

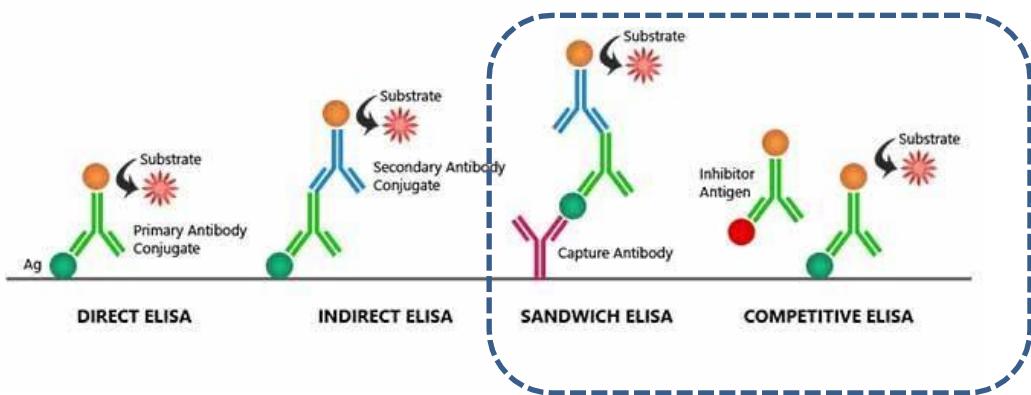
CONCENTRATION MOLES/LITER	METABOLITES/IONS	THERAPEUTIC DRUGS	STEROID AND AMINO ACID HORMONES	PROTEIN/POPPYPEPTIDE HORMONES	ANTIBODIES
$10^{-1}$	SODIUM CHLORIDE				
$10^{-2}$		ETHANOL			
(mM)	GLUCOSE UREA				
$10^{-3}$	CHOLESTEROL CALCIUM TRIGLYCERIDES	SALICYLATE ACETAMINOPHEN			IgG (Total)
$10^{-4}$	PHENYLANINE	THEOPHYLLINE			
$10^{-5}$	AMMONIA IRON BILIRUBIN	GENTAMICIN			
( $\mu$ M)					
$10^{-6}$			CORTISOL $T_4$ (Total)	THYROXINE BINDING GLOBULIN PLACENTAL LACTOGEN	IgM (Total)
$10^{-7}$					IgG (SPECIFIC)
$10^{-8}$	DIGOXIN	CORTICOSTERONE $T_3$ (Total)	ESTRIOL PROGESTERONE	PROLACTIN HCG	SYphilis RUBELLA ETC.
(nM)					
$10^{-9}$			$T_4$ (Free)	INSULIN	IgE (Total)
$10^{-10}$		ALDOSTERONE		PARATHYROID HORMONE	
$10^{-11}$		TSI (Thyroxine Stim. Hormone)		HGH (Growth Hormone)	
(pM)				LH (Luteinizing Hormone)	
$10^{-12}$	ANGIOTENSIN OXYTOCIN VASOPRESFIN				

FIGURE 2. CLASSES OF CLINICALLY SIGNIFICANT ANALYTES AS A FUNCTION OF CONCENTRATION IN THE SAMPLE.

# Enzyme Linked Immuno-Sorbent Assay (ELISA)

	ELISA	HPLC	LC-MS/MS
Price	Low	Medium	High
No. analytes per run	1 target	Multiple target	Up to 650
Accuracy	Screening	Reference method, highly sensitive and precise	Reference method (accredited results), highly sensitive and precise

## Types of ELISA



Used for food and clinical analysis

An Enzyme-Linked Immunosorbent Assay (ELISA) is a widely used laboratory technique designed to detect and quantify soluble substances such as peptides, proteins, antibodies, and hormones. ELISA tests are based on the principle of antigen-antibody interaction and are commonly used in diagnostics, research, and various industries for their sensitivity and specificity.

### **Key Components of ELISA:**

- 1. Antigen:** The substance to be detected or measured.
- 2. Antibody:** A specific protein that binds to the antigen.
- 3. Enzyme:** An enzyme linked to an antibody or antigen that produces a measurable signal, often a color change, when a specific substrate is added.
- 4. Substrate:** A chemical that the enzyme acts upon to produce a detectable signal.

### **Types of ELISA:**

- 1. Direct ELISA:** Involves the direct attachment of the antigen to a plate, followed by the application of an enzyme-linked antibody.
- 2. Indirect ELISA:** Uses a two-step process where a primary antibody binds to the antigen, and a secondary enzyme-linked antibody binds to the primary antibody.
- 3. Sandwich ELISA:** Requires the antigen to be captured between two antibodies – a capture antibody bound to the plate and a detection antibody that binds to another site on the antigen.
- 4. Competitive ELISA:** Measures the concentration of an antigen by detecting its ability to compete with a labeled antigen for binding to an antibody.

## **Procedure:**

- 1. Coating:** The wells of a microplate are coated with the antigen or antibody.
- 2. Blocking:** A blocking agent is used to prevent nonspecific binding.
- 3. Incubation:** The sample containing the antigen or antibody is added and incubated to allow binding.
- 4. Washing:** Unbound substances are washed away.
- 5. Detection:** An enzyme-linked antibody or antigen is added, followed by a substrate to produce a measurable signal.
- 6. Measurement:** The signal, often a color change, is measured using a spectrophotometer or other appropriate device.

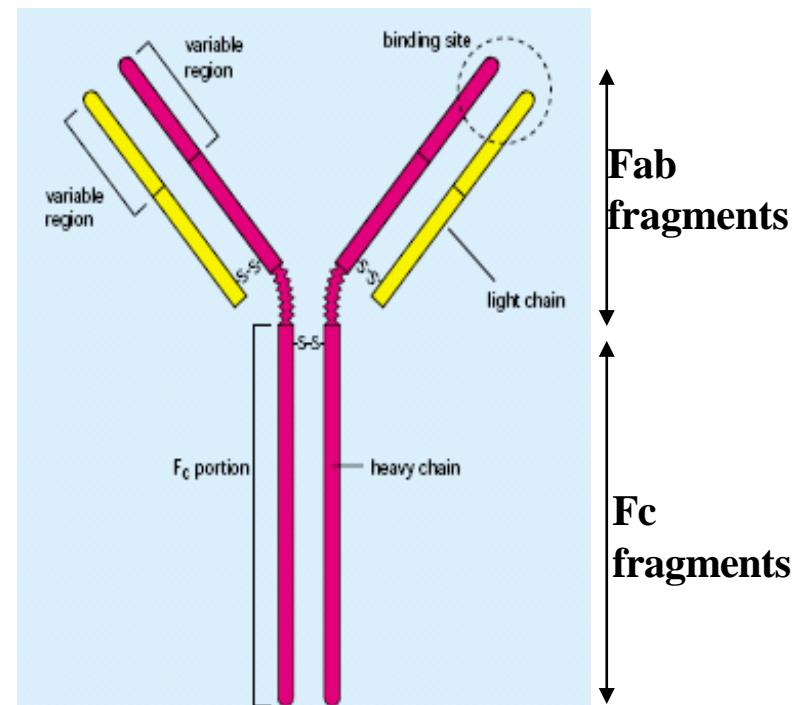
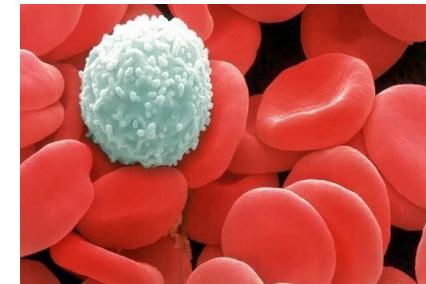
## **Applications:**

- **Medical Diagnostics:** Detection of diseases, including HIV, hepatitis, and COVID-19.
- **Food Industry:** Detection of allergens or pathogens.
- **Research:** Quantification of cytokines, hormones, and other biomarkers.

ELISA is valued for its accuracy, ability to process multiple samples simultaneously, and relatively

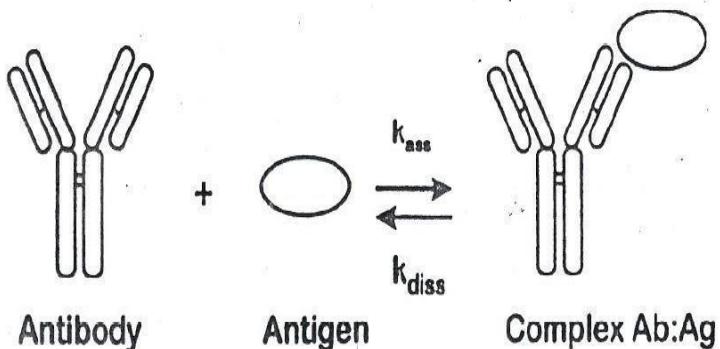
# Antibodies

- Proteins secreted by B-lymphocytes (type of white blood cell), in vertebrates.
- Recognise and bind to molecules (**antigens**) on foreign particles, marking them for destruction by T-lymphocytes.
- Each antigen may generate several antibodies for different sites (**epitopes**) on antigen.



IgG molecule

## Antibody - Antigen interaction



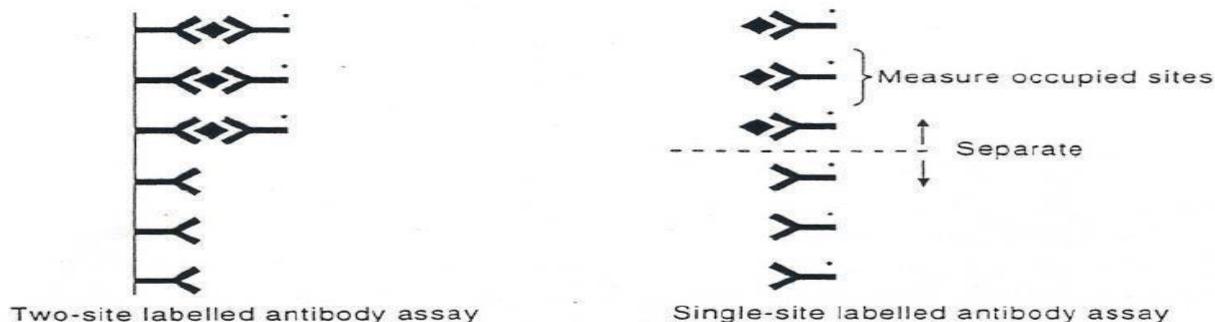
$$v = \frac{d[Ab:Ag]}{dt} = k_{ass} [Ab][Ag] - k_{diss} [Ab:Ag]$$

Equilibrium:  $\frac{d[Ab:Ag]}{dt} = 0$  and  $K_{aff} = \frac{k_{ass}}{k_{diss}} = \frac{[Ab:Ag]}{[Ab][Ag]}$

- Non - covalent
- Highly specific
- $k_{ass} \approx 10^6 - 10^8 \text{ M}^{-1}\text{s}^{-1}$
- $k_{diss} \approx 10 - 10^{-4} \text{ s}^{-1}$
- $K_{aff} \approx 10^6 - 10^{12} \text{ M}^{-1}$

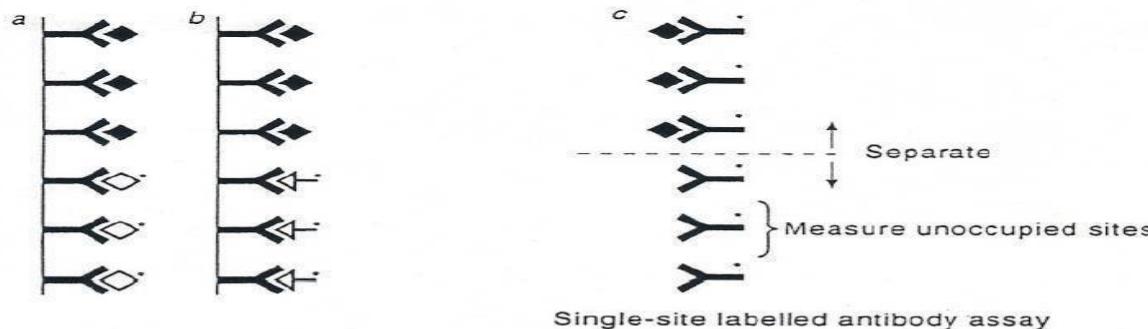
**A Noncompetitive immunoassay**  $Ab \rightarrow \infty$  for maximum sensitivity

Measurement of occupied sites



**B Competitive immunoassay**  $Ab \rightarrow 0$  for maximum sensitivity

Measurement of unoccupied sites

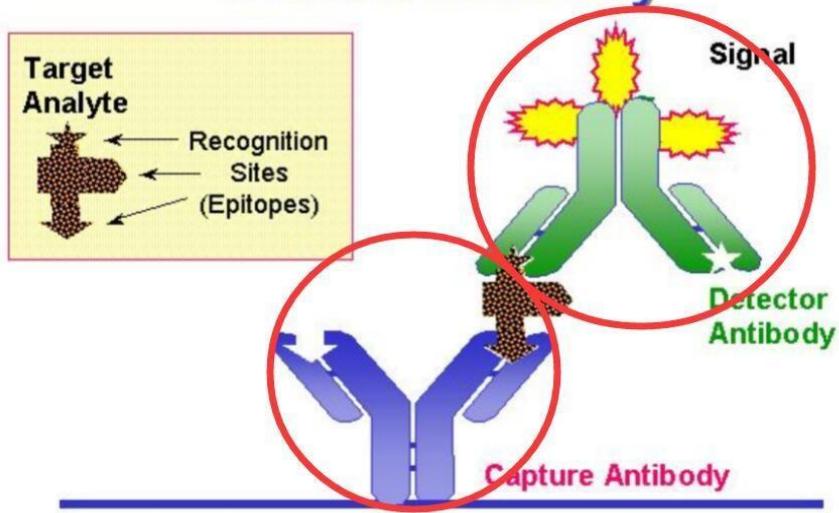


**Key**

- ◇ Labelled antigen
- ◀ Labelled anti-idiotypic antibody
- Y Labelled antibody
- ◆ Analyte

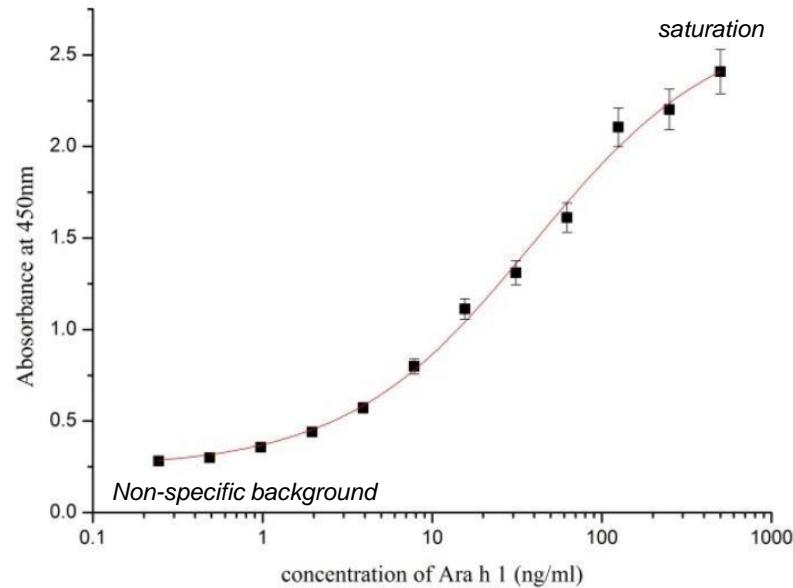
**Figure 6** Basic competitive and noncompetitive immunoassay designs. The distinction between noncompetitive (a) and competitive (b) reflects the way in which antibody binding site occupancy is observed. Labelled antibody methods are noncompetitive if occupied sites of the (labelled) antibody are directly measured, but are competitive (Bb) when unoccupied sites are measured. Labelled antigen (Ba) or labelled anti-idiotypic antibody methods (Bb) rely on measurement of sites unoccupied by analyte, and are therefore invariably of competitive design.

# Double Antibody Sandwich Immunoassay



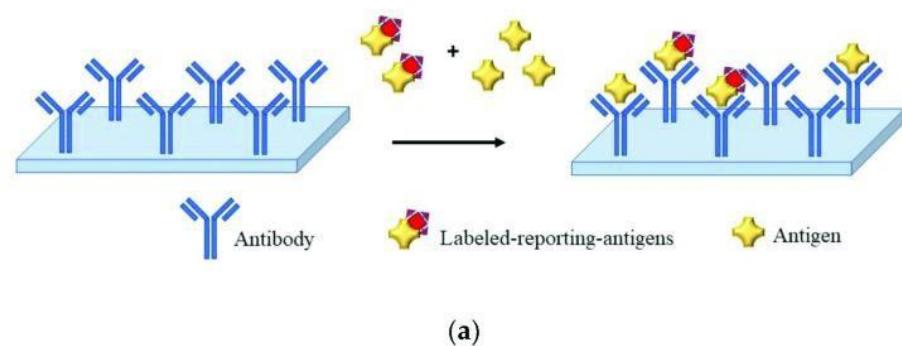
*Excess Antibody LOG SCALE!!*

Dosaggio di un allergene (proteina) di arachide

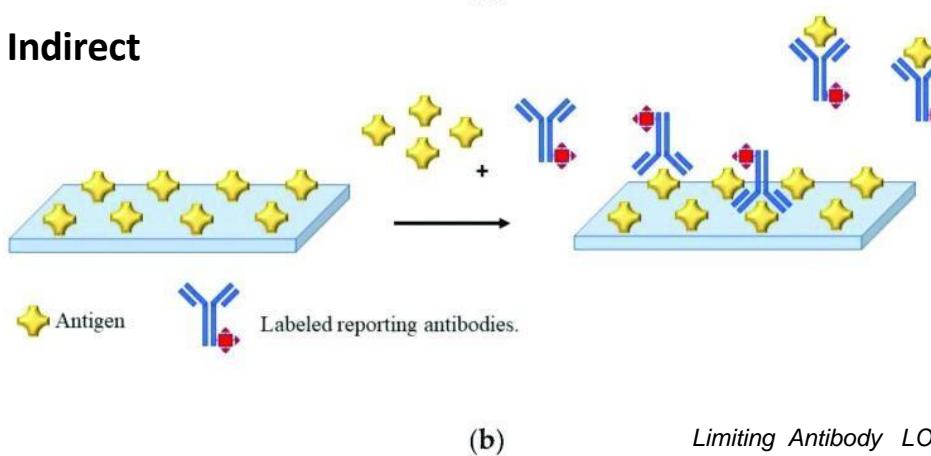


# Competitive immunoassays

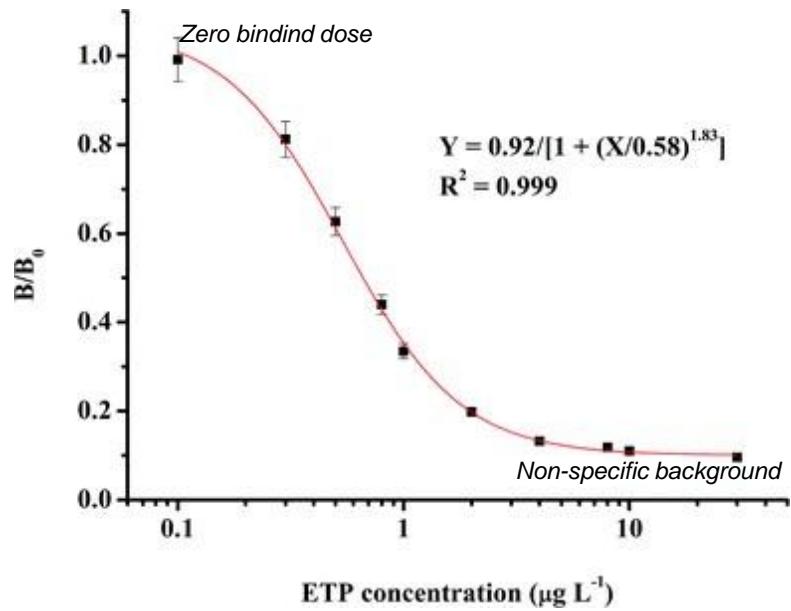
## Direct



## Indirect



Calibration curve for ethopabate (veterinary drug)



Limiting Antibody LOG SCALE!!

**TABLE 17.1**  
**Overview of Immunoassay Techniques**

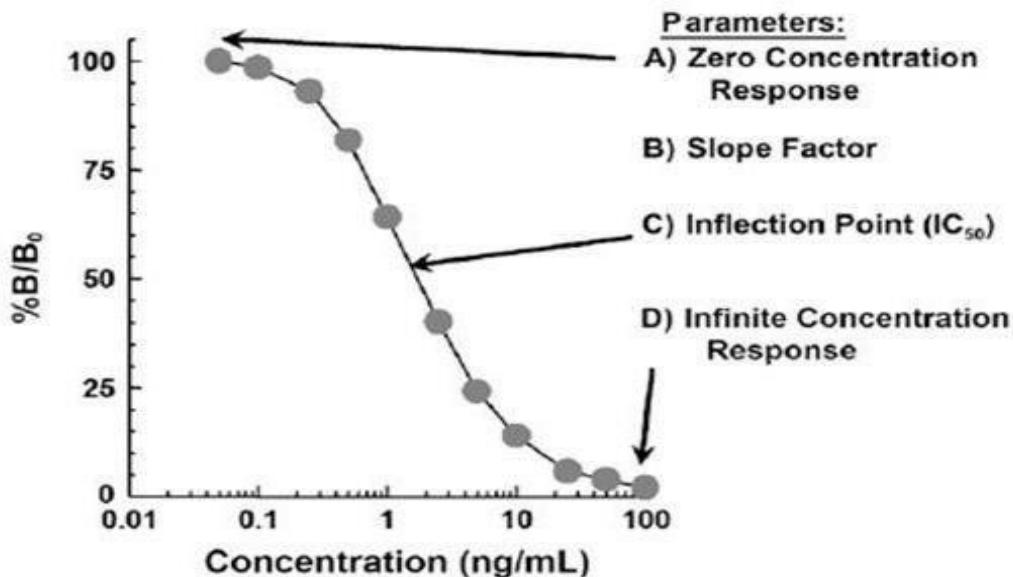
Assay method	Label	Detected	Detector
RIA (radioimmunoassay)	$^{125}\text{I}$ , $^{3}\text{H}$ , $^{14}\text{C}$	Radiation	Scintillation counter
EIA (enzyme immunoassay)	HRP AP $\beta$ -D-galactosidase HRP, AP, galactosidase HRP HRP, AP, GOD, catalase	Color change (absorbance) Fluorescence Luminescence Current Fluorescence	Photometer Fluorimeter Luminometer Amperometric electrode Fluorimeter
FrIA (fluoroimmunoassay)	Fluorescein rhodamines, dansyl chloride, cumarines, phycoerythrin, also liposomes		
TR-FrIA (time-resolved FIA)	Lanthanoid cations: $\text{Eu}^{3+}$ , $\text{Tb}^{3+}$ , $\text{Sm}^{3+}$	Delayed fluorescence	Time-resolved fluorimeter
LIA (luminescence immunoassay)	Acridinium esters Dioxetanes Peroxyoxalates Luminol Luciferase/luciferin Peroxidase Pyrene	Chemi- and bioluminescence Electroluminescence	Luminometer Electrode luminometer
Electrochemical immunoassays	Metallocenes Metals GOD, catalase  Urease Liposomes	Current	DPP (differential pulse polarograph) DPASV (differential pulse anodic stripping voltammetry) Potentiometric electrode
		Ions (potential change)	

*Note:* AP: alkaline phosphatase, GOD: glucose oxidase, HRP: horseradish peroxidase.

$$y = \frac{a-d}{(1+(x/c)^b)} + d$$

a = (theoretical) response at low concentration/dilution  
b = absolute value of the slope at the inflection point  
c = value of x at inflection point  
d = (theoretical) response at high concentration/dilution  
x = concentration or dilution  
y = response (OD)

## Typical calibration curve for immunoassay

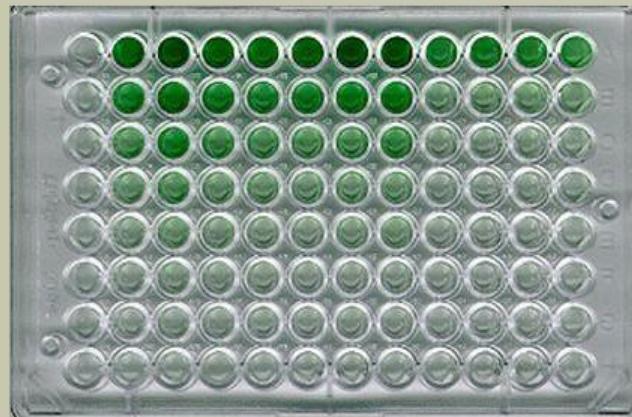


**Figure 1.** Typical 4-parameter logistic graph for a competitive-format immunoassay.

# ELISA



- **Enzyme-linked immunosorbent assay (ELISA)** is a test that uses antibodies and color change to identify a substance.





