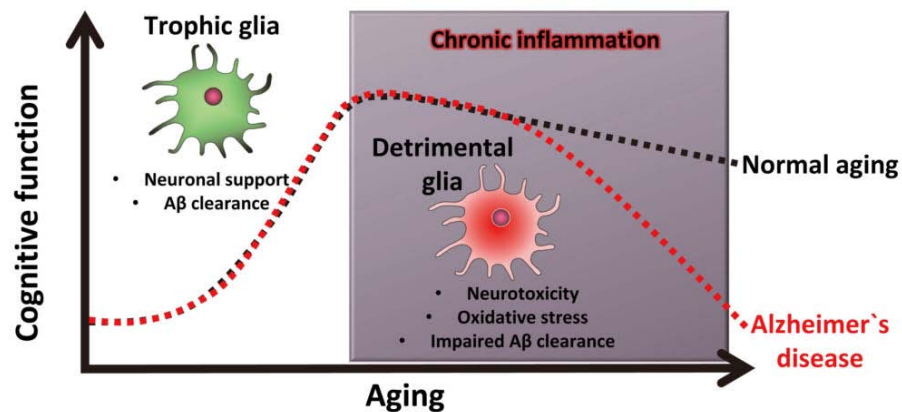
DCA



The Metabolic trajectory in aging and Alzheimer's Disease

Driving Biological Question:

- The Epigenetic Oxidative Redox Shift (EORS) theory of aging proposes that a sedentary low-energy state with aging triggers an oxidative shift and mitochondrial dysfunction* and further causes metabolic disturbances

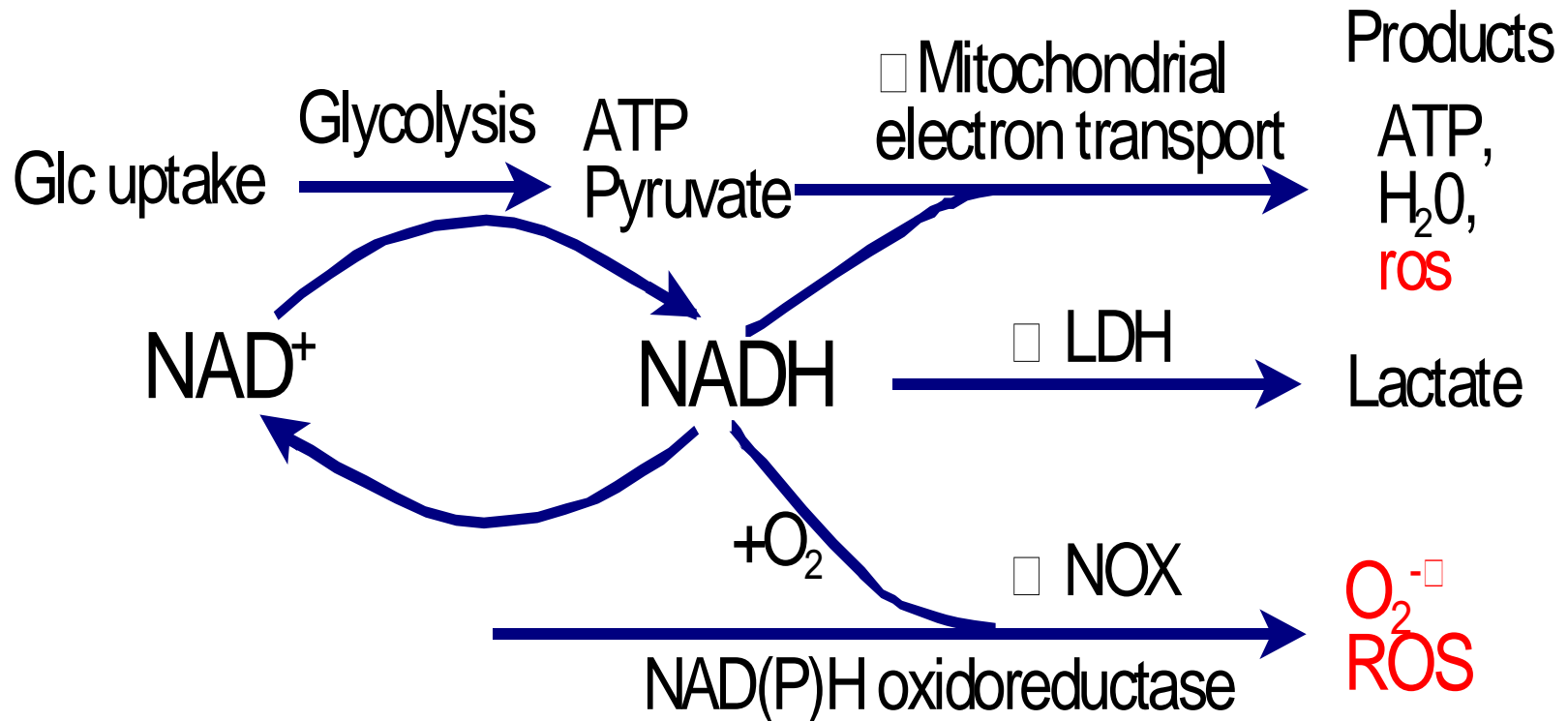


Aging is associated with chronic inflammation, which has been identified as both a key contributor to the initiation and progression of Alzheimer's disease

- Our hypothesis: free NADH in neurons from old mice is lower than the levels in young mice and even lower in neurons from the 3xTg-AD Alzheimer's disease (AD) mouse model

*due to the accumulation of free radical damage over time

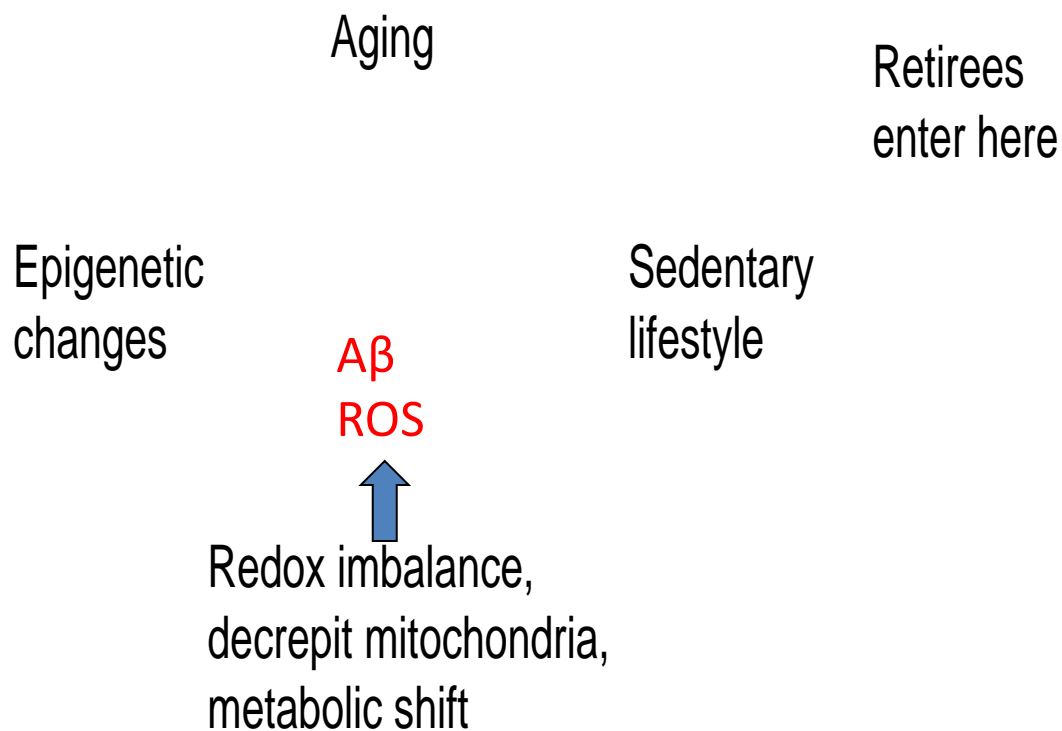
Three alternative mechanisms for energy production that maintain NAD⁺ levels for glycolysis and redox balance



Brewer 2010

Dilemmas of current aging theories		
Aging Paradigm/Observations	Impact	Dilemmas
1) Oxyradical damage to macromolecules (Harman, 1952)	Cumulative cellular damage leads to organ failure, shortened life-span	Exercise extends lifespan but produces ROS (Powers SK 2008) Ignores insulin signaling SOD, catalase Xgenics(Perez VI 2009)
2) Genetic reductions in the insulin signaling path (Carter CS 2002) or caloric restriction (Weindruch R 2001)	Increased life-span, may decrease metabolism = “slower” rate of living (Pearl R 1928)	Metabolism actually increases (Westbrook R 2009) Ignores role of ROS
3) Both		Ignore healthy aging, energetics and ROS signaling vs. ROS inflammation

EORS, a vicious cycle of aging enforced by an Epigenetic
Oxidized Redox Shift
unifies the free radical and insulin signaling theories



- Aging and AD share some common characteristics such as:
 - oxidative stress
 - mitochondrial impairment
 - bioenergetic deficits

Clinical features of Alzheimer's Disease and Frontotemporal Dementia




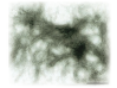



- | | | |
|---|--|-----|
| <ul style="list-style-type: none">• Neuronal/synapse loss at specific predilection sites• Early memory deficits• Subdivided into behavioral and language variants• 14% parkinsonism• Memory loss in advanced stages | | AD |
| | | FTD |

Histopathologically, AD is characterized by $A\beta$ plaques & neurofibrillary lesions.

$A\beta$ in the plaques is fibrillar.

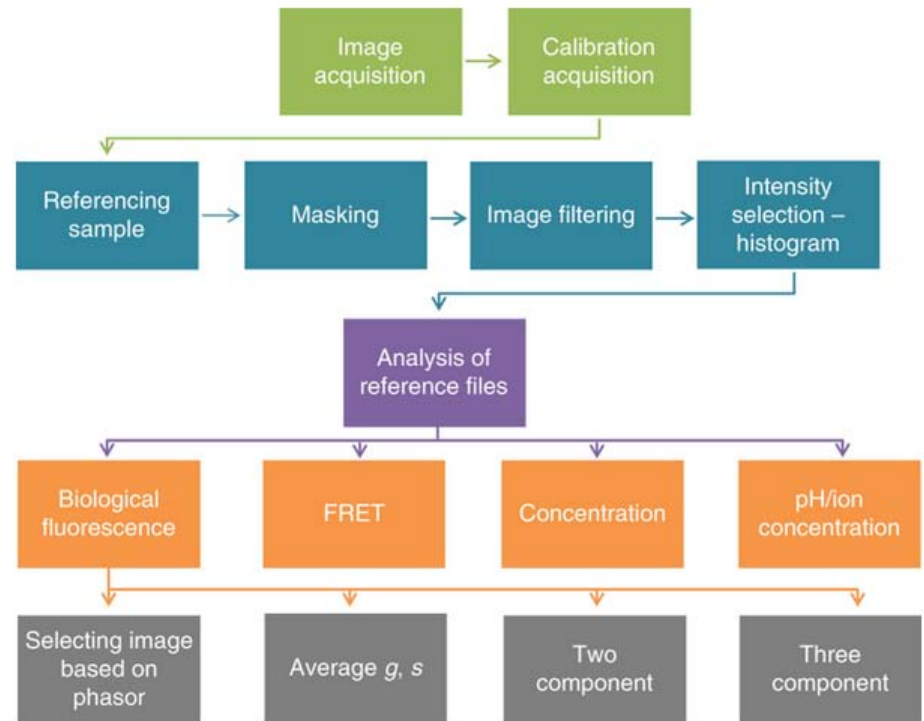
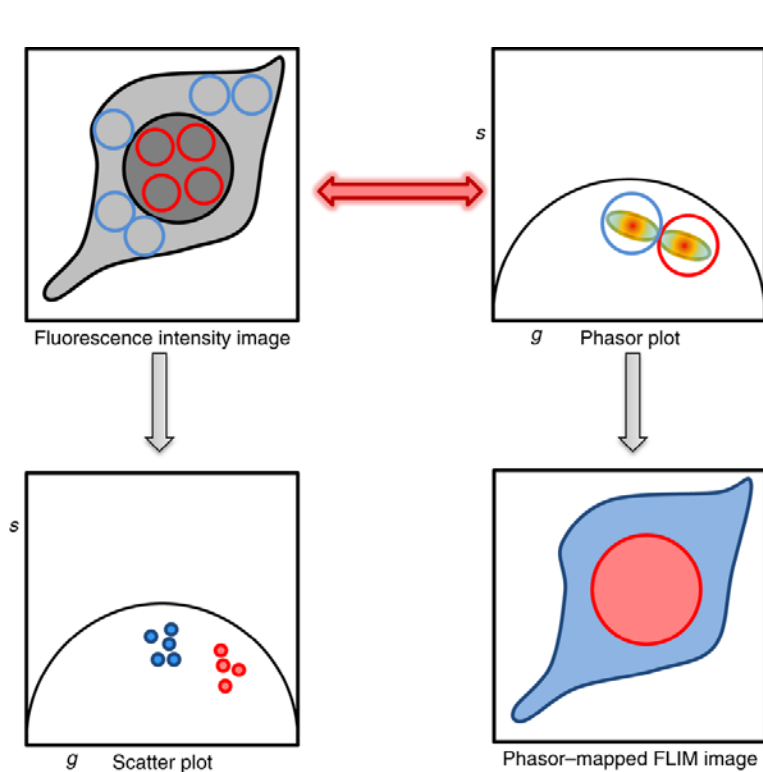
The 3xTg-AD mouse model: combined A β plaque and NFT (neurofibrillary tangles) pathology

Tau	X	A β	Observations	References
A. JNPL3 (P301L)		Tg2576 (A β^{sw}) 	<ul style="list-style-type: none"> • 7x NFT induction • No increase in Aβ pathology 	Lewis et al., Science 2001
B. pR5 (P301L)		i.c. A β 	<ul style="list-style-type: none"> • 5x NFT induction 	Gotz et al., Science 2001
C. 3xtg-AD (P301L/APP ^{sw} /PSEN1 ^{M146/-} ki)			<ul style="list-style-type: none"> • NFTs and plaques preceded by synaptic/LTP deficits • Correlated with intra-neuronal Aβ • Blocking Aβ delays onset/progression of tau pathology 	Oddo et al., Neuron 2003; Oddo et al., J Neurosci 2008; Billings et al., Neuron 2005

- Decreased mitochondrial numeric density (number of mitochondria/ μm^3 of cytoplasm) in 10-month-old mice but the volume and average length increased
- decreased mitochondrial number with age in the hippocampus along with abnormal mitochondrial morphology with elongation, swelling and disorganized cristae
- decline in the motility of mitochondria in the due to changes in Drp1 levels for fission

*these mitochondrial structural changes reflect functional deficits of free NADH as a bioenergetic substrate for oxidative phosphorylation (OXPHOS) that initiates a vicious cycle involving redox imbalance, energetic shortage, and mitochondrial dysfunction.

