

Raman Spectroscopy in Biogeology

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Terre Planètes Environnement

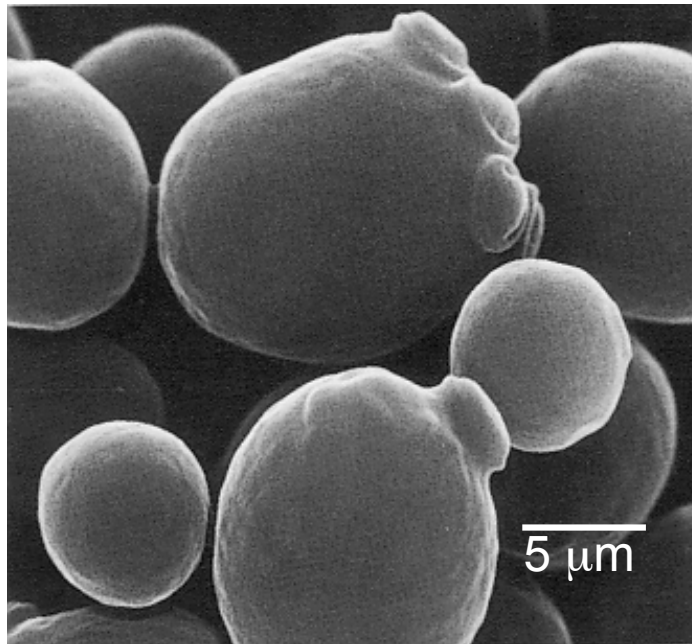


Outline

- Quantitative Raman spectroscopic analysis of microbial metabolic activity
- Raman spectroscopy for probing live cells
- Measurements of minute amount of biological and biotic products

Quantitative Raman spectroscopic analysis of microbial metabolic activity

The fermentation
by the baker yeast *Saccharomyces cerevisiae*
as a function of pressure



The yeast *S. cerevisiae* at high hydrostatic pressure

0.1 MPa
Optimal growth pressure



20-50 MPa
Cell cycle arrest



40-60 MPa
Internal acidification
Induction of stress
transcriptional factors



70-200 MPa
Induction of stress
transcriptional profile

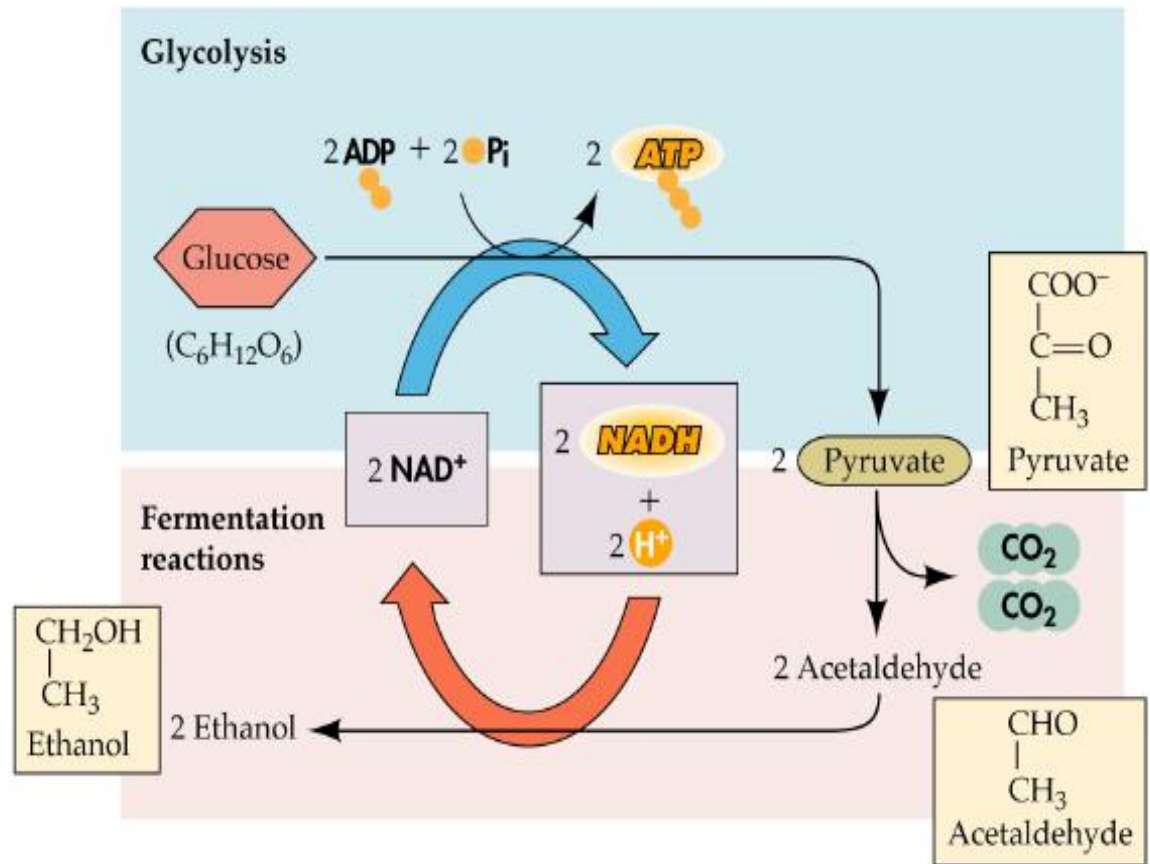
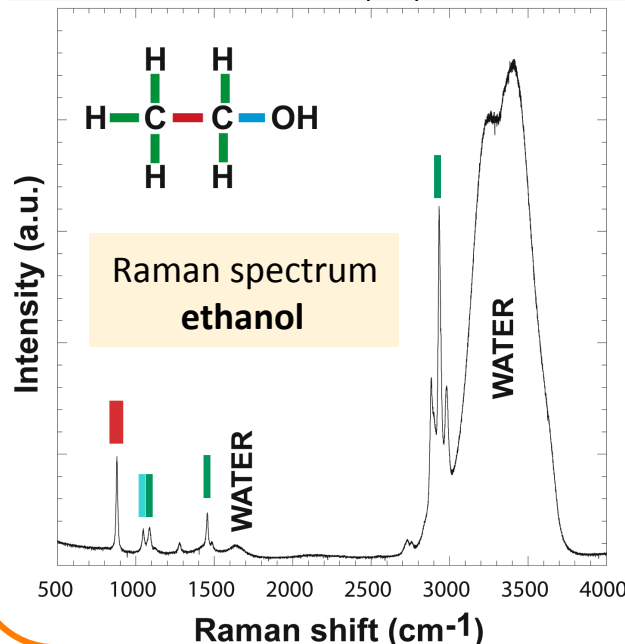
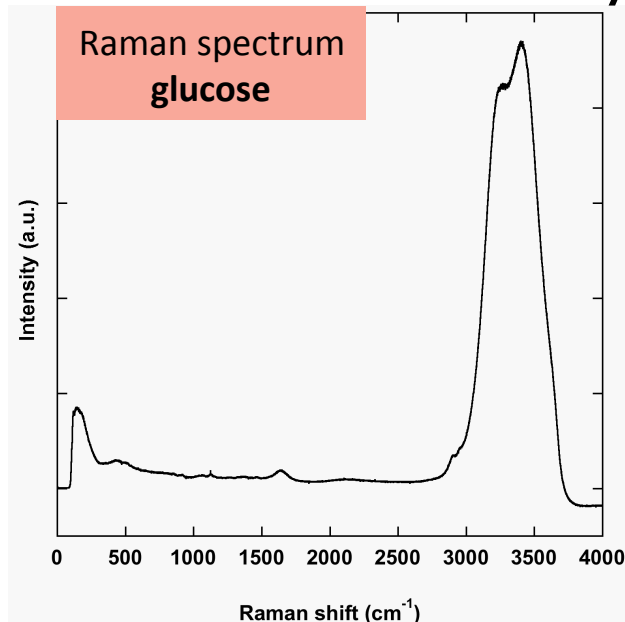


≥220 MPa
Death



- The **eukaryotic model** for high-pressure studies
- No metabolic data for *S. cerevisiae* under high pressure
- Alcoholic fermentation well constrained at ambient pressure
- Ethanol easily detectable by Raman spectroscopy
- Arrest of alcoholic fermentation predicted at ca. 50 MPa (Abe *et al.* 2004)

Characterization of alcoholic fermentation by Raman spectroscopy

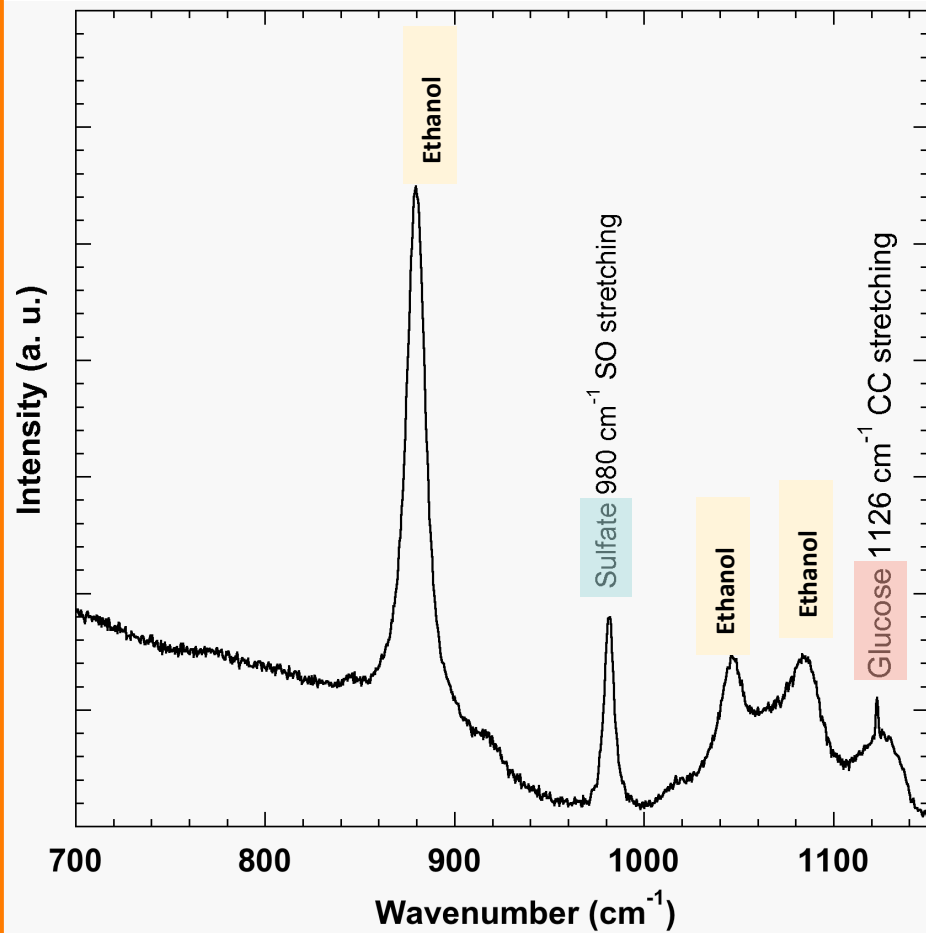


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Symmetric C---C stretching mode of ethanol
@ 883 cm⁻¹

Quantification of ethanol by Raman spectroscopy

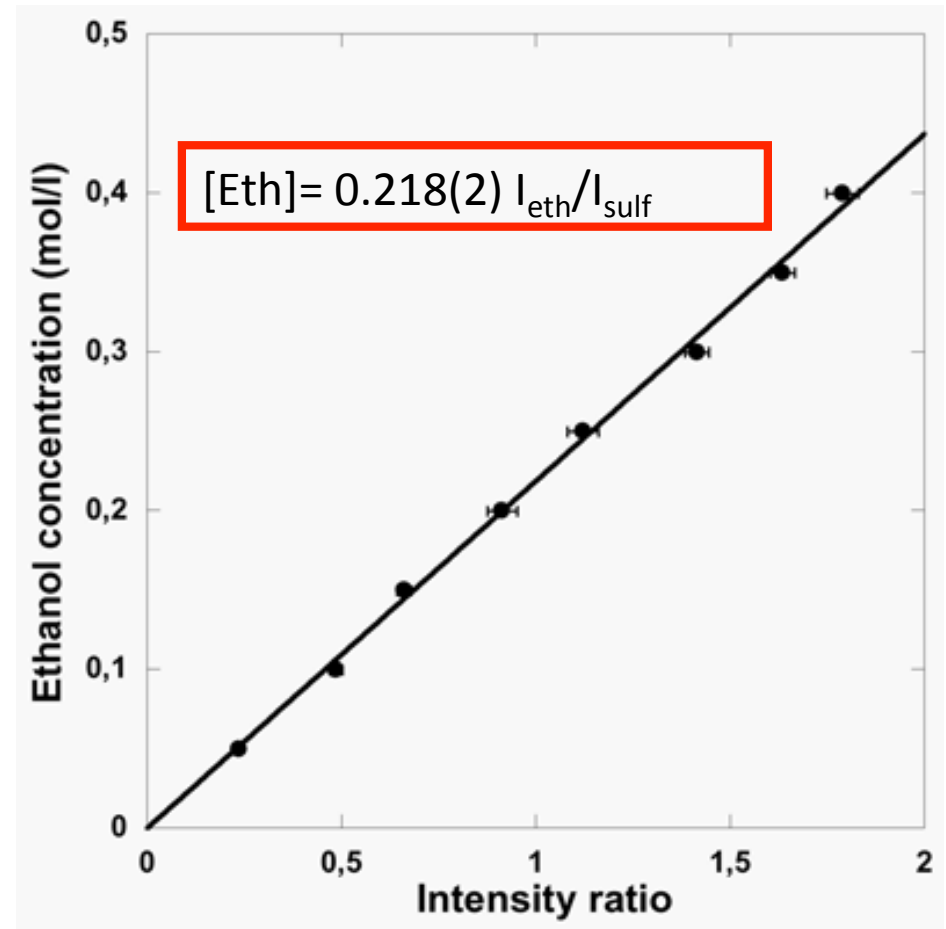
... in the low-fluorescence culture medium



Experimental details:

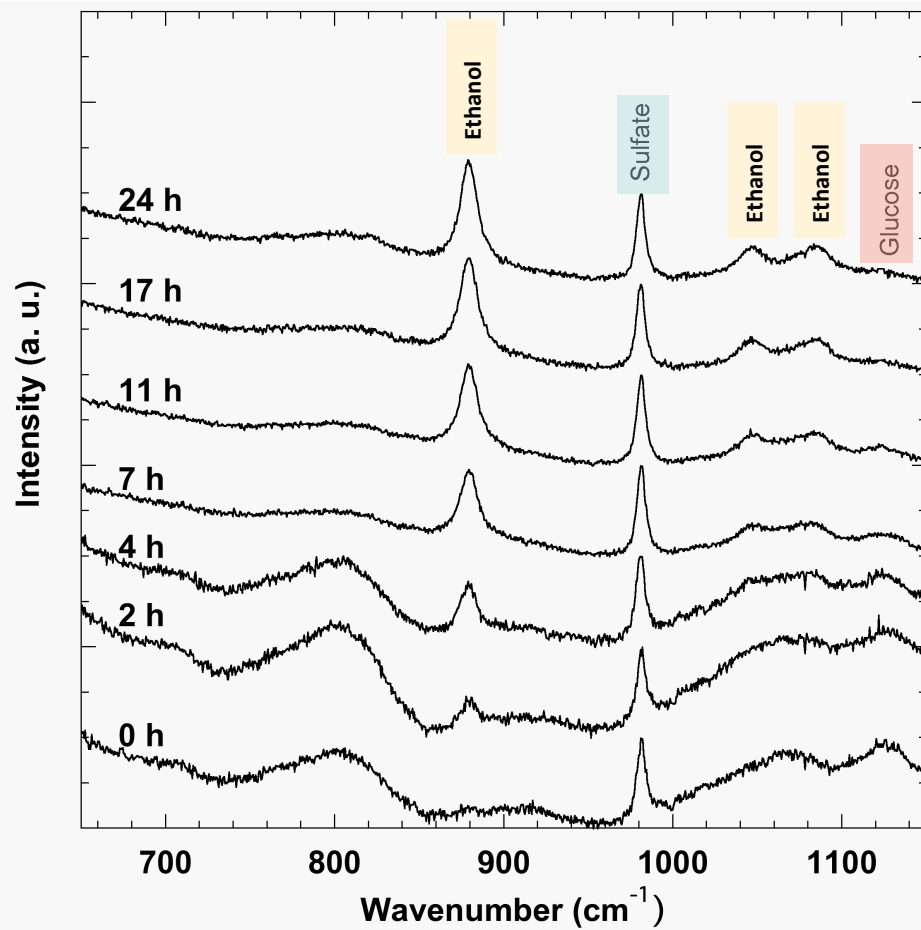
- Jobin Yvon® HR800 spectrometer
- Ar laser, 514.53 nm, 40-50 mW
- 10x20 s acquisitions