# DIAGNOSTIC AUTOMATION, INC.

IMMUNODIAGNOSTICS



## AccuDiag™ ELISA Aflatoxin B1 (In Food)

REF 5120-8

96 Tests

LOT AFB-132

8°C

2012-02

2°C



DIAGNOSTIC AUTOMATION, INC.

23961 Craftsman Road, Ste E/F, Calabasas, CA 91302 U.S.A.

Tel: 1 (818) 591-3030 Fax: 1 (818) 591-8383

Website: <http://www.rapidtest.com>

Email: [onestep@rapidtest.com](mailto:onestep@rapidtest.com)

Made in USA

961 CRAFTSMAN ROAD, SUITE E/F • CALABASAS, CA 91302, U.S.A.

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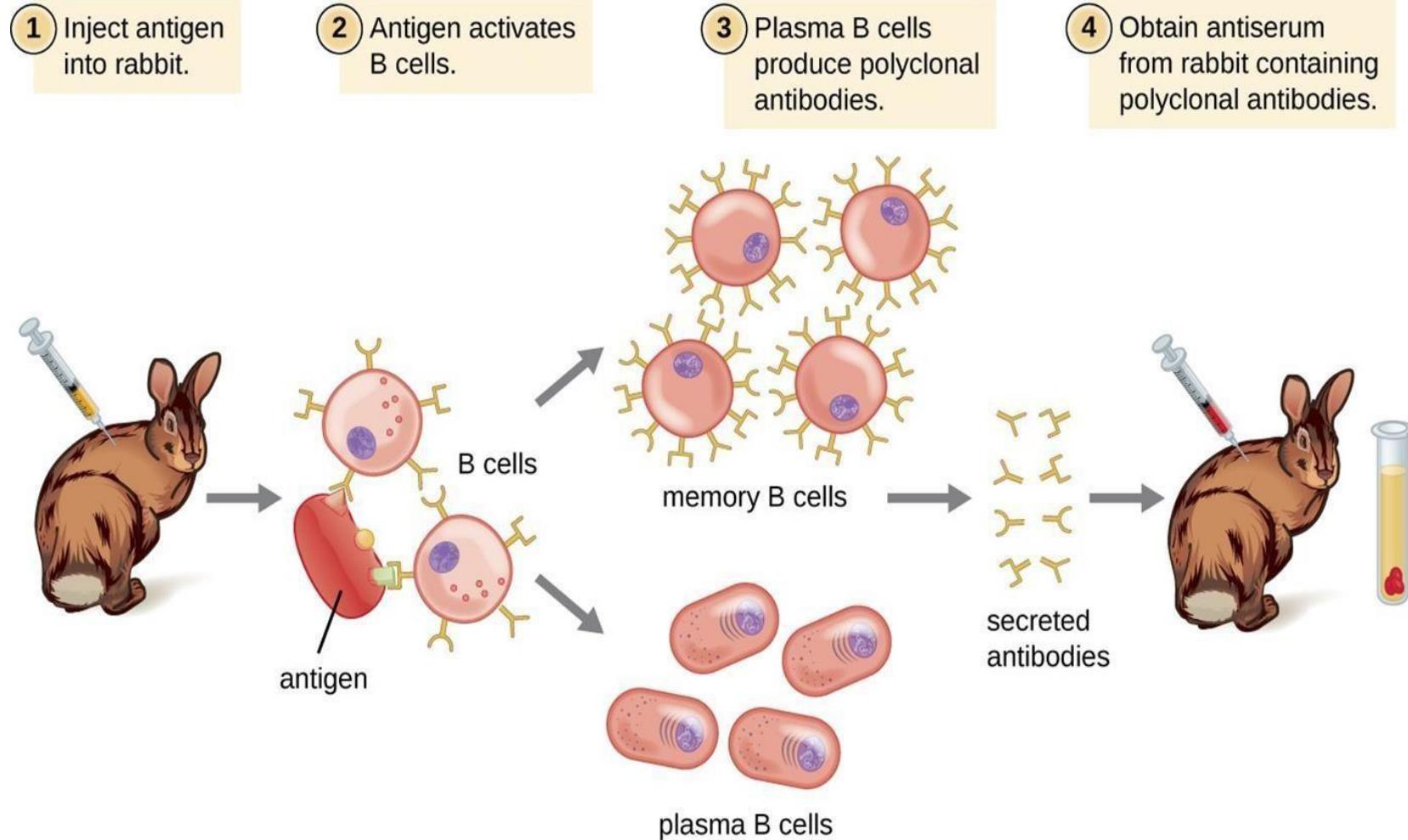




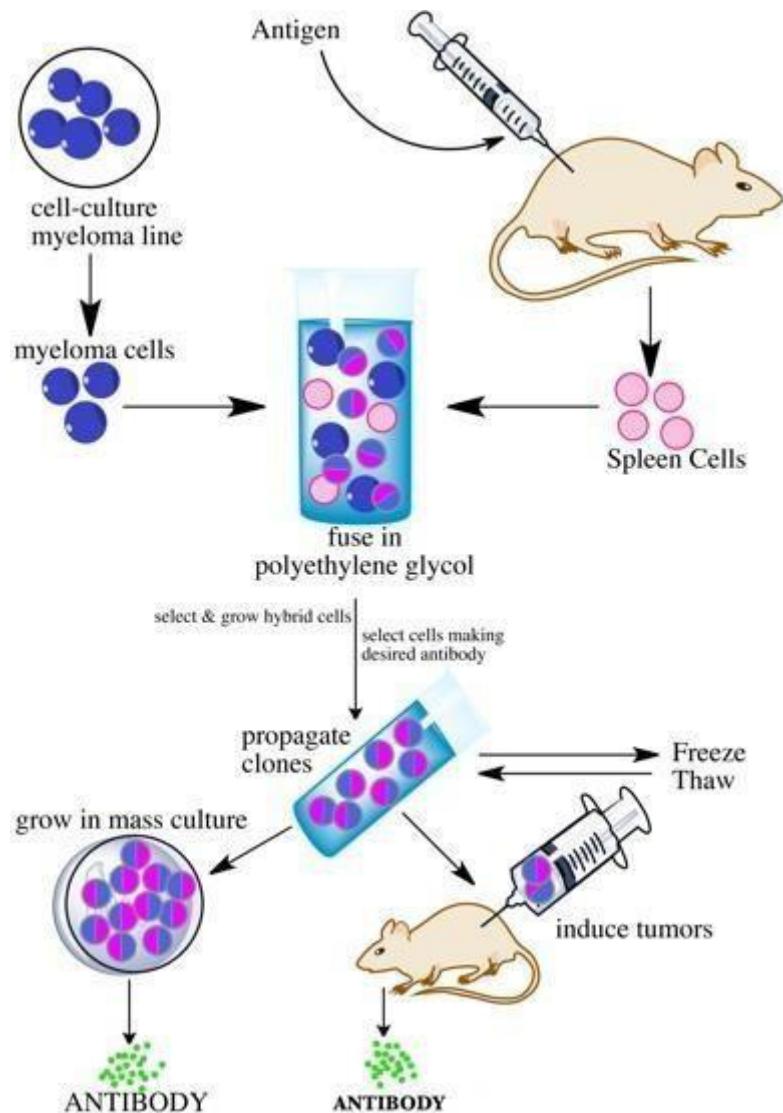
## À la carte ELISA Systems

Almond	Beta-Lactoglobulin	Buckwheat	Casein	Crustacean
Egg	Gluten	Hazelnut	Lupin	Mustard
Peanut	Sesame	Soy		

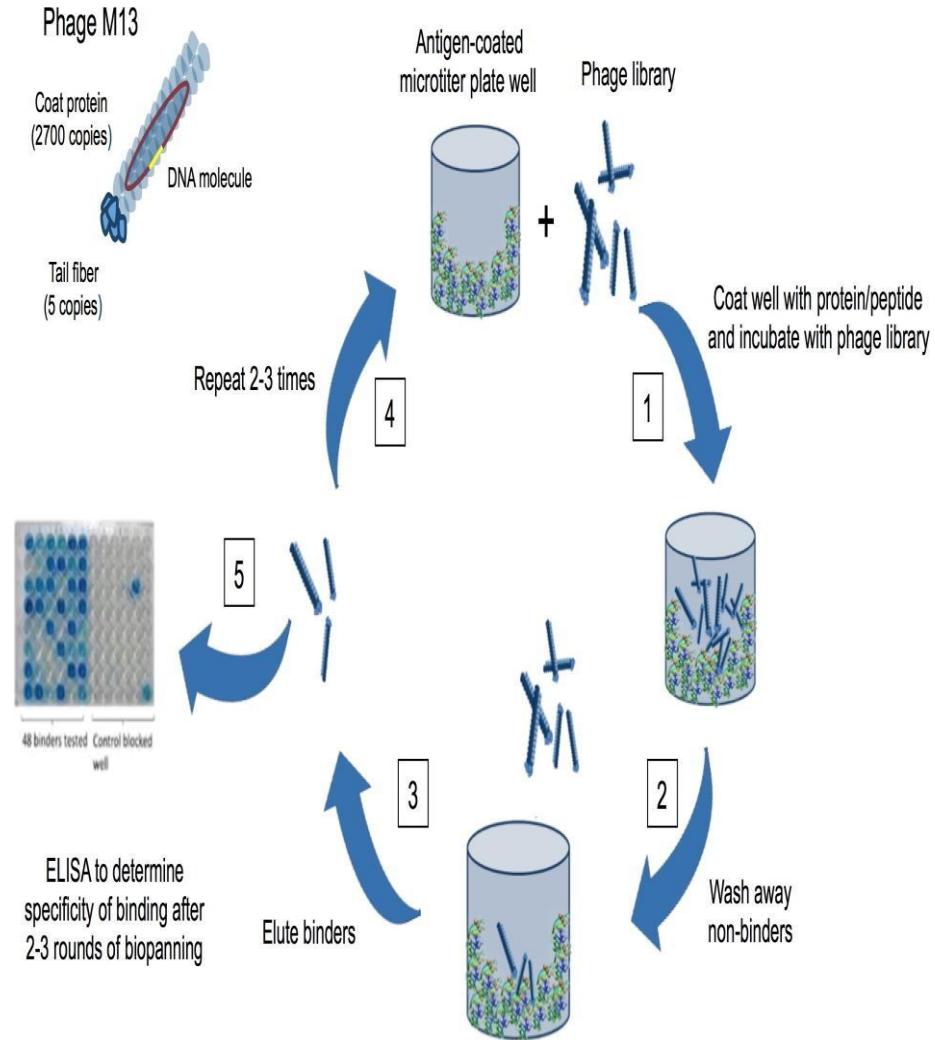
# Polyclonal antibodies production



# Monoclonal antibodies

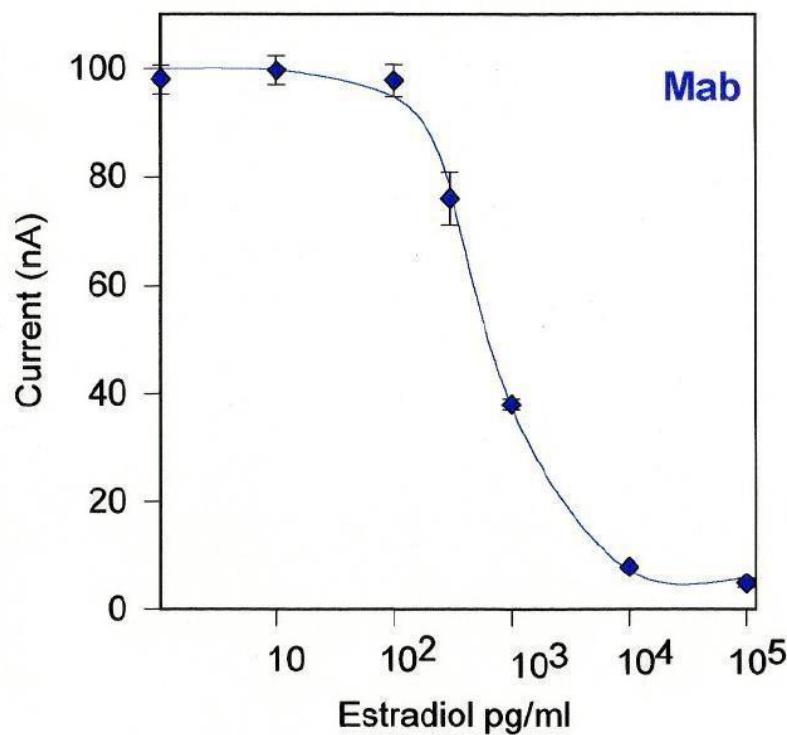
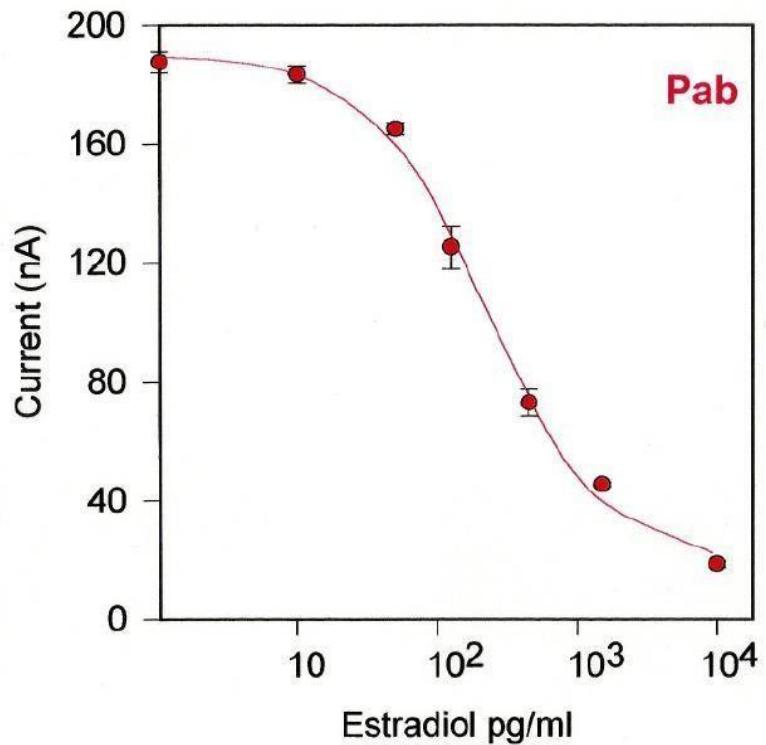


# Recombinant antibodies



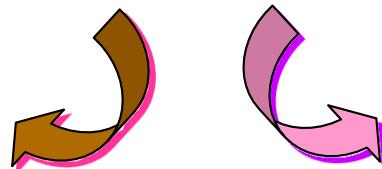
# Enzime Linked Immuno-Sorbent Assay

## ELISA elettrochimico



## ❖ Electrochemical detection:

Chronoamperometry



Differential pulse  
voltammetry (DPV)

## ❖ Enzymes and substrates:

Alkaline  
phosphatase



1-naphthyl-phosphate

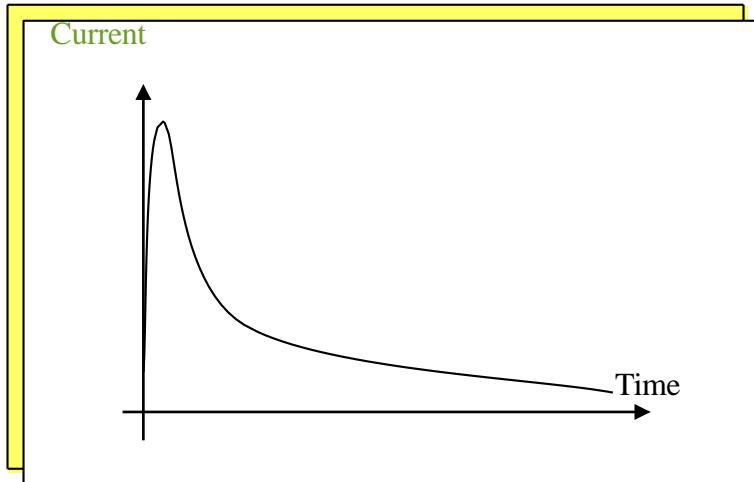
Horseradish peroxidase



- TetramethylBenzidine +  $H_2O_2$
- $[K_4Fe(CN)_6]$  +  $H_2O_2$

electrochemical detection:  
electrochemical detection:

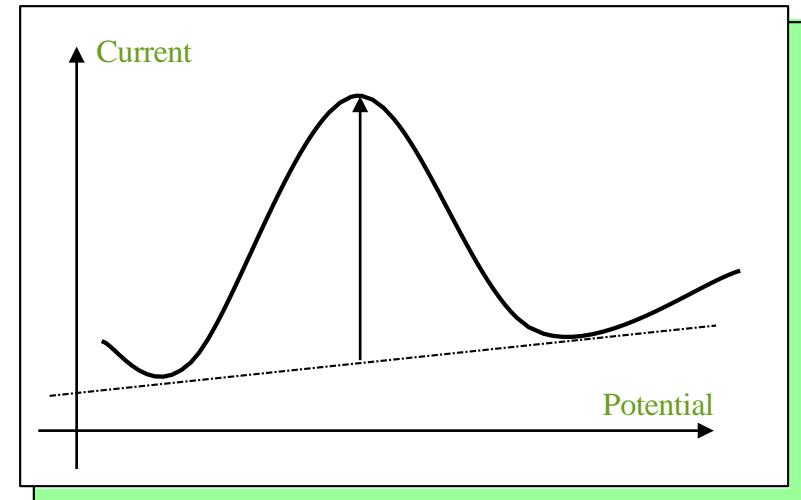
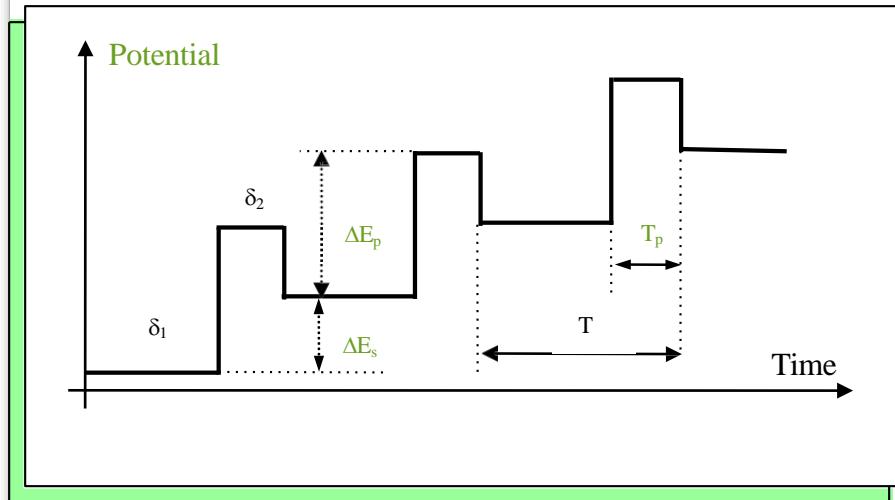
chronoamperometry and differential pulse voltammetry (DPV) :  
chronoamperometry and differential pulse voltammetry (DPV) :

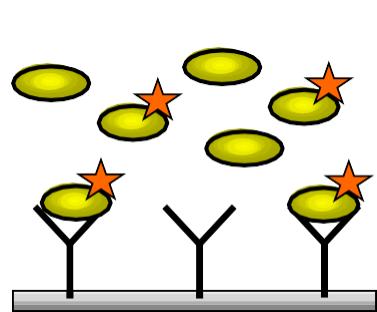


$$\Delta E = \text{const.}$$

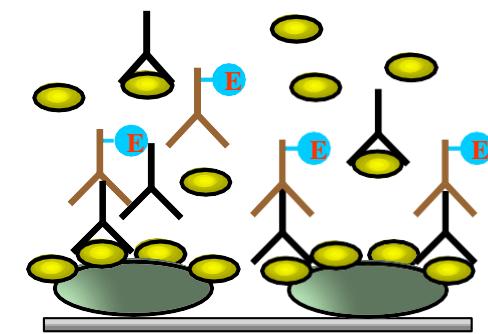
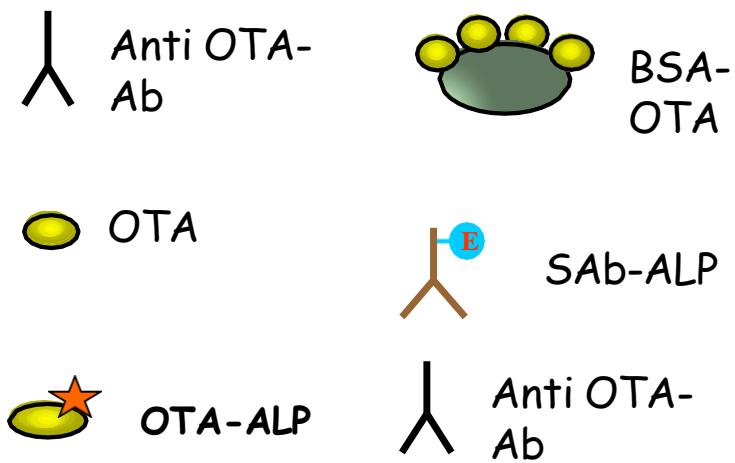
$$\delta_2 = 60 \text{ ms}$$

$$\Delta E_p = 5-100 \text{ mV}$$

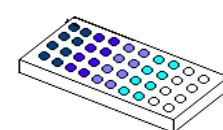
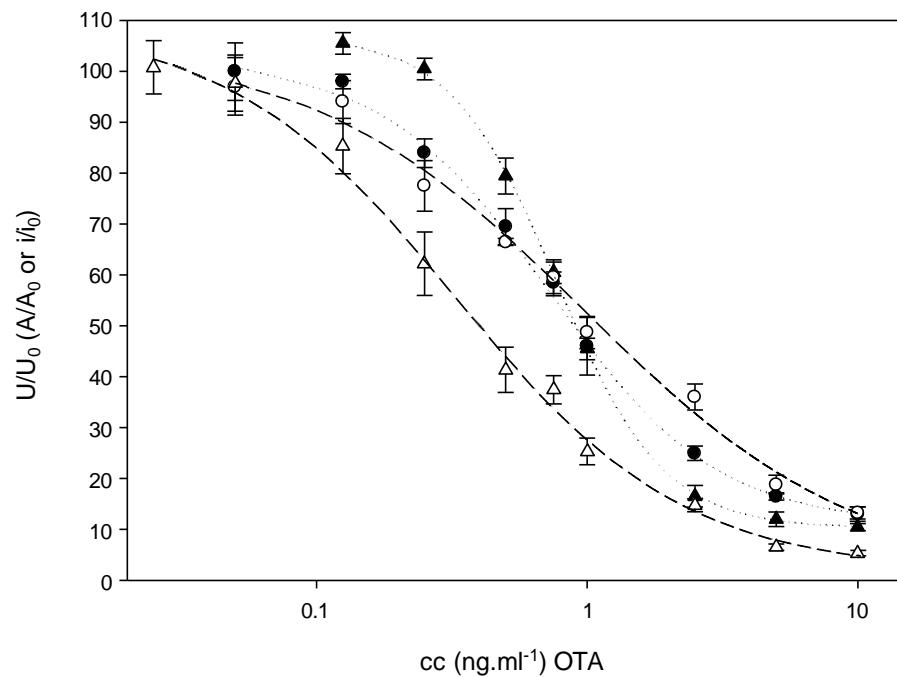
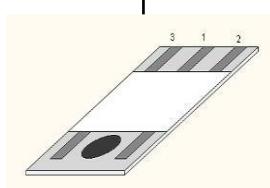




Direct



indirect



$$f(x) = \{ (a - d) / [1 + (x/c)b] \} + d$$

	Competition curve parameters				Linear regression
	$a$ (A or nA)	$b$ (nA.ng.ml <sup>-1</sup> )	$c$ (ng.ml <sup>-1</sup> )	$d$ (A or nA)	
ic spettr	1.220 ± 0.053	1.40 ± 0.50	0.80 ± 0.22	0.129 ± 0.077	$f(x) = 49.3 (\pm 0.8) - 57.5 (\pm 0.1) x$ [ $r = 0.991$ ]
ic amp.	6019 ± 118	0.90 ± 0.22	0.93 ± 0.10	176 ± 30	$f(x) = 52.5 (\pm 0.4) - 43.7 (\pm 0.5) x$ [ $r = 0.994$ ]
dc spettr	1.392 ± 0.061	2.17 ± 0.15	0.80 ± 0.14	0.132 ± 0.071	$f(x) = 47.4 (\pm 0.7) - 86.0 (\pm 0.3) x$ [ $r = 0.993$ ]
dc amp.	707 ± 56	1.10 ± 0.10	0.35 ± 0.04	16 ± 13	$f(x) = 34.9 (\pm 0.6) - 52.2 (\pm 0.9) x$ [ $r = 0.992$ ]