Image segmentation and phasor compartment average:

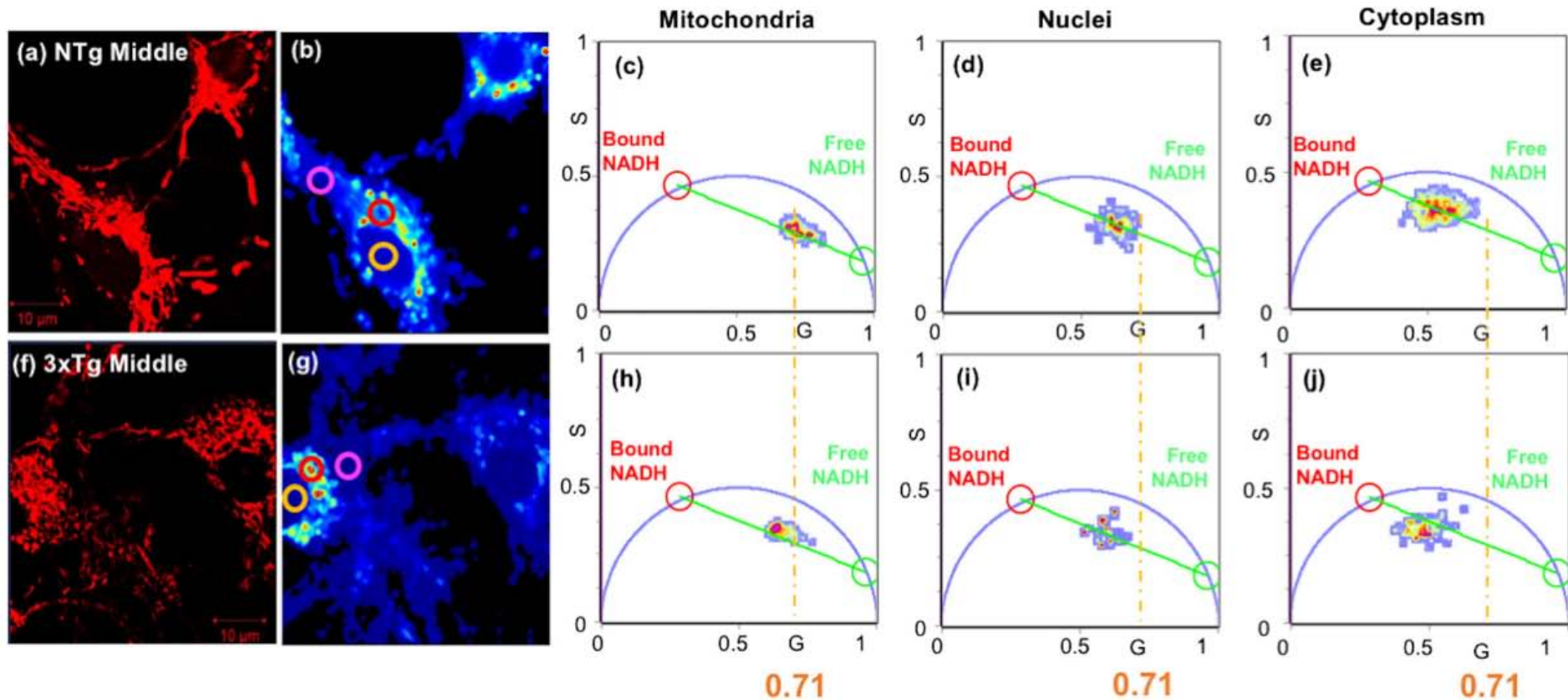


Schematic overview of the steps involving FLIM analyses according to the phasor approach.

Fit-free analysis of fluorescence lifetime imaging data using the phasor approach

Ranjit et al, Nature Protocols 3, pages1979–2004 (2018)

AD genotype shows lower free NADH fraction than the corresponding compartment levels of *NTg control* mice neurons.

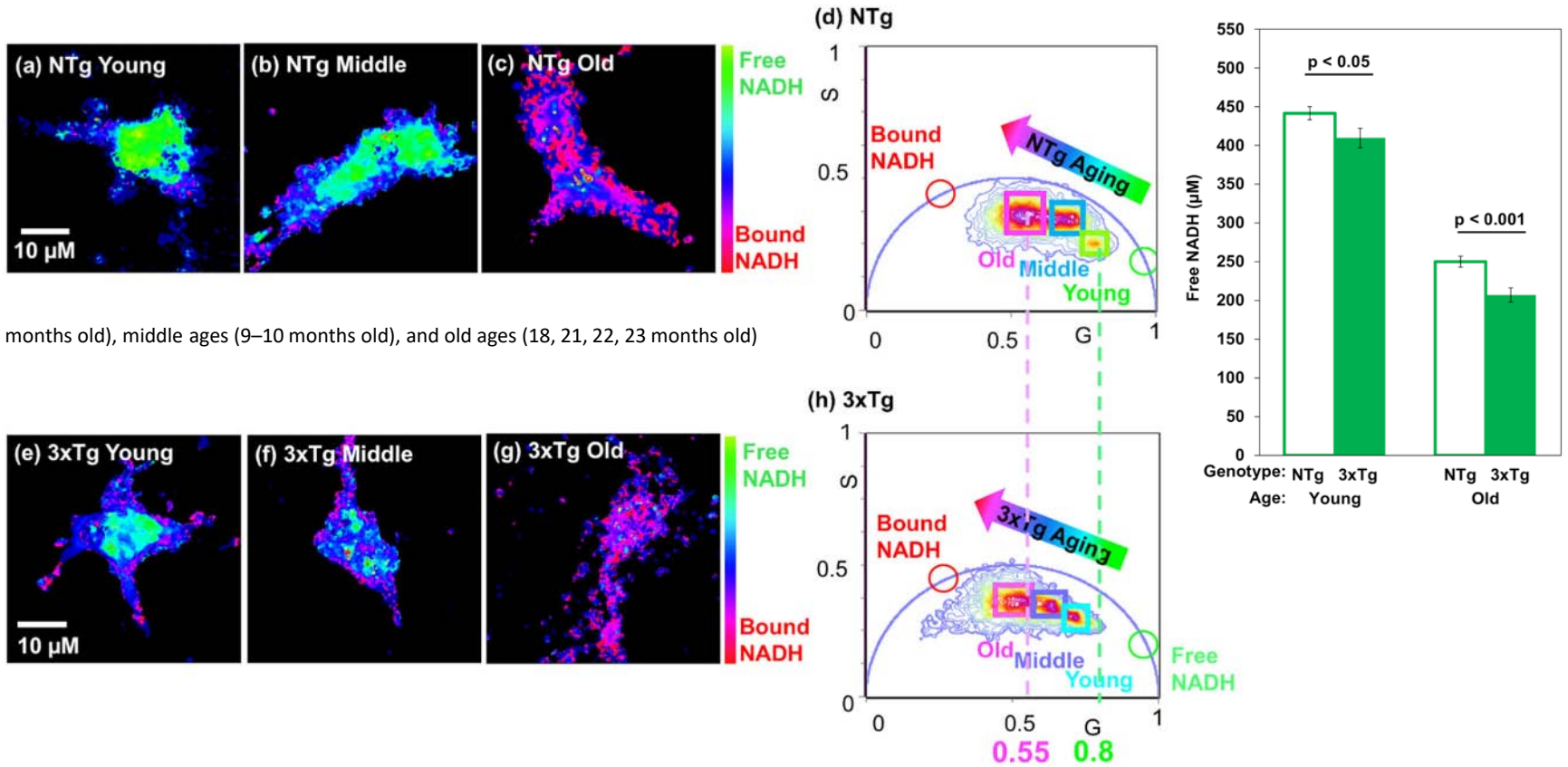


The vertical orange lines are centered on the NTg young mitochondrial distribution for reference at $G=0.71$

Free NADH highest in mitochondria > nuclei > cytoplasm

Free NADH levels lower in 3xTg-AD genotype neurons than NTg

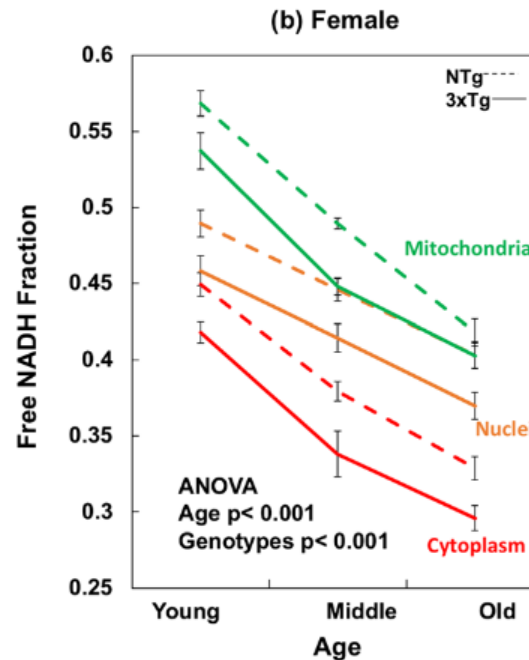
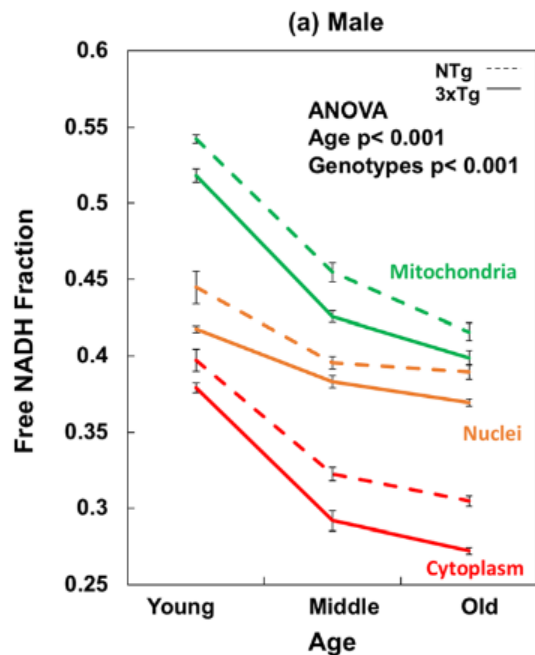
Aging promotes shifts toward more bound and less free NADH in NTg control and 3xTg-AD neurons



young (3–4 months old), middle ages (9–10 months old), and old ages (18, 21, 22, 23 months old)

NTg 43% and 3xTg 50% drop in mitochondria free NADH available for oxidative phosphorylation with aging

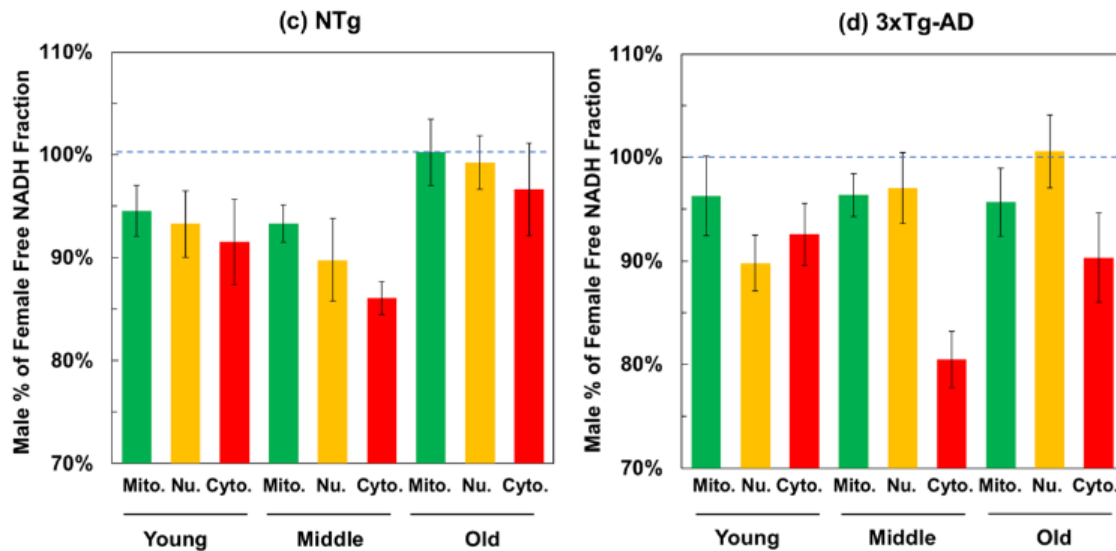
Aging strongly drove a decline in the free NADH fractions in all subcellular compartments in both genotypes **and both genders**.