Calibration of
the normalised intensity of ν_s ethanol
as a function of ethanol titration

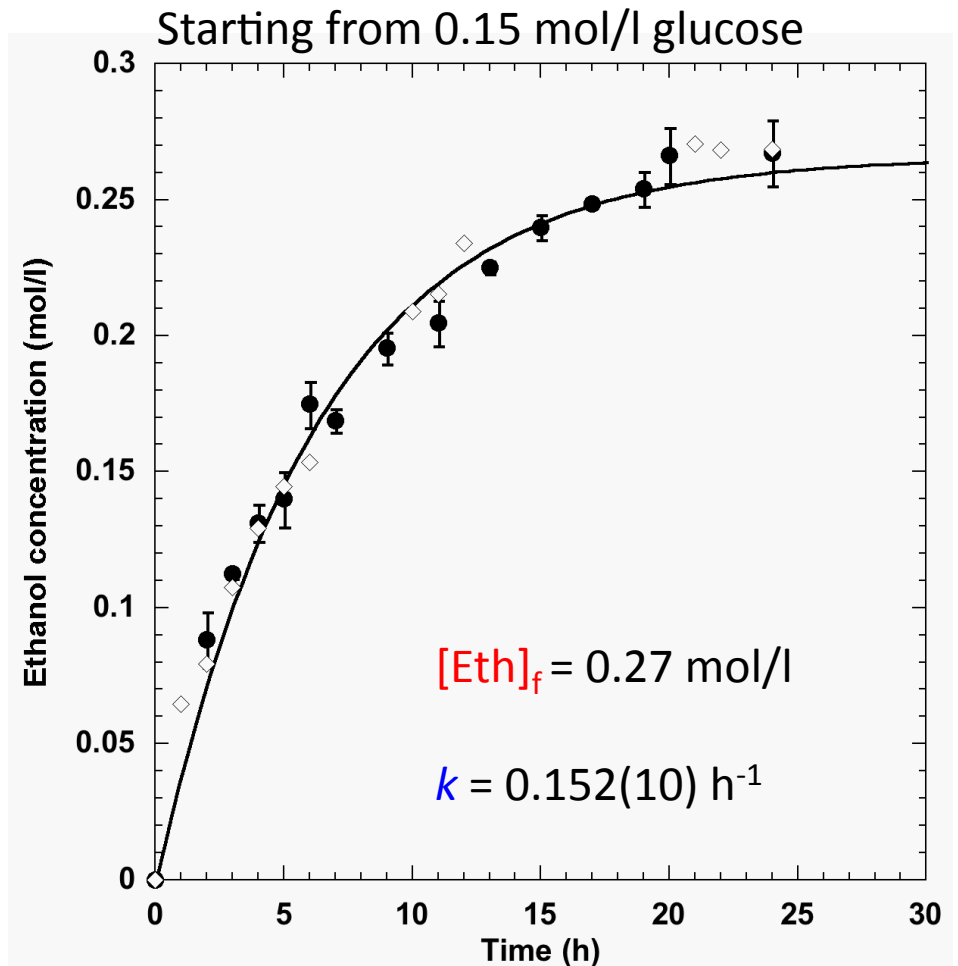


$$I_{eth, 883} / I_{sulf, 980}$$

Fermentation of *S. cerevisiae*, at ambient pressure



First order kinetic reaction
 $[\text{Eth}] = 2[\text{Glc}](1 - e^{-kt})$



Raman spectroscopy in a 'low'-pressure DAC



Coll. with J.C. Chervin, at IMPMC, Paris

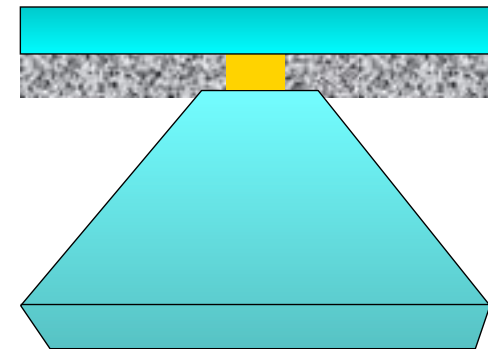
Pressure and temperature range

1.5 GPa

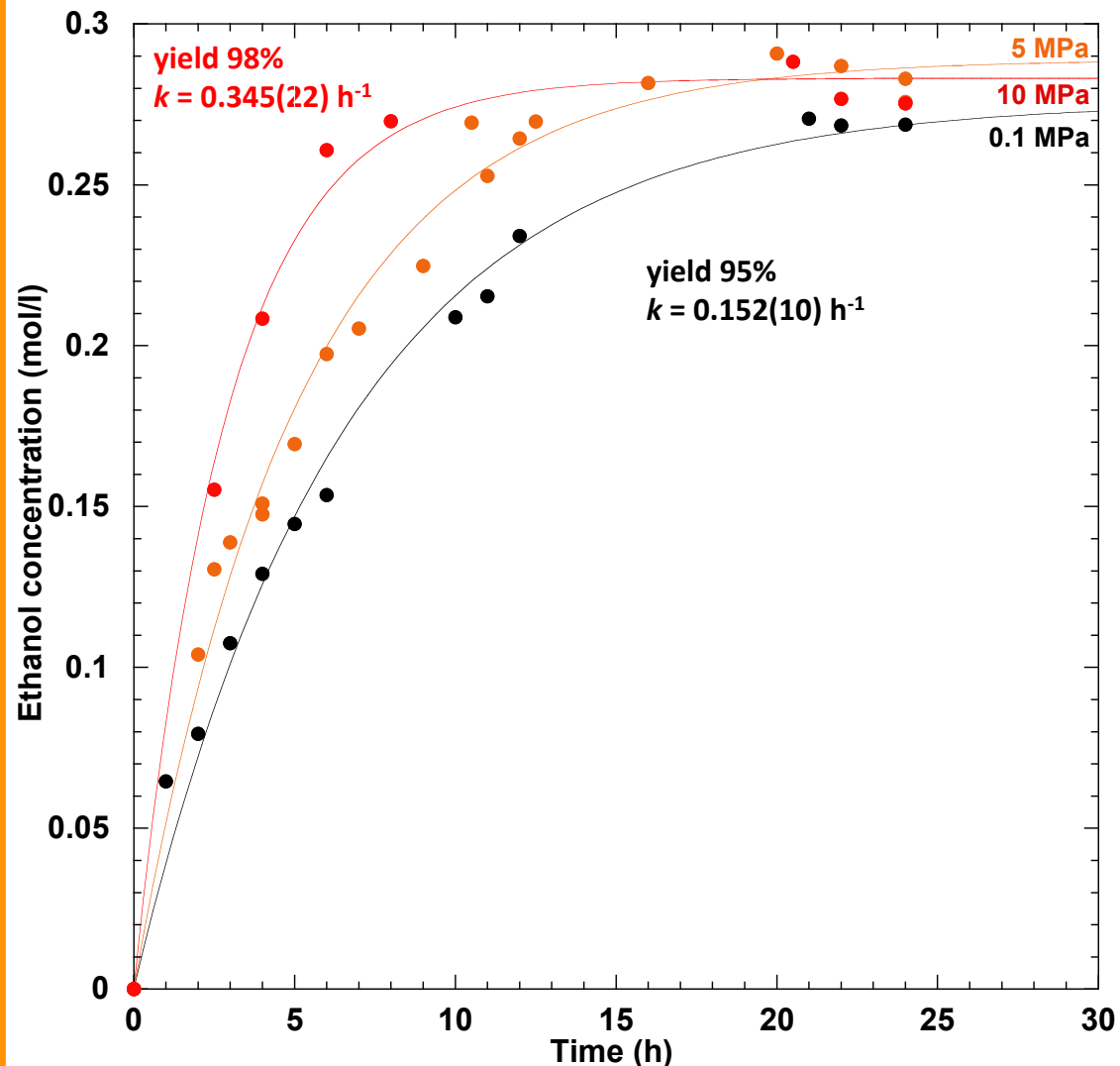
300 °C

sample 60 nl

- Diamond window 0.4 - 0.6 mm thick
- Ni gasket, 0.3 mm thick, 0.5 mm across
- Diamond anvil 2.2 mm thick, 1.4 mm culet



Fermentation of *S. cerevisiae* in the DAC, as a function of pressure



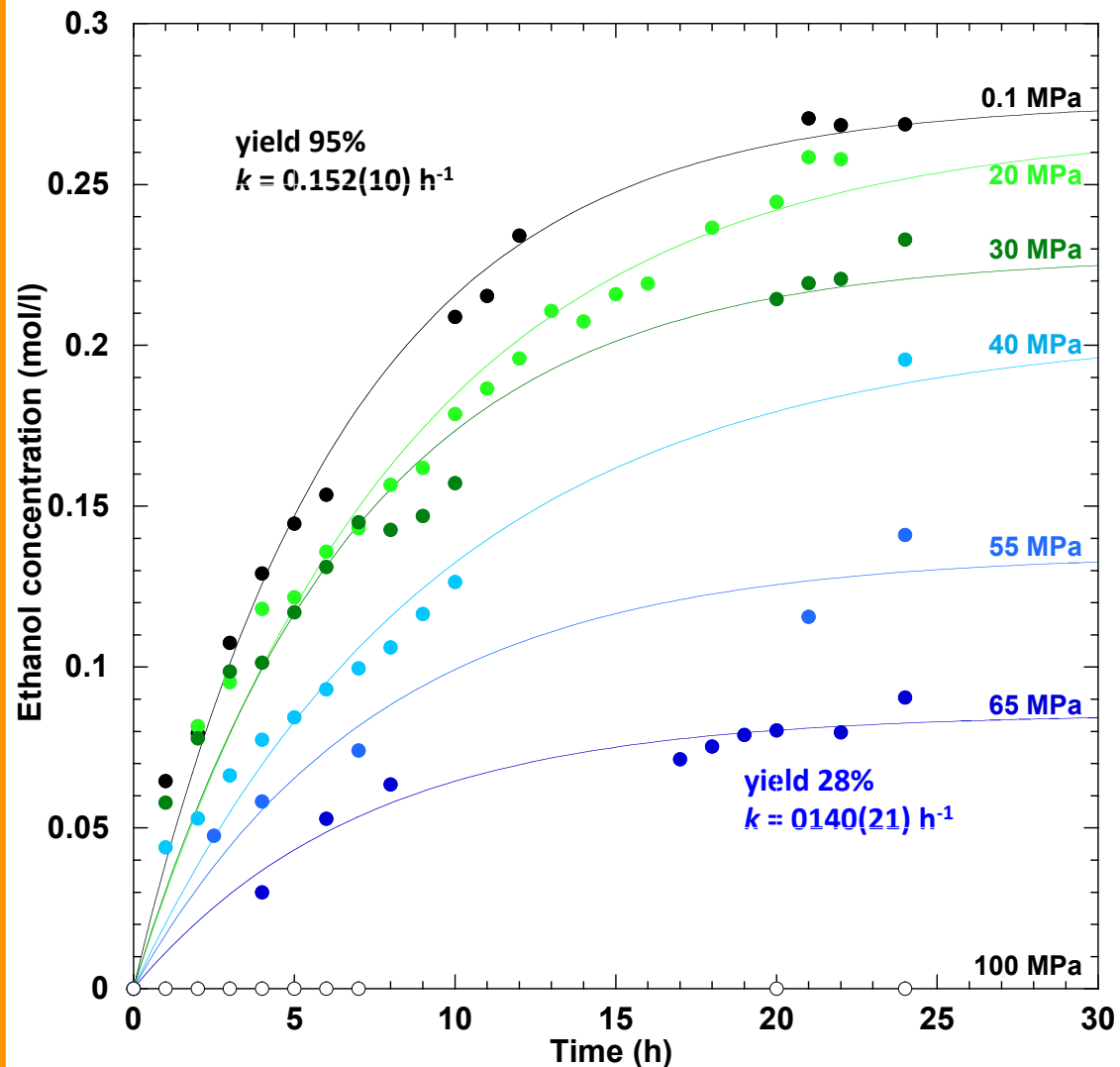
Observations : ambient to 10 MPa

- reaction twice-thrice faster
- yield almost at the theoretical limit

Interpretation

- enhanced uptake of glucose
- enhanced activity of one/several enzymes of the glycolysis and/or fermentation pathways.
- no measurable lag phase tends to exclude pressure-induced increase in protein synthesis.
- more efficient expellation ethanol from the cell under pressure, due to an increase of passive diffusion.

Fermentation of *S. cerevisiae* in the DAC, as a function of pressure



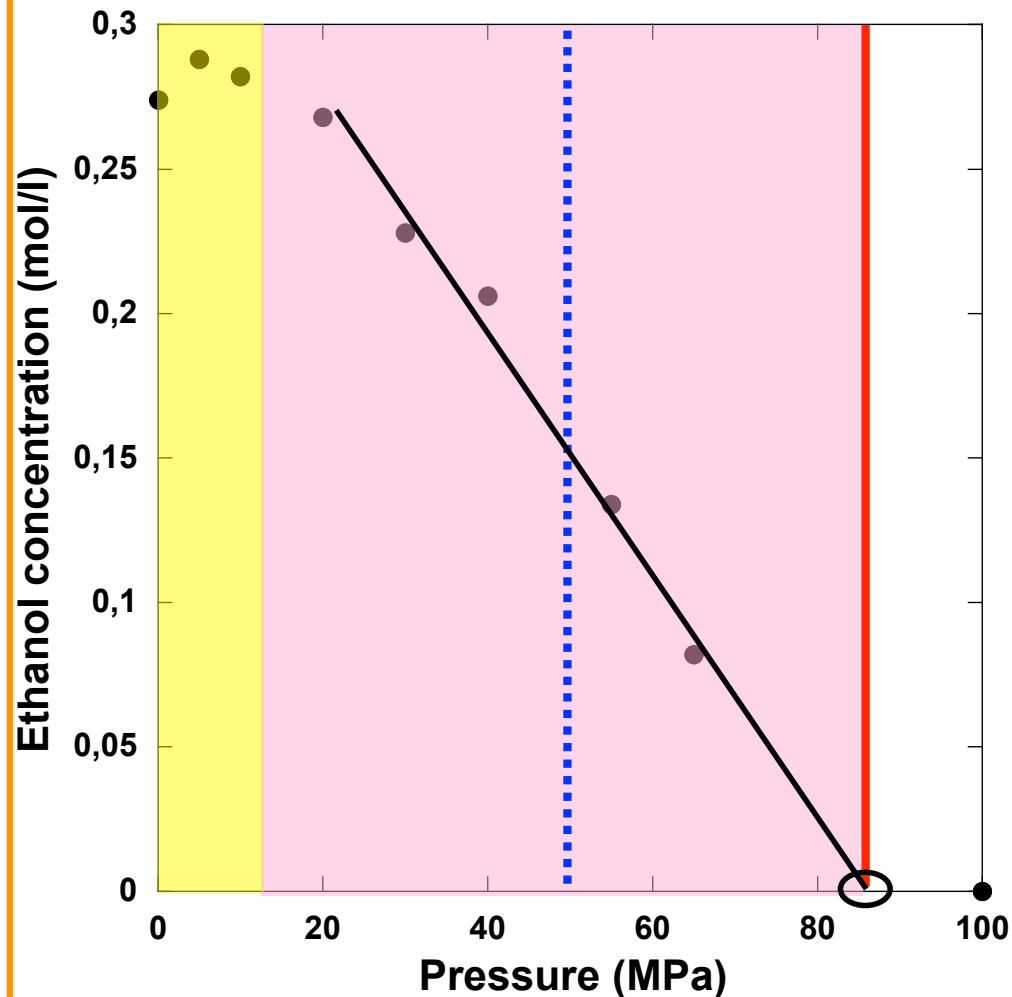
Observations : above 10 MPa,

- yield decreases
- reaction rate almost constant

At 40 MPa, yield of 68%
similar to Abe & Horikoshi (1997)

Alcoholic fermentation stops between 65
and 100 MPa

Fermentation of *S. cerevisiae* in the DAC, as a function of pressure



Maximal pressure for ethanol fermentation calculated at **87±7 MPa**

37 MPa higher than the predicted value by Abe *et al.* (2004) and than pressure limit for growth

0-10 MPa

Activated steps of ethanol fermentation:

- Increased glucose import?
- Activation of glycolysis or fermentation pathway enzymes?
- Facilitated excretion of ethanol?