Immunoassay	Working Range (ng/ml)	L.O.D. (Blank – 3 $\sigma$ ) (ng/ml)
ic spettr	0.20 – 2.5	0.150
ic amp.	<b>0.10 – 7.5</b>	<b>0.120</b>
dc spettr	0.10 – 10	0.080
dc amp.	<b>0.05 – 2.5</b>	<b>0.060</b>

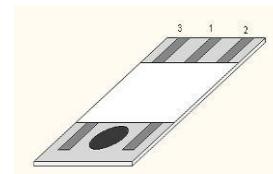
## Immunosensor procedure:

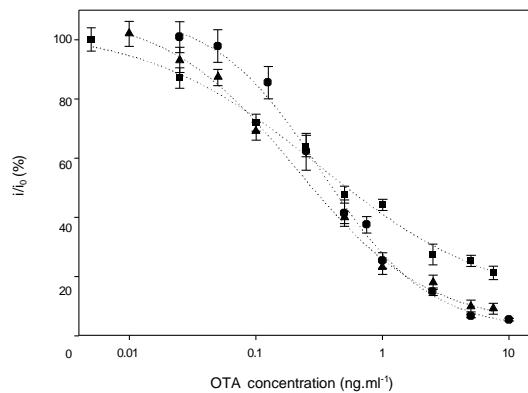
- ✓ Pre-coating: 6 µl of rabbit anti IgG (4°C overnight)
- ✓ Blocking: 6 µl of 1 % PVA (polyvinyl alcohol) (30 min)
- ✓ Coating: 6 µl of anti-OTA Ab (1 h)
- ✓ Competition: 6 µl of OTA-AP + standard/sample (30 min)
- ✓ Detection: 100 µl of 5 mg/ml 1-Naphtylphosphate (2 min) + DPV washings: 150 µl phosphate buffer pH 7.4

## effect of extraction solvent

Activity of an electrode modified with IgG-ALP after 30 min incubation with 1:9 - 9:1 solutions (1:1 in DPBS) acetonitrile:water → 95-108%

Sensitivity of the calibration curve ~ 50%





25 g in 100 mL di  
ACN:H<sub>2</sub>O

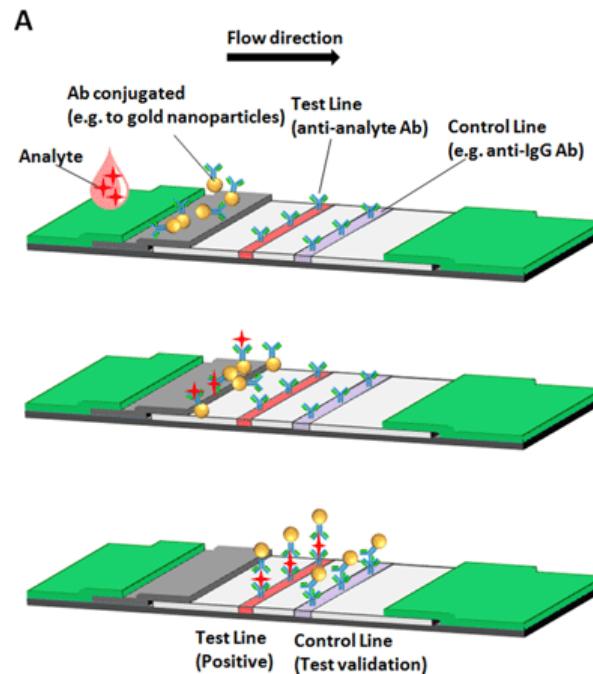
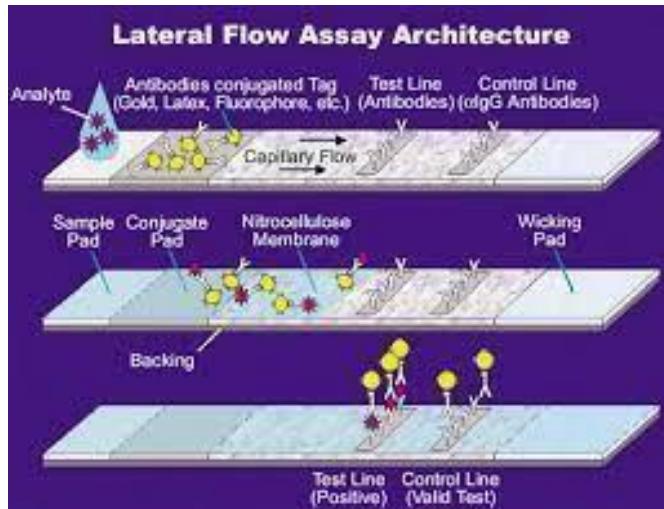
Final dilution 1:8

MRL = 3 ng/g

I<sub>50</sub> = 1.6 ng/g

Parameters		0.1 PBS Buffer ●	ACN:H <sub>2</sub> O (6:4) ▲	Wheat Extract (blank) ■
<i>a</i>	( nA)	707 ( $\pm$ 56)	260 ( $\pm$ 12)	408 ( $\pm$ 72)
<i>b</i>	(nA.ng.ml <sup>-1</sup> )	1.1 ( $\pm$ 0.1)	0.62 ( $\pm$ 0.03)	0.8 ( $\pm$ 0.1)
<i>c</i> (I <sub>50</sub> )	(ng.ml <sup>-1</sup> )	0.35 ( $\pm$ 0.04)	0.32 ( $\pm$ 0.02)	0.20 ( $\pm$ 0.03)
<i>d</i>	(nA)	16 ( $\pm$ 13)	24 ( $\pm$ 8)	13 ( $\pm$ 15)
w.r.	(ng.ml <sup>-1</sup> )	0.05 – 2.5	0.02 – 5.0	0.05 – 2.5
L.O.D.	(ng.ml <sup>-1</sup> )	0.06	0.015	0.05
Lin.	Reg.	30.9 ( $\pm$ 0.6) – 52.2 ( $\pm$ 0.9) <i>x</i>	42.3 ( $\pm$ 0.3) – 25.4 ( $\pm$ 0.6) <i>x</i>	23.5 ( $\pm$ 0.1) – 41.1 ( $\pm$ 0.5) <i>x</i>

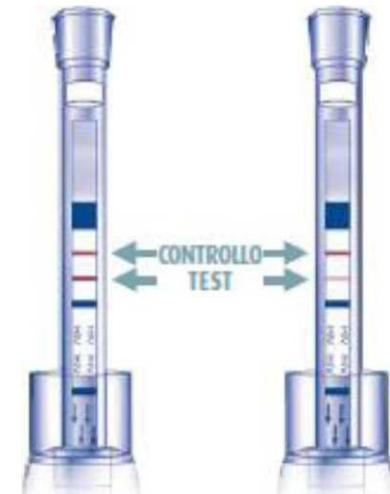
# Lateral flow immunoassays



## Human corionic gonadotropin in urine assay



First Commercialised LFIA by Unipath, 1988

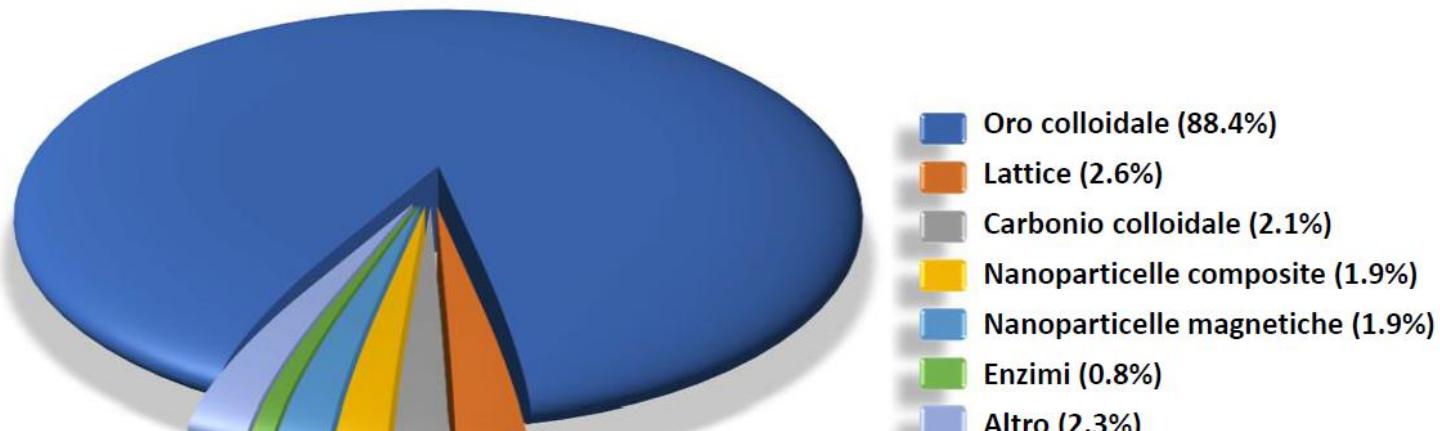


## Drugs of Abuse



## Marcatori colorimetrici in LFIA

Marcatori colorimetrici più utilizzati in LFIA (pubblicazioni scientifiche dal 2010 al 2019 compresi):



\* Di Nardo, F.; Chiarello, M.; Cavalera, S.; Baggiani, C.; Anfossi, L. *Sensors* 2021, 21, 5185.

## LFIA FORMATS

---

NON-COMPETITIVE

COMPETITIVE

**Negative**



**Positive**



**Negative**

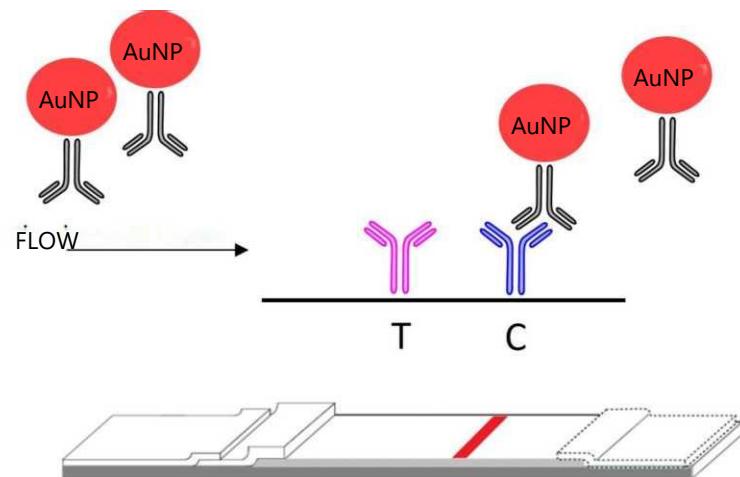


**Positive**

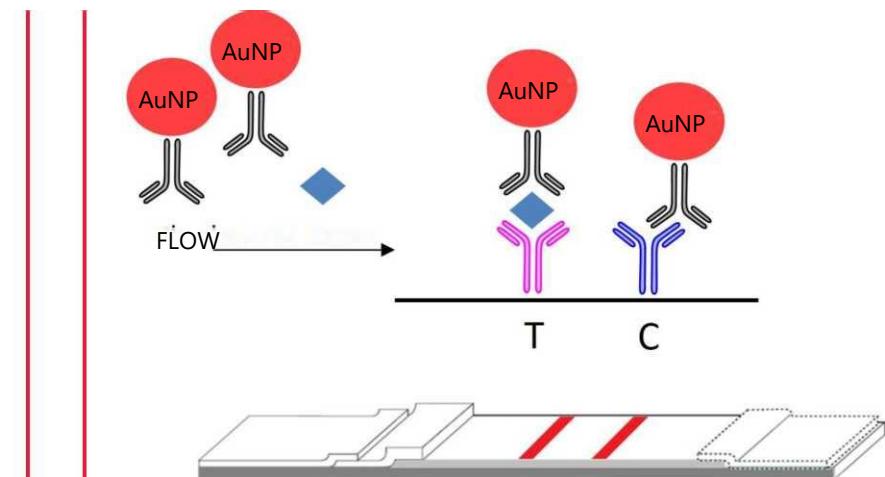


## non competitive (sandwich)

NO TARGET



TARGET PRESENT



Ab di rivelazione  
marcato con AuNPs



Ab di cattura  
anti-analita



Ab anti anticorpo  
di rivelazione



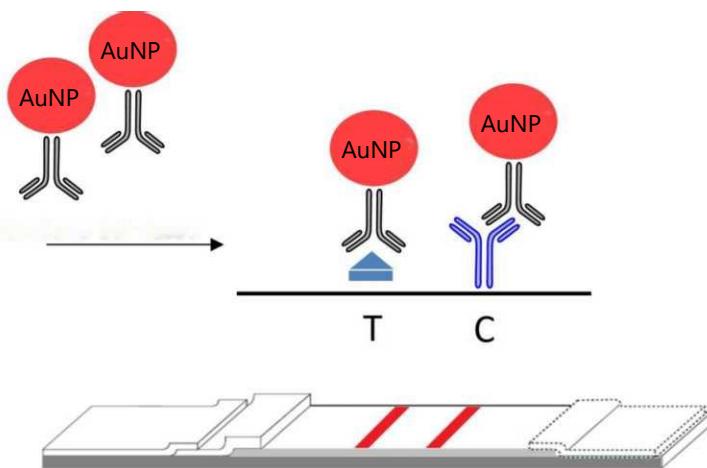
Analita



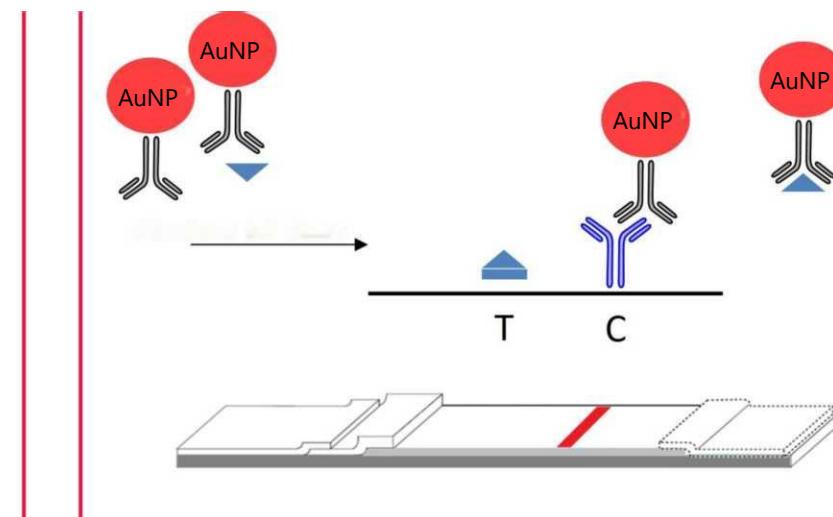
Fabio Di Nardo - Università di Torino

## competitive indirect

### NO TARGET



### TARGET PRESENT



 Ab di rivelazione  
marcato con AuNPs

 Coniugato proteico  
dell'analita

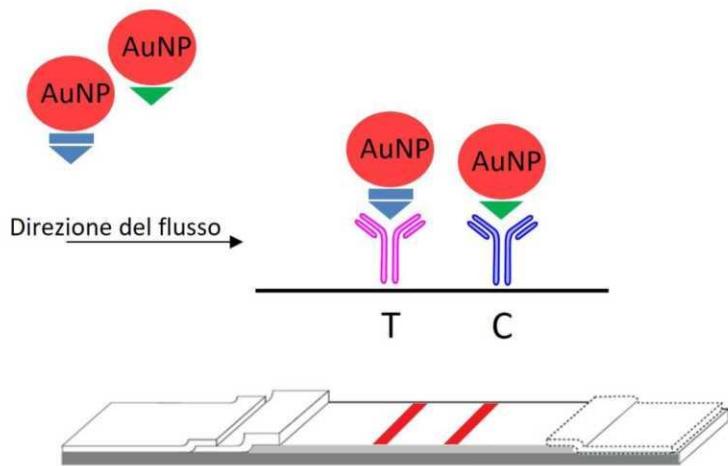
 Ab anti anticorpo  
di rivelazione

 analita

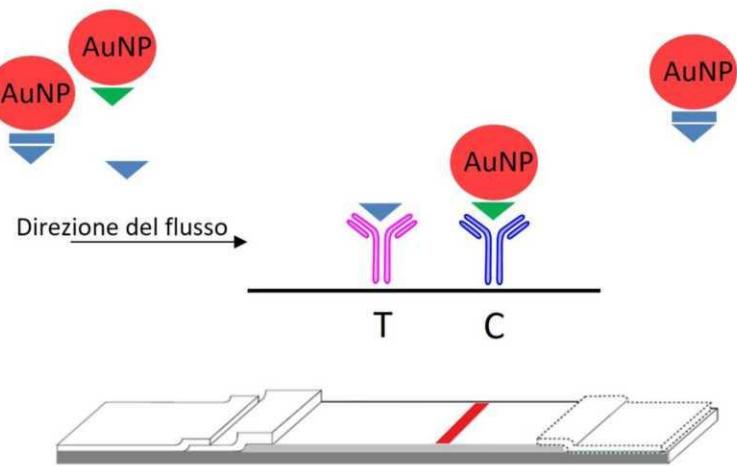


## competitivo direct

### NO TARGET



### ! TARGET PRESENT



Omologo dell'analita  
marcato con AuNPs



Antigene non-target  
marcato con ANPs



Ab di cattura  
anti-analita



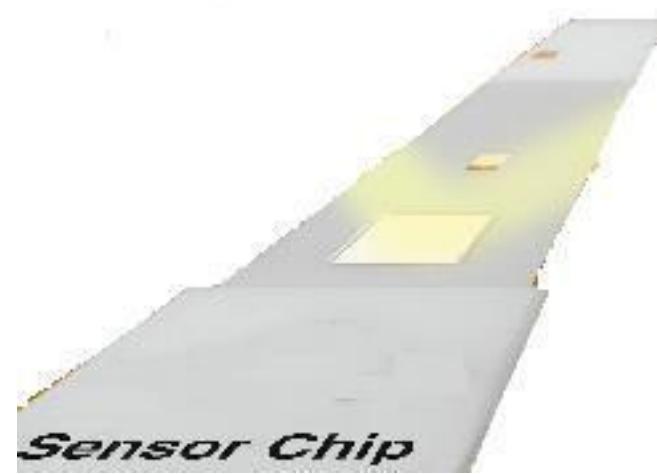
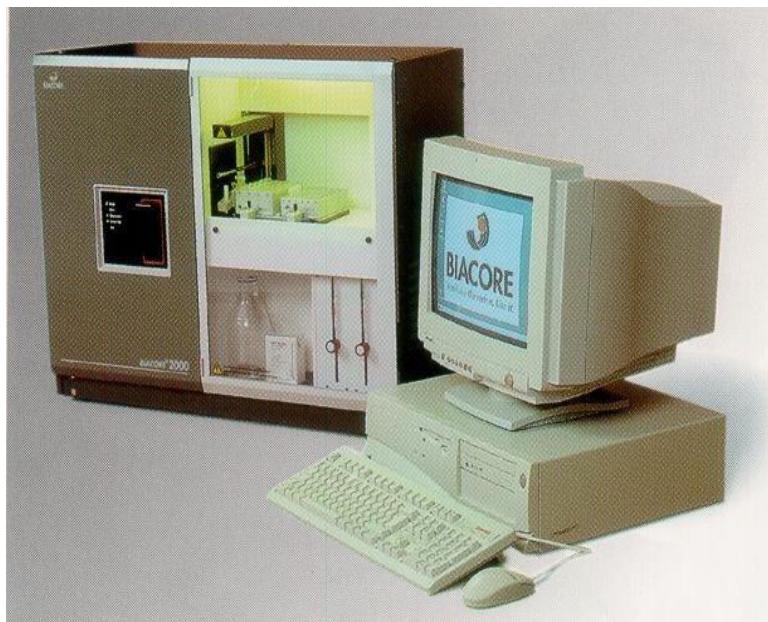
Ab di cattura anti  
antigene non target