Mitochondria

- The relative order of compartmental free to bound NADH redox states was independent of age and genotype.
- Overall, the largest changes in free NADH were associated with age, then compartment, genotype, and gender
- The highest free NADH fraction in mitochondria suggests that mitochondria have the highest capacity for free NADH production compared to nuclei and cytoplasm.

Aging depleted free NADH levels in all compartments of both genotypes: Free NADH fraction only 5-10% in male



- 1) By gender: male neurons were lower in free NADH fraction than neurons at young and middle ages, but the multiple comparison results indicated no significant differences in old ages between female and male neurons
- 2) There is gender differences in the young and middle age of all compartments, but no significant gender differences at old ages. This suggests that higher reductive capacity in females than males is gradually lost in old age.

Summary and Conclusions

Age- and AD-dependent decline in free NADH

- The lower free NADH with age and further decrease with AD genetic load can be due to a decrease in enzymes concentration which produce NADH in the TCA cycle. AD cells are known to have declines in these dehydrogenases that would lead to lower free NADH production in mitochondria.
- As compensatory effects, pathways may redirect fluxes to replenish the decreasing NADH levels. Nicotinamide nucleotide transhydrogenase reverses the direction of reaction to generate more NADH at the cost of NADPH. This begins a vicious cycle because NADPH is needed to regenerate GSH as the major redox buffer, when GSH levels are decreased by age-associated oxidative stress. Age- and AD-related oxidative shifts stimulate up-regulation of glycolytic pathway to feed the NADH & energetic demand.
- We found that levels of free NADH in male neurons were significantly lower than those of age and genotype matched female mouse brains, particularly in the cytoplasmic compartment. This protective effect of higher free NADH levels in female neurons appears to lessen with age, ending in similar free NADH levels in old neurons of female and male.

GeroScience (2019) 41:51–67

<https://doi.org/10.1007/s11357-019-00052-8>



CrossMark

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ORIGINAL ARTICLE

Age- and AD-related redox state of NADH in subcellular compartments by fluorescence lifetime imaging microscopy

the
tela

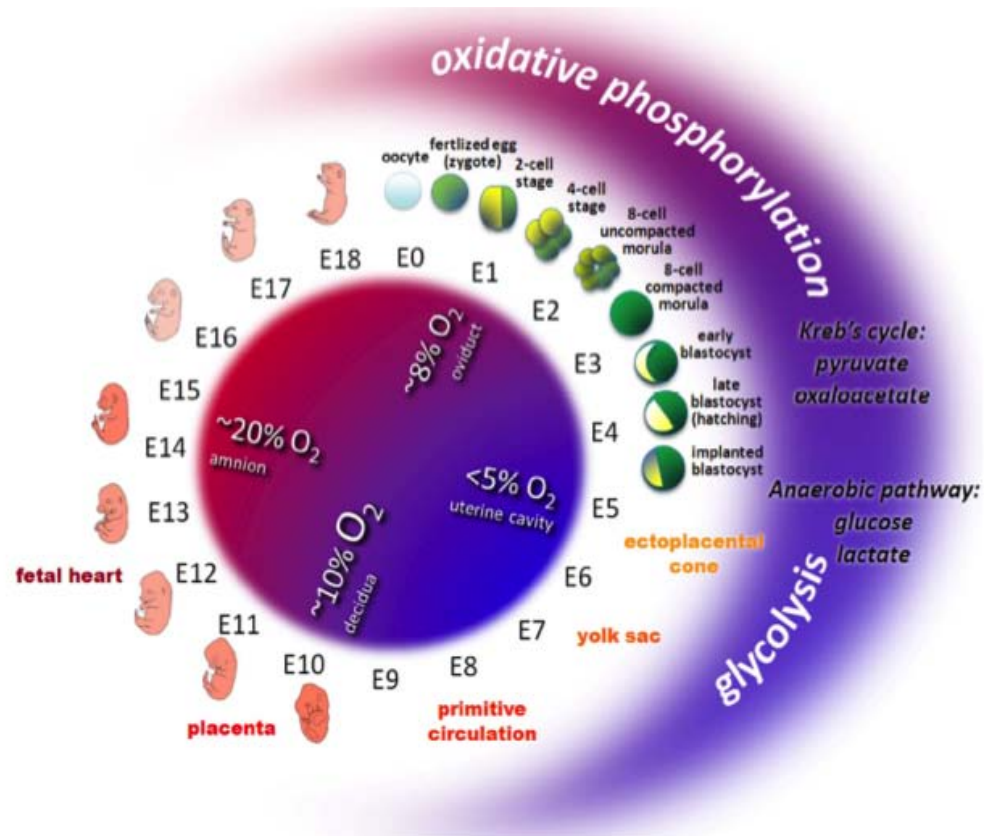
Yue Dong • Michelle A. Digman • Gregory J. Brewer

Future Directions

1. How could age-related deficits in intracellular NADH be assessed in the elderly non-invasively and cost-effectively? (red blood cells transdermal?)
2. How could intracellular NADH be elevated orally and can they be quantitatively controlled?
3. Can elevated intracellular NADH slow progression of AD, stall conversion from MCI to AD or improve cognitive function in MCI subjects?
4. Will supplementary intracellular NADH rescue the observed age- and AD-related metabolic shifts implicated in AD?

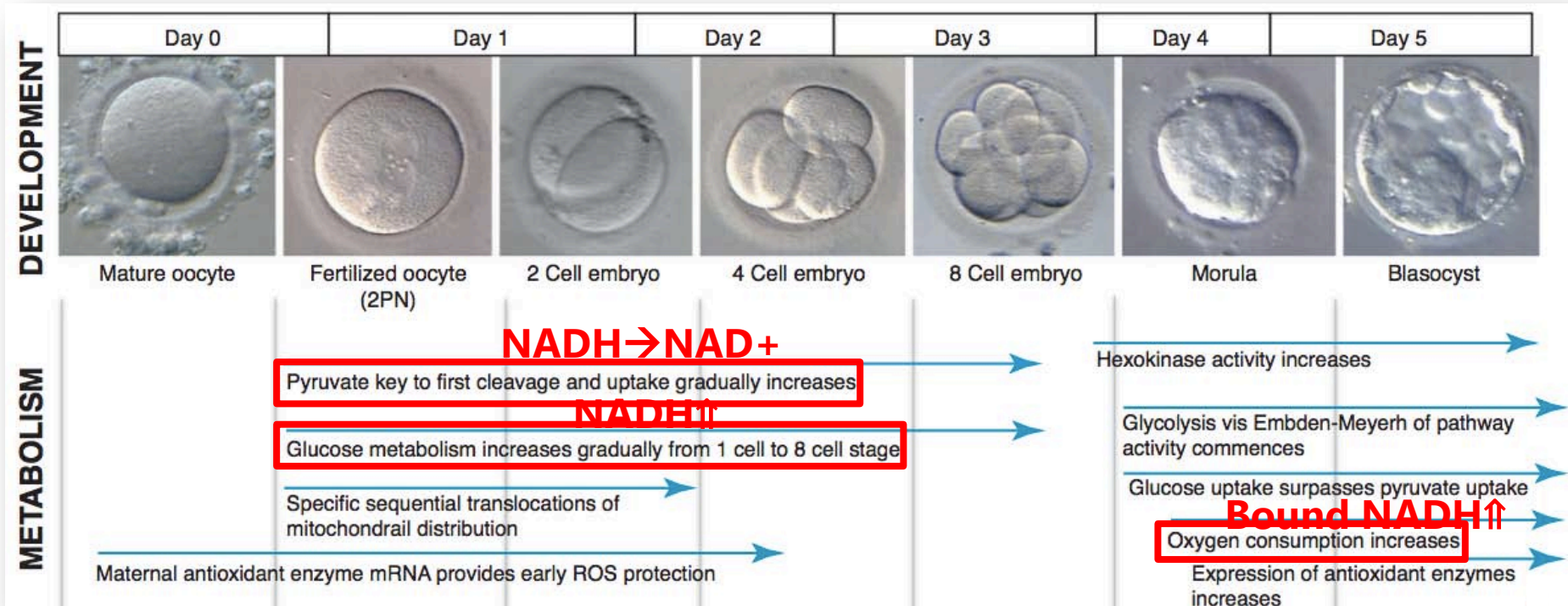
Metabolism During Mouse Embryo Development

1. From the 2-8 cells stage metabolism is pyruvate based (oxidative phosphorylation).
2. Hexokinase is inhibited in the first stage of embryo development then around the morula stage there is an increase in glucose uptake and increase activity of hexokinase thereby making glycolysis the major source of energy production



Ufer and Wang. Front. Mol. Neurosci., 28 July 2011

Activity of Metabolic Cofactor NADH during Pre-implantation Embryo Development



- Nicotinamide adenine dinucleotide(NADH)
- Endogenous fluorescent
- Plays a key role in production of energy in cells and tissues
- Oxidized form (NAD⁺) of the coenzyme does not fluoresce

Metabolism was found to be associated with viability

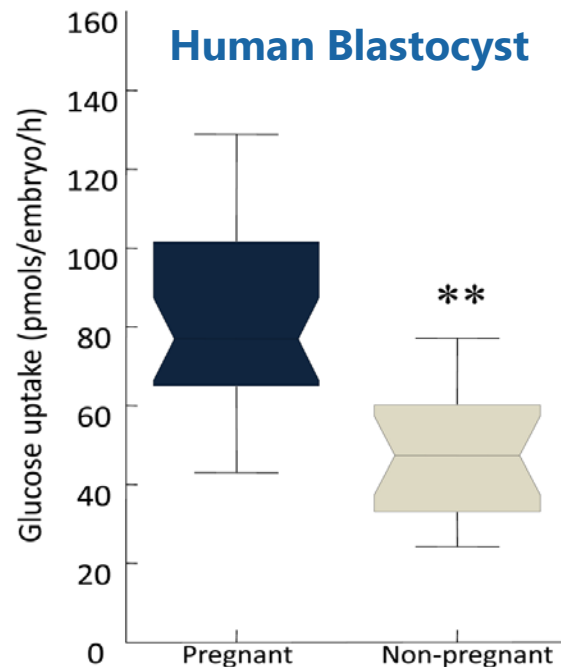
- Studies of the embryonic developmental in multiple mammalian species shows glucose uptake correlates to embryo health

Mouse Blastocyst

TABLE 1. The uptake of glucose (1 mM) by single mouse blastocysts prior to transfer into pseudopregnant recipients^a

Glucose uptake (pmol/embryo/hour)		
Male	Female	Nonviable
4.37 ± 0.22 ^b	4.92 ± 0.36	3.57 ± 0.46

Gardner and Leese, 1987



Gardner, 2015

Bovine Blastocyst

Development of cultured bovine embryos after single cervical transfer

	Embryos showing glucose uptake	Embryos with no glucose uptake	Total
No. of recipients	13	14	27
No. of pregnancies at Day 50 (%)	9* (69.2) ⁺	2* (14.2)	11 (40.7)

* Significantly different ($\chi^2=6.30$; $P < 0.025$).

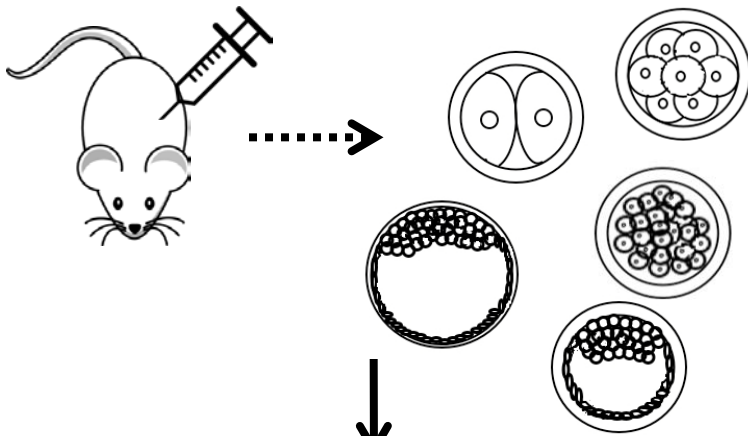
+ Including 1 dead fetus

Renard et al., 1980

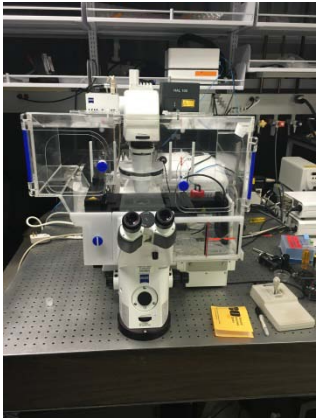
Imaging Strategy:

Non-invasive Phasor-FLIM mouse preimplantation embryo model

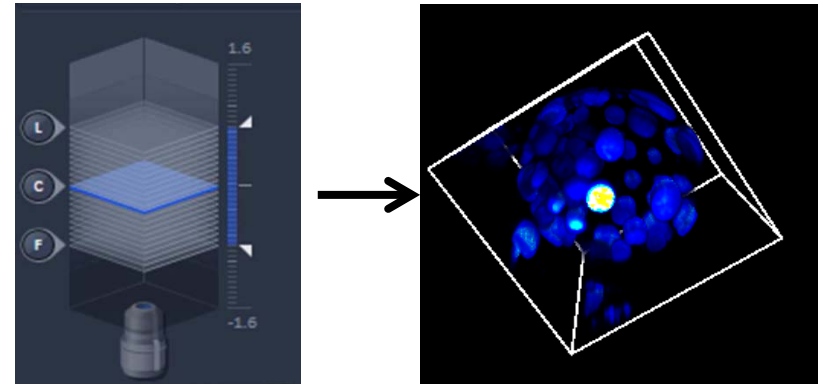
1. Superovulation and Embryo Collection



2. FLIM imaging

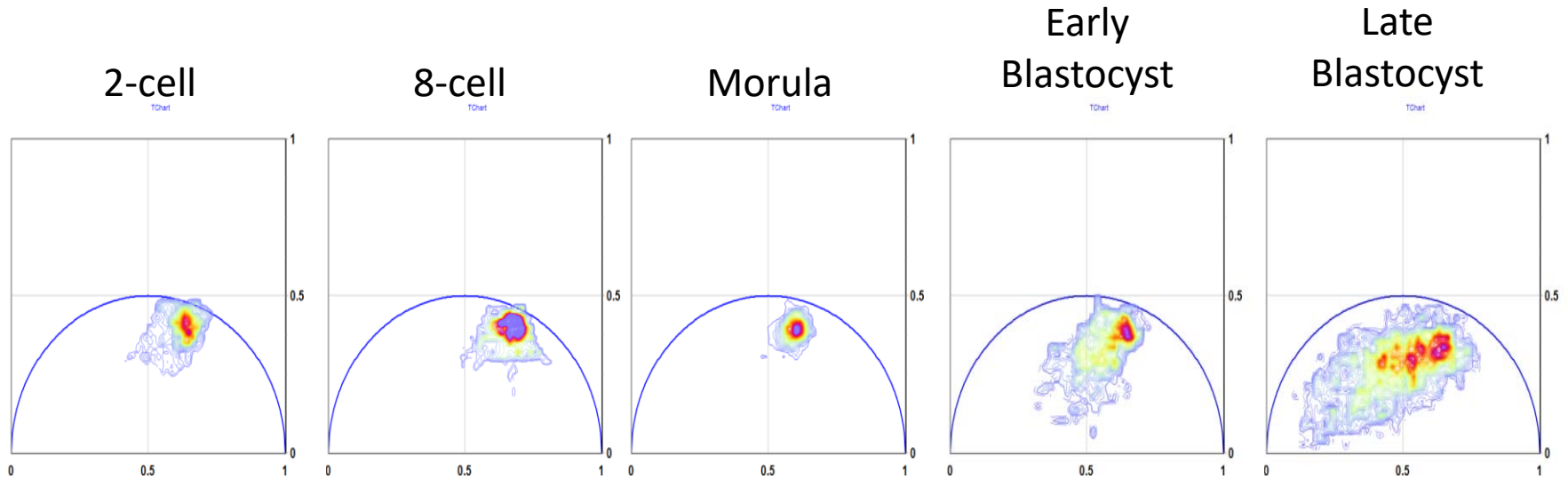


- Non-inbred CD-1 and Inbred C57BL/6
- 24-day old females
- Superovulation with PMSG and hCG



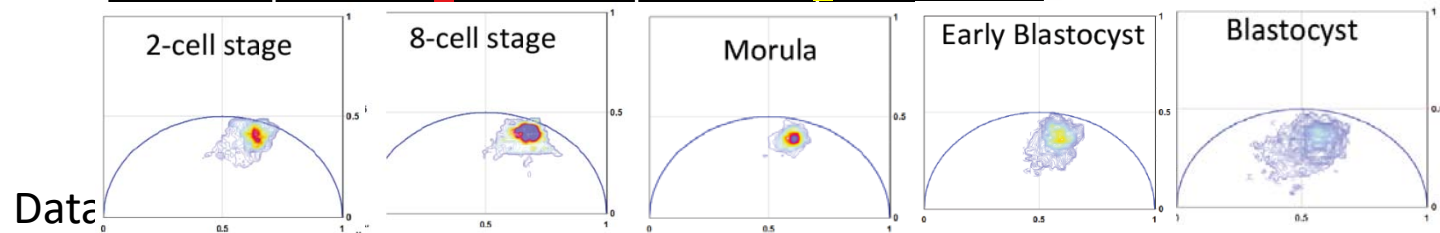
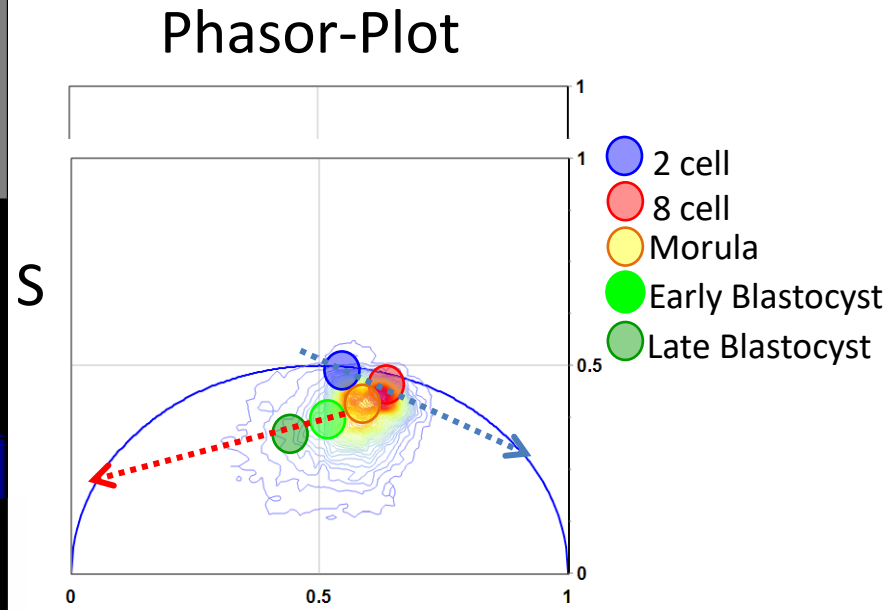
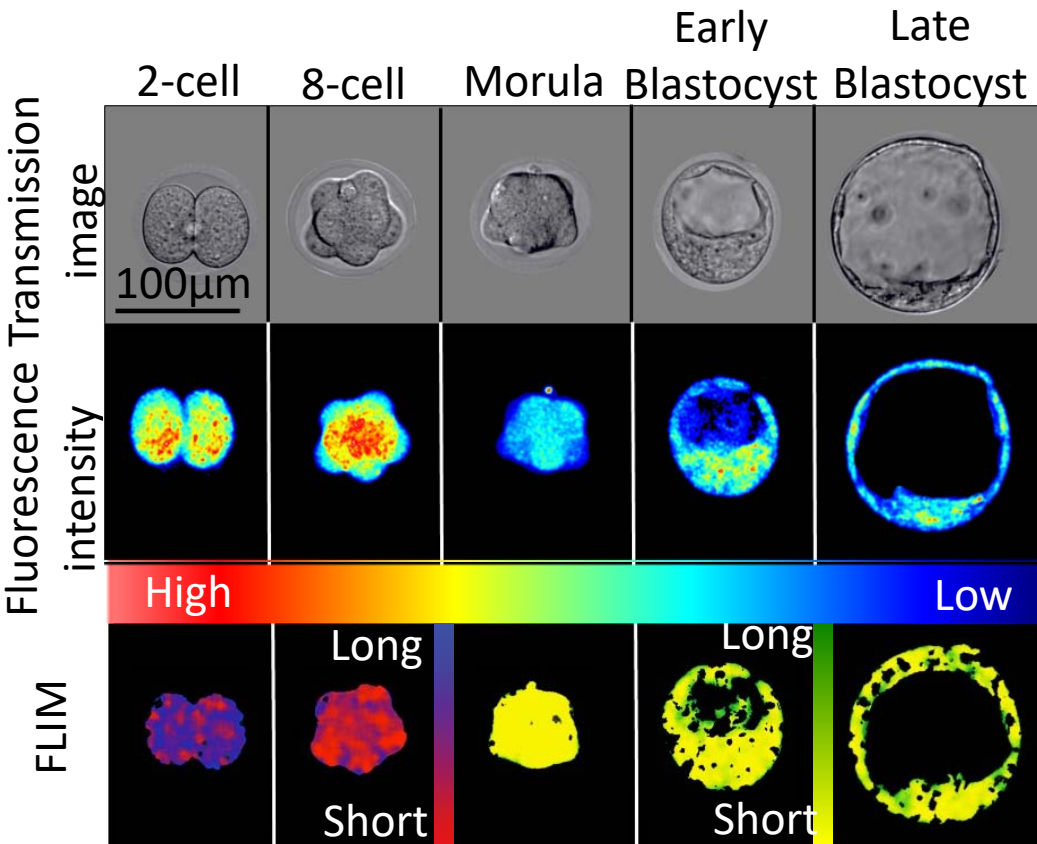
- Zeiss LSM710 + Fast FLIM data acquisition card
- Zeiss EC Plan-Neofluar 20x/0.5 NA objective
- Excitation/Emission: 740nm/420-500nm
- ~3.5mW laser power
- 50 frames z-stack integration (~3 minute)
- 37 °C, 5% CO₂

Progression of ROS production shifts the lifetime towards the oxidized lipid lifetime



- In bovine embryos by 8-cell stage, ROS levels increase, and then they decrease by the blastocyst stage (Dalvit, 2005).
- Production of ROS, or increase of free radicals, is a normal result of cellular metabolism responsible for maintaining a good redox balance

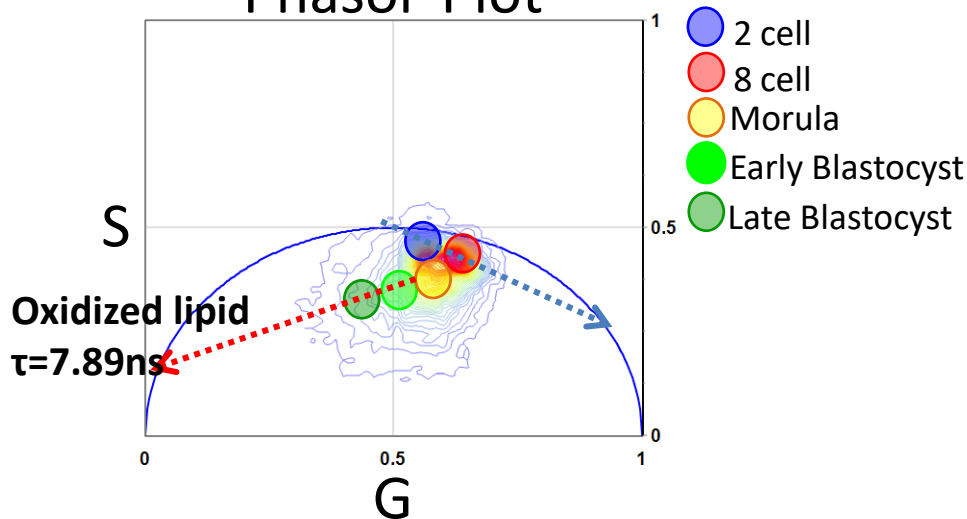
Lifetime population trajectory of healthy developing embryos



Lifetime population trajectory of healthy developing embryos

Stage	Major metabolism in literature	Expected NADH signature
2-cell	Pyruvate uptake (Biggers et al. ,1967, Whitten et al., 1957, Brinster et al., 1965)	free NADH↓
Late 4-cell/8-cell	Glycolysis (Brinster et al. ,1966)	free NADH↑
Blastocyst	Oxidative phosphorylation (Gardner et al, 2010, Lane et al. 1996, Gardner et al. 1987)	Bound NADH↑

Phasor-Plot

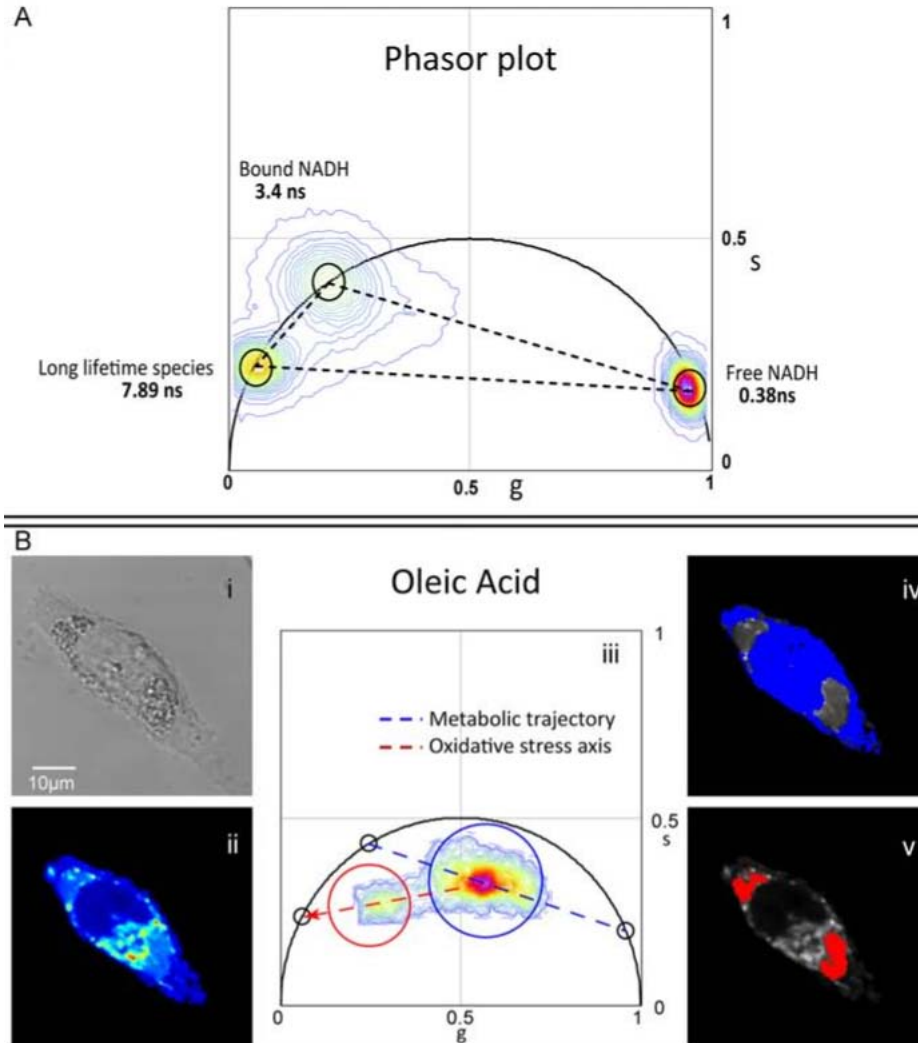


Lifetime:

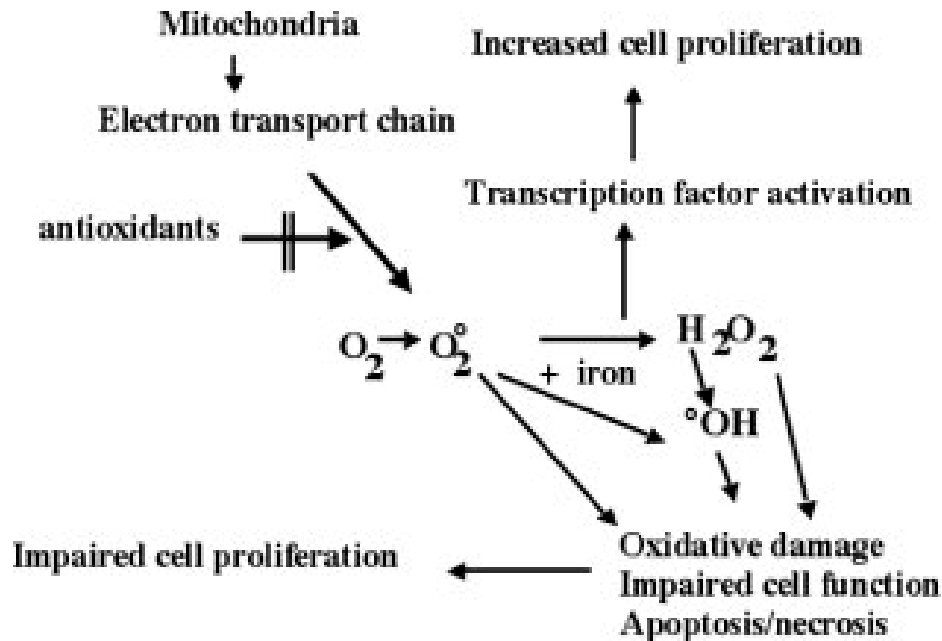
Long \rightarrow Short \rightarrow Long

Phasor-FLIM can measure metabolism during embryonic development in pre-implantation embryos

Long lifetime species (LLS): FLIM signature and a new oxidative stress axis on phasor



Why is ROS production needed for embryonic development?



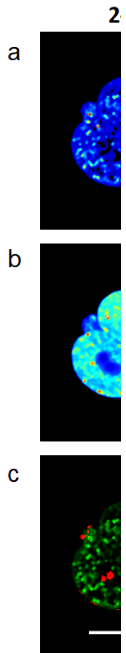
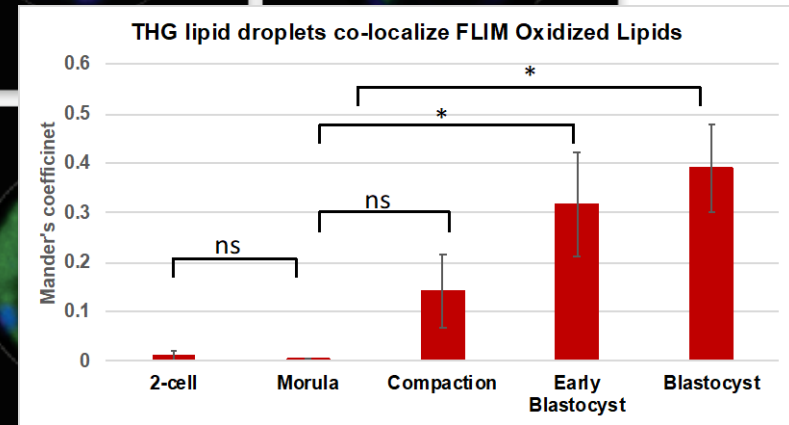
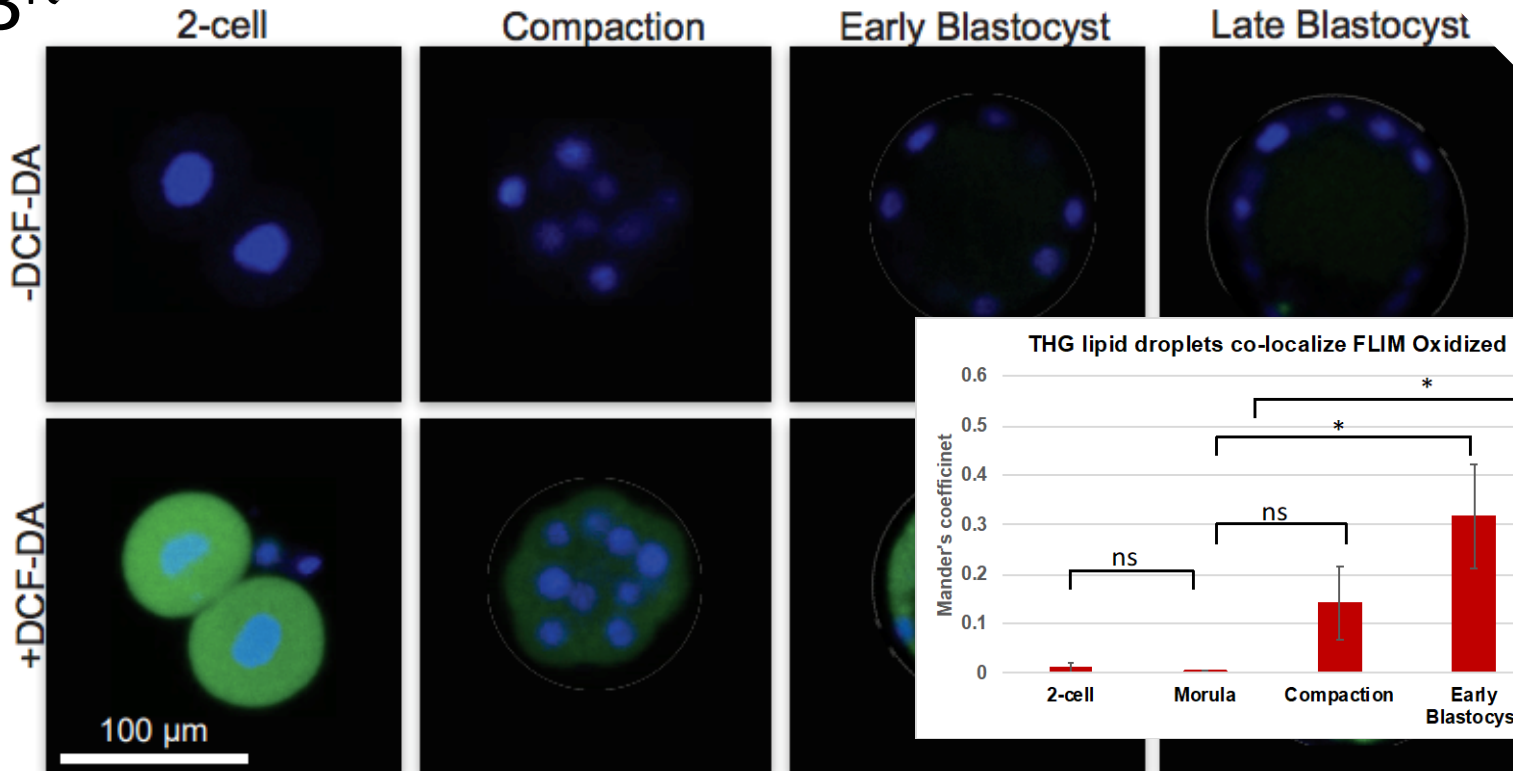
- In bovine embryos by 8-cell stage, ROS levels increase, and then they decrease by the blastocyst stage . Production of ROS, or increase of free radicals, is a normal result of cellular metabolism responsible for maintaining a good redox balance
- ROS are important in normal development through cellular signaling and control of cellular fate.

Reactive Oxygen Species (ROS) detection

- DCF-DA: Reactive Oxygen Species

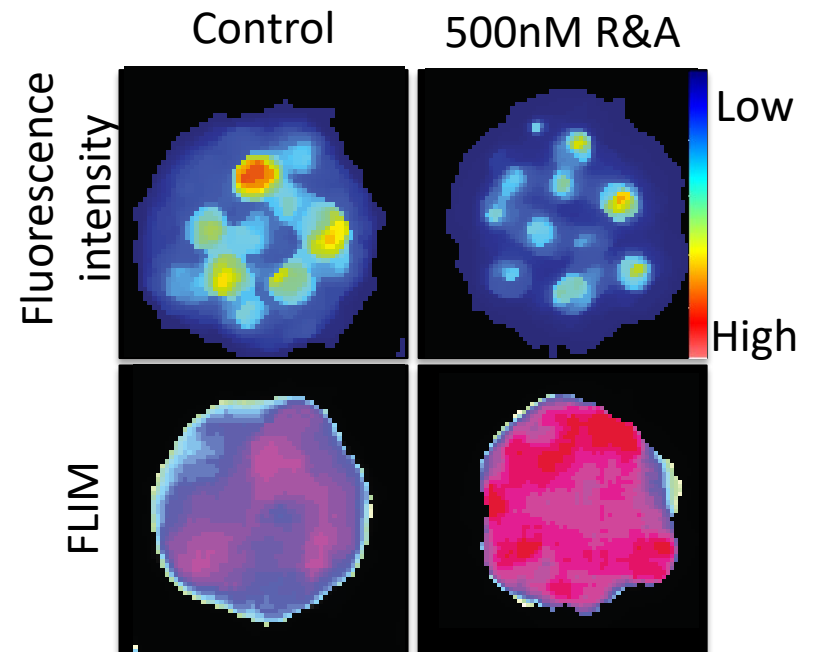
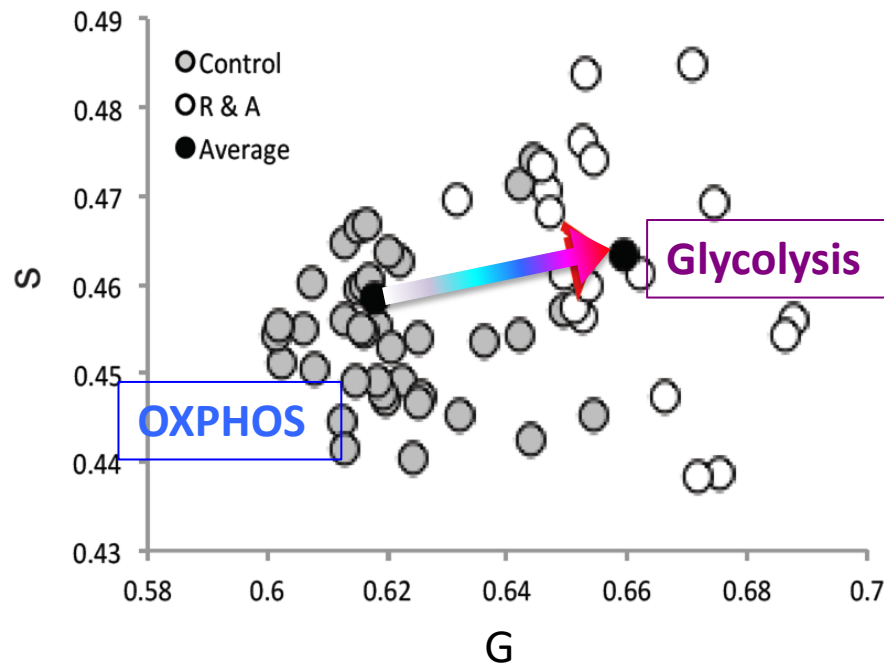
Blue: DAPI
Green: DCF-DA

- 3rd ...