20-87 MPa

Decrease of final ethanol production:

- Loss of only 1 log after 24 hours at 70 MPa
- Progressive inhibition of enzymes?

Uncoupling of growth and metabolism

Raman spectroscopy for probing live cells

- CARS

Coherent anti-Stokes Raman Scattering

- Resonant Raman Scattering

- SERS

Surface Enhanced Raman Scattering

For 2D mapping

Coherent anti-Stokes Raman Scattering

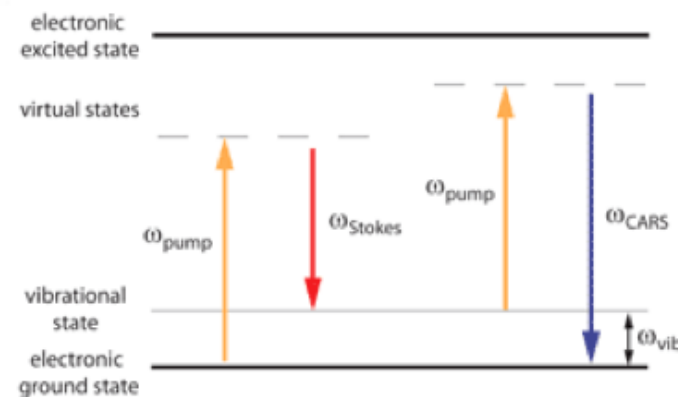
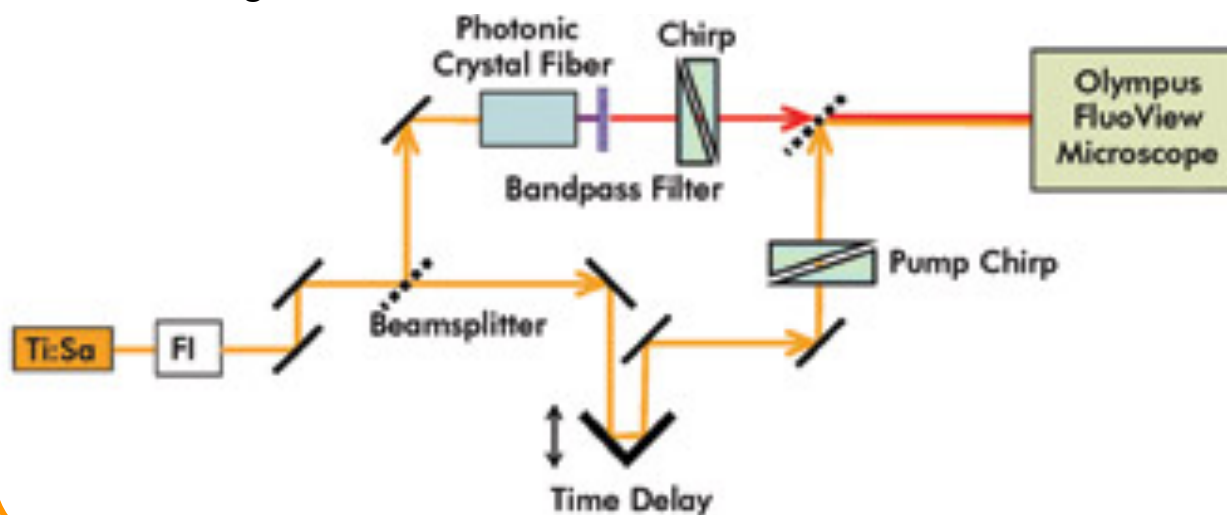
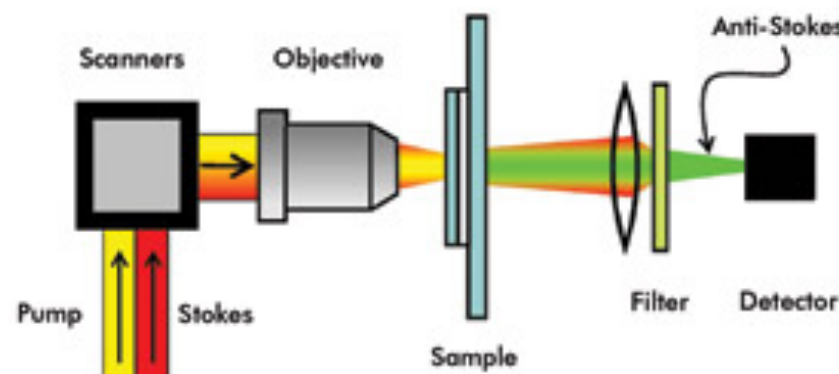
CARS microscopy provides label-free imaging

Non linear spectroscopy

A simple CARS microscope.

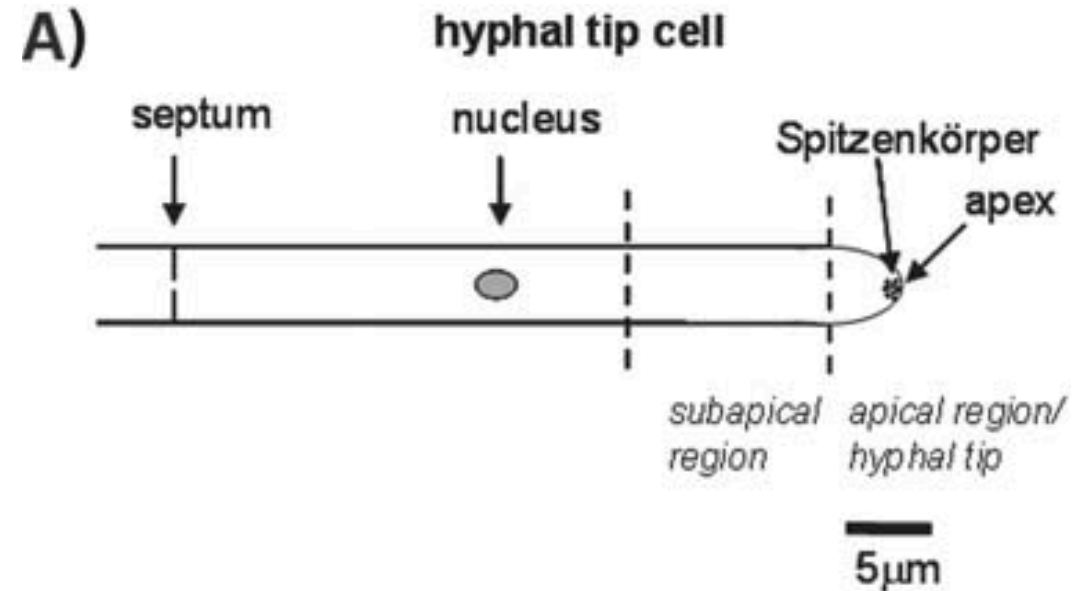
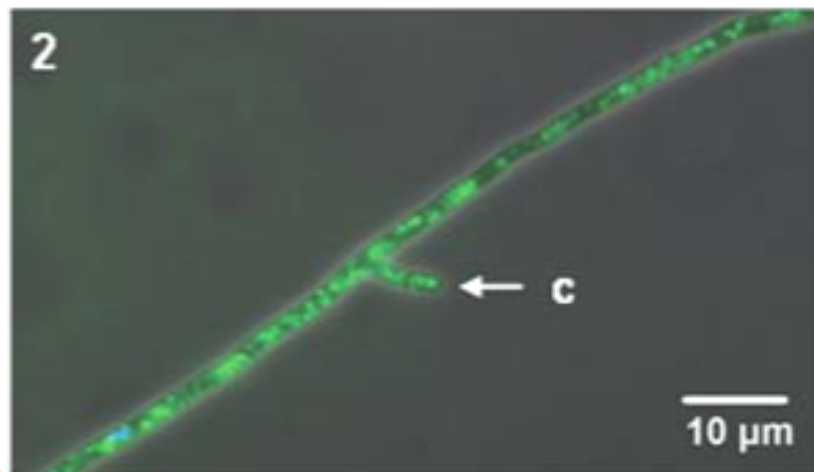
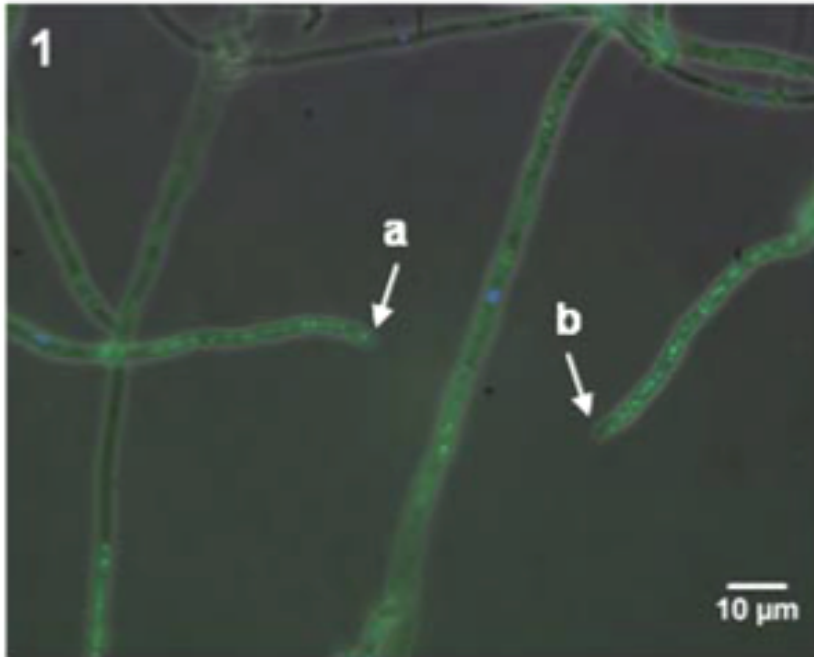
A beamsplitter splits pulses from an isolated femto-second Ti:sapphire or Nd:vandate laser.

Half goes to a photonic crystal fiber to generate the Stokes pulses, followed by a bandpass filter, before being recombined on a dichroic mirror.



$$\omega_{CARS} = 2\omega_{pump} - \omega_{Stokes}$$

Analysis of cytochrome distribution

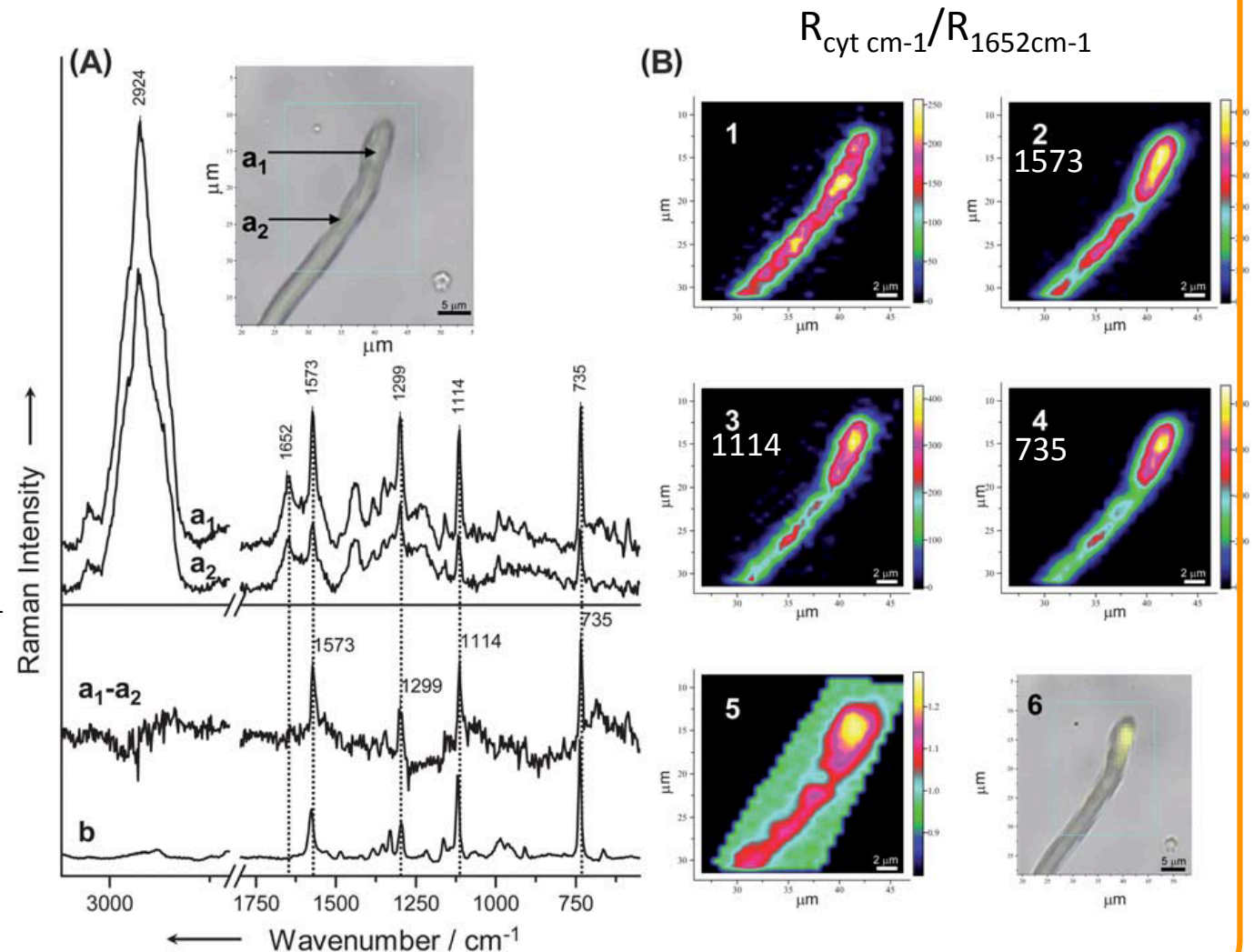


The cytochrome distribution in hyphal tip cells of the fungi *Schizophyllum commune*