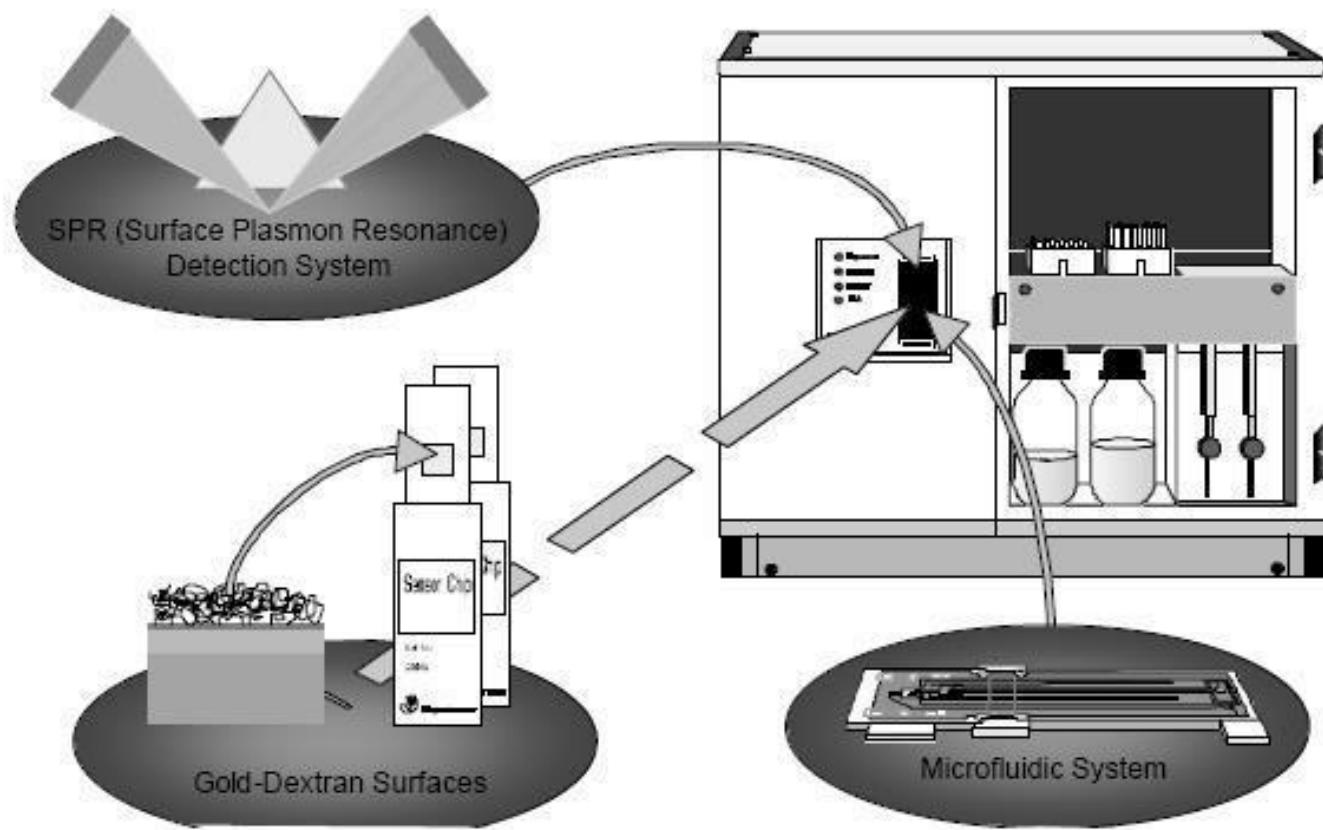


Analita

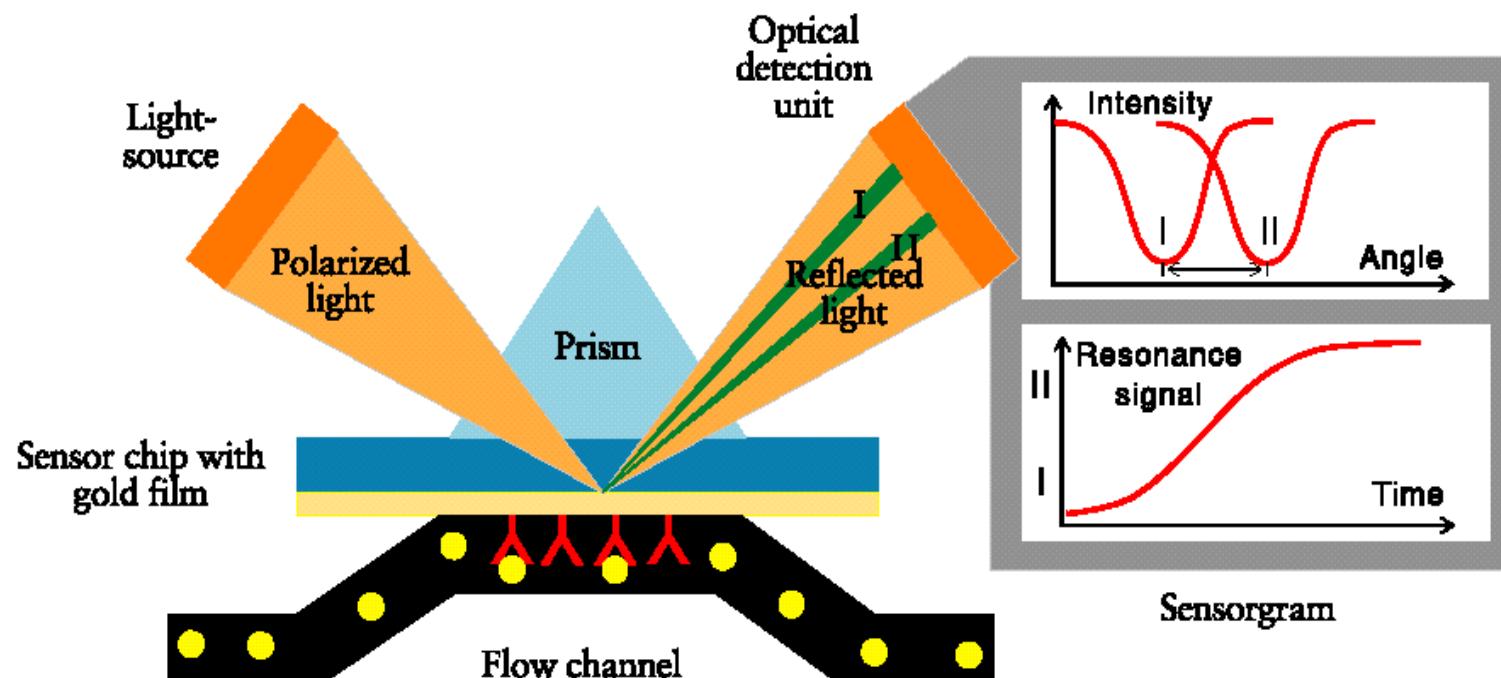


# Biacore

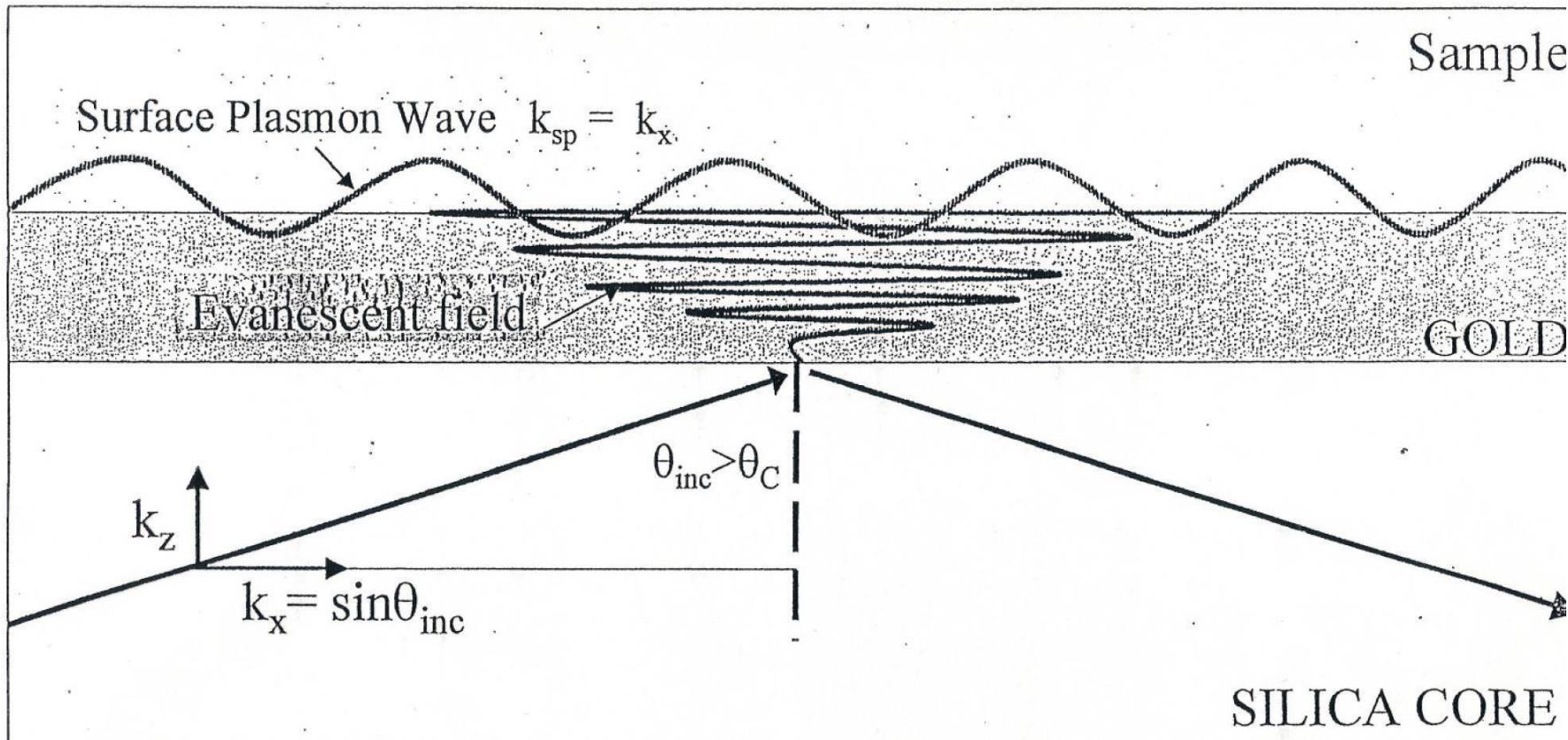




# SPR Biosensors



# Surface Plasmon Resonance



$\theta_{inc}$  - angle of incident light

$\lambda_{inc}$  - wavelength of incident light

$n_{glass}$  - Refractive Index of glass

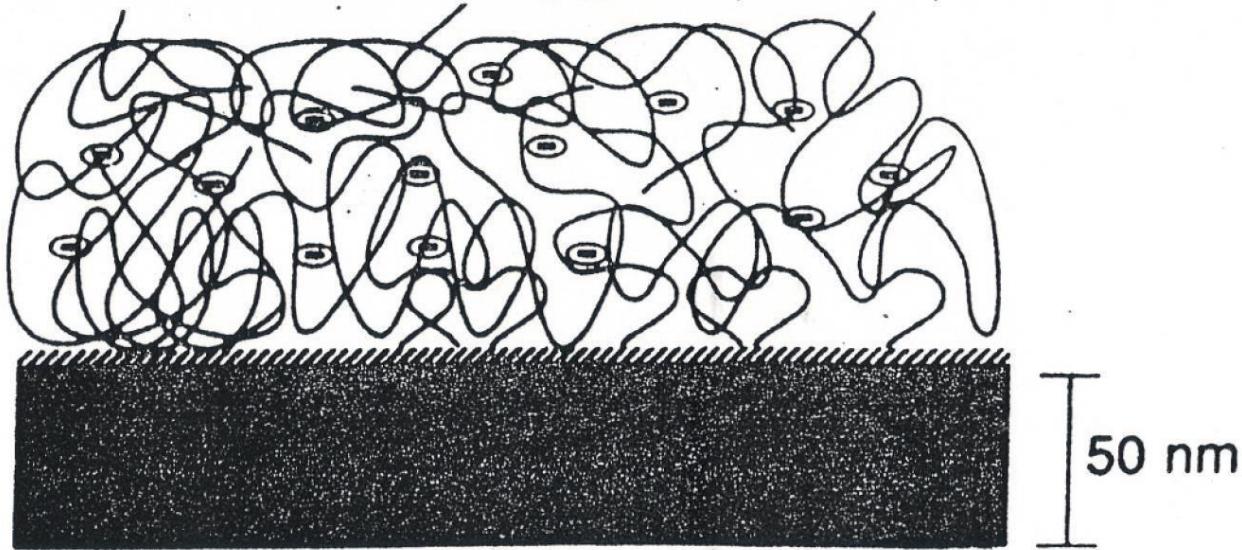
$n_{metal}$  - Refractive Index of metal

$n_{sample}$  - Refractive index of sample

$t_{metal}$  - Thickness of metal

# BIACORE approach

Carboxylated  
dextran  
Linker layer  
Gold film



**dextran hydrogel**

**open structure (good accessibility)**

**no denaturation**

- enhancement of the capacity of the interaction layer
- stagnant layer / mass transport flow needed ( $\mu\text{l}/\text{min}$ )
- negative charge
- regenerable (up to 100 x)

## SPR principles

Surface plasmon resonance (SPR) arises when light is reflected under certain conditions from a conducting film at the interface between two media of different refractive index. The media are the sample and the glass of the sensor chip, and the conducting film is a thin layer of gold on the chip surface. SPR causes a reduction in the intensity of reflected light at a specific angle of reflection. This angle varies with the refractive index close to the surface on the side opposite from the reflected light.

When molecules in the sample bind to the sensor surface, the concentration and therefore the refractive index at the surface changes and an SPR response is detected. Plotting the response against time during the course of an interaction provides a quantitative measure of the progress of the interaction. This plot is called a sensogram.

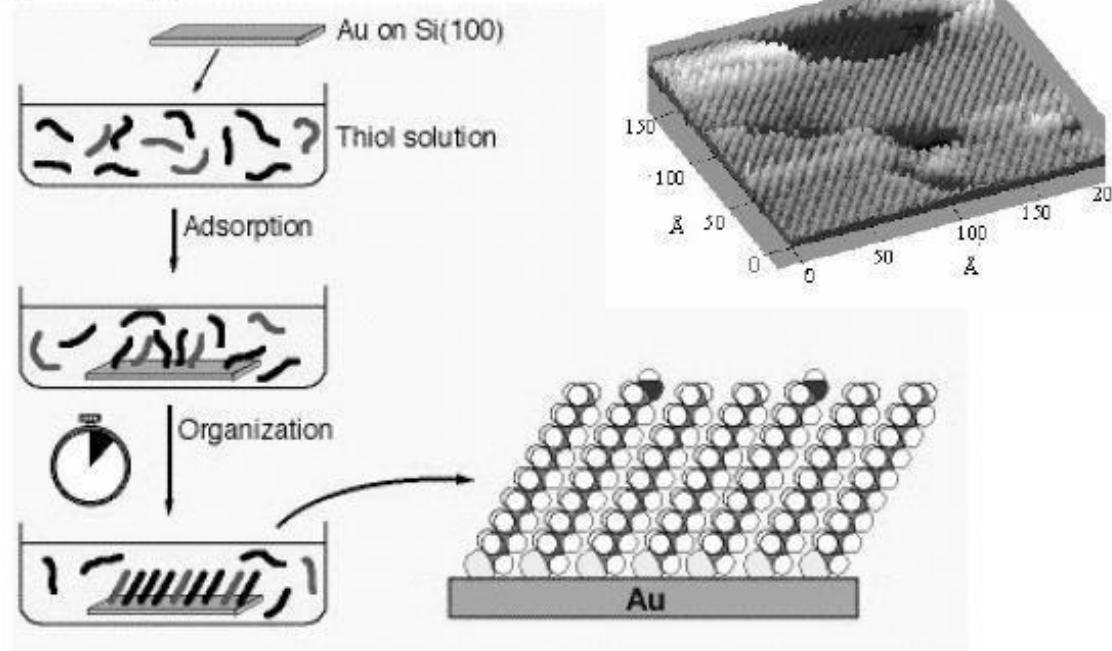
What Biacore actually measures is the angle of minimum reflected light intensity. The light is not absorbed by the sample: instead the light energy is dissipated through SPR in the gold film. Thus the light used to detect interaction processes never enters the sample.

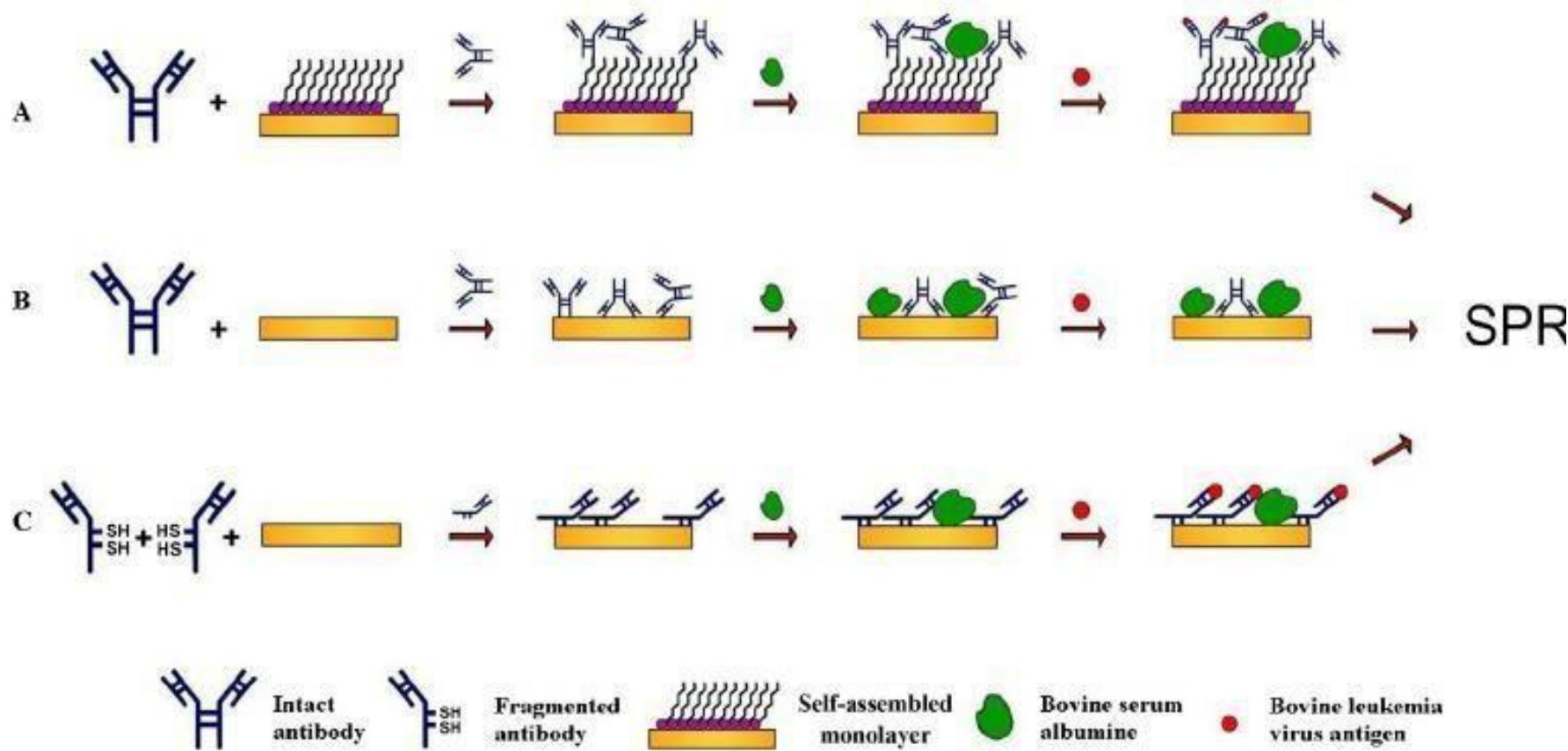
SPR response values are expressed in resonance units (RU). One RU represents a change of  $0.0001^\circ$  in the angle of the intensity minimum. For most proteins, this is roughly equivalent to a change in concentration of about  $1 \text{ pg/mm}^2$  on the sensor surface. The exact conversion factor between RU and surface concentration depends on properties of the sensor surface and the nature of the molecule responsible for the concentration change.

# Immobilisation of organic molecules on gold

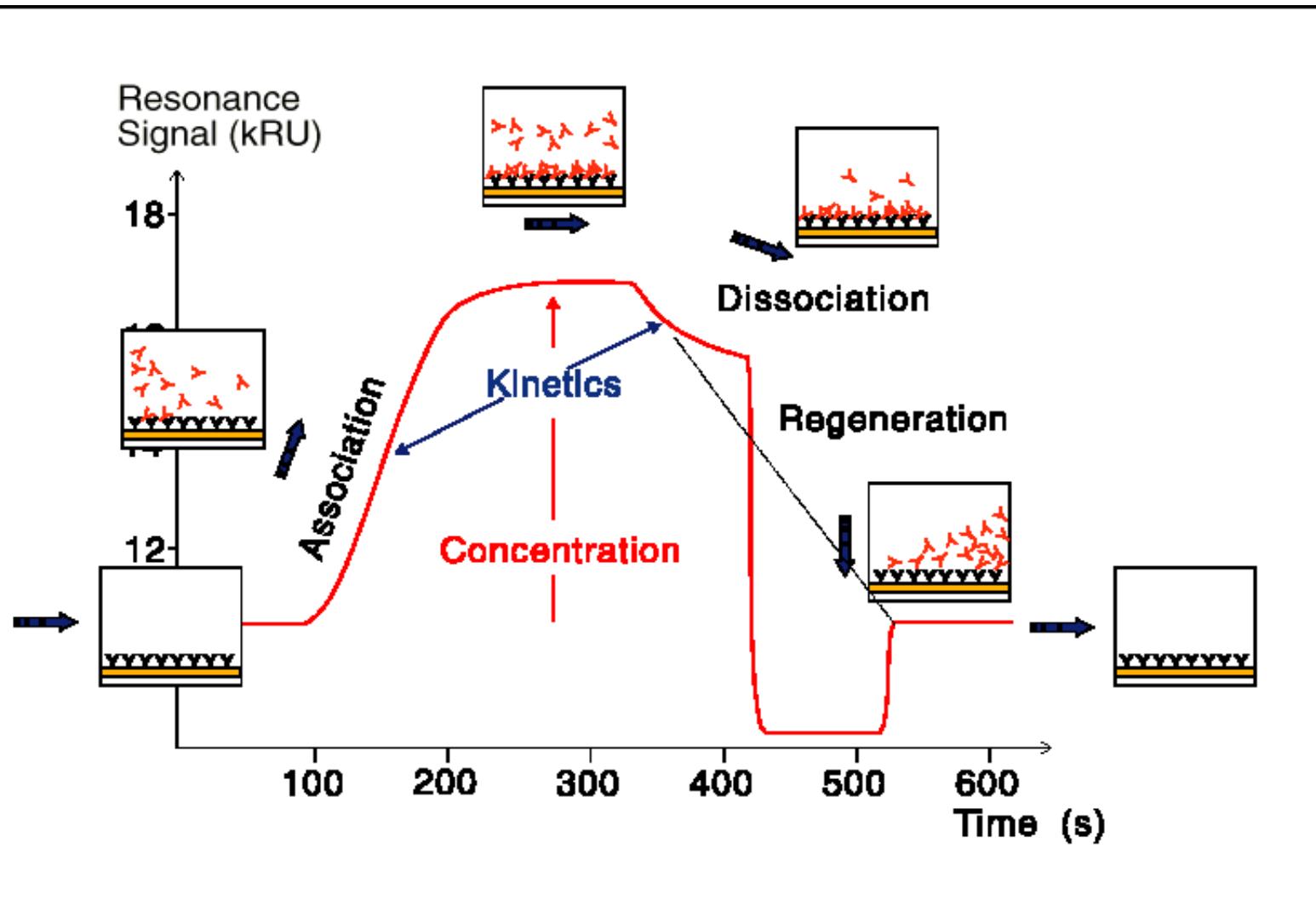
## Self Assembled Monolayers (SAM)

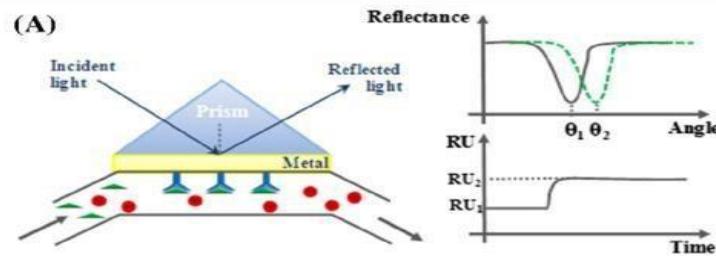
**Disulphides (R-S-S-R)**  
**Sulphides (R-S-R)**  
**Thiols (R-SH)**





# Sensorgram





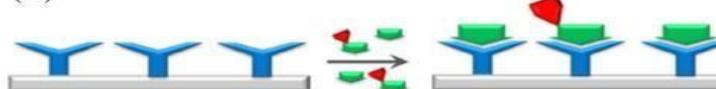
**(B) DIRECT**



**(C) SANDWICH**



**(D) COMPETITIVE**

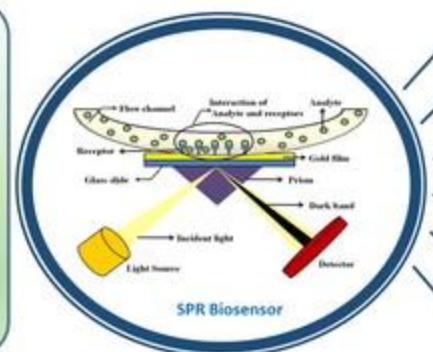


**(E) INHIBITION**



### Why SPR biosensor

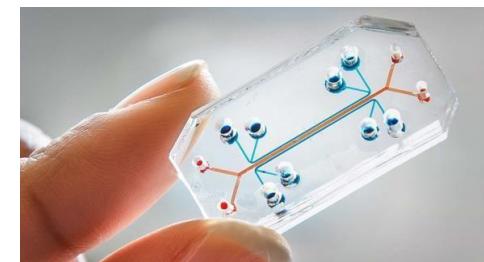
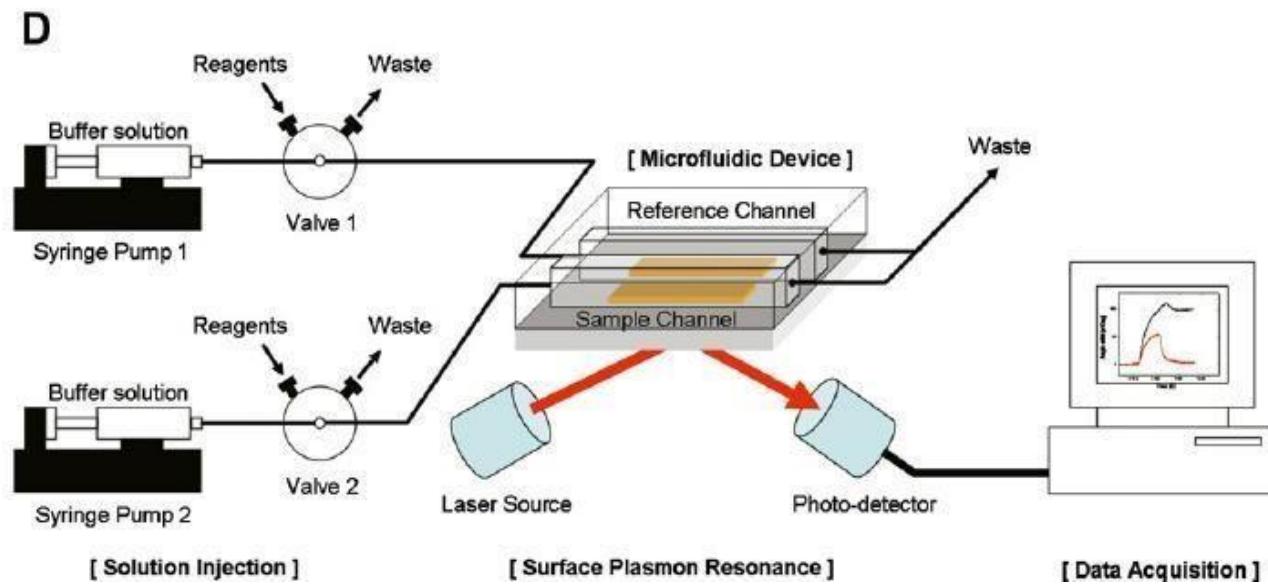
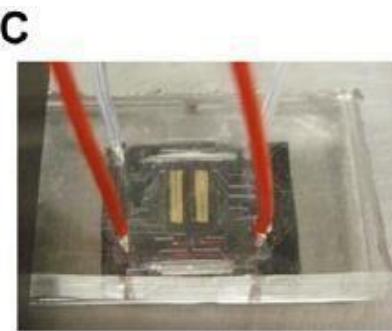
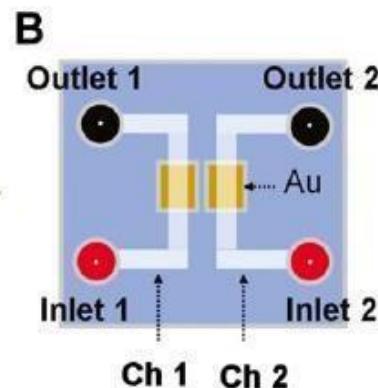
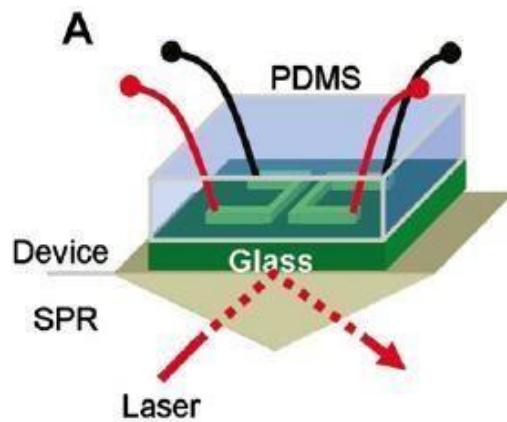
- Optical Biosensor
- High specificity
- Rapid detection
- Real time and label free assay
- Compact design
- Light weight
- Can be coupled with modern technology.