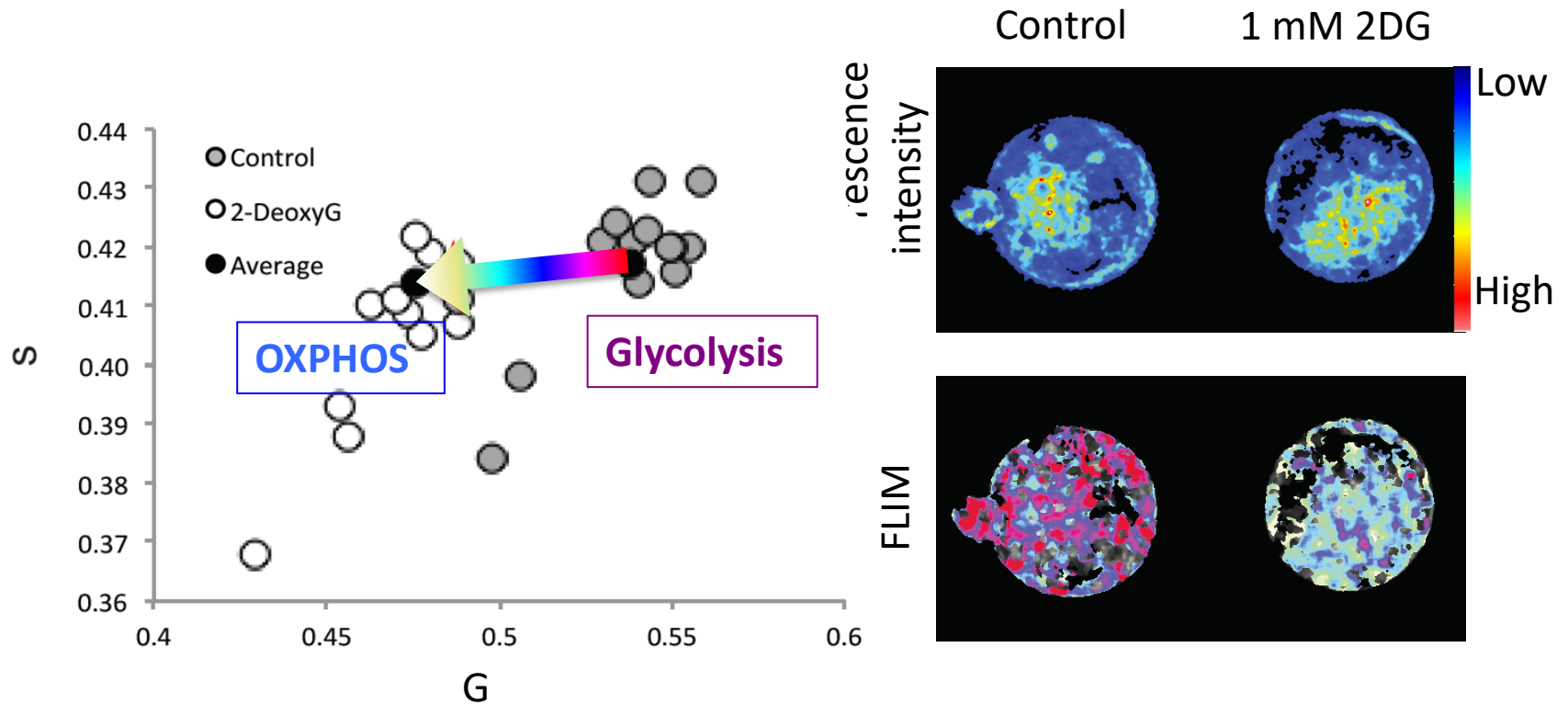
Manipulation of embryo metabolism

- [Rotenone and Antimycin A \(R&A\): Oxidative Phosphorylation inhibitor](#)



Manipulation of embryo metabolism

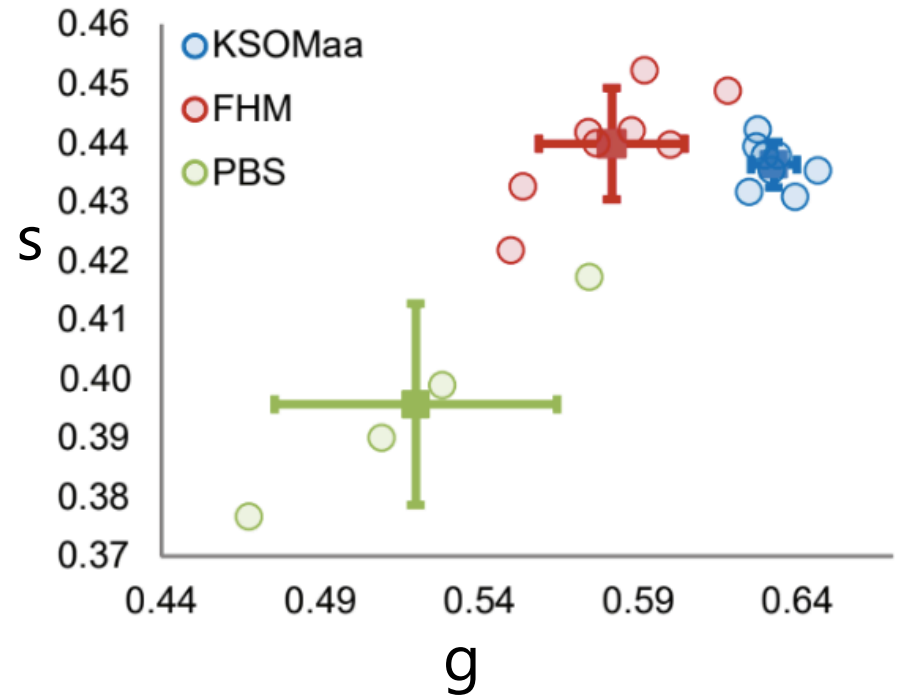
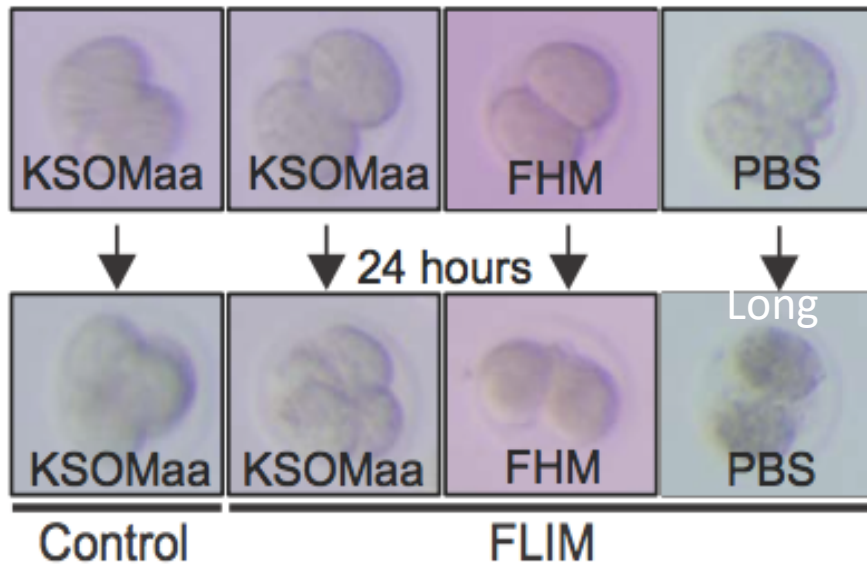
- [2 Deoxy-D-Glucose \(2DG\): Glycolysis inhibitor](#)



FLIM can distinguish healthy embryos and the embryos under “high stress” conditions

After ~~24 hours~~ treatment

2-cell stage embryos



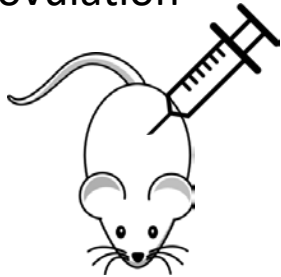
KSOMaa: standard culture media

FHM: nutrient lacking flushing and holding media

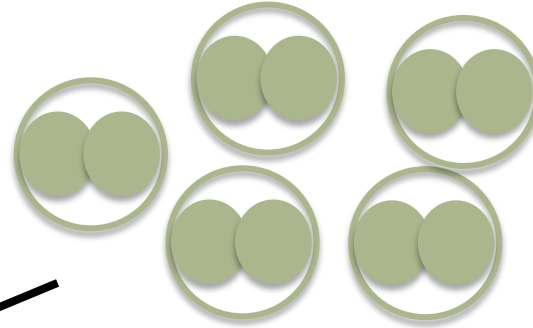
PBS: saline solution

Can we use the phasor/FLIM method to predict the best developmental potential for each embryo?

1. Superovulation



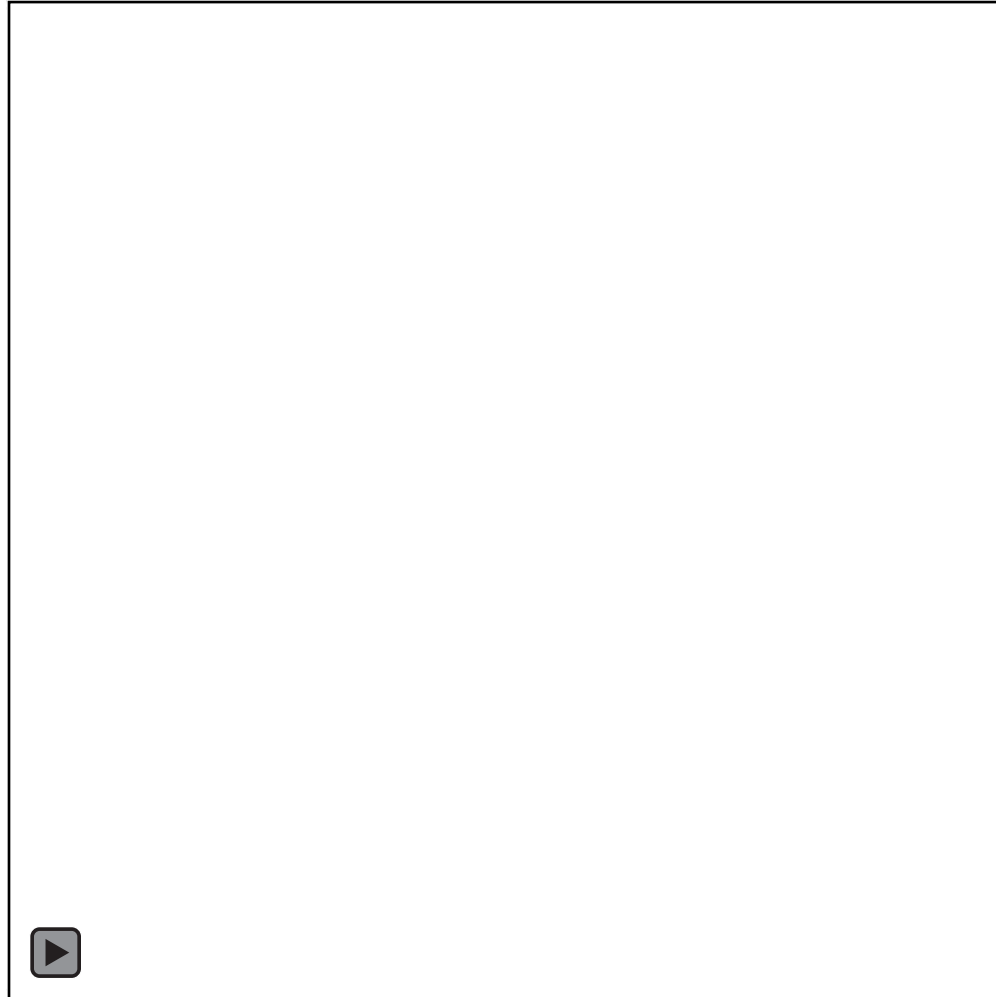
2. Embryo Collection (2-cell stage)
40~180 embryos for each experiment



3. FLIM imaging and Time-lapse
Imaging for ~60 hours



Time-lapse imaging of a healthy (H) and an “unhealthy” (UH) preimplantation embryo



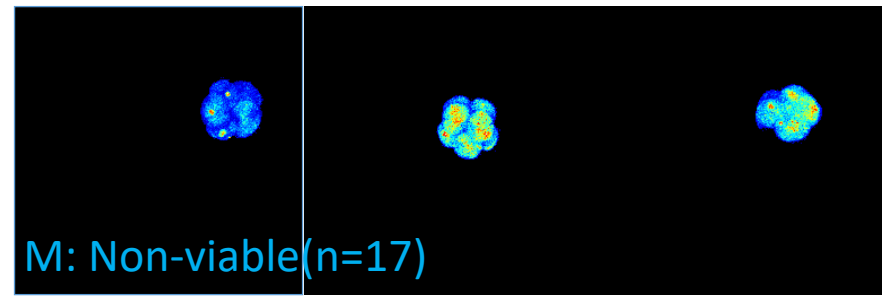
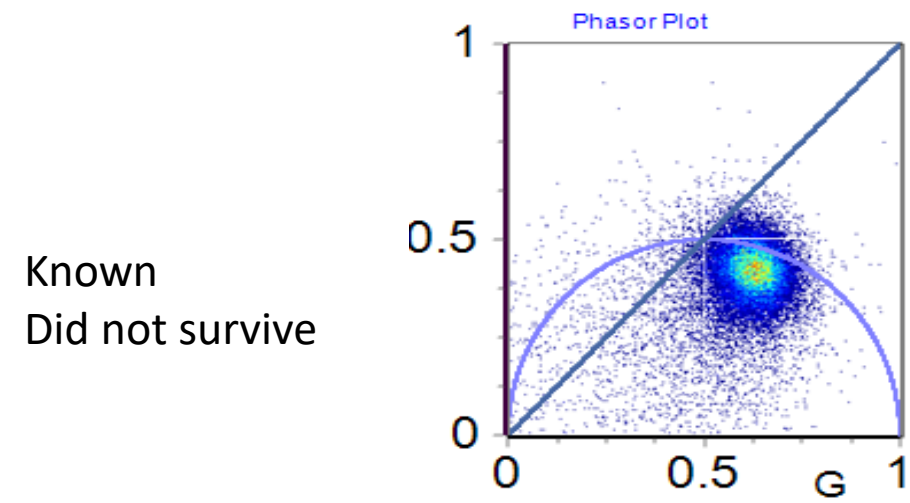
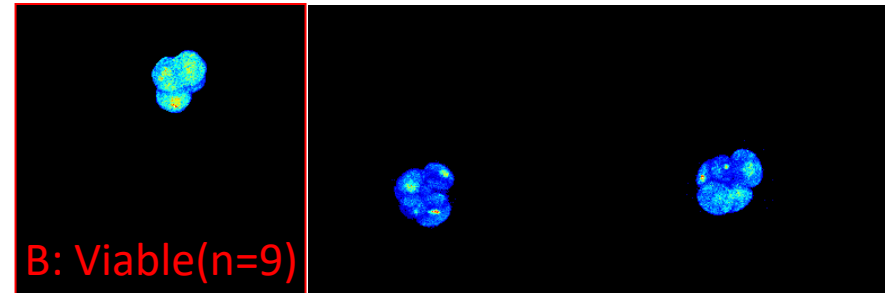
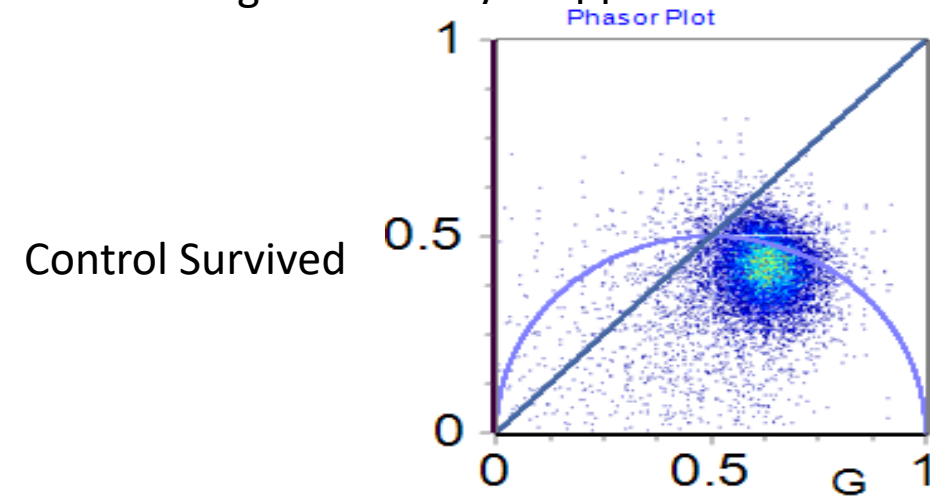
H: Embryos developed to blastocyst