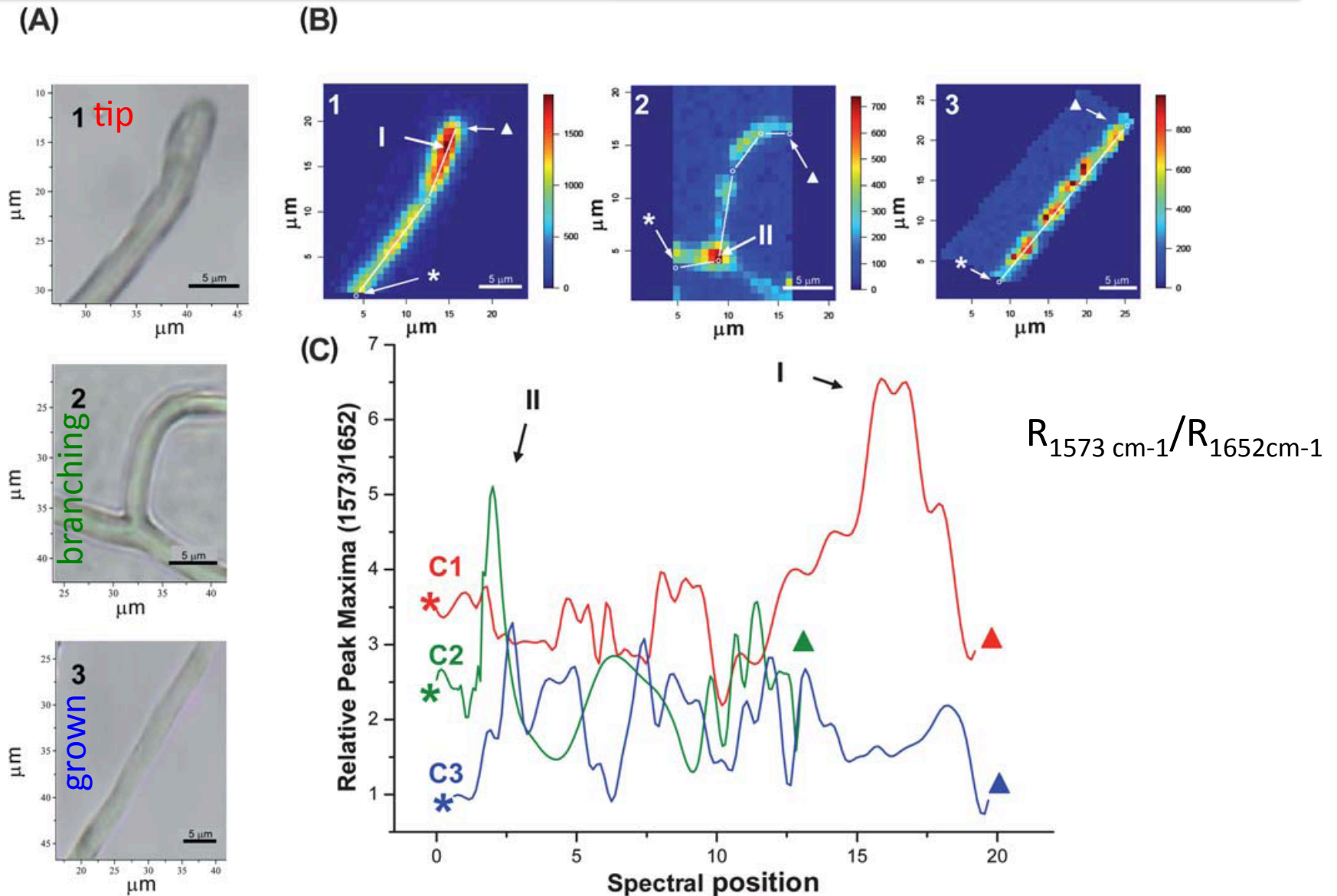


Resonant Raman spectra of a hyphal tip cell of *S. commune*

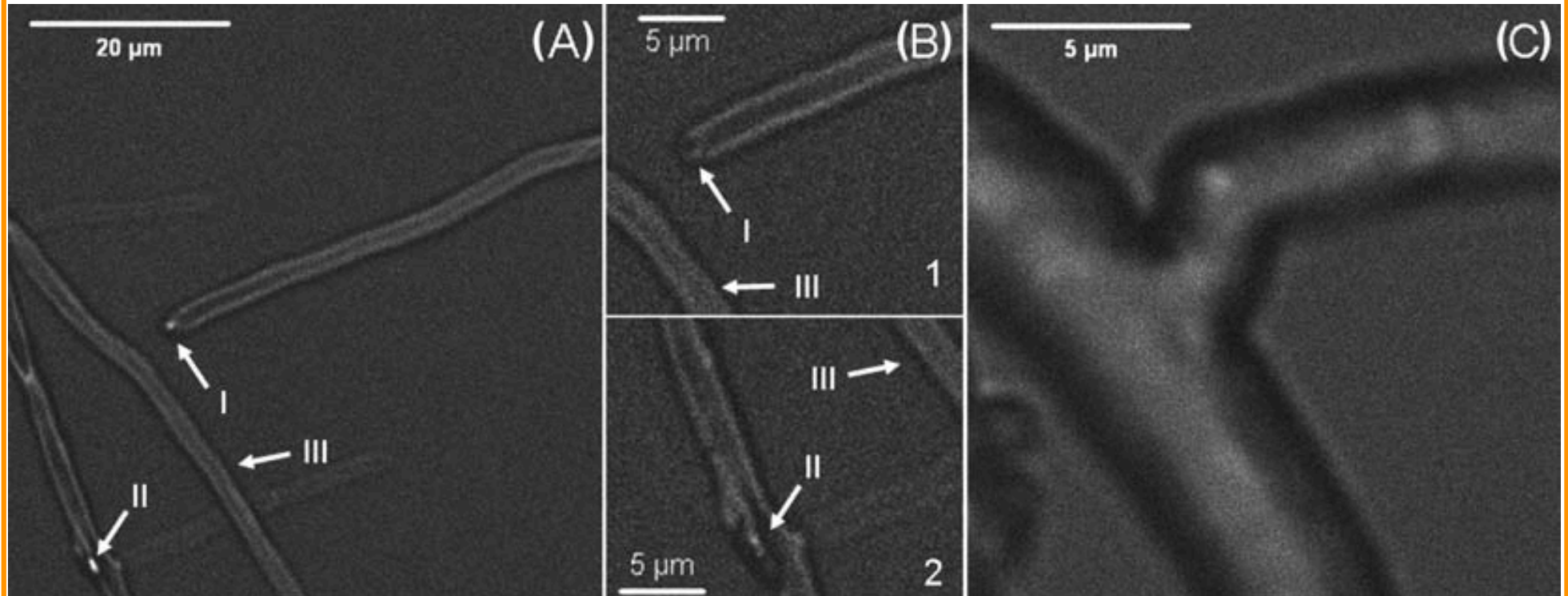
- 2924 cm^{-1}
asymmetric CH-stretching
vibration of methylene
groups from all cell
constituents
- 1652 cm^{-1}
protein, lipid and
polysaccharide vibrations
- 1573, 1299, 1114, 735 cm^{-1}
cytochrome vibrations
- Spatial resolution 0.7 μm
 $\lambda_0 = 532 \text{ nm}$ resonant with
the electronic absorption of
cytochrome



Resonant Raman spectra of a hypha of *S. commune*



CARS images of the fungal hyphae



@1572 cm⁻¹
cytochrome marker band
spec. resolution 20 cm⁻¹

@1552 cm⁻¹ under non
resonant conditions

@2990 cm⁻¹, CH vibration
spatial resolution 30x30 nm²
spec. resolution 110 cm⁻¹

In less than 25 s

Measurements of minute amount of biological and biotic products

- by **SERS**

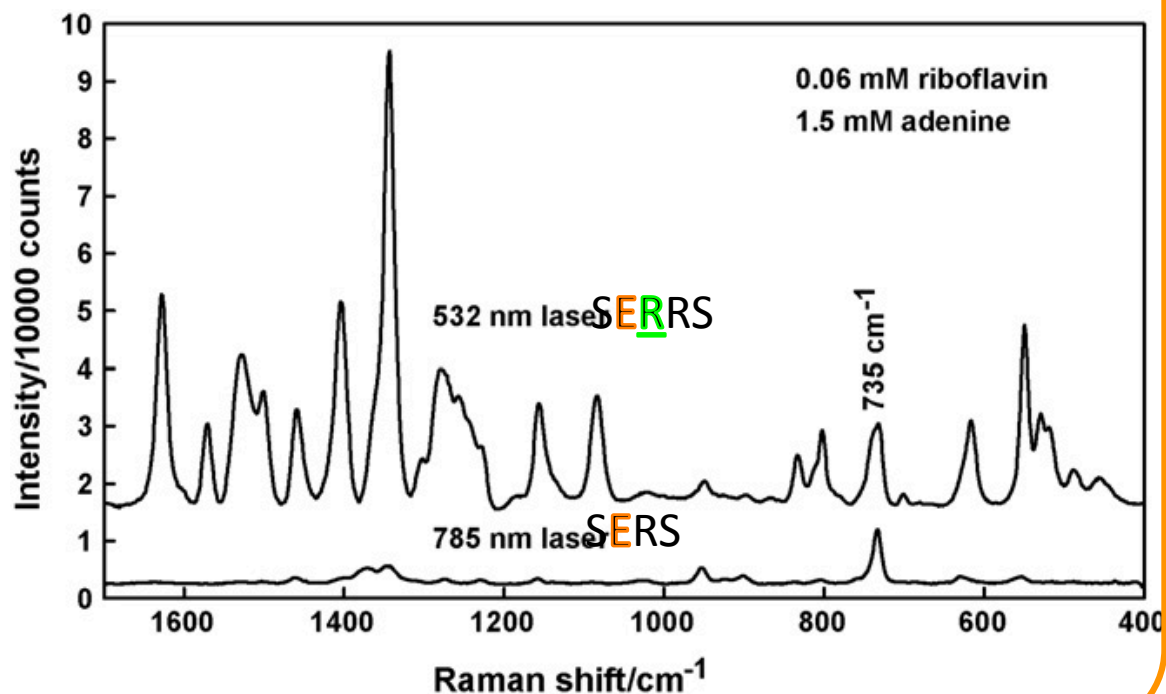
Surface Enhanced Raman Spectroscopy

- by **SERS**

Surface Enhanced Resonant Raman Spectroscopy

relies on the enhancement of the EM field by a substrate

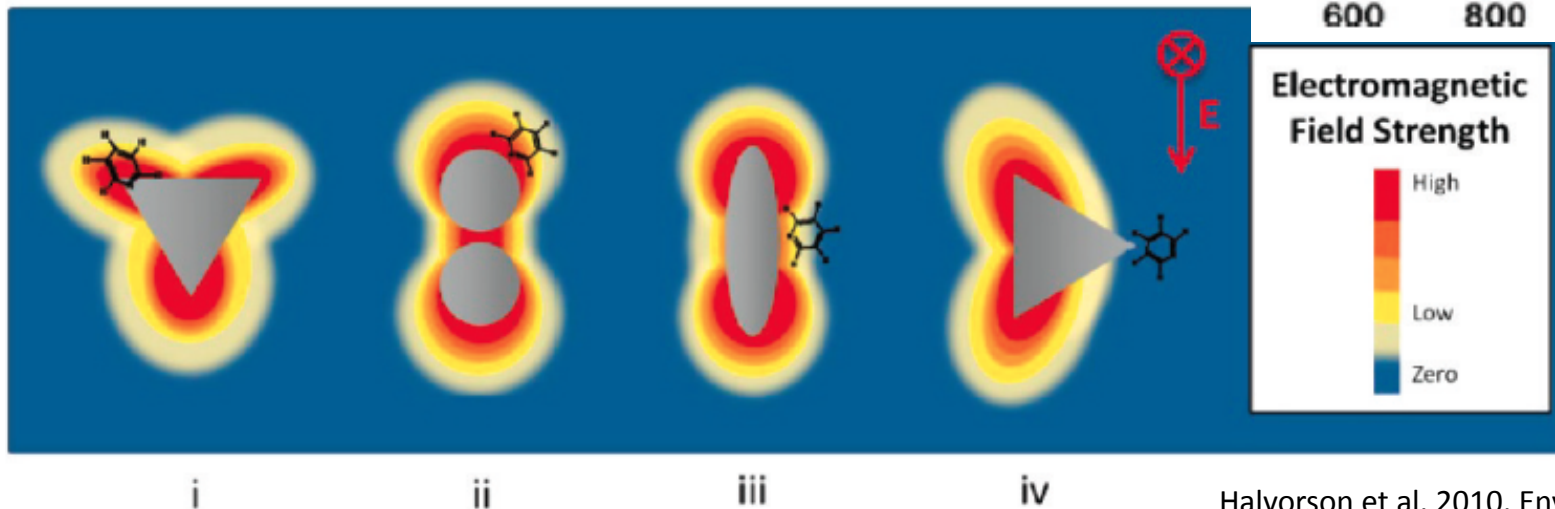
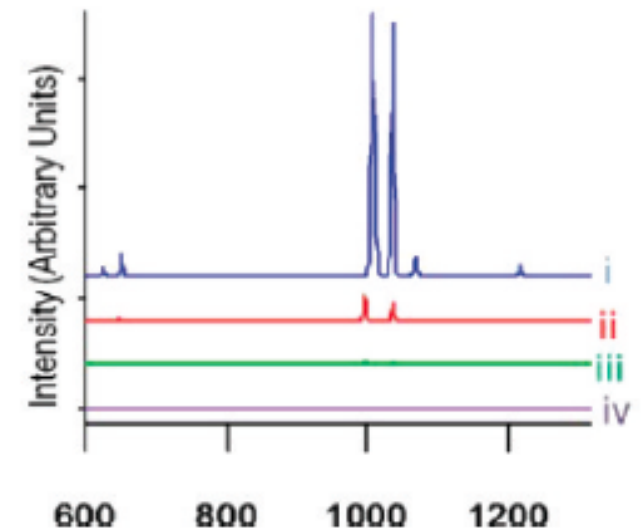
... requires contact between the analyte and metal nanoparticles



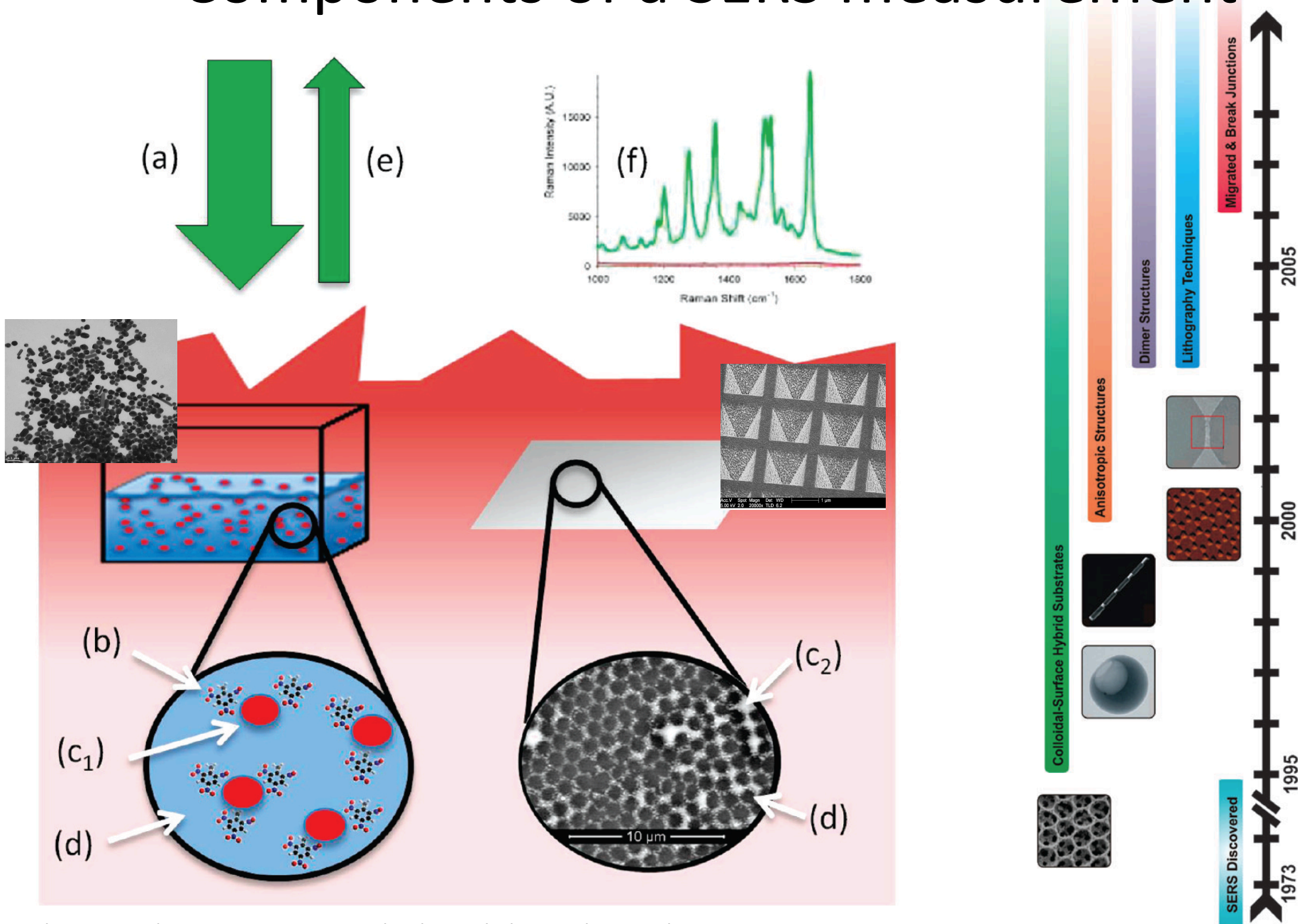
SERS & SERRS

- ✓ EM enhancement occurs when the incident laser excites surface plasmons (electrons at the metal surface that collectively oscillate upon excitation) thereby creating an electromagnetic field extending up to 20 nm from the metal enhancement $\times 10^4$ up to 10^{11}
- ✓ CT enhancement when transfer of electrons between the analyte and metal, $\times 10$ - 100
- ✓ resonance enhancement if the laser wavelength falls near an absorption wavelength of the sample