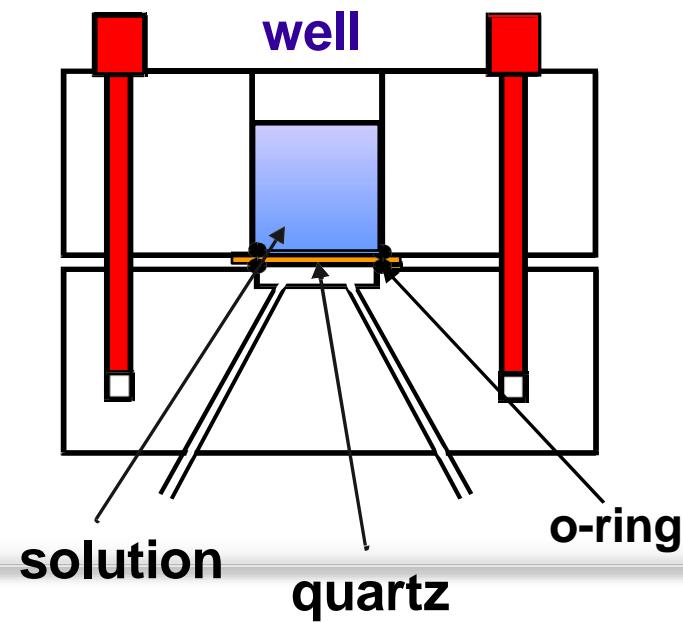
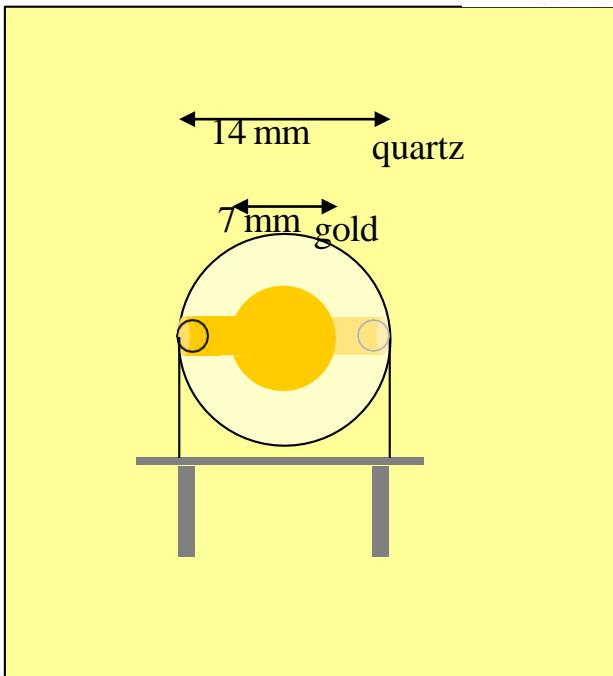
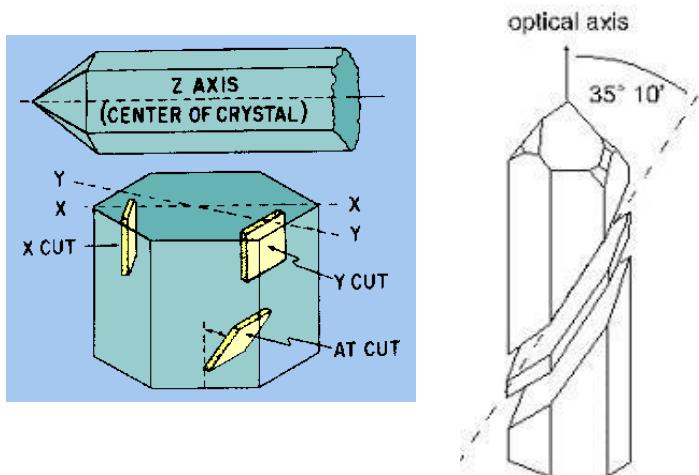
### Food Applications

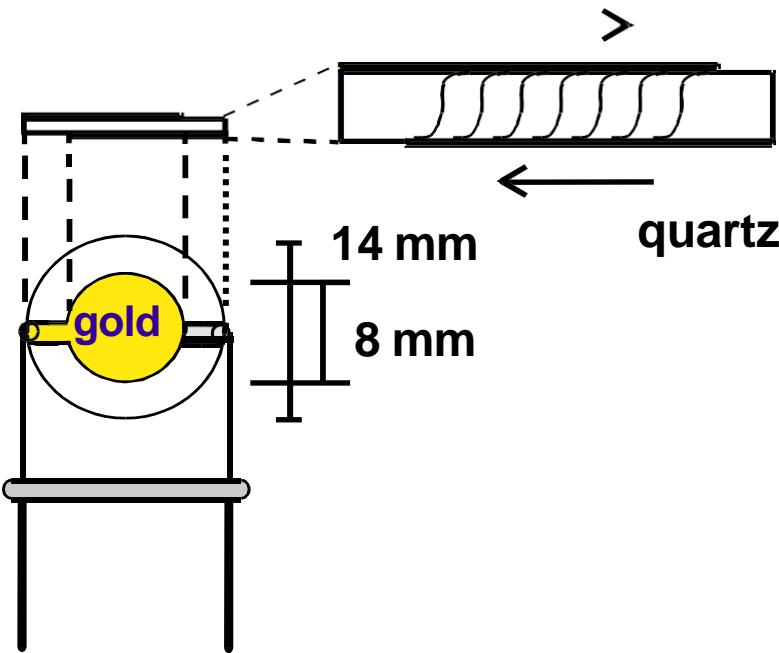
- Antibiotics detection
- Adulteration detection
- Biomolecules detection
- Microbes and Toxins detection
- Pesticide detection
- Genetically modified food detection

# Spr sensors and microfluidics



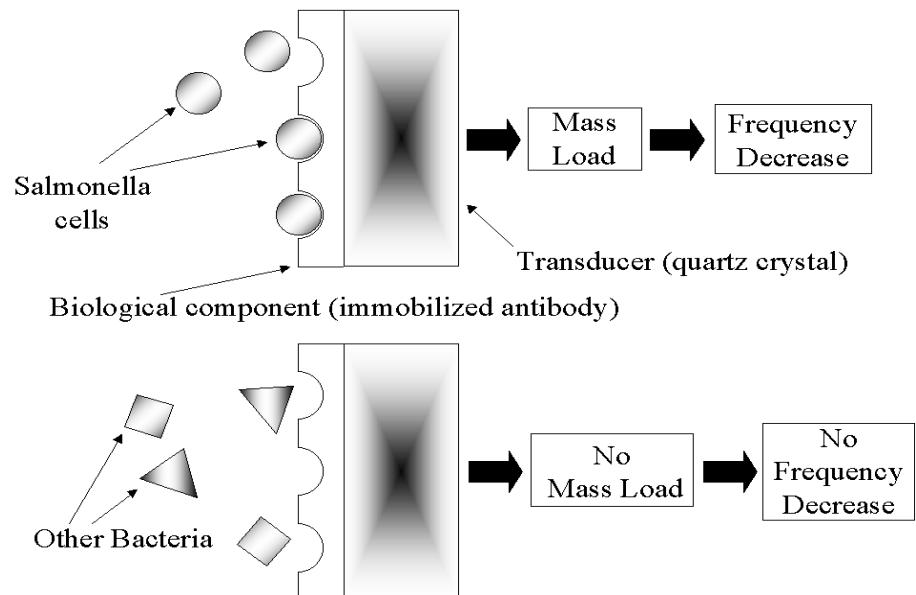
# Piezoelectric Biosensors





The standard QCM measures the mass of a material deposited on a quartz crystal surface as a linear function of a change in the oscillating crystal resonant frequency.

The mass-loading frequency effects of the transducer are based upon Sauerbrey's equation



$\Delta F$  (Hz) = frequency shift of the coated crystal

$F$  (Hz) = resonance frequency of the crystal

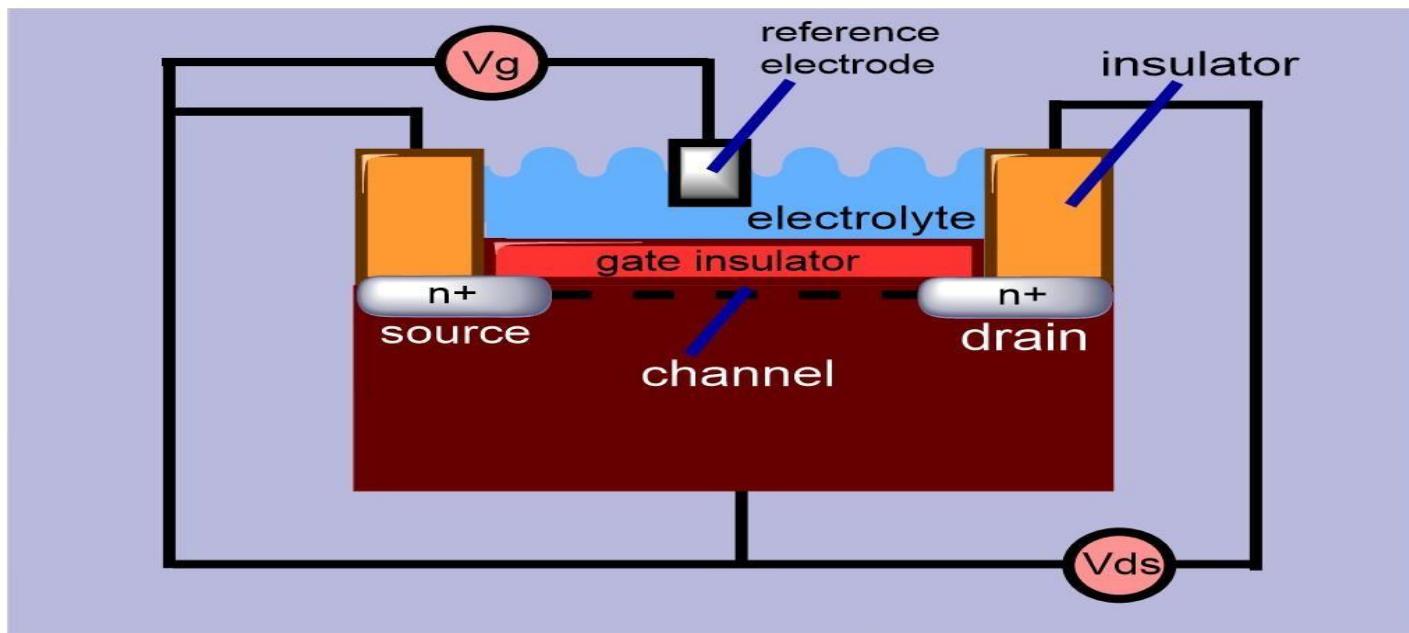
$\Delta M$  (g) = increase in mass loading

$A$  ( $\text{cm}^2$ ) = area of the coated crystal

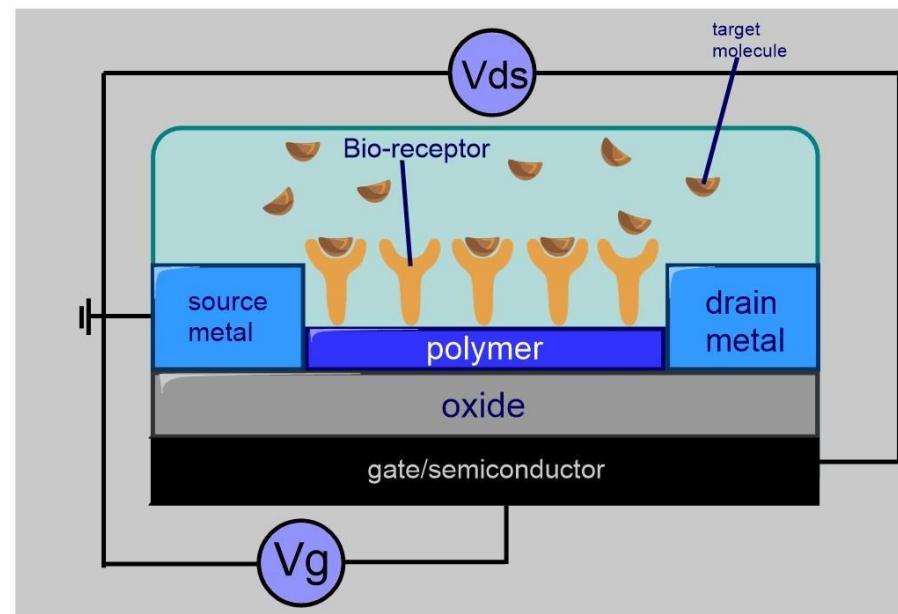
$$\Delta F = (-2.3 \times 10^{-6}) F^2 \Delta M / A$$

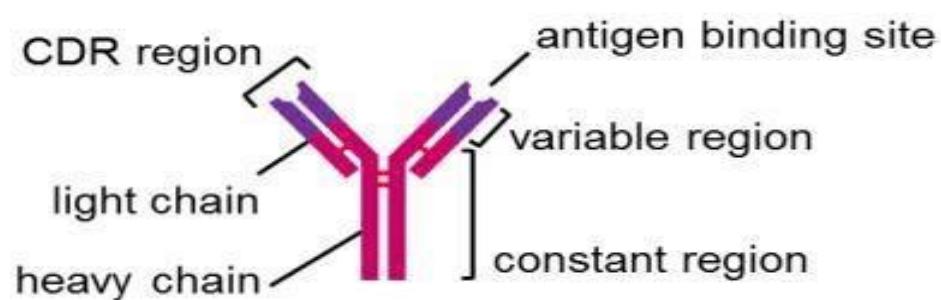
# BIOFETs

An **ion-sensitive field-effect transistor (ISFET)** is a field-effect transistor used for measuring ion concentrations in solution; when the ion concentration (such as  $\text{H}^+$ , see pH scale) changes, the current through the transistor will change accordingly. Here, the solution is used as the gate electrode. A voltage between substrate and oxide surfaces arises due to an ion sheath. It is a special type of MOSFET (metal-oxide-semiconductor field-effect transistor),<sup>[1]</sup> and shares the same basic structure, but with the metal gate replaced by an ion-sensitive membrane, electrolyte solution and reference electrode.<sup>[2]</sup> Invented in 1970, the ISFET was the first biosensor FET (BioFET) source wikipedia

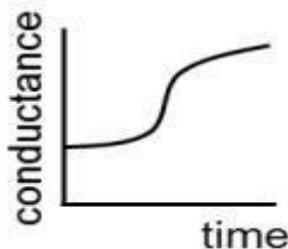
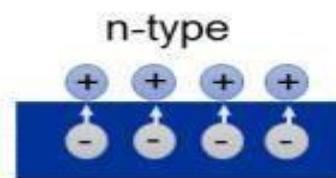
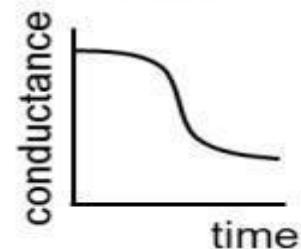
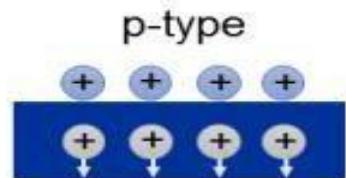
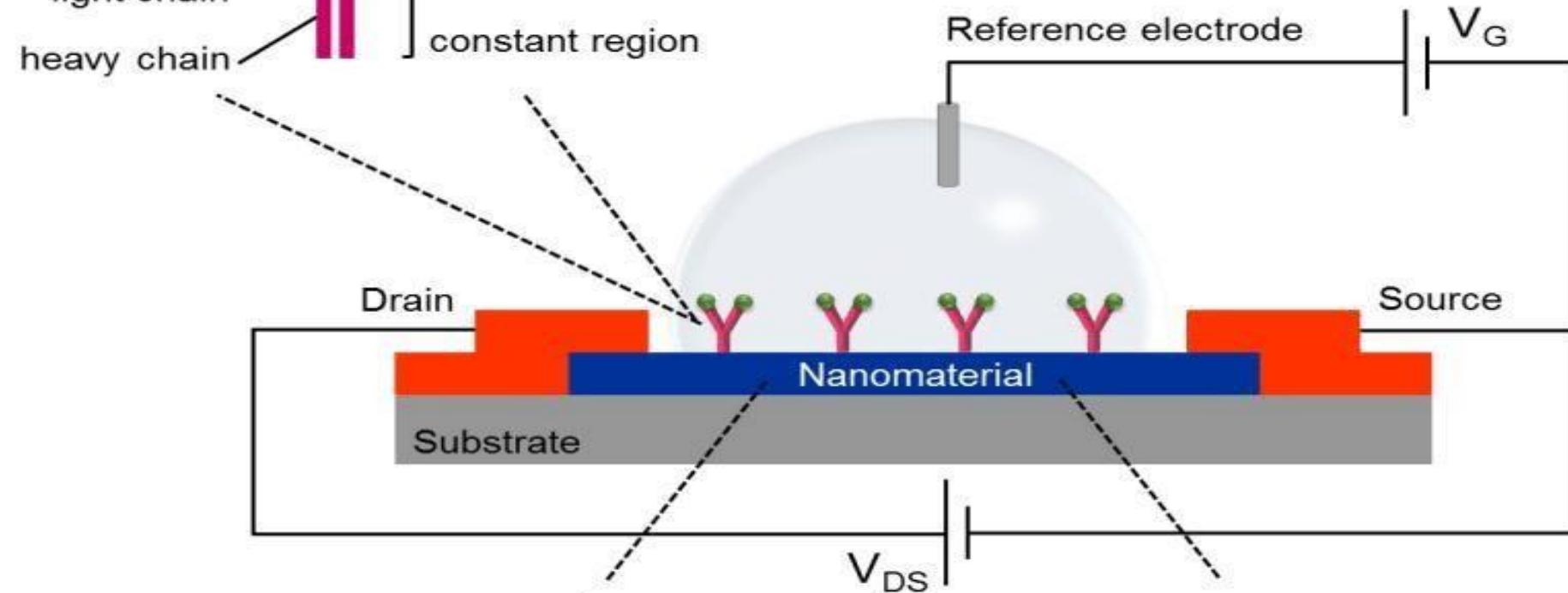


Bio-FETs couple a transistor device with a bio-sensitive layer that can specifically detect bio-molecules such as nucleic acids and proteins. A Bio-FET system consists of a semiconducting field-effect transistor that acts as a transducer separated by an insulator layer (e.g.  $\text{SiO}_2$ ) from the biological recognition element (e.g. receptors or probe molecules) which are selective to the target molecule called analyte.<sup>[8]</sup> Once the analyte binds to the recognition element, the charge distribution at the surface changes with a corresponding change in the electrostatic surface potential of the semiconductor. This change in the surface potential of the semiconductor acts like a gate voltage would in a traditional MOSFET, i.e. changing the amount of current that can flow between the source and drain electrodes.<sup>[9]</sup> This change in current (or conductance) can be measured, thus the binding of the analyte can be detected. The precise relationship between the current and analyte concentration depends upon the region of transistor operation (source Wikipedia)

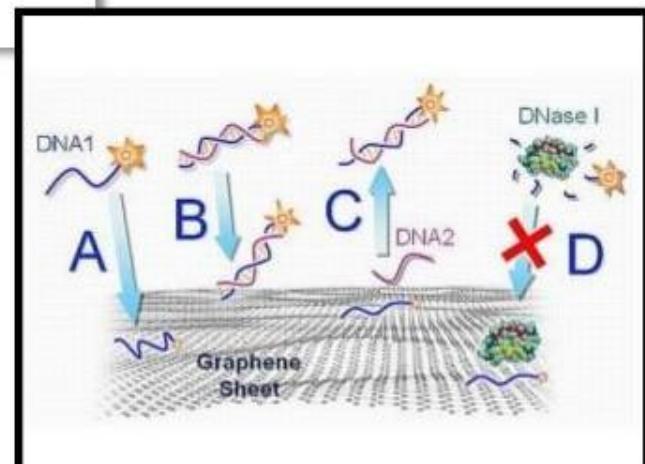
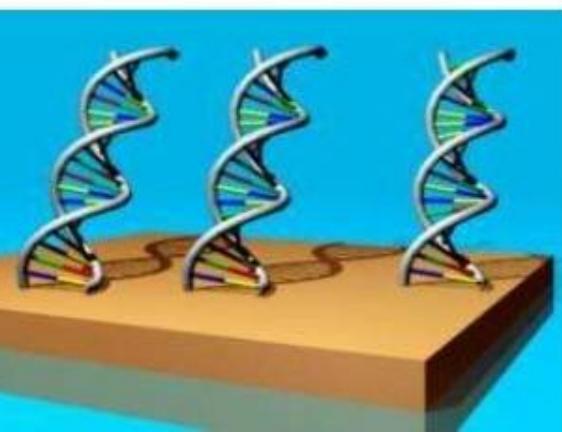
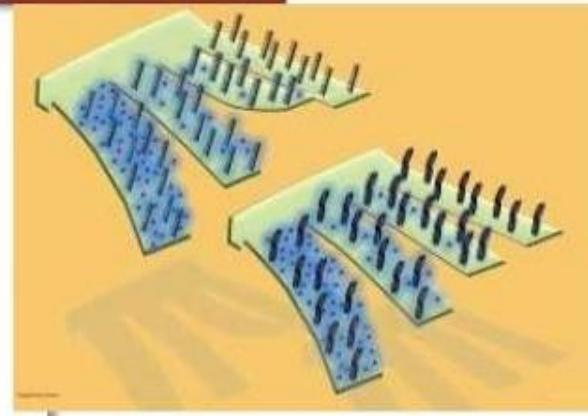
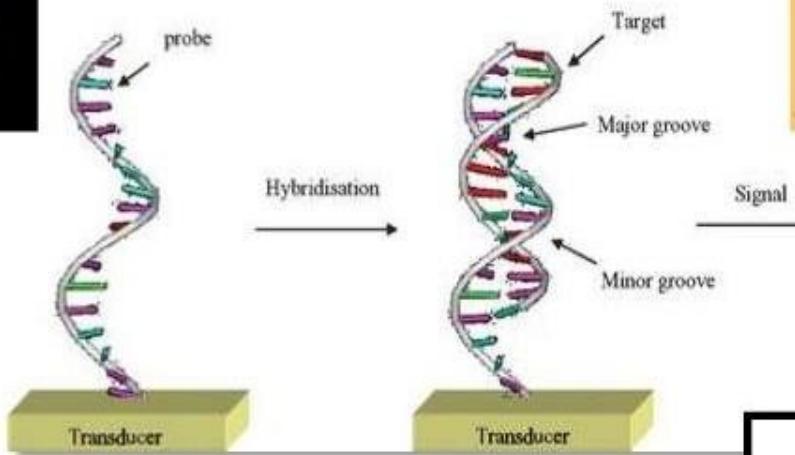
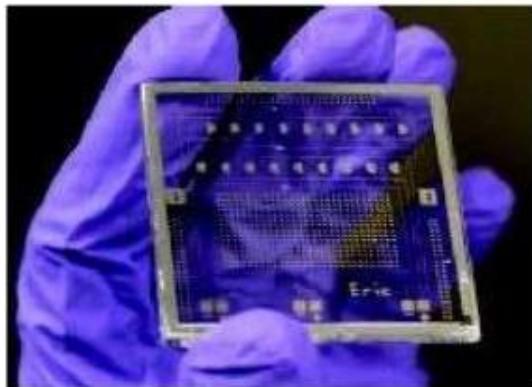