UH: Embryos arrested before blastocyst at morula stage

The “Distance” analysis approach of the phasor plot

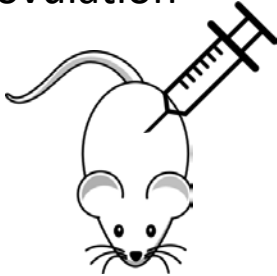
Premise: the phasor plot contains about 10,000 points for each image and generally we analyze 10 to 100 images all together

At 8 cell stage the embryos appear to be indistinguishable through 2D phasor

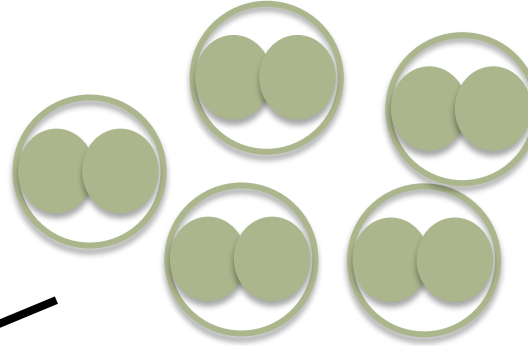


Combination of FLIM imaging and time-lapse morphological tracking work flow

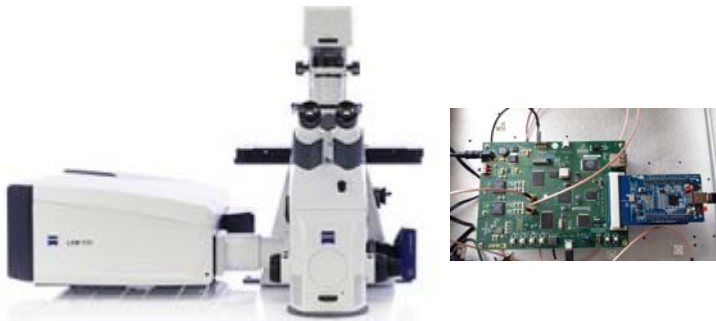
1. Superovulation



2. Embryo Collection (E1.5)

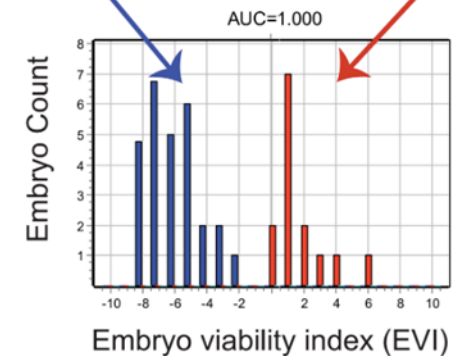


3. FLIM imaging and Time-lapse Imaging for ~60 hours

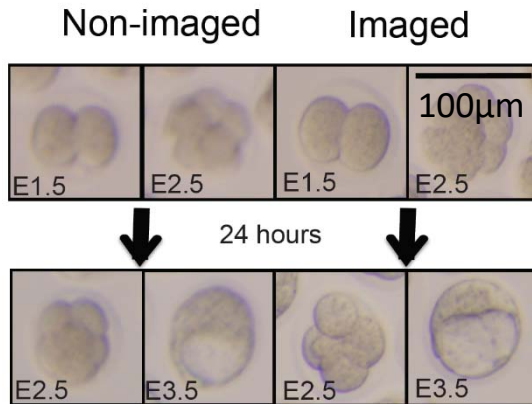


4. Distance-

Healthy embryos (EVI < 0) Unhealthy embryos (EVI > 0)

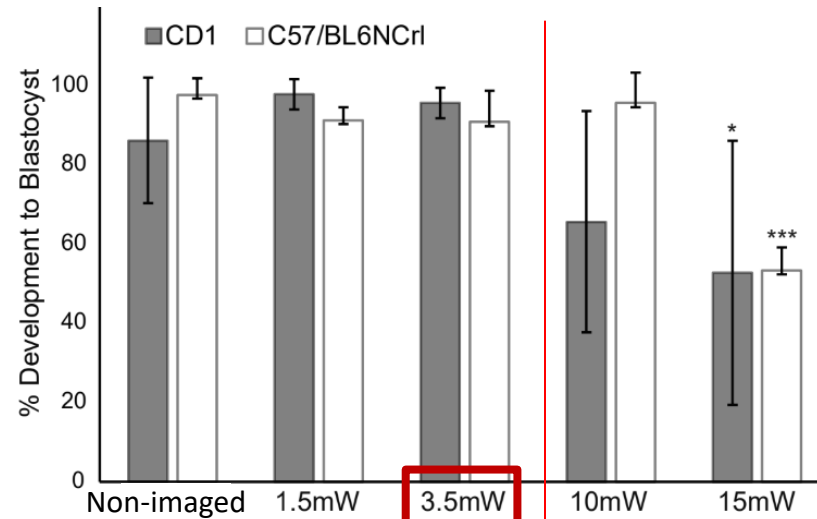


FLIM does not cause DNA damage

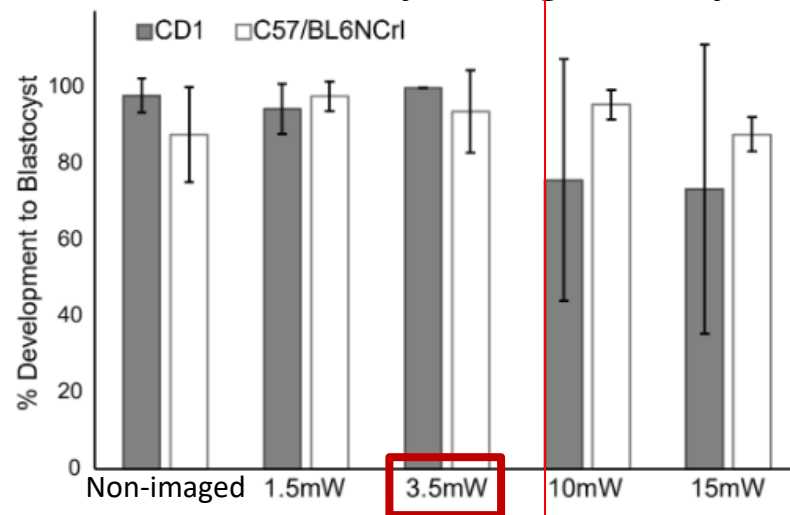


3.5mW: average laser power for embryo imaging (50 frames)

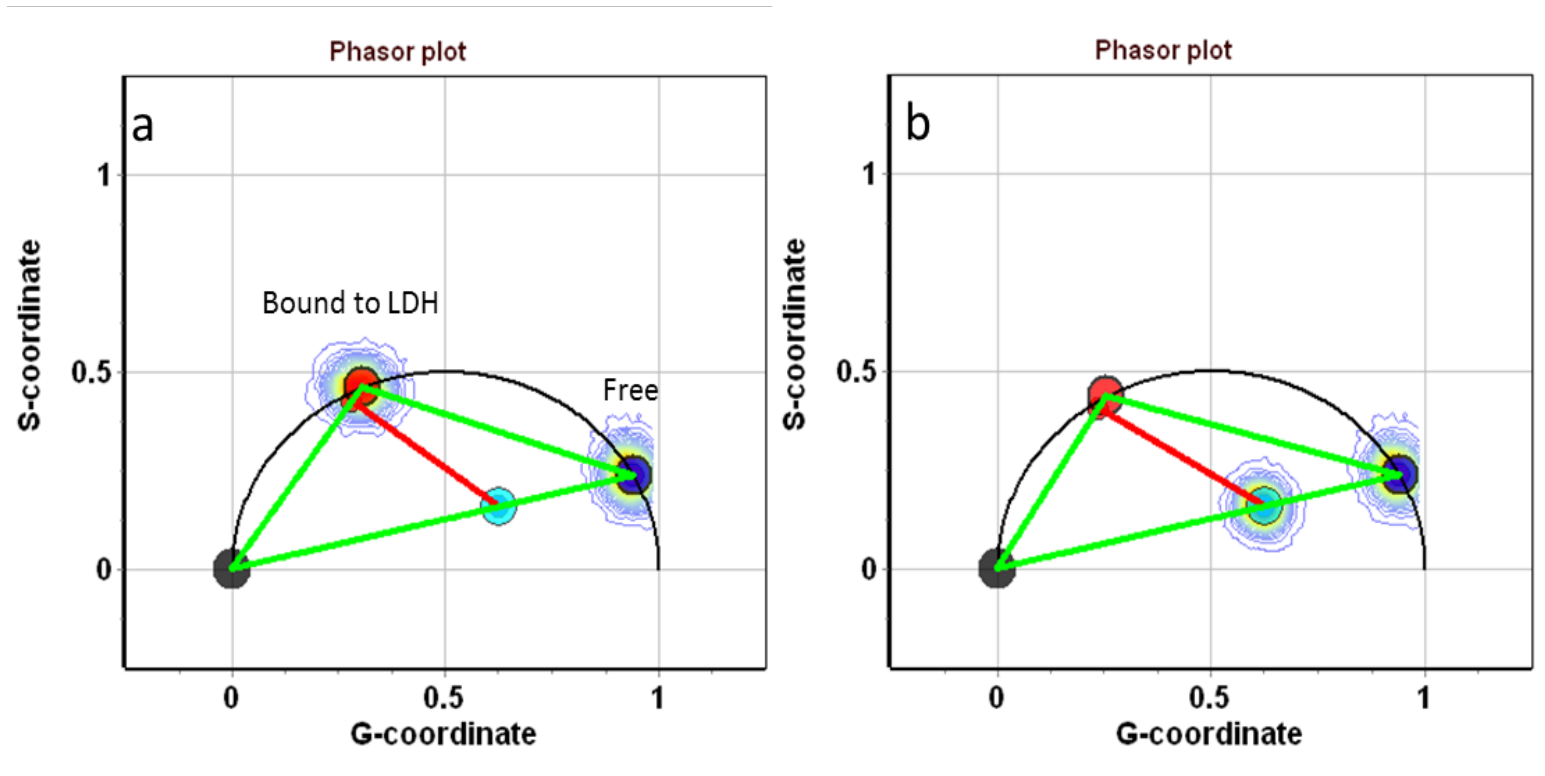
2-cell to 4-cell stage embryos



Morula to Blastocyst stage embryos



Principle of absolute NADH concentration measurement



$$M = \frac{L}{L + F}$$

g: experimental g

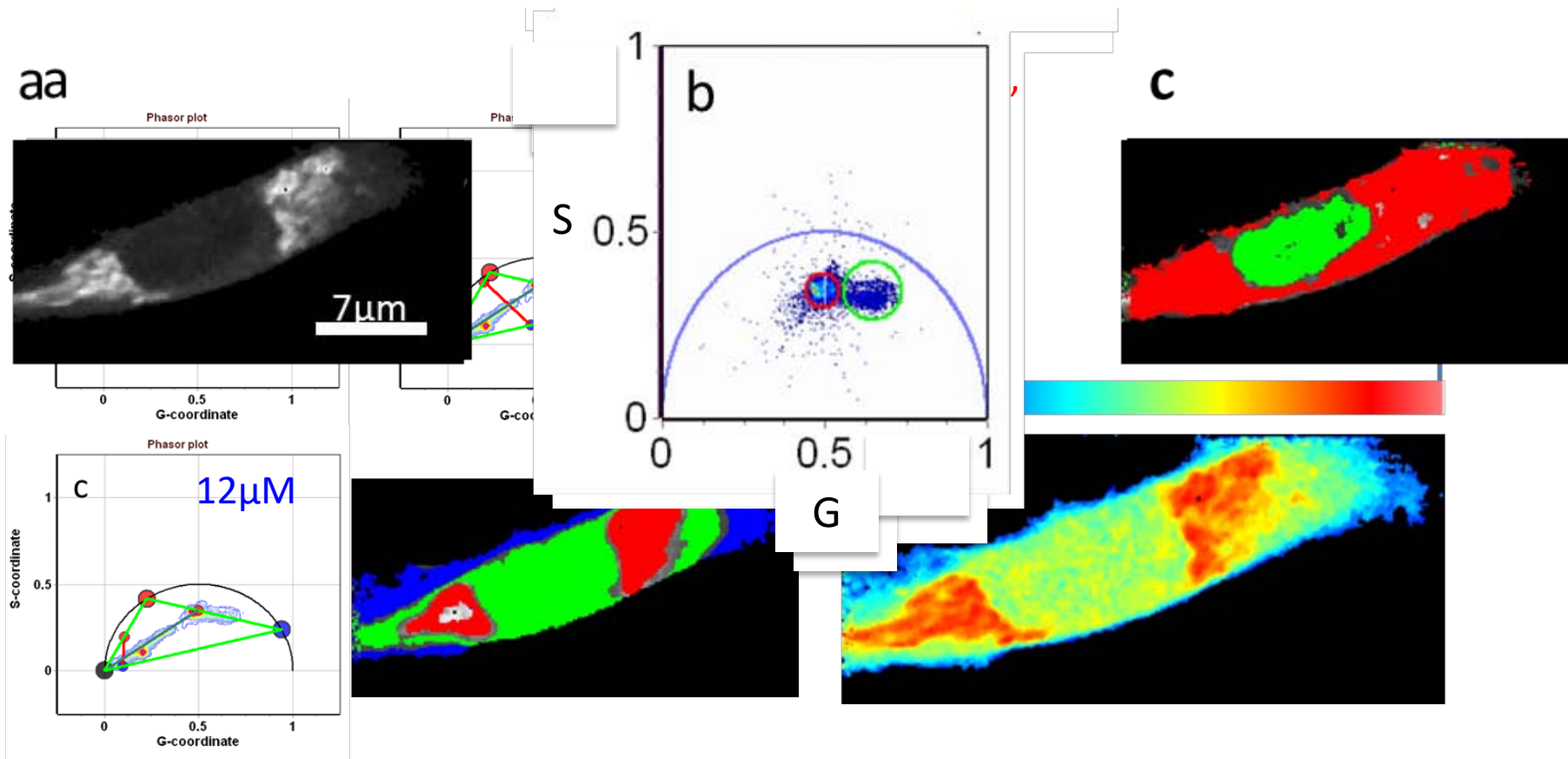
M_0 : g coordinate of the calibration solution without