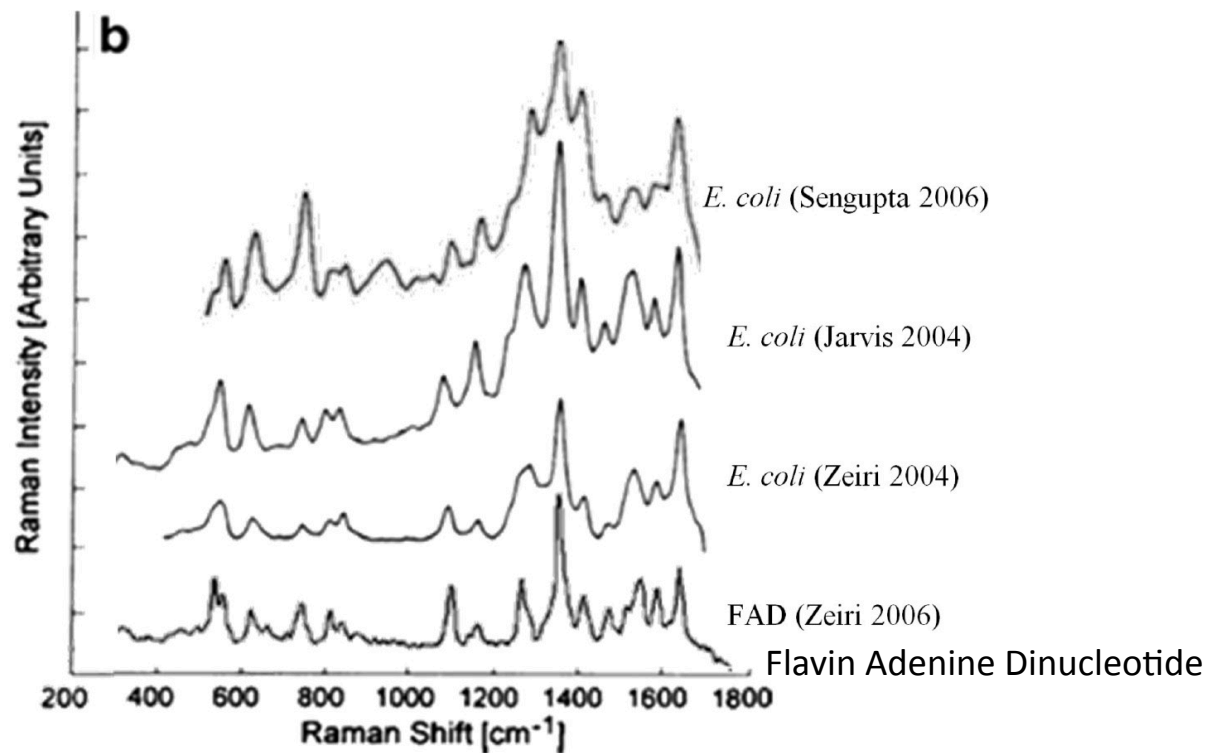
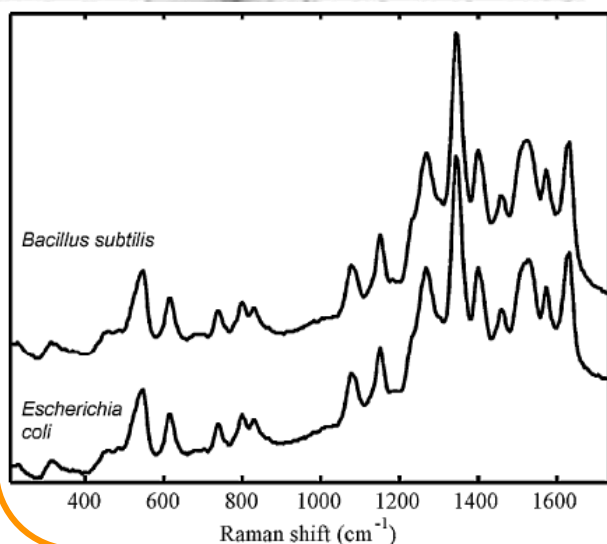
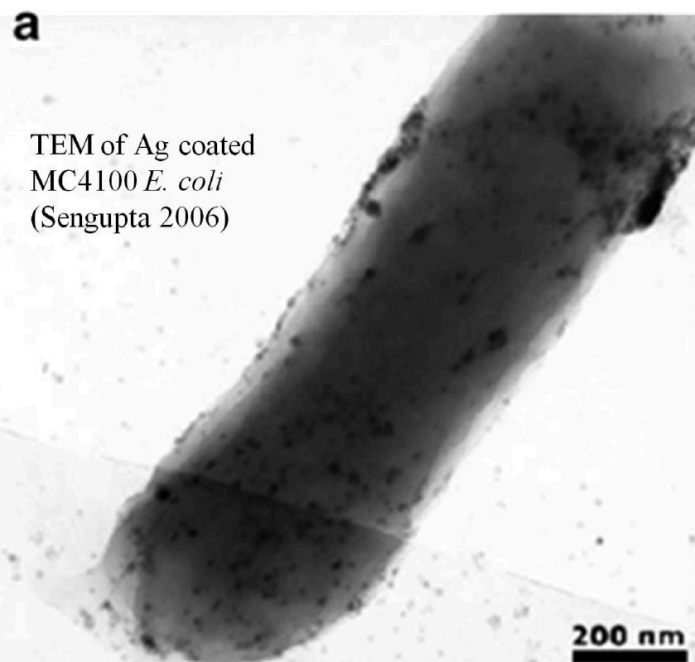
Expected SERS spectra for pyridine on nanostructures i, ii, iii, and iv



Components of a SERS measurement

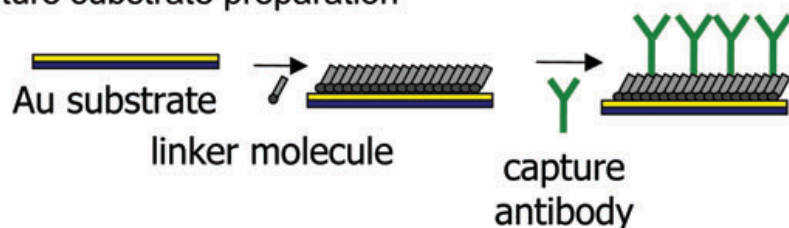


SER spectra of bacteria @332 cm⁻¹



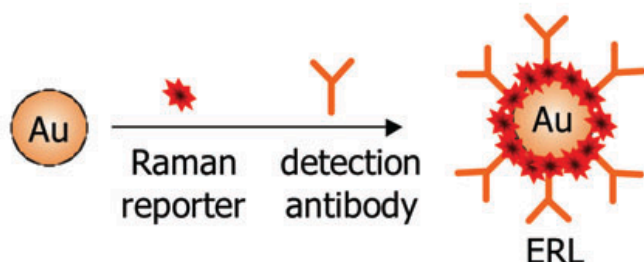
SERS-based immunoassays

Capture substrate preparation

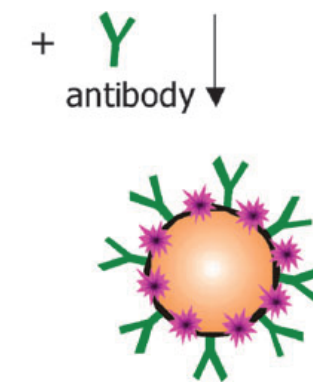
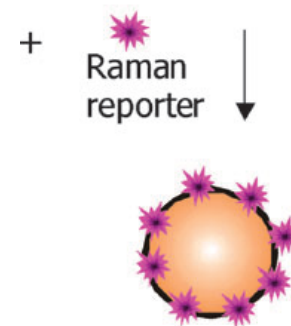
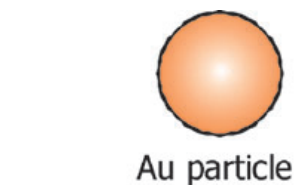


With *femtomolar* detection of the analyte

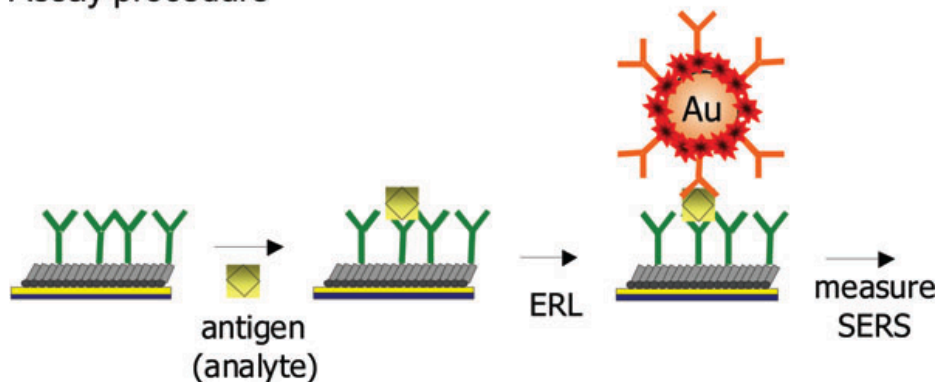
Extrinsic Raman Label (ERL) preparation



With 10^3 - 10^5 reporters on each particle



Assay procedure

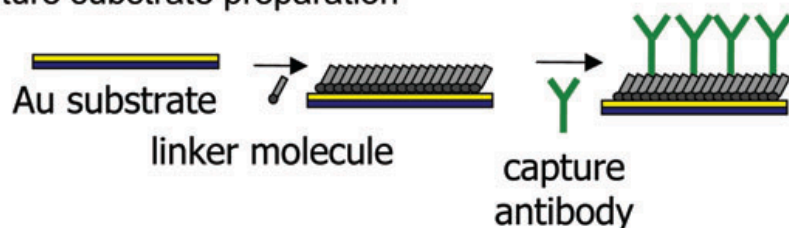


SER(R)S of the **extrinsic Raman labels (ERLs)**

Immunoassay platform

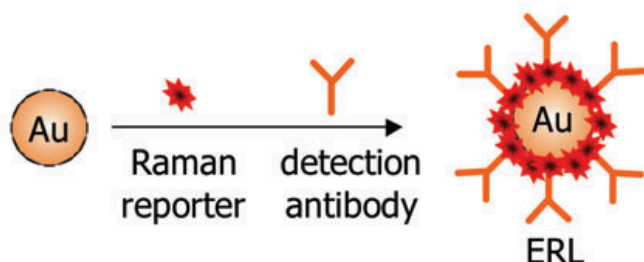
SERS-based immunoassays

Capture substrate preparation



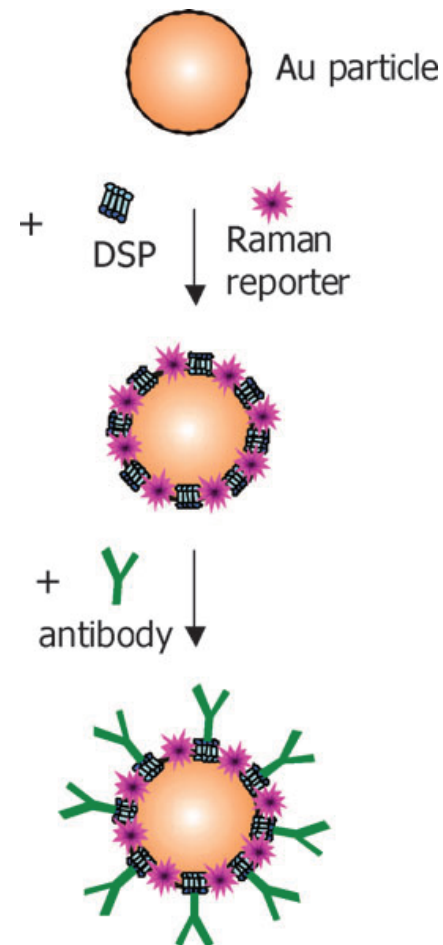
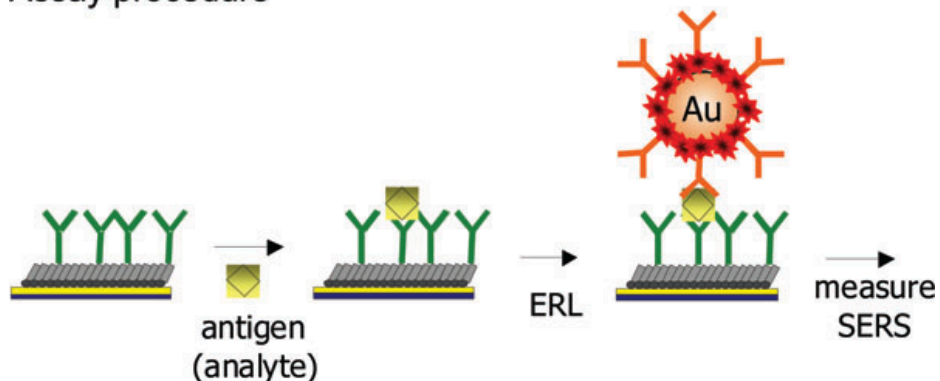
With *femtomolar* detection of the analyte

Extrinsic Raman Label (ERL) preparation



With 10^3 - 10^5 reporters on each particle

Assay procedure



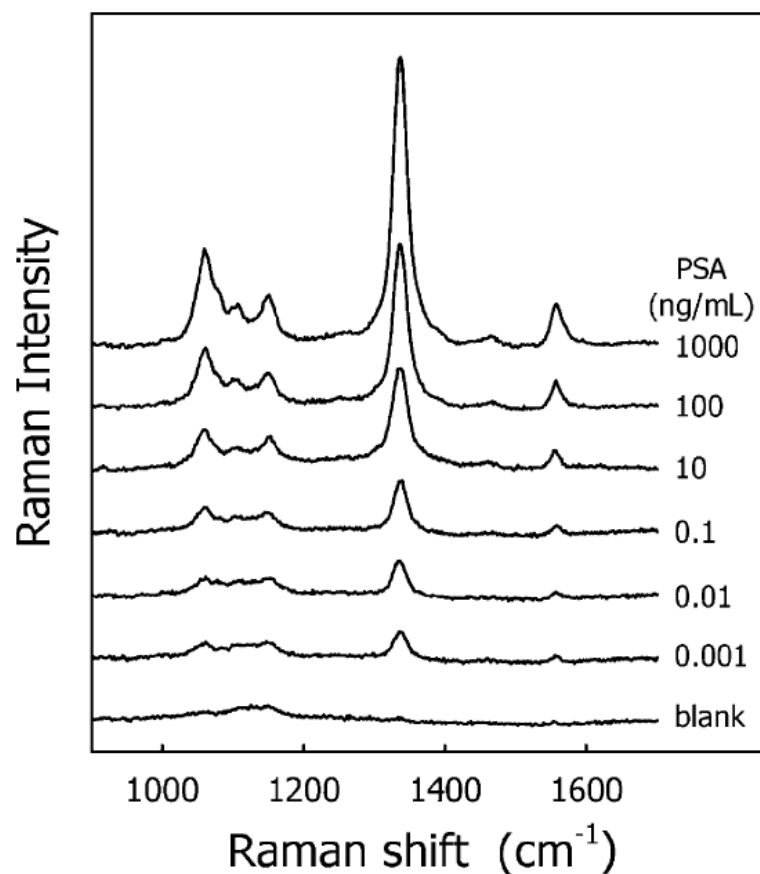
SER(R)S of the **extrinsic Raman labels (ERLs)**

Immunoassay platform

SERS-based immunoassays, examples

early disease diagnosis

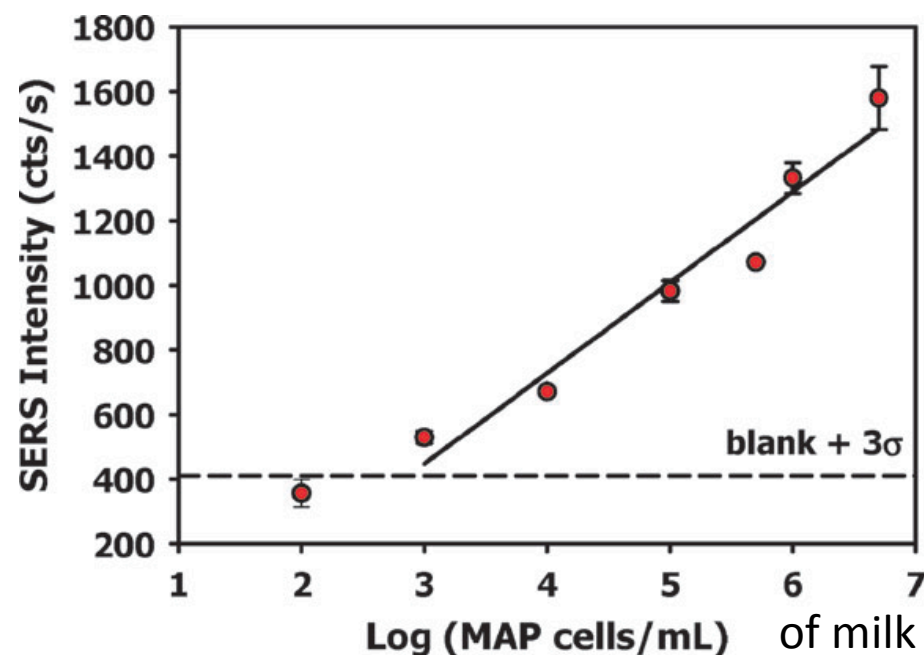
detection of prostate specific antigen (PSA)
a 33 kDa glycoprotein