● Antigen  
Y Antibody



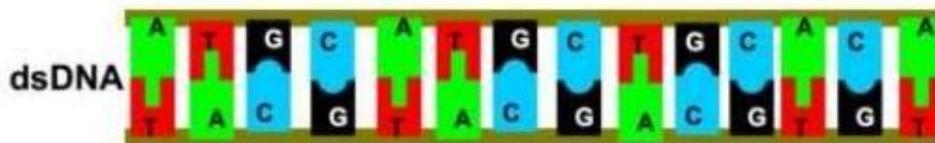
# DNA biosensors



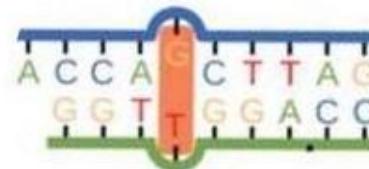
# Principles of DNA biosensors

## ❖ Nucleic acid hybridization

- **Perfect match**  
stable dsDNA, strong hybridization



- **One or more base mismatches**  
weak hybridization



## ❖ Forms of DNA Biosensors

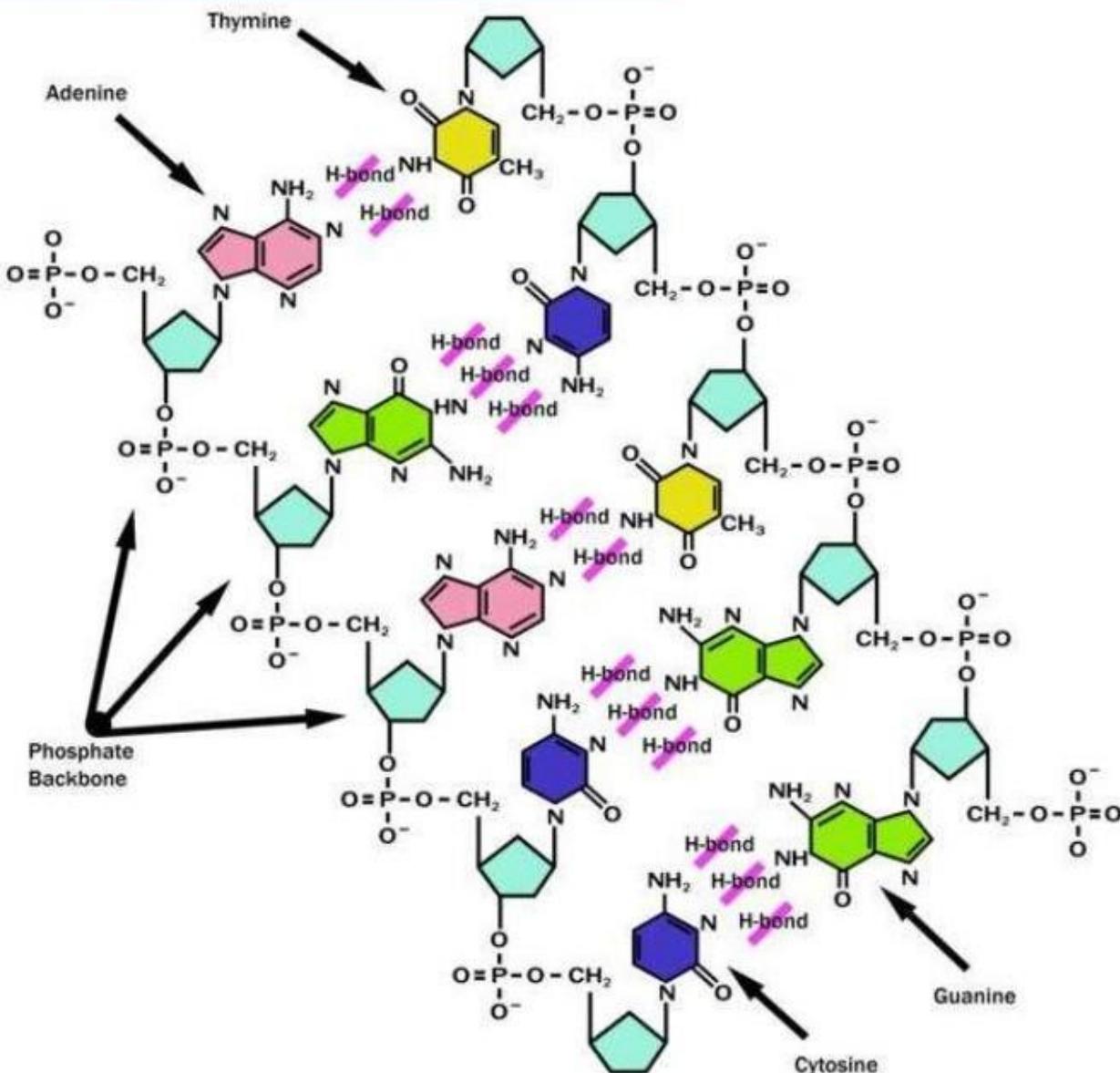
- Electrodes
- Chips
- Crystals

## ❖ Types of DNA Based Biosensors

- Optical
- Electrochemical
- Piezoelectric

# Immobilization of DNA Probe onto Transducer Surface

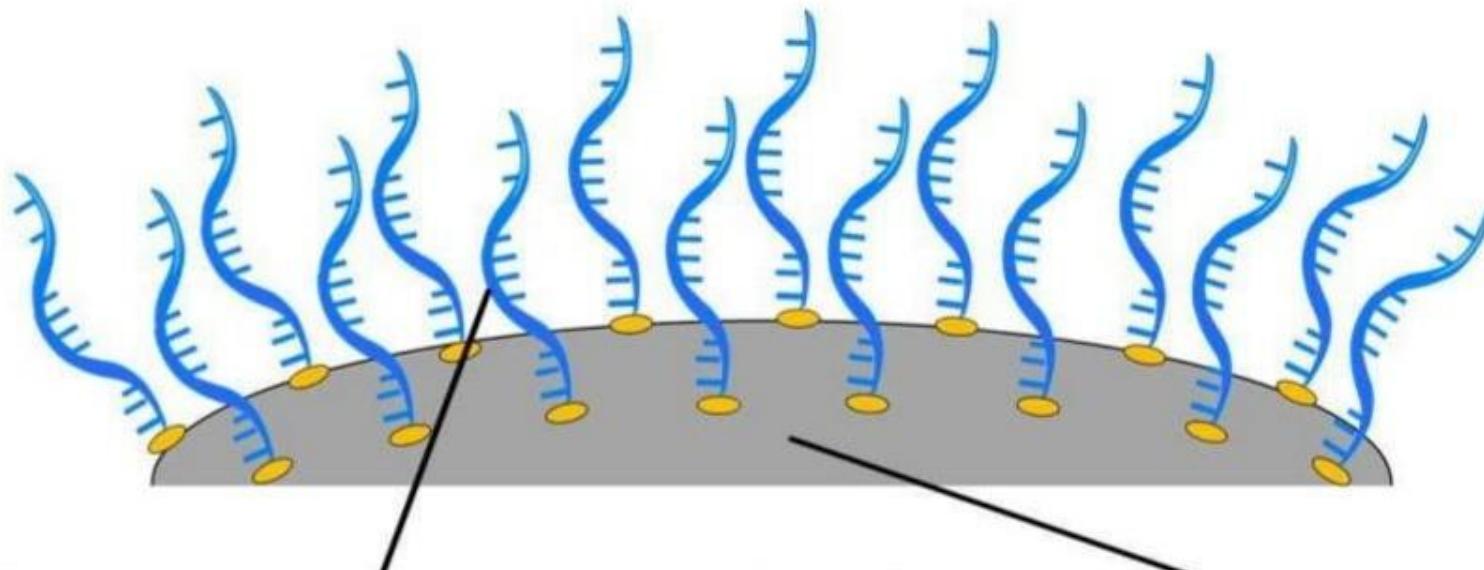
- simple adsorption onto carbon surfaces



## Immobilization of DNA Probe onto Transducer Surface

- Thiolated DNA for self assembly onto gold (or platinum) transducers

### SAM conjugation

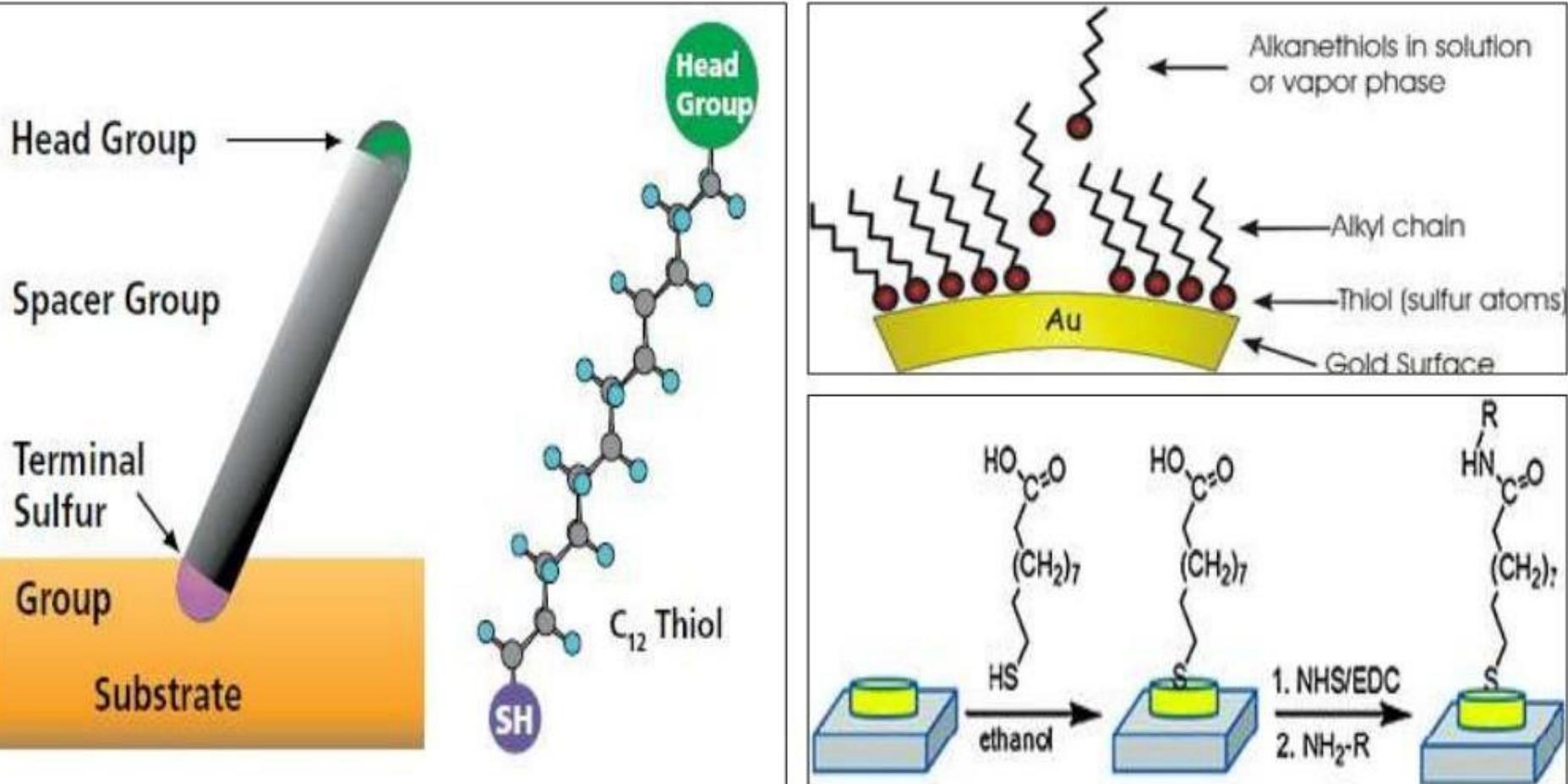


**Thiol modified DNA**

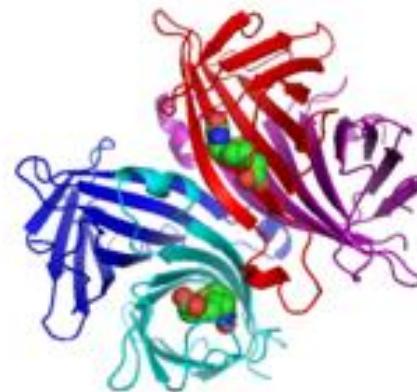
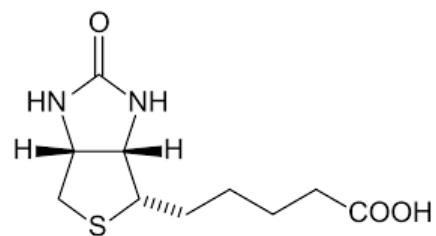
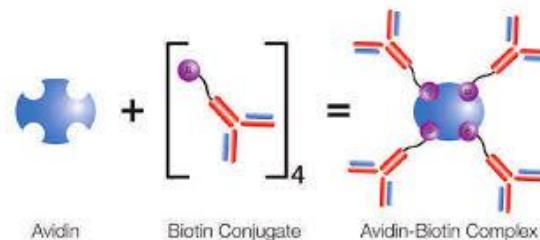
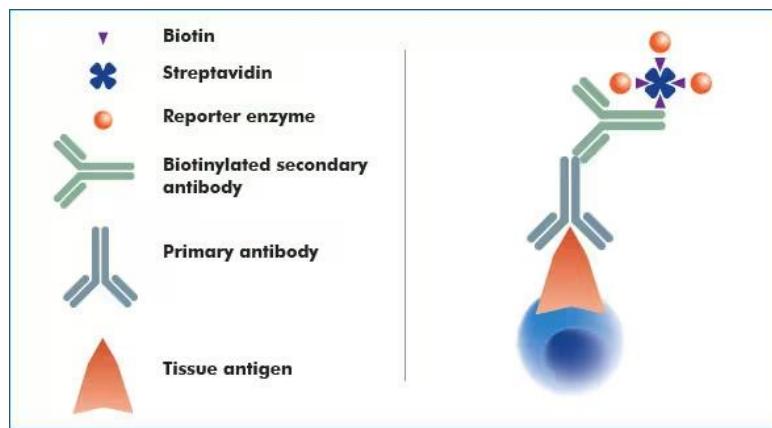
**Au or Pt Surface**

## Immobilization of DNA Probe onto Transducer Surface

- Covalent linkage to the gold surface via functional **alkanethiol-based** monolayers

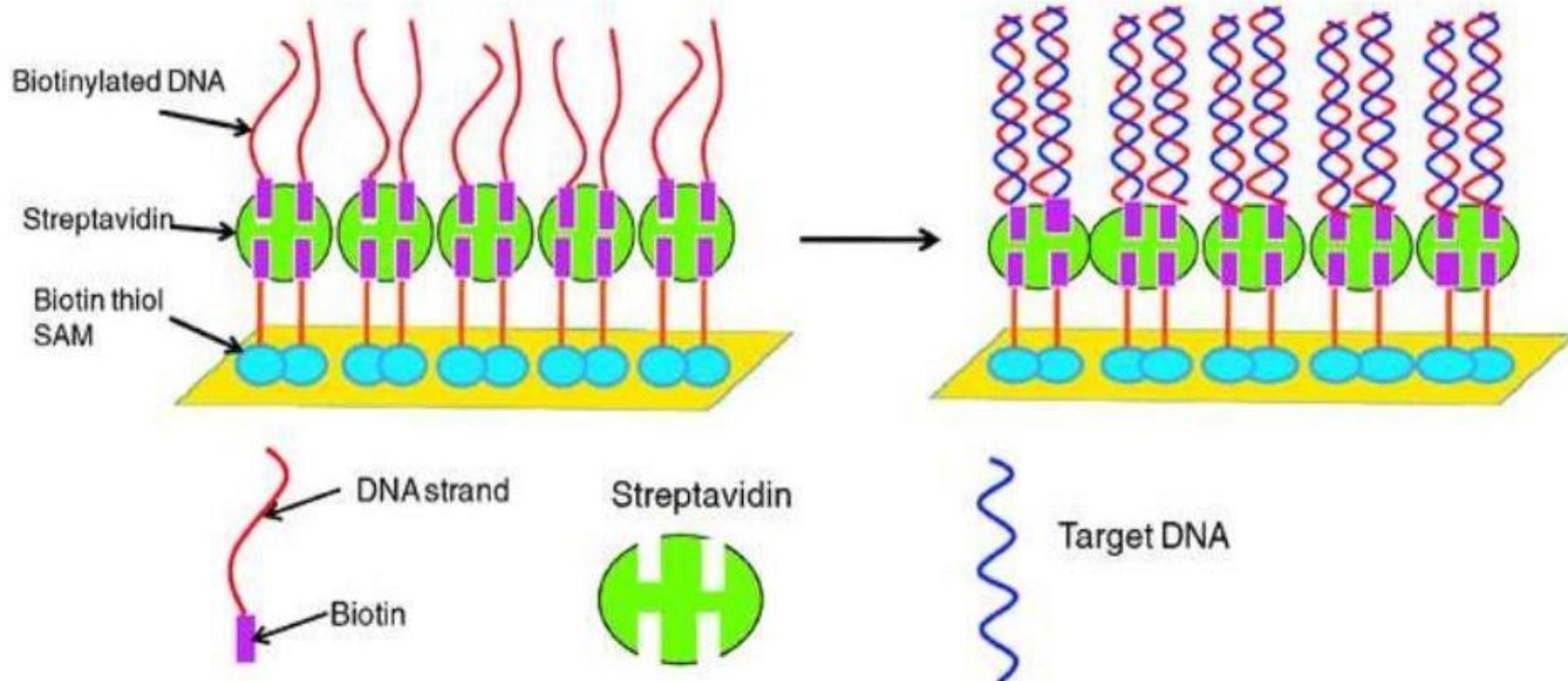


Streptavidin- Biotin  $K_D = 10^{-15}$



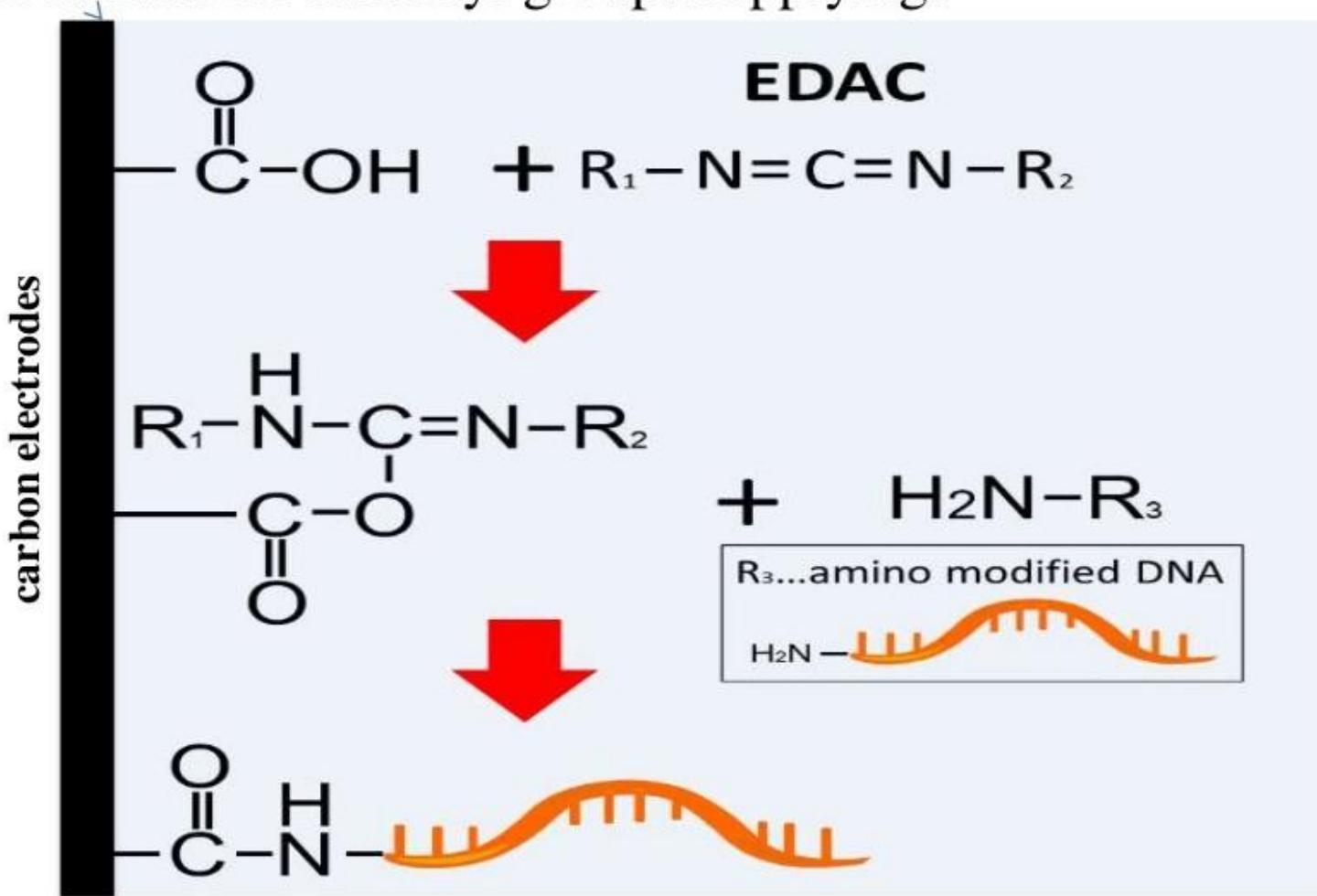
## Immobilization of DNA Probe onto Transducer Surface

- Use of biotinylated DNA for complex formation with a surface-confined avidin or streptavidin



## Immobilization of DNA Probe onto Transducer Surface

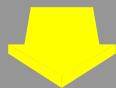
- Covalent (carbodiimide) coupling to functional groups on carbon electrodes for carboxyl groups supplying.



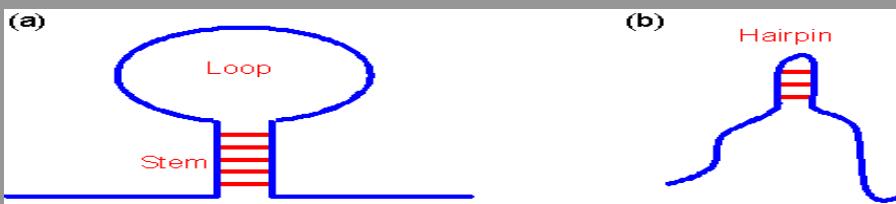
# Immobilization method of DNA probes on functionalized surfaces.