external light added

L: intensity of unmodulated light

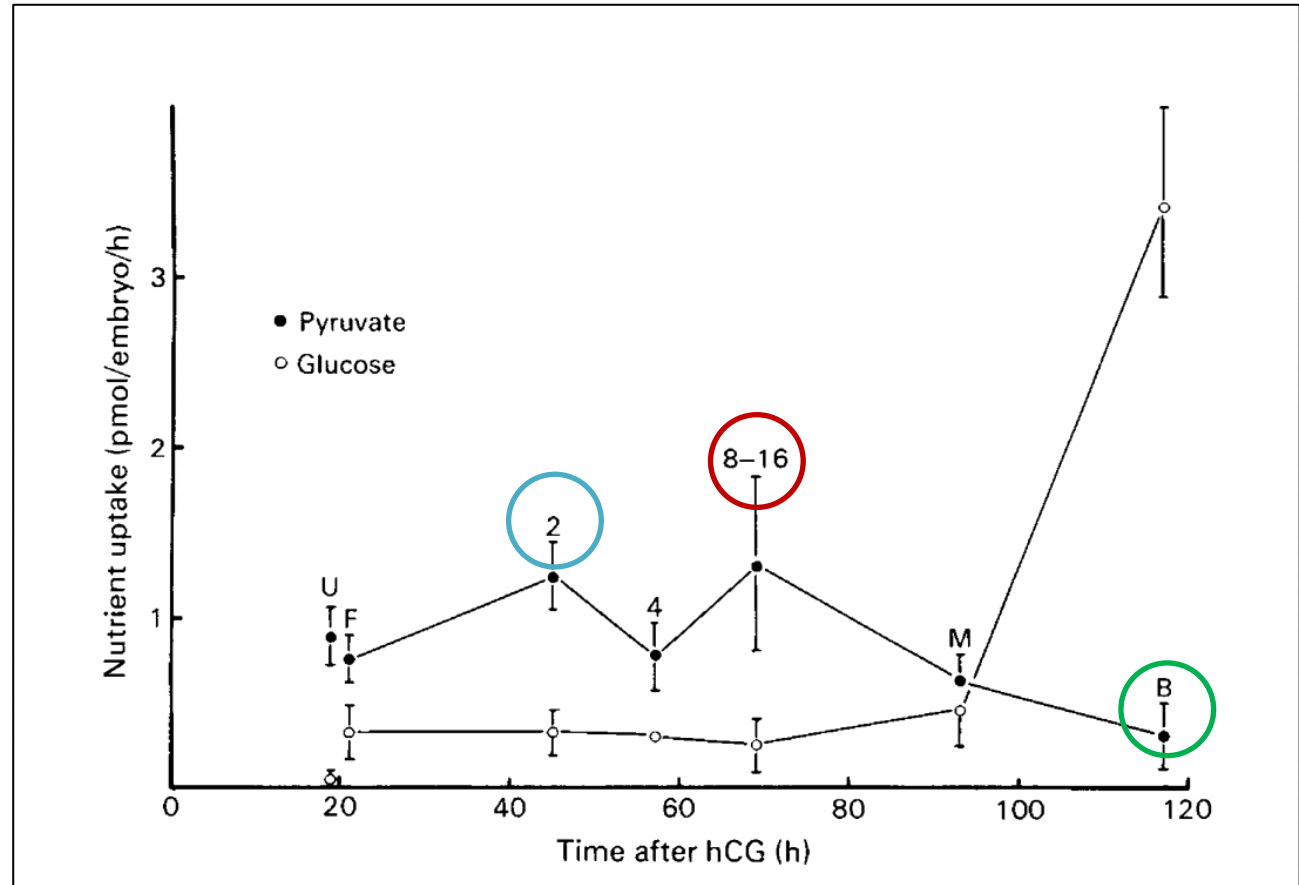
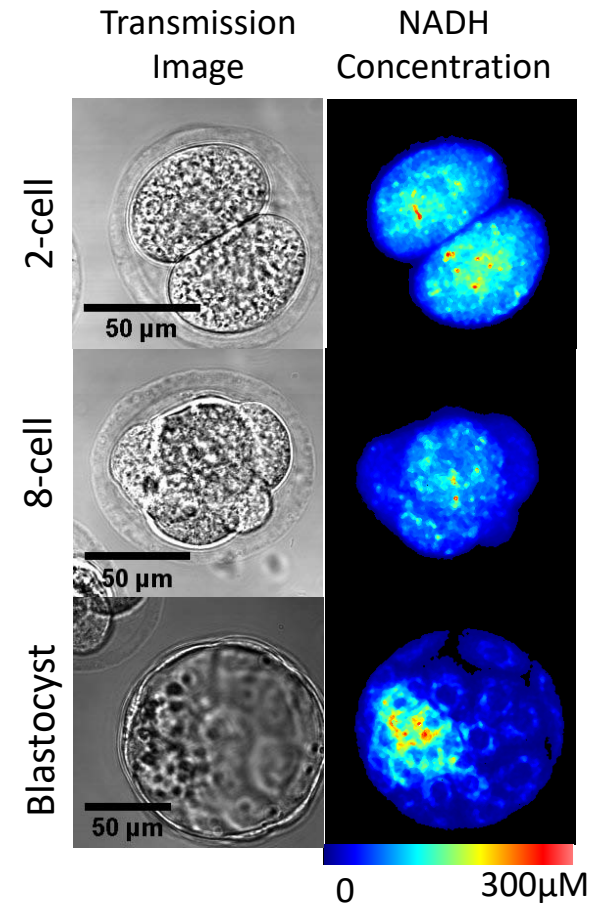
F: intensity of fluorescence of the specimen

Absolute NADH concentration measurement in Cho-K1 cells

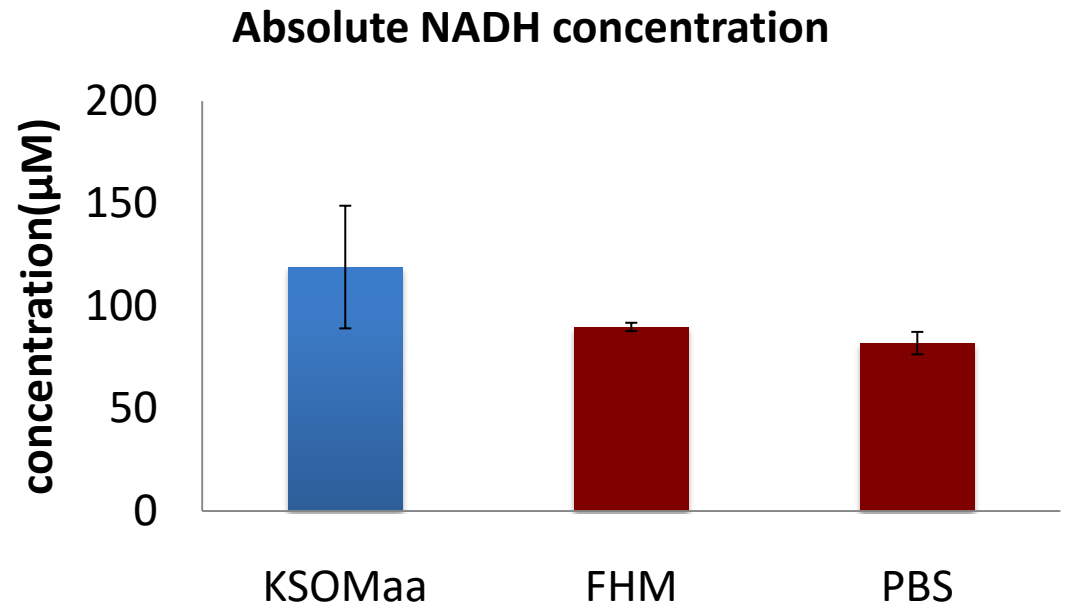
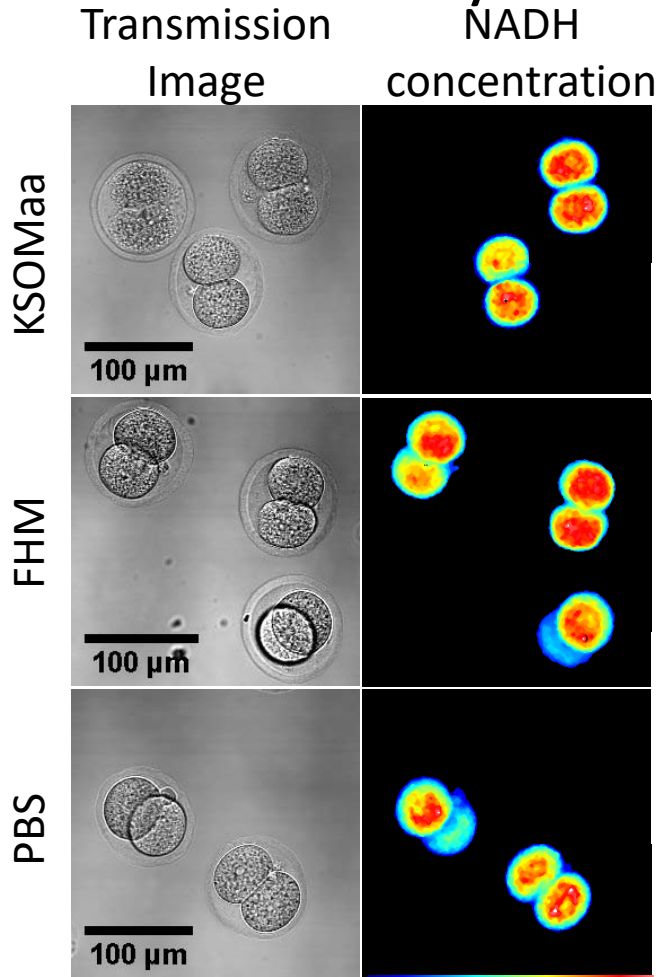


Previous measurement of NADH concentration in normal cells and cancer cells are in the range 97µM-168µM, the values demonstrate our method can derive absolute NADH concentration in cells (Klaidman et al. 1995, Yu et al., 2009)

Preliminary data for NADH absolute concentration during embryonic development



Comparing absolute NADH concentration in healthy embryos and embryos under “high stress”



Embryo culture media Nutrient lacking media

The viable embryos appeared to have higher NADH level.

Embryo viability prediction Pipeline

MENU ▾

SCIENTIFIC REPORTS

Article | [Open Access](#) | Published: 13 September 2019

Label-free assessment of pre-implantation embryo quality by the Fluorescence Lifetime Imaging Microscopy (FLIM)-phasor approach

Ning Ma, Nabora Reyes de Mochel, Paula Duyen Pham, Tae Yeon Yoo, Ken W. Y. Cho

✉ & Michelle A. Digman ✉

Scientific Reports **9**, Article number: 13206 (2019) | [Cite this article](#)

677 Accesses | **4** Altmetric | [Metrics](#)

Abstract

Summary

- Mouse pre-implantation embryo has unique lifetime trajectory during development which can be detected by Phasor-FLIM noninvasively and quantitatively
- 8-cell stage is the optimal stage for the prediction
- FLIM can be a useful tool to detect embryo viability

The Digman Lab

Metabolic Alterations

Embryo Development
Neurodegeneration
Cancer

Mechanobiology and the
Influence of ECM on
Metabolism in Cancer Cells

Mitochondrial recruitment
Tumor organoids

Imaging Technologies:
Phasor/FLIM
Multiphoton excitation
Fluorescence Correlation Spectroscopy
N&B
RICS
pCF

DNA Damage and Protein
Phase Transitions in
Health and Disease

Protein dynamics and
distinct physical states

Signal transduction
dynamics during
epidermal wound healing
and
partial Epithelial-to-
Mesenchymal Transition

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Angel Balam Benitez Mata



Yu-Kai Huang



Dr. Ning Ma

COLLABORATORS:



Greg Brewer



Yue Dong



DGE-1839285
(Austin Lefebvre)



Career Award #1847005
(Dr. Digman)



Ken Cho



Soledad Mochel



Paula Pham



Tommy Yoon