ERL: 30 nm gold part. with a monolayer of DSNB
30s acq; LOD= 1 pg.mL⁻¹ or 30 fg

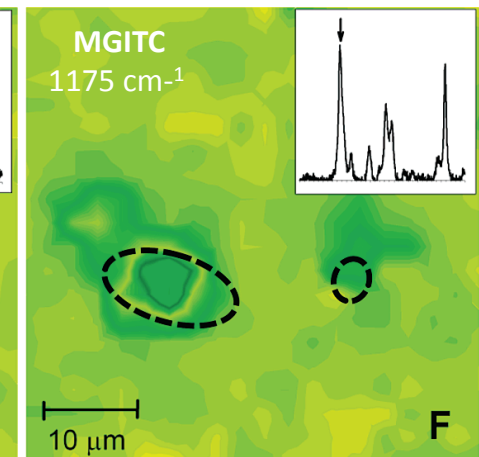
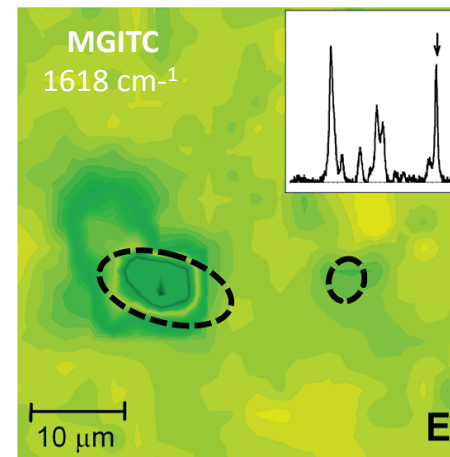
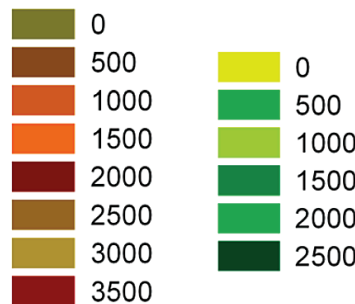
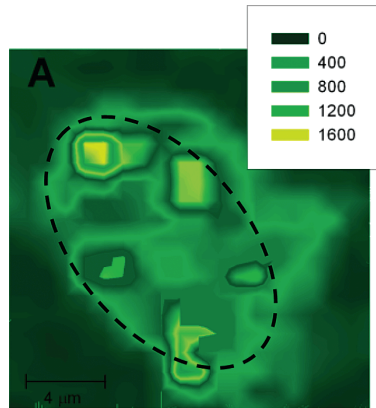
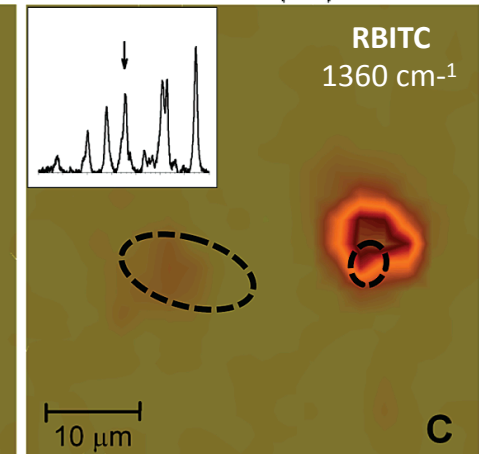
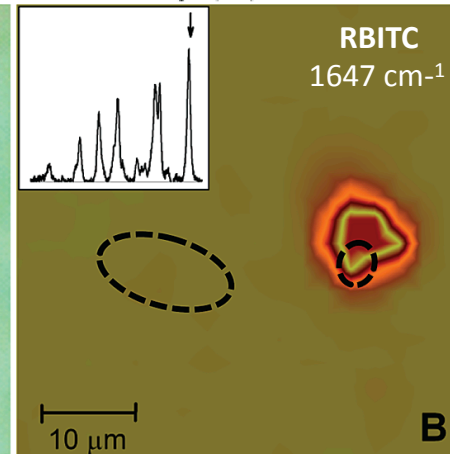
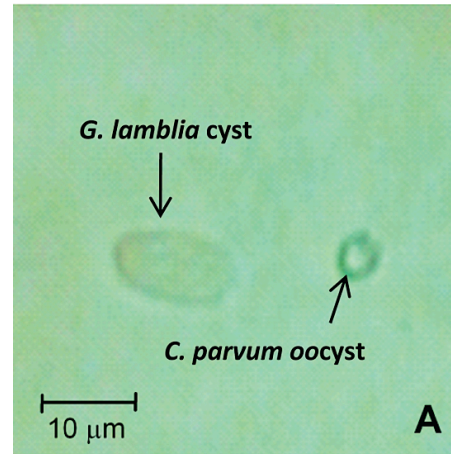
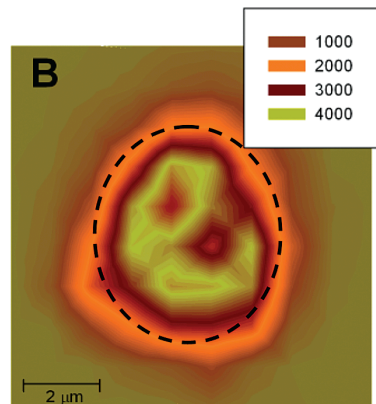
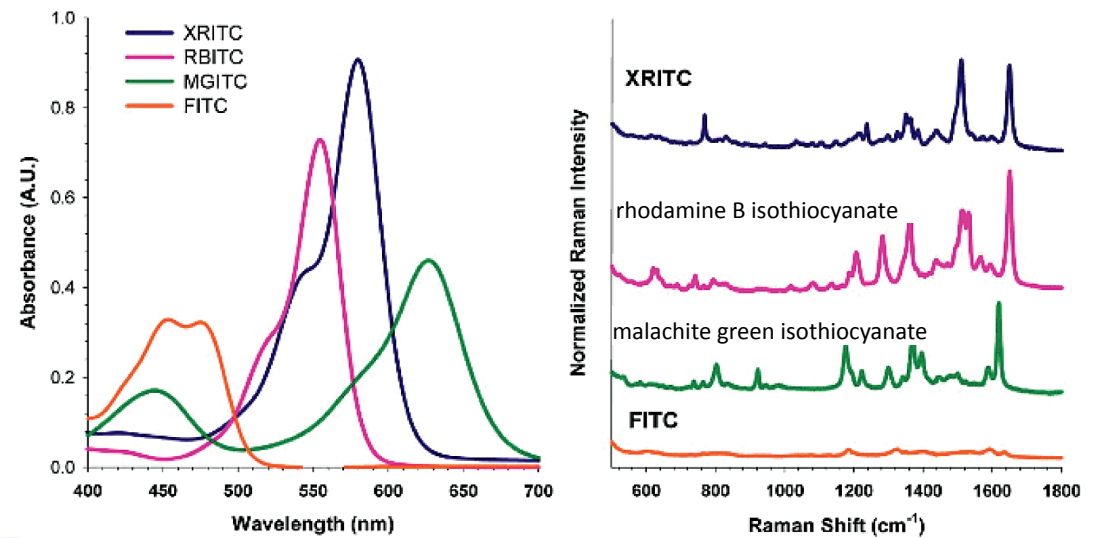
Pathogen detection

Mycobacterium avium subsp. paratuberculosis
(MAP)



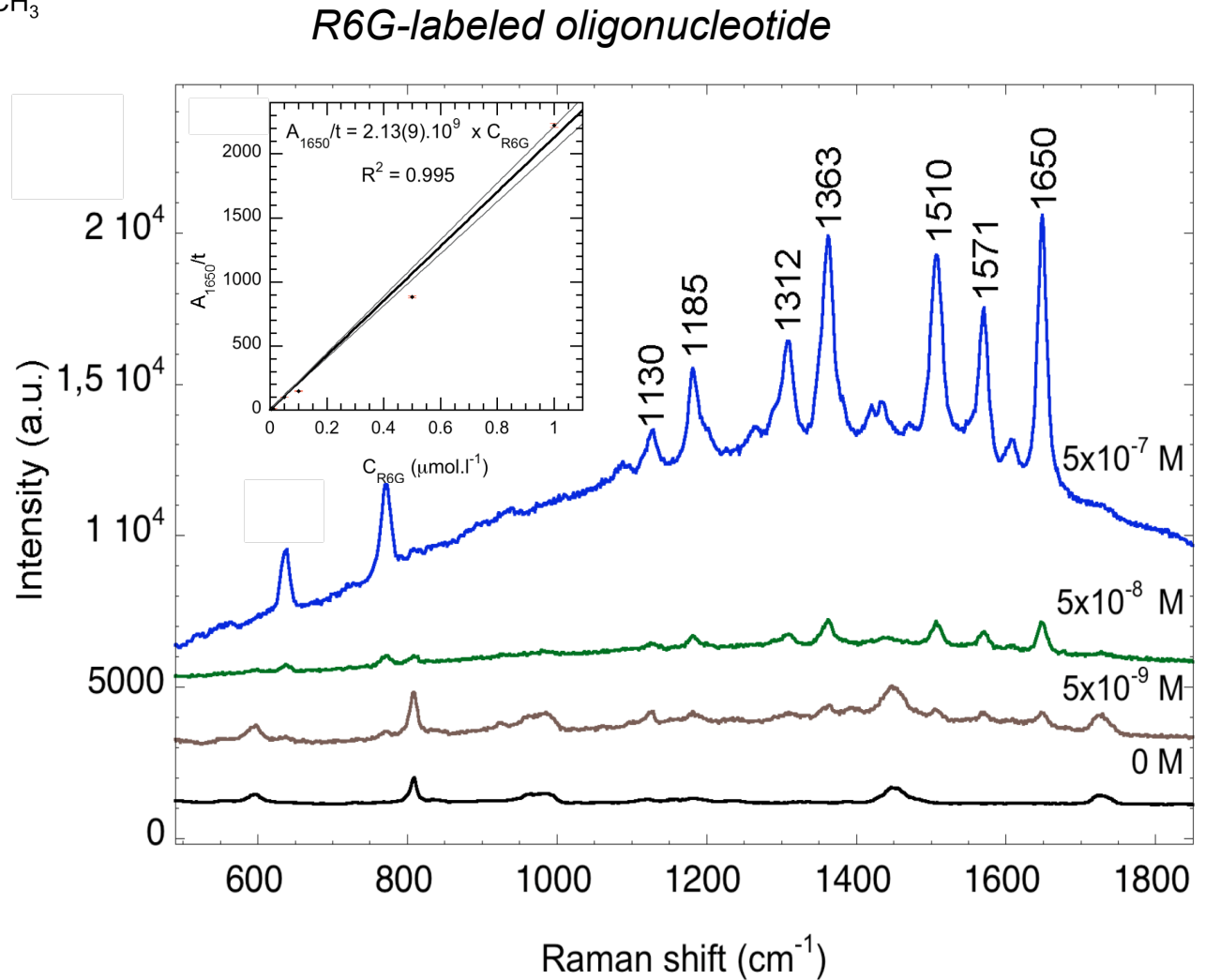
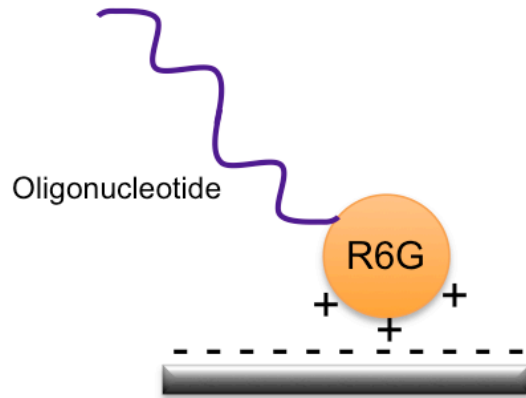
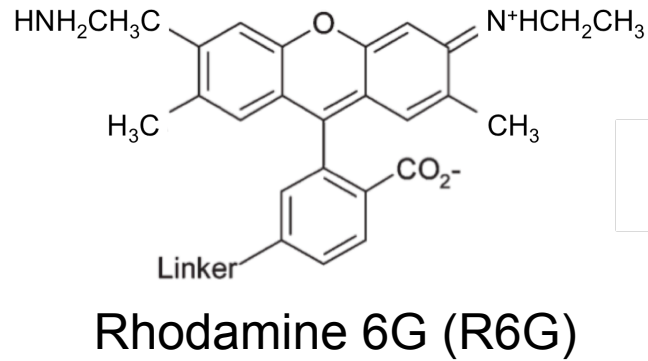
Classical culture method: 12-14 weeks
SERS (sample preparation, antigen extraction,
ERL incubation, and readout): 24hrs
LOD 1000 MAP mL⁻¹

An example of SERRS immunogold labeling for aquatic pathogens



20x20 pixels, 2 μm steps, 1s acquisition time
 $\lambda_e = 632.8$ nm, He-Ne laser

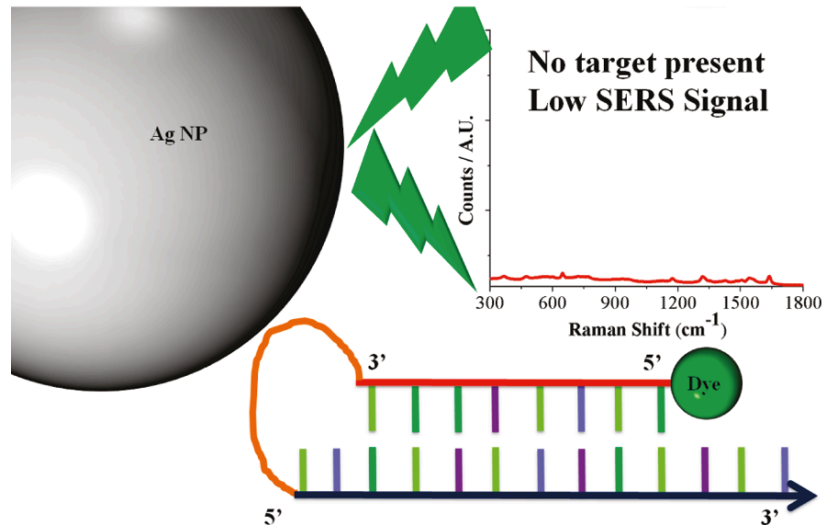
SERRS detection of minute amount of DNA



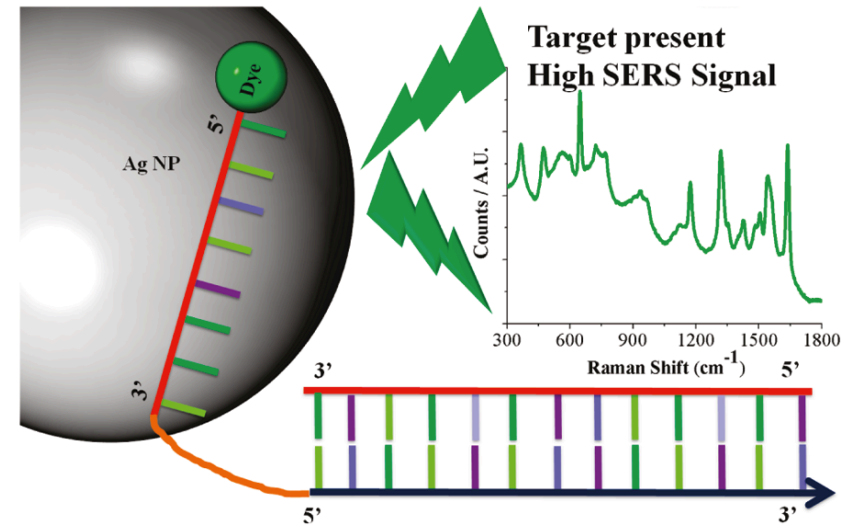
SERRS on single-stranded DNA

Van Lierop et al. 2011, Anal. Chem.