Immobilization Method	Interaction or Reaction	Advantages	Drawbacks
<b>Physical Adsorption</b>		Simple	Desorption by change of ionic strength or pH
	Charge-charge interaction or	Fast	Random orientation
	Hydrophobic interaction	Direct method (no linker molecules)	Desorption by detergent
		Suitable to DNA, RNA, and PNA	Problem of crowding effect and poor reproducibility
<b>Covalent bonding</b>	Chemical bonding	Good stability	Use of linker molecules
		High binding strength	Slow, Irreversible
		Use during long term	Problem of crowding effect
			Island formation
<b>Streptavidin-Biotin interactions</b>	Specific Streptavidin-Biotin interaction	Improved orientation	Expensive, Slow
		High specificity and functionality	Problem of crowding effect
		Well-controlled	Use of biocompatible linker
		Reversible	Poor reproducibility

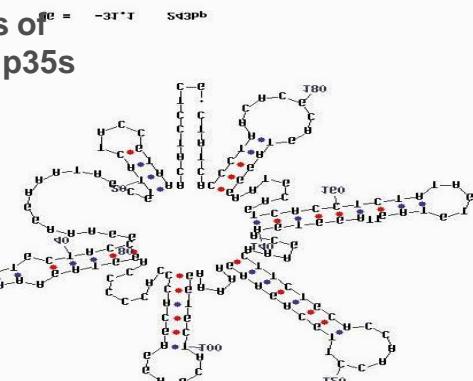
ssDNA strands can form intra-strand base pairs and secondary structures



Limitation of the available ssDNA target for hybridisation  
with the immobilised probe



Secondary structures of  
single stranded DNA p35s

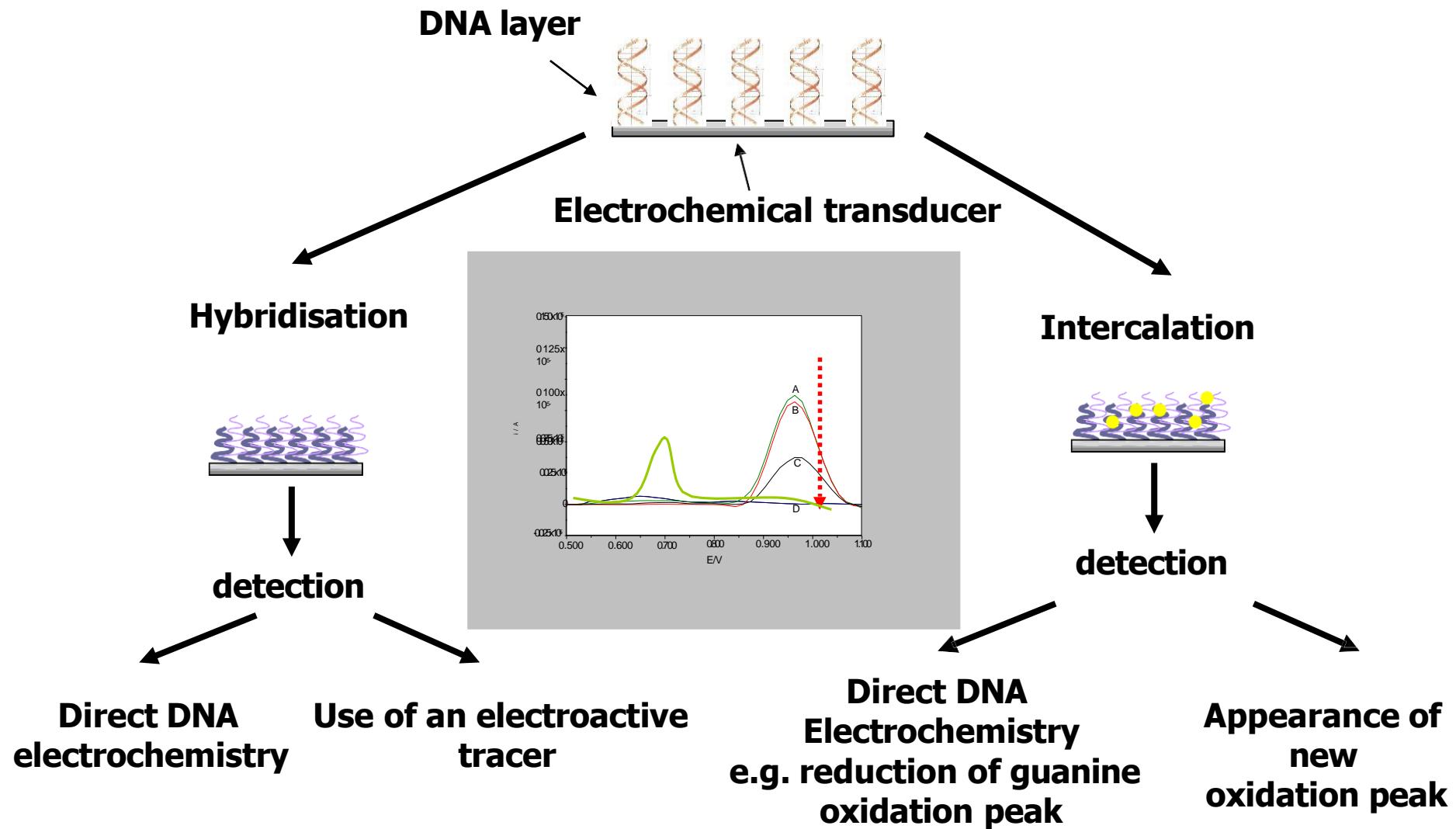


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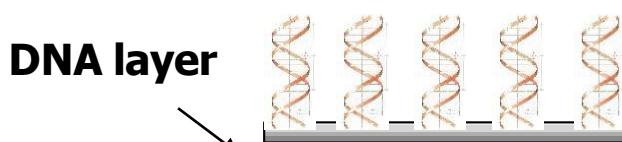
To avoid this: proper denaturation!

Aim: prolongate the ssDNA life time as much as possible, preventing dsDNA formation

# DNA electrochemical biosensors



# DNA electrochemical biosensors application



**Hybridisation:**

**Molecular diagnostic application:**

**microorganism**

**gene single-point mutation**

**species identification**

**GMO**

**Intercalation**

**Antitumoral drugs**

**Genotoxic compounds:**

**stable DNA complex**

**adduct formation**

# Sensori elettrochimici a DNA

Table 1  
Categories of electrochemical NA sensing principles.

	Label-free, reagent-less	Label-free, reagent-dependent	Labeled, reagent-less	Labeled, reagent-dependent
Heterogeneous detection	<ul style="list-style-type: none"><li>• Change in capacitance</li><li>• Change in impedance</li><li>• Field-effect</li></ul>	<ul style="list-style-type: none"><li>• Intercalation</li><li>• Groove-binding</li><li>• Electrostatic binding</li><li>• Electrostatic repulsion</li></ul>	<ul style="list-style-type: none"><li>• Labeled capture probes</li><li>• Labeled signaling probes</li><li>• Labeled nucleotides</li></ul>	<ul style="list-style-type: none"><li>• Enzyme labels</li></ul>
Homogeneous detection	<ul style="list-style-type: none"><li>• Detection of NA amplification by-product with ISFET</li></ul>	<ul style="list-style-type: none"><li>• Consumption of electroactive molecules by interaction with NA</li></ul>	<ul style="list-style-type: none"><li>• Release of electroactive molecules</li></ul>	<ul style="list-style-type: none"><li>• No principle known</li></ul>
Advantages	<ul style="list-style-type: none"><li>• Cost-effective reagents possible</li></ul>	<ul style="list-style-type: none"><li>• Enhanced signals</li></ul>	<ul style="list-style-type: none"><li>• Specific signal</li></ul>	<ul style="list-style-type: none"><li>• Signal amplification</li></ul>
Drawbacks	<ul style="list-style-type: none"><li>• Low signals</li><li>• Risk of unspecific signal changes</li></ul>	<ul style="list-style-type: none"><li>• Risk of unspecific signal changes</li></ul>	<ul style="list-style-type: none"><li>• Modification of oligonucleotide increases costs</li></ul>	<ul style="list-style-type: none"><li>• Additional process steps complicate automation</li></ul>

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Review: Electrochemical DNA sensing – Principles, commercial systems, and applications

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<sup>b</sup> Laboratory for MEMS Applications, Department of Microsystems Engineering – IMTEK, University of Freiburg, Georges-Koehler-Allee 103, 79110, Freiburg, Germany

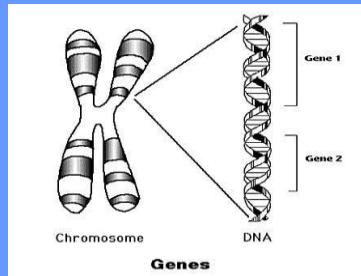
<sup>c</sup> Faculty of EECS, Chair of Sensor and Actuator Systems, TU Berlin, Einsteinstr 17, 10587, Berlin, Germany

# Detection of DNA hybridisation in PCR samples



**Fig. 1.** Images of some of the reviewed commercial electrochemical NA detection systems in the order of their appearance in the text. A: Cubed Laboratories' NESDEP instrument (copyright: Cubed Laboratories) B: Canon's Genelyzer II instrument (copyright: Canon Medical Systems Corp.) C: GenMark's ePlex instrument (four tower version shown – the device can also be equipped with fewer towers, copyright GenMark Diagnostics Inc.) D: Friz Biochem's envisioned Cycle device (copyright: Friz Biochem GmbH) E: CustomArray's ElectraSense reader (copyright: Custom Array Inc.) F: Elice's Leo instrument (copyright: Easy Life Science) G: Binx's io instrument (copyright: Binx Health Inc.). The images are not to scale. All images are published with the permission of the respective companies.

- *Development of an Hybridisation sensor*



**Synthesis of a DNA fragment (probe, bioreceptor) containing the sequence of interest (analytical problem)**

**Immobilisation of the probe onto the solid support of the sensor (surface) (thiol/dextran/streptavidin/biotinylated probe)**

**Extraction of the DNA from the real sample (blood, water, food) and amplification of the sequence of interest (sample pretreatment)**

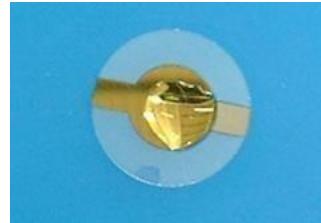
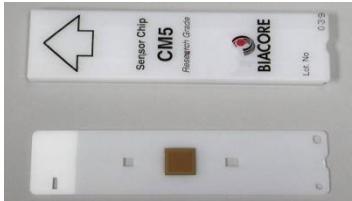
**Denaturation of the dsDNA (amplified fragment or genomic) to obtain a single-stranded DNA (sample pretreatment)**

**Hybridisation of the obtained ssDNA with the immobilised probe**

**Changes in the physicochemical parameters of the layer formed on the transducer (quartz crystal or gold –glass chip)**

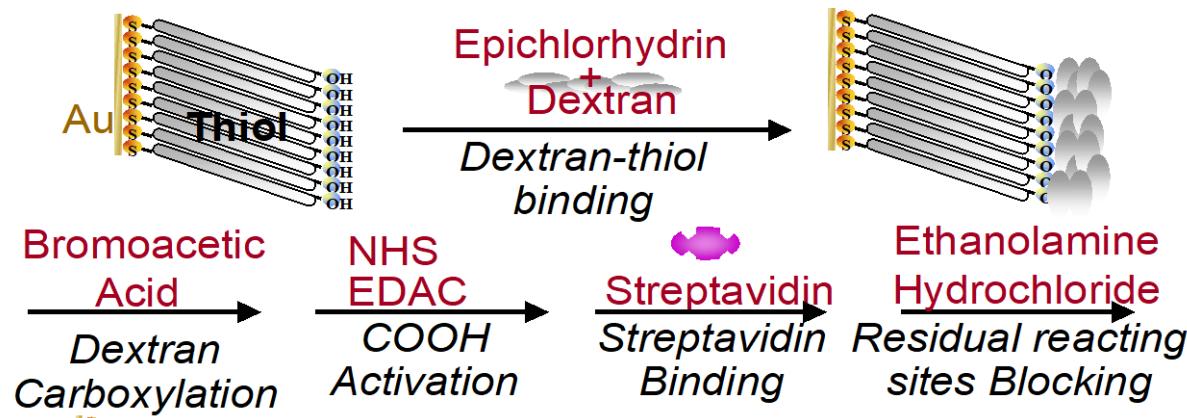
# Probe immobilisation on gold film

optical

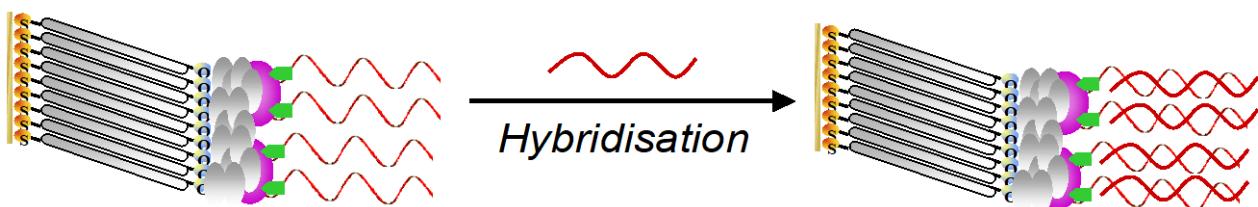


piezoelectric

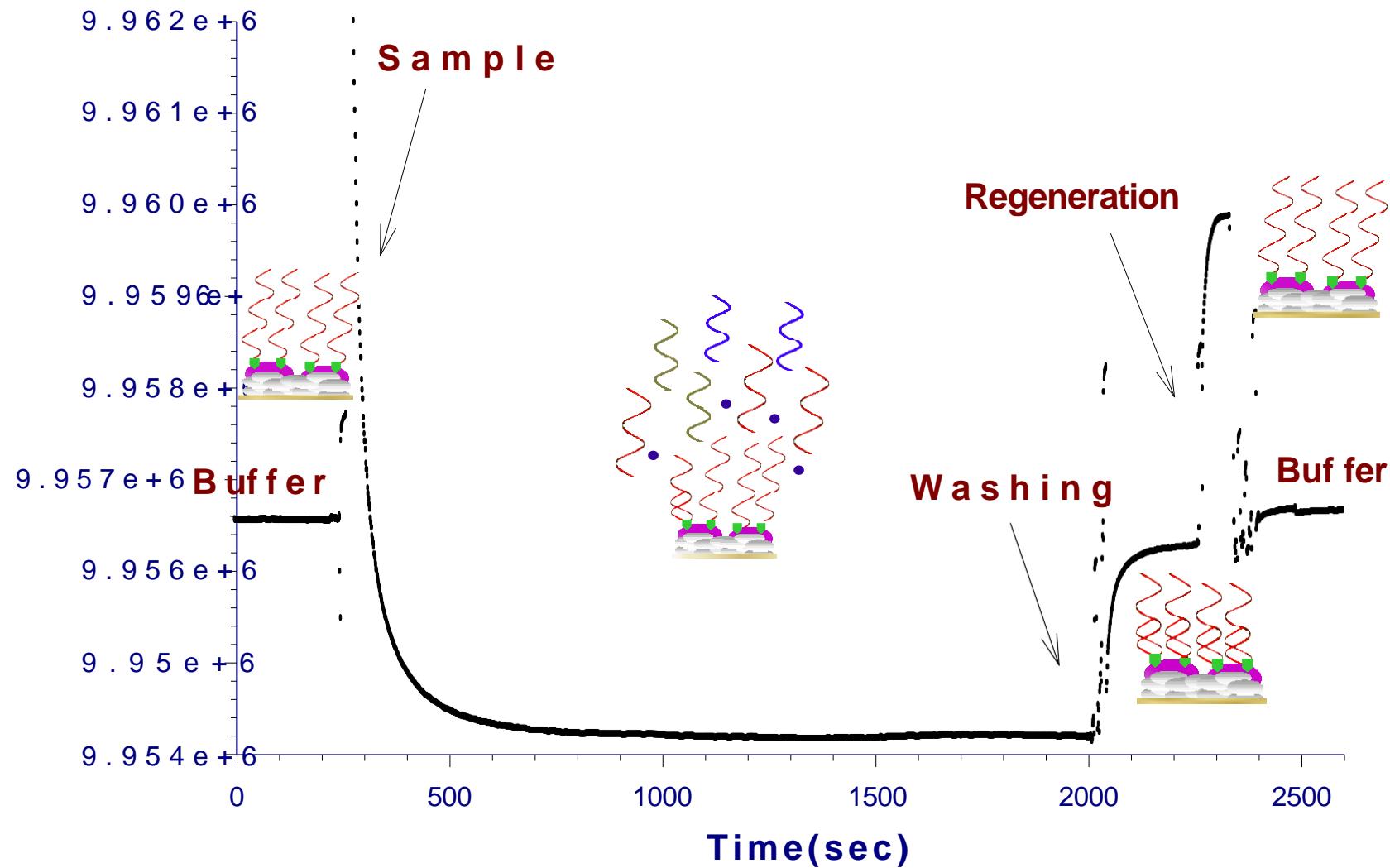
## *thiol/dextran/streptavidin/biotinylated probe*



Specificity,  
no aspecific  
adsorption,  
stability,  
multi-use



# Hybridization-Regeneration Cycle



# Detection of DNA target sequence in real matrices

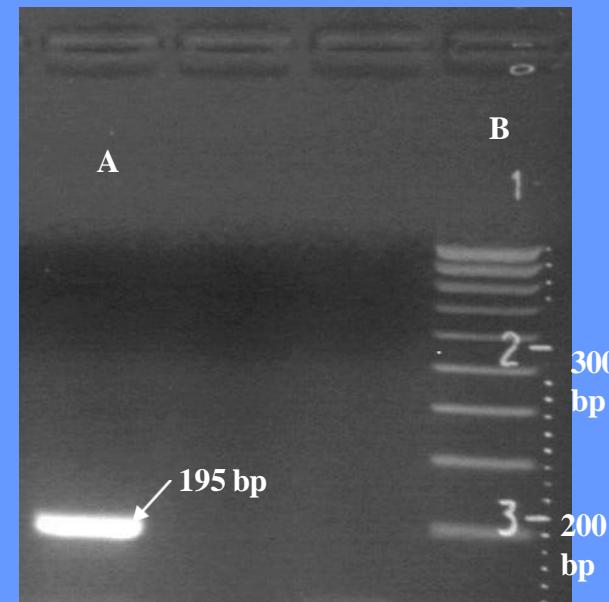
- Certified Reference Materials (CRM) -Fluka
- Processed food (dietetic snaks, soy crackers, soft drinks)

Control: Post PCR  
Electrophoresis

## 1. PCR amplified DNA

### *Processing the sample*

- Extraction of DNA from samples (CTAB method)
- Amplification of DNA by PCR (Pietsh K. et al. 1997)
- Dilution with *Hybridisation buffer*: NaCl 150 mM, Na<sub>2</sub>HPO<sub>4</sub> 20 mM, EDTA 0.1 mM, pH 7.4
- Denaturation to obtain ssDNA from amplified dsDNA

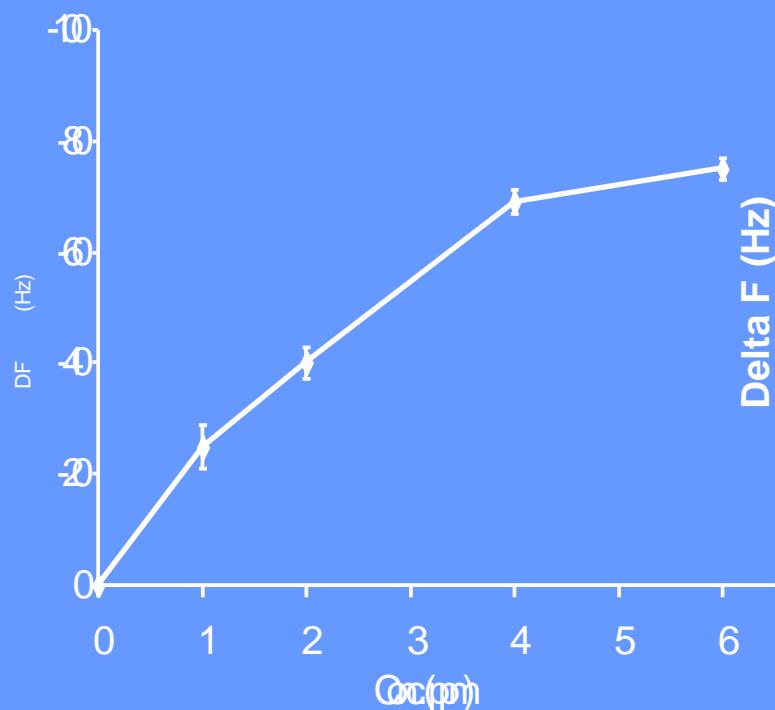


A: amplified fragment  
(Promoter 35S 195 bp)

B: Standard length fragments

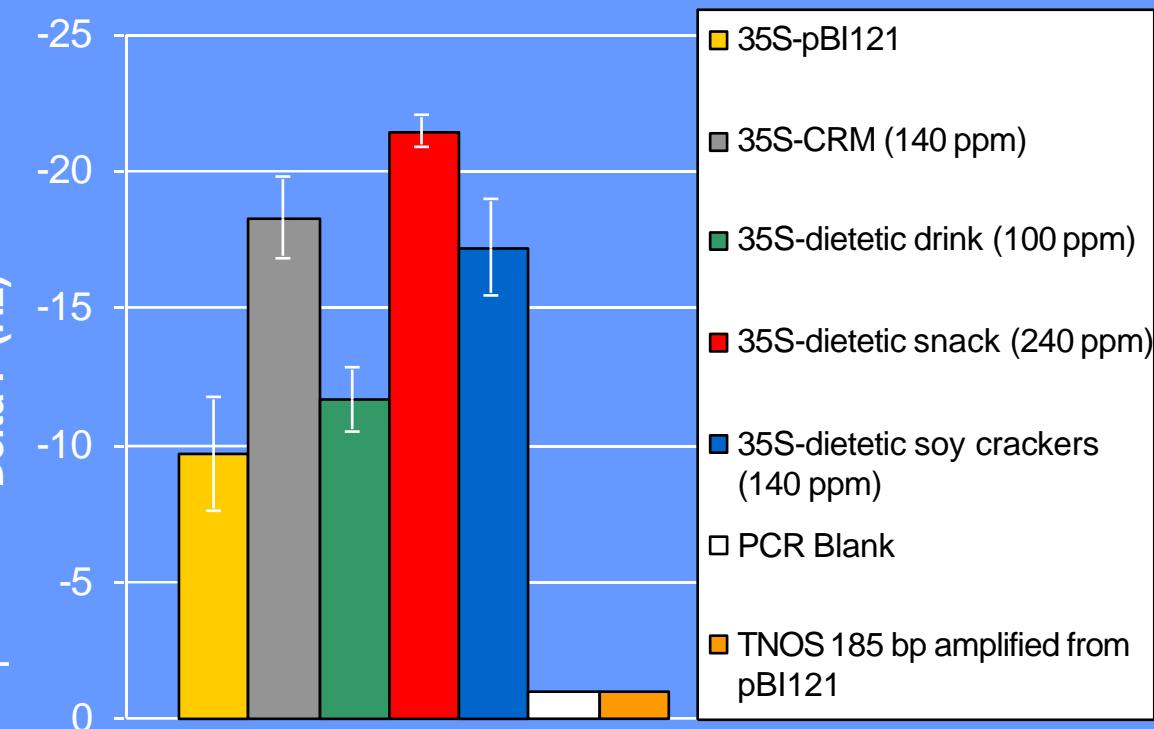
# Piezoelectric sensor, CRM 2% samples and processed food samples

Sample pre-treatment: PCR amplified DNA, thermal denaturation



DL: 0,3 ppm

CV% 6 (n=3)



5'-BIOT-ggc cat cgt tga aga tgc ctc tgc c-3'      probe 35S  
3'- ccg gat gca act tct acg gag acg g-5      target 35S

# Biomimetic receptors

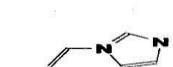
Obtained via combinatorial chemistry and/or molecular modelling

MIP (Molecularly imprinted polymers)

Peptides

Aptamers

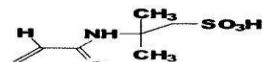
## Functional Monomer Database



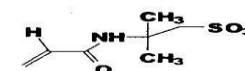
F1 1-VINYLMIDAZOLE



F2 2-VINYLPYRIDINE



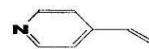
F3 ACRYLAMIDO-2-METHYL-1-PROPANESULFONIC ACID (AMPSA)



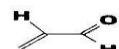
F3a ACRYLAMIDO-2-METHYL-1-PROPANESULFONIC ACID (AMPSA)



F4 2-HYDROXYETHYL METHACRYLATE



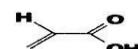
F5 4-VINYLPYRIDINE



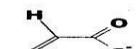
F6 ACROLEIN



F7 ACRYLAMIDE



F8 ACRYLIC ACID



F8a ACRYLIC ACID



F9 ACRYLONITRILE



F10 ALLYLAMINE



F10a ALLYLAMINE



F11 p-DIVINYLBENZENE



F12 ETHYLENE GLYCOL DIMETHACRYLATE (EGDMA)



F13 UROCANIC ACID ETHYL ESTER



F14 ITACONIC ACID



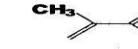
F14a ITACONIC ACID



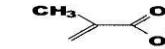
F15 m-DIVINYLBENZENE



F16 N,N-METHYLENE BIS ACRYLAMIDE



F17 METHACRYLIC ACID



F17a METHACRYLIC ACID



F18 STYRENE



F19 UROCANIC ACID



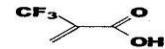
F19a UROCANIC ACID



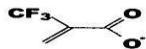
F20 N,N-DIETHYLAMINO ETHYL METHACRYLATE (DEAEM)



F20a N,N-DIETHYLAMINO METHACRYLAMIDE (DEAEM)