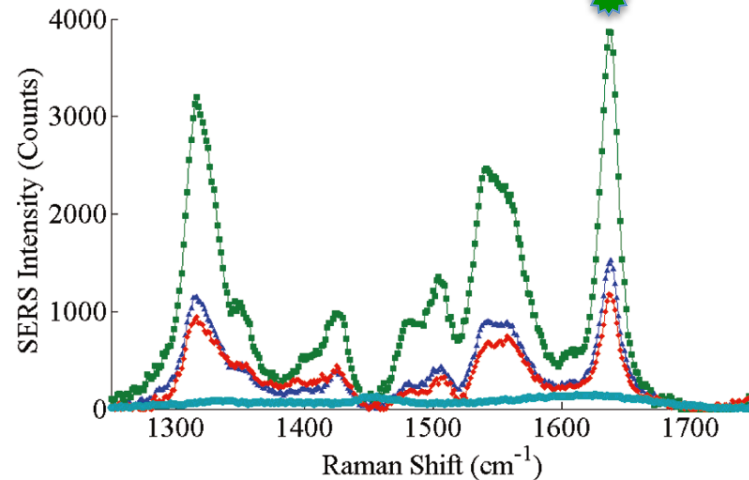
(A)



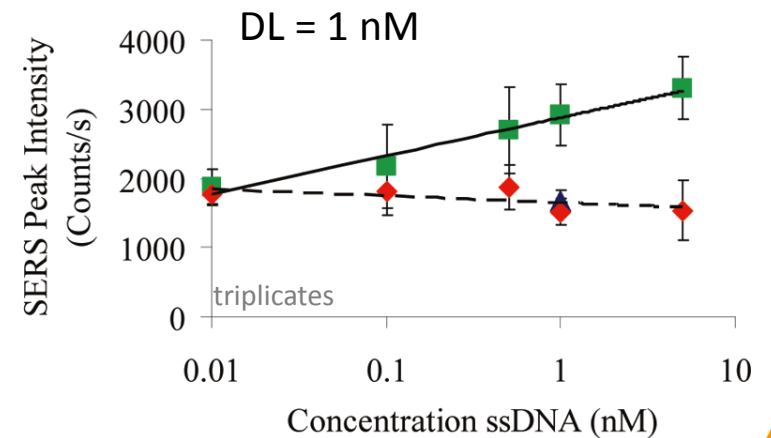
(B)



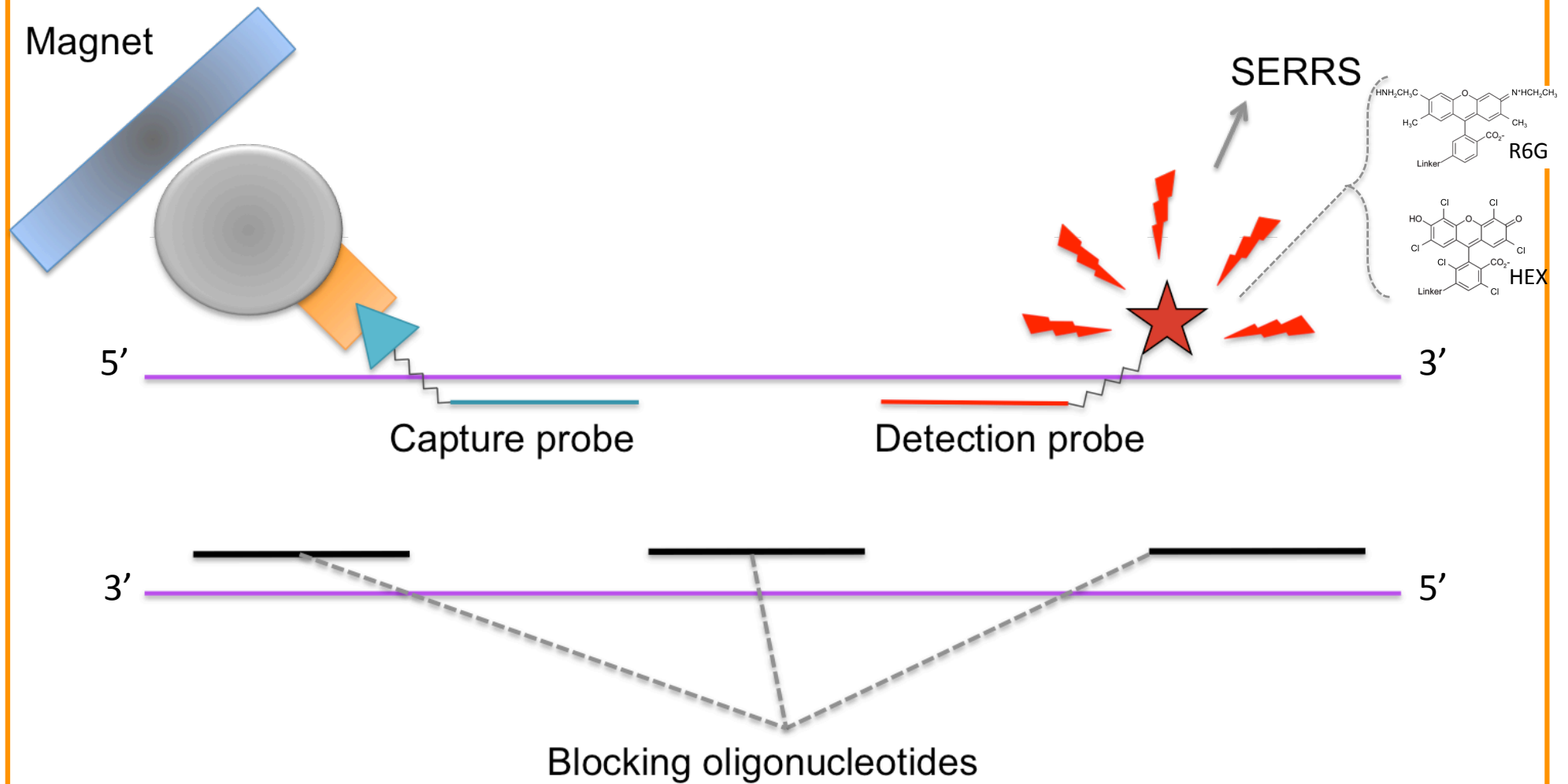
(A)



(B)

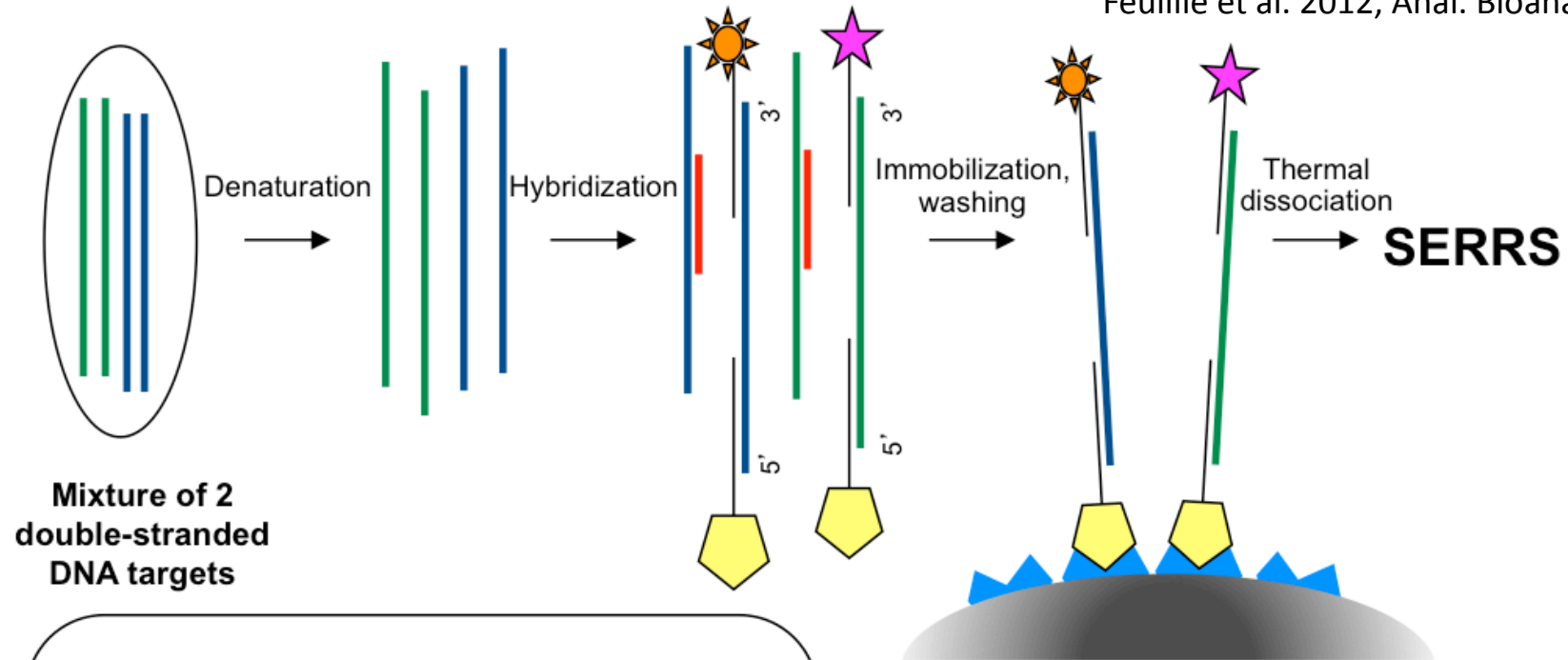


SERRS on double-stranded DNA

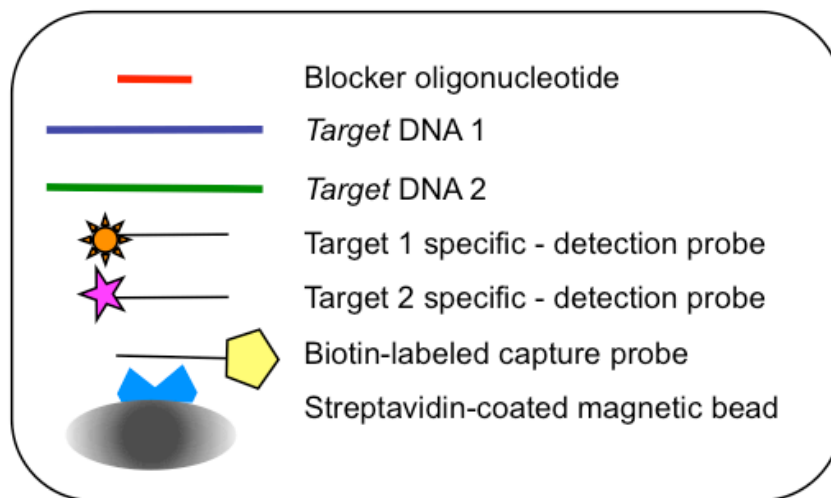


A SERRS hybridization assay for ds DNA

Feuillie et al. 2012, Anal. Bioanal. Chem.



Mixture of 2 double-stranded DNA targets

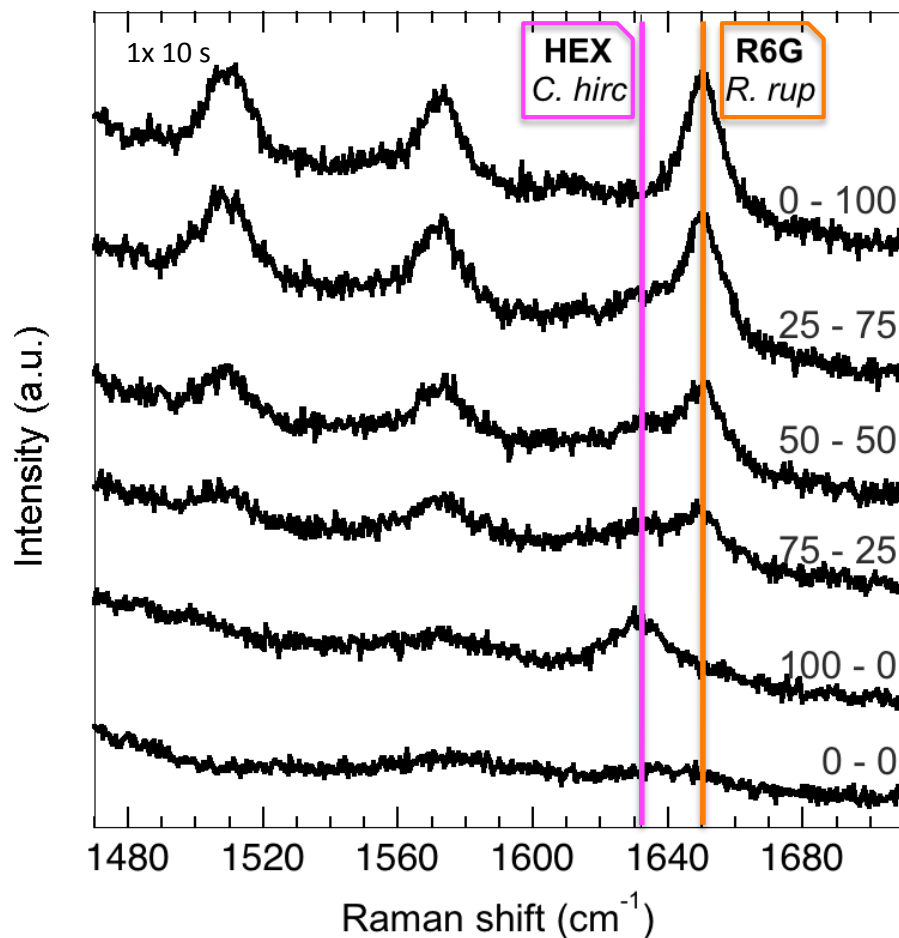


R6G

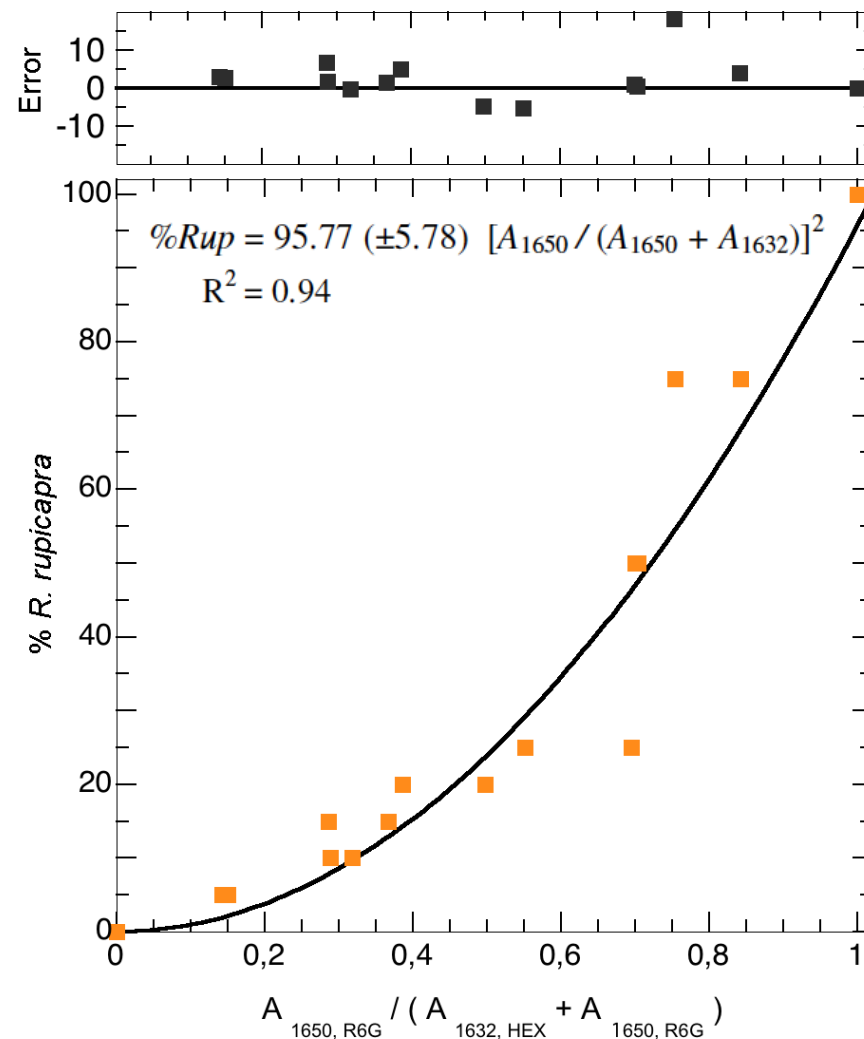
HEX



Simultaneous detection of 2 species



Identification of both sequences
 Total DNA amount = $5 \cdot 10^{-8}$ M
 DOL = $4 \cdot 10^{-10}$ M



Quantification of their relative amount

Detection of degraded DNA

Nomenclature

N5' / N3'

N5' / I3'

II5' / N3'

N5' / IV3'

V5' / N3'

I5' / I3'

II5' / I3'

I5' / IV3'

V5' / I3'

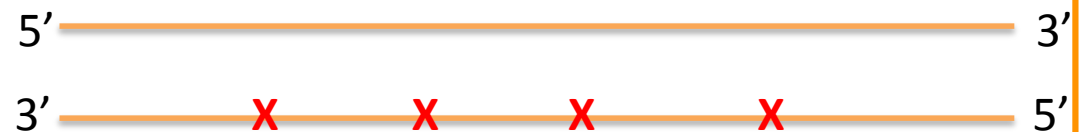
II5' / IV3'

V5' / IV3'

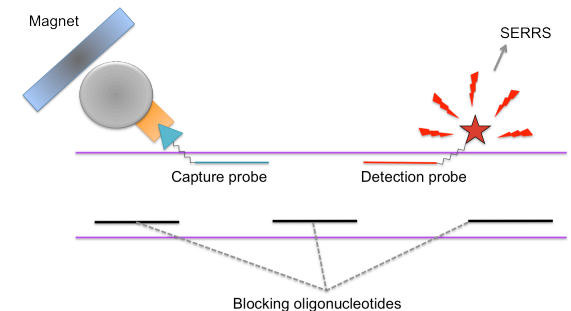
- Double-stranded DNA molecules of 139 bp
- Model blocking lesion : abasic sites
- Lesions distributed on both strands

N = no abasic site
I = 1 abasic site
II = 2 abasic sites ...

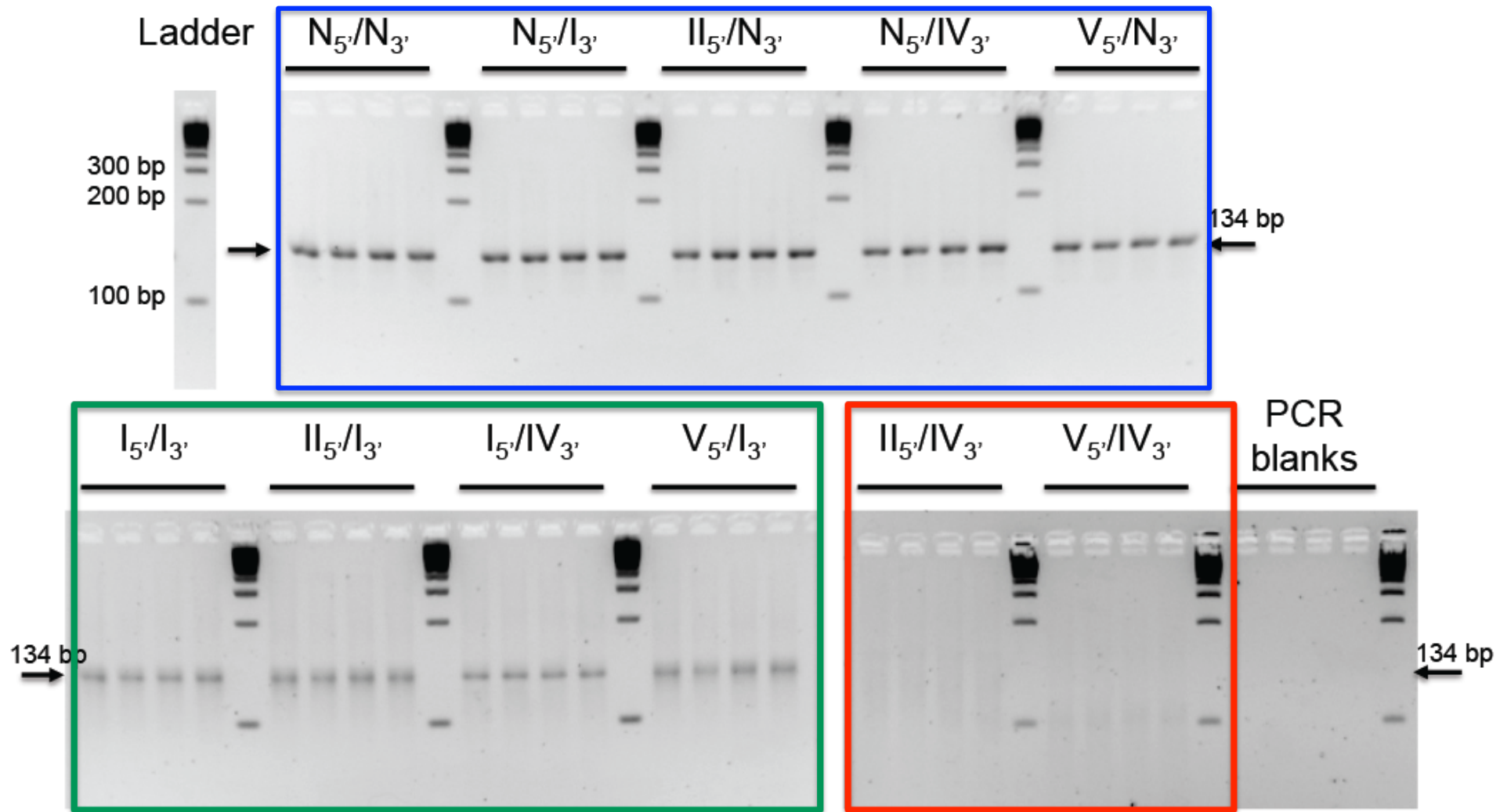
Example : N5' / IV3'



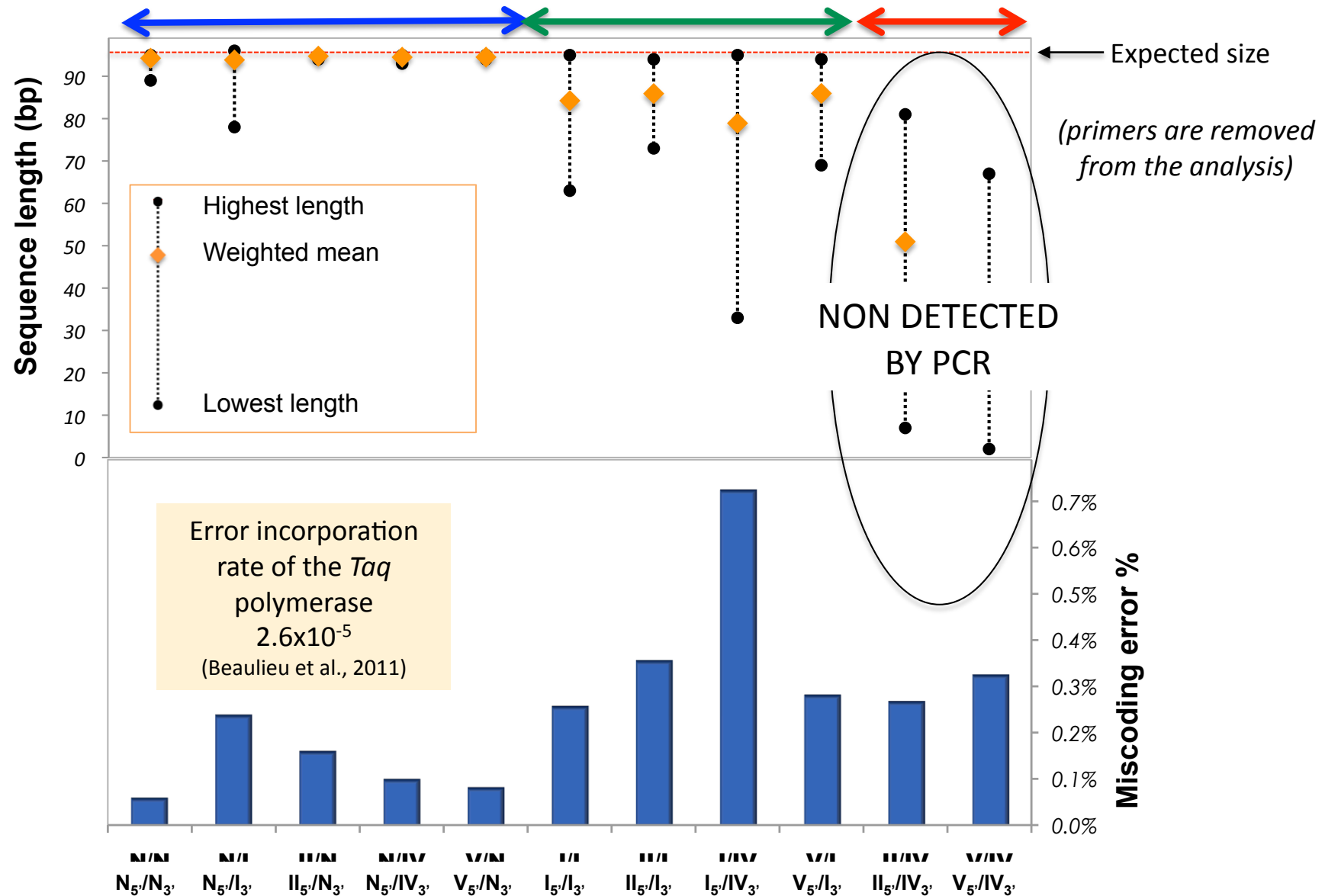
PCR vs SERRS-hybridization assay



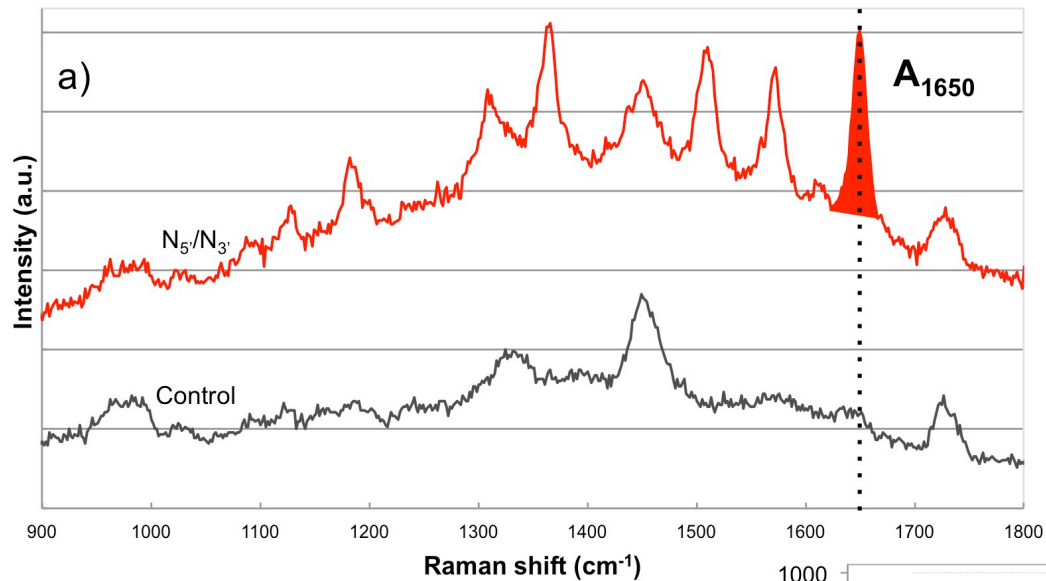
Detection of damaged DNA by PCR



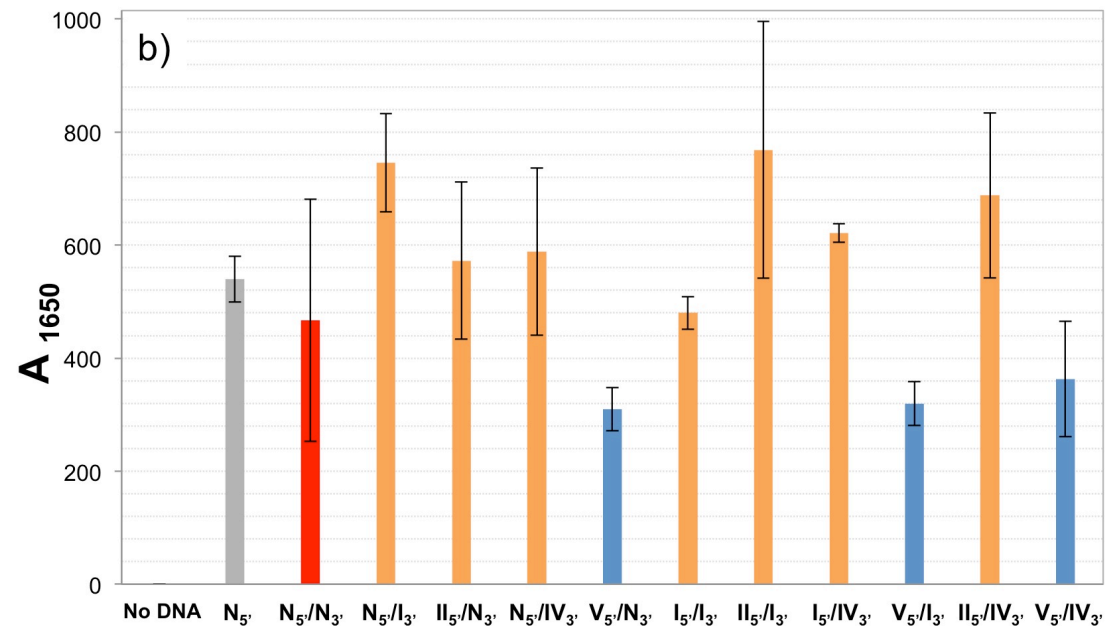
Detection of damaged DNA by PCR



Detection of damaged DNA by SERRS

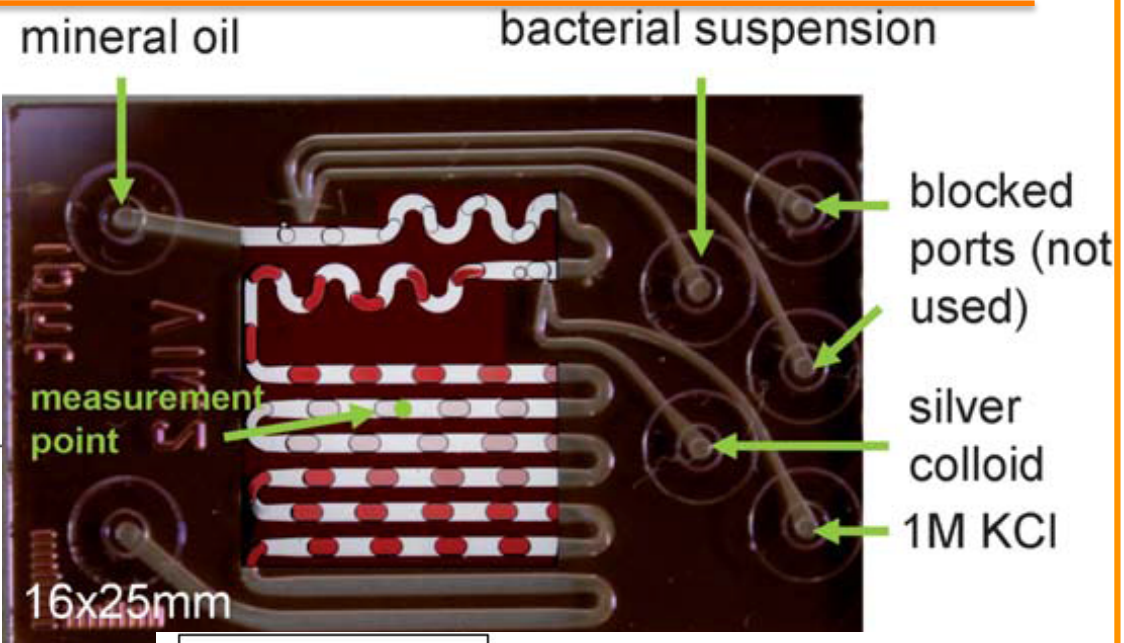


All molecules are detected
by the
SERRS-hybridization assay



On a Lab-on-a-chip device...

Walter et al. 2011, Lab Chip



Highly reproducible SER spectra of a strain of *E. coli*