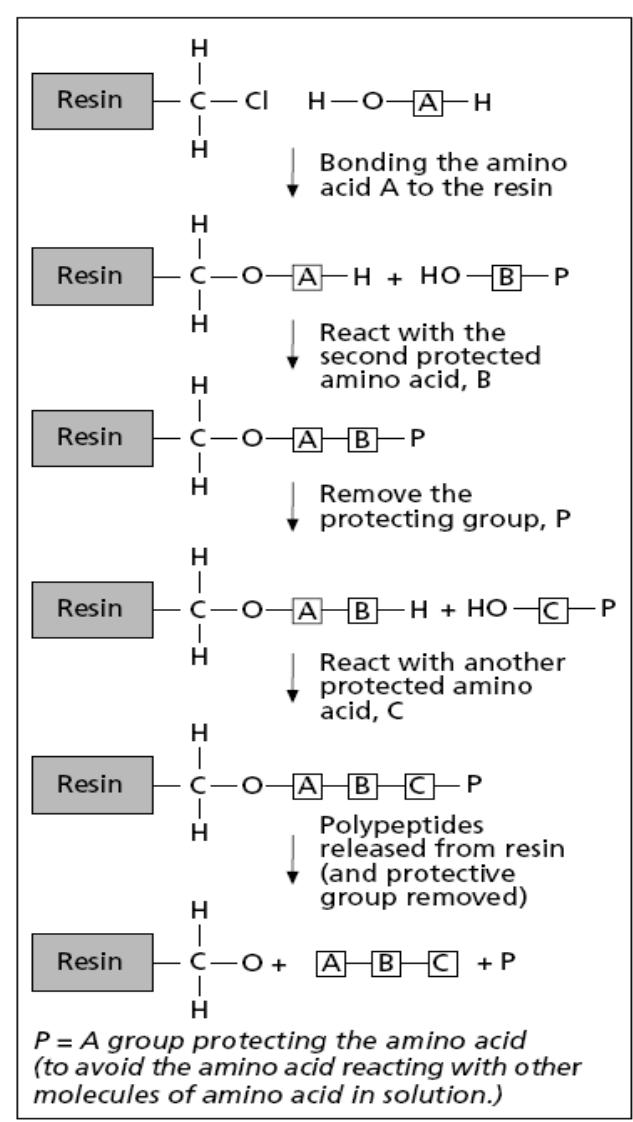


F17 TRIFLUOROMETHACRYLIC ACID



F17a TRIFLUOROMETHACRYLIC ACID

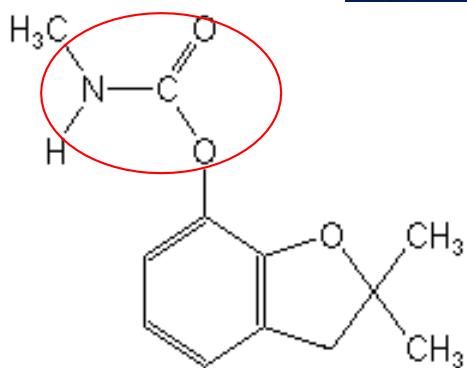
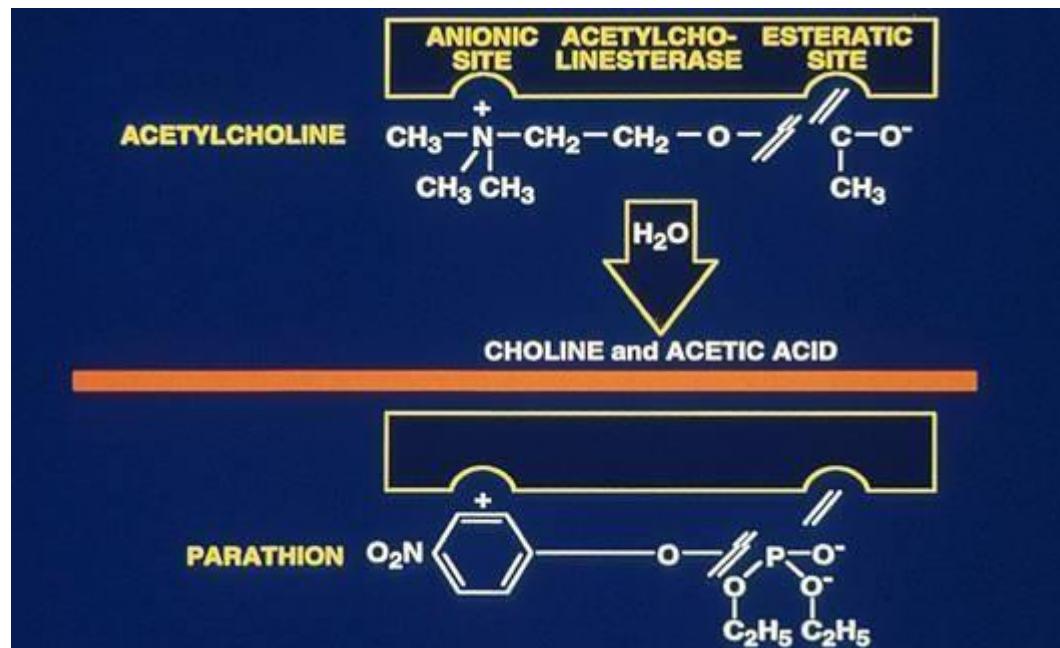
# Combinatorial chemistry approach: Synthesys of aminocids via split and mix



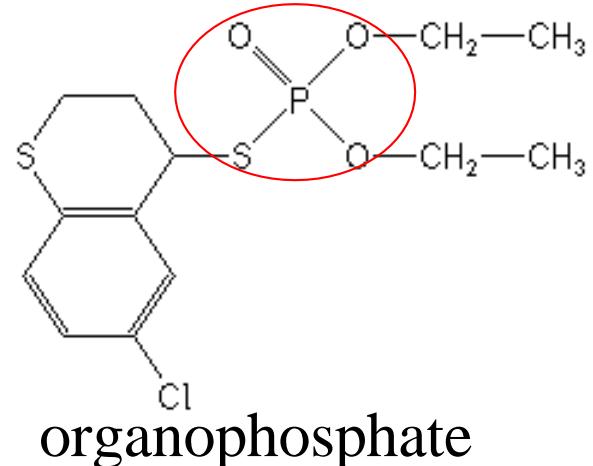
*Split synthesis*

Stage	Reaction vessel 1 (A)	Reaction vessel 1 (B)	Reaction vessel 1 (C)	
1	Resin + A	Resin + B	Resin + C	3 compounds
			MIX	
2	Resin-A+A Resin-B+A Resin-C+A	Resin-A+B Resin-B+B Resin-C+B	?	9 compounds
			MIX	
3	Resin-A-A+A Resin-B-A+A Resin-C-A+A Resin-A-B+A Resin-B-B+A Resin-C-B+A Resin-A-C+A Resin-B-C+A Resin-C-C+A	Resin-A-A+B Resin-B-A+B Resin-C-A+B Resin-A-B+B Resin-B-B+B Resin-C-B+B Resin-A-C+B Resin-B-C+B Resin-C-C+B	Resin-A-A+C Resin-B-A+C Resin-C-A+C Resin-A-B+C Resin-B-B+C Resin-C-B+C Resin-A-C+C Resin-B-C+C Resin-C-C+C	27 compounds
			MIX	

# BIOMIMETIC RECEPTORS FOR PESTICIDES



Carbamate



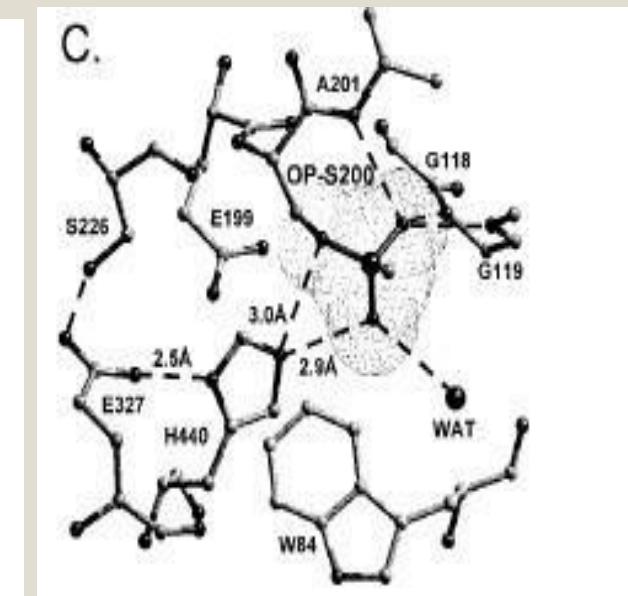
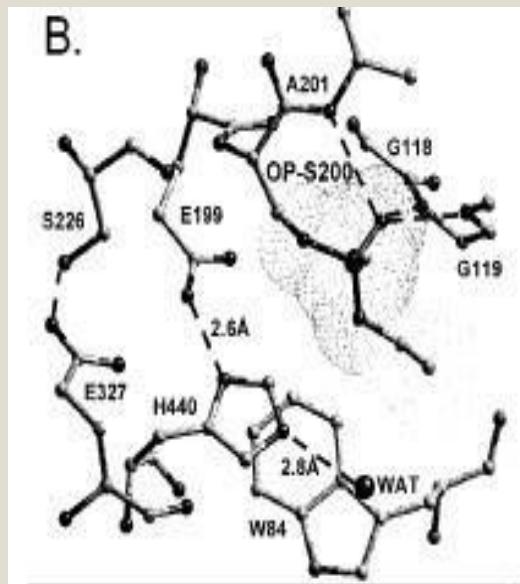
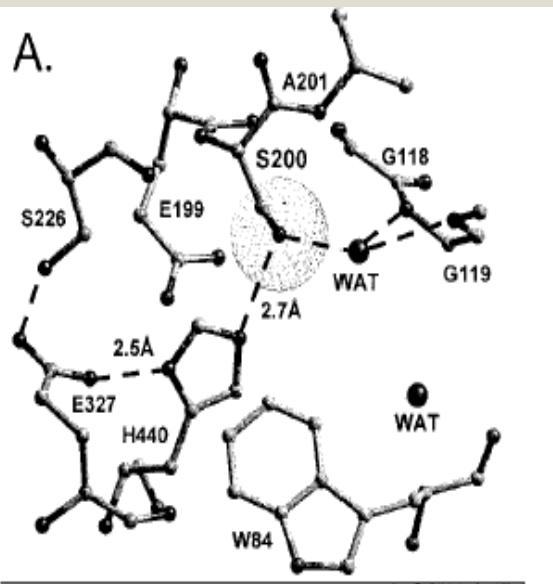
organophosphate

# ✓ Mechanism of AChE inhibition

**AChE**, the target enzyme of pesticides, **is an efficient serine hydrolase** that catalyzes the breakdown of acetylcholine (ACh)



## How pesticides work

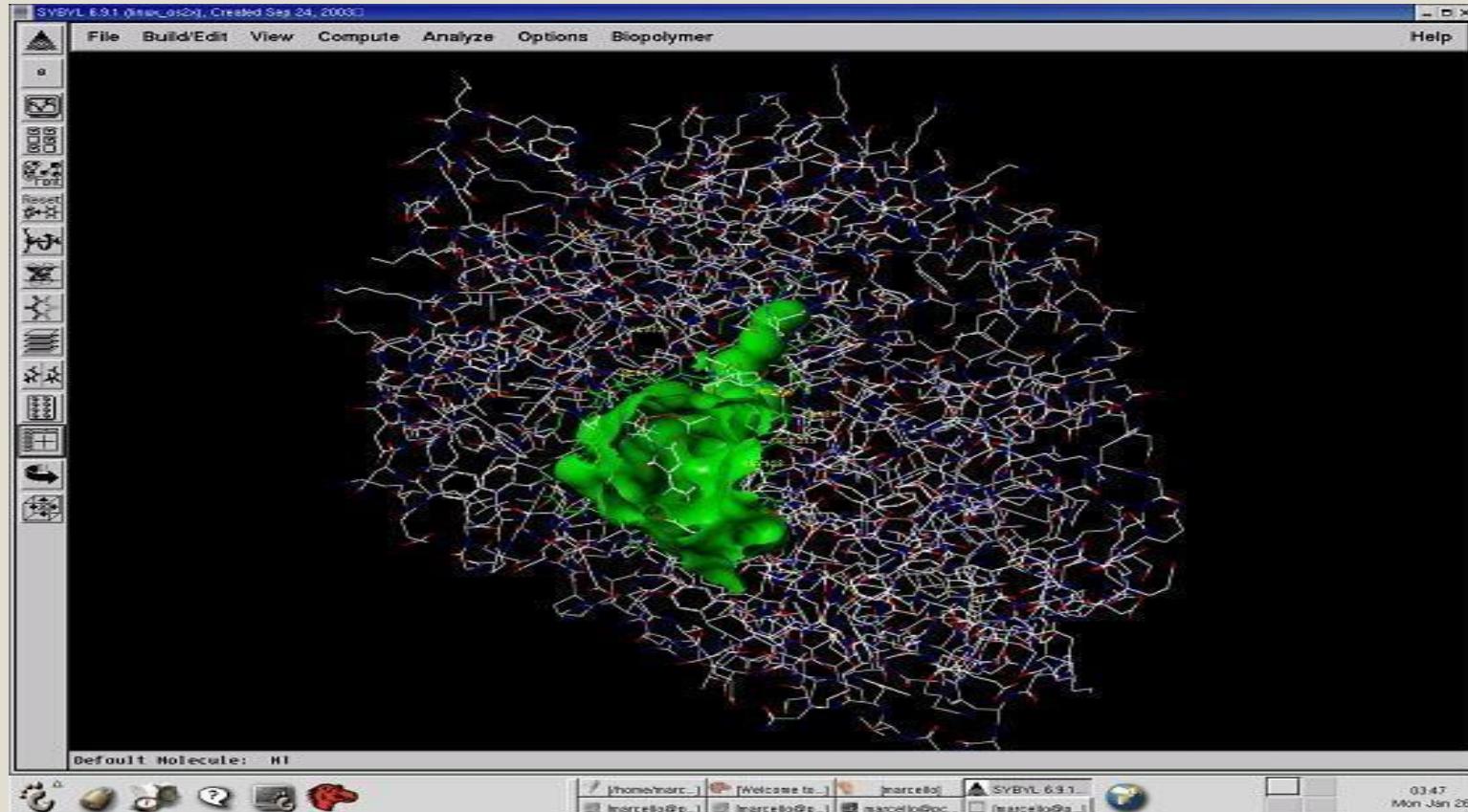


From Millard *et al* J.Am.Chem.Soc. 121, (1999)

## ❖ Computational screening

✓ AChE-OP crystallographic structure (PDB ID: 1VXO)

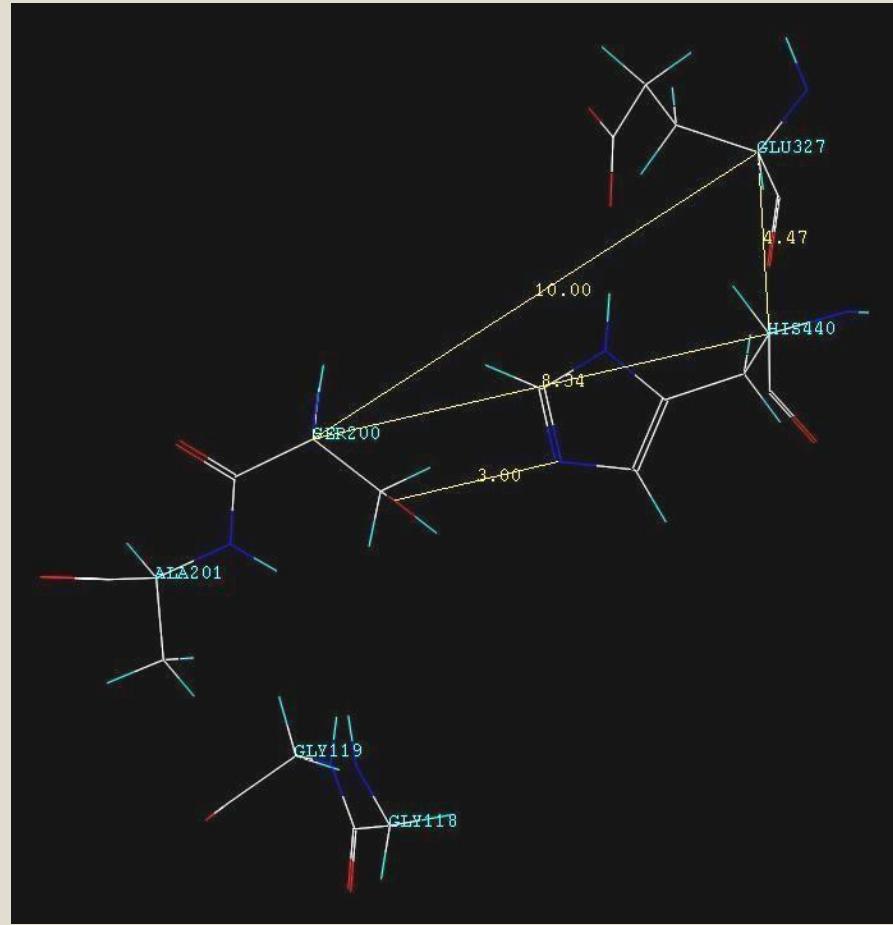
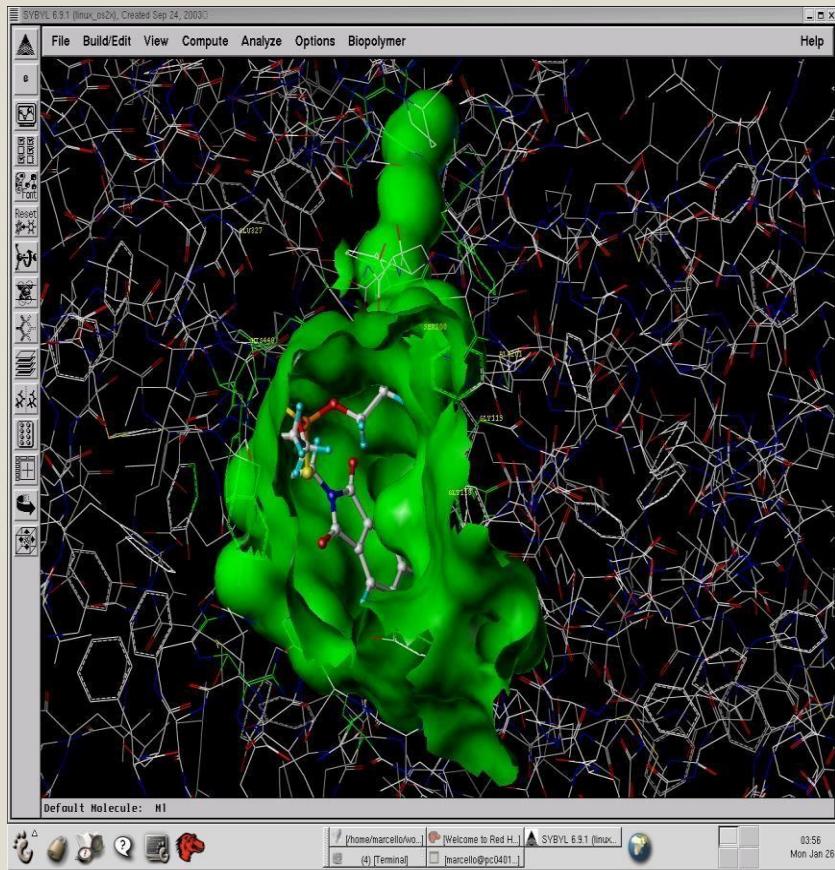
Methylphosphonylated Acetylcholinesterase (Aged) Obtained By Reaction With O-Ethyl-S-[2-[Bis(1-Methylethyl) Amino]Ethyl] Methylphosphonothioate (Vx) conventional X-ray crystallography resolution [Å]: 2.40



In green the molecular electrostatic potential distribution on the surface of the enzyme binding pocket

# ✓ Design of the oligopeptides library as possible receptors

The geometry of the binding pocket was investigated to create oligopeptides library



Three dimensional coordinates of the asymmetric carbon ( $C\alpha$ ) of each aminoacid involved in the binding pocket were calculated in order to reproduce the geometry observed

## ✓ Tetrapeptides library

- easy to synthesise
- more possibility to preserve in solution the secondary structure predicted
- A series of tetrapeptides, containing the possible combinations of the catalytic triad (SER 200, HIS 440, GLU 327) and the catalytic oxyanion hole (GLY 118 GLY 119 ALA 201) was drawn
- The proper geometry of binding pocket was achieved using alternatively a GLY or a PRO residue

Library (24 tetrapeptides)	
Ser-Gly-His-Glu	Glu-Gly-Ser-Ala
Ser-Gly-Glu-His	His-Gly-Ser-Ala
His-Glu-Gly-Ser	Gly-Pro-Ser-Ala
Glu-His-Gly-Ser	Ser-Ala-Pro-Glu
Ser-Pro-His-Glu	Ser-Ala-Pro-His
Ser-Pro-Glu-His	Ser-Ala-Pro-Gly
His-Glu-Pro-Ser	Glu-Pro-Ser-Ala
Glu-His-Pro-Ser	His-Pro-Ser-Ala
Gly-Gly-Ser-Ala	Gly-Ser-Gly-Ala
Ser-Ala-Gly-Glu	Ala-Gly-Ser-Gly
Ser-Ala-Gly-His	Ser-Gly-Pro-Ala
Ser-Ala-Gly-Gly	Ala-Pro-Gly-Ser

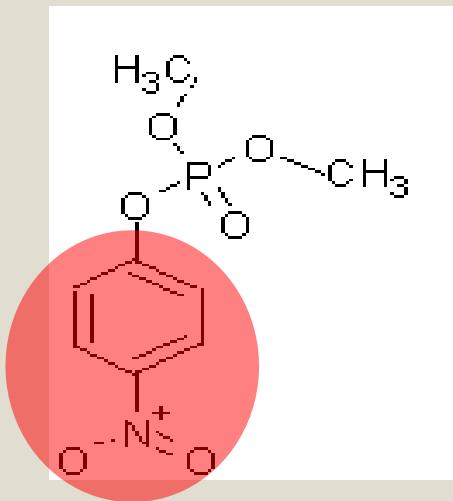
✓ Simulated binding results vs paraoxon of the tetrapeptides selected for experimental screening

	A	B	C	D
	Ser-Ala-Gly-Glu	His-Gly-Ser-Ala	Glu-Pro-Ser-Ala	His-Glu-Pro-Ser
Binding Score (kJ/mol)	38	73	21	93

Negative control (NC): Glu-His-Ser-Gly

Primary sequence of AChE catalytic triad

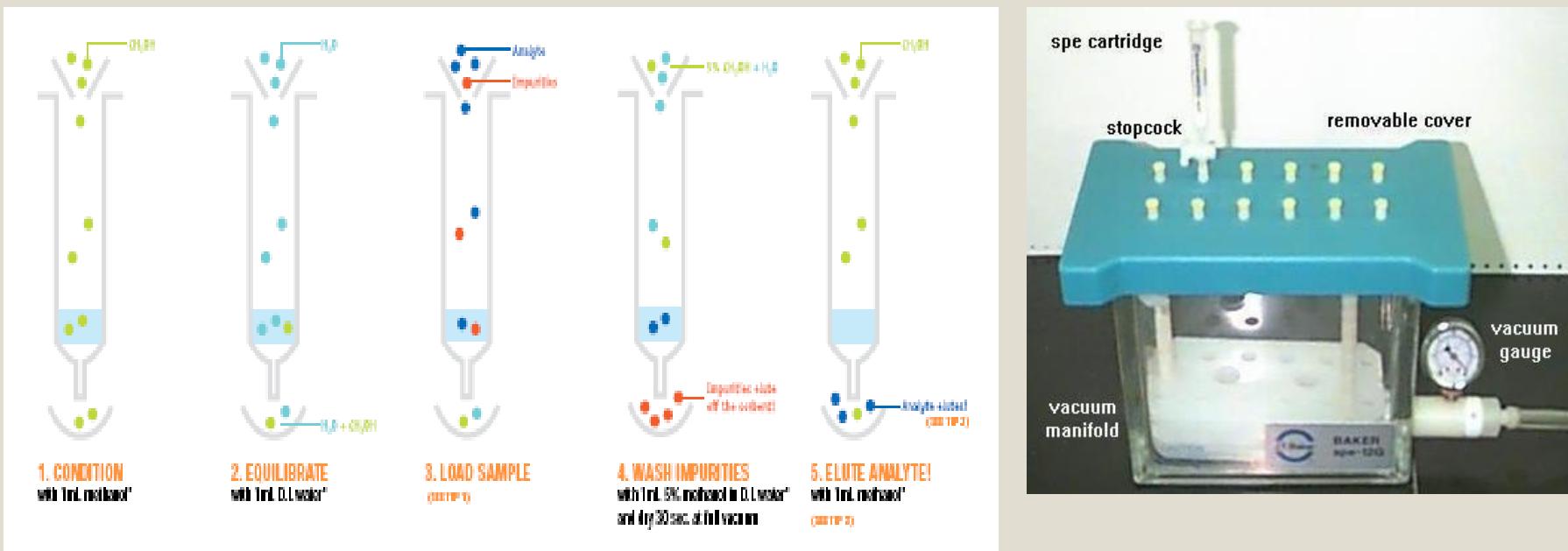
PARAOXON



- A. Ser-Ala-Gly-Glu
- B. His-Gly-Ser-Ala
- C. Glu-Pro-Ser-Ala
- D. His-Glu-Pro-Ser

NC

## ✓ Pre-analytical applications: selective affinity columns (Extraction or purification)



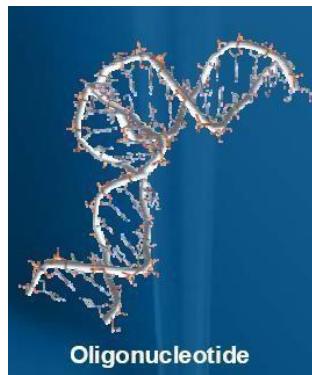
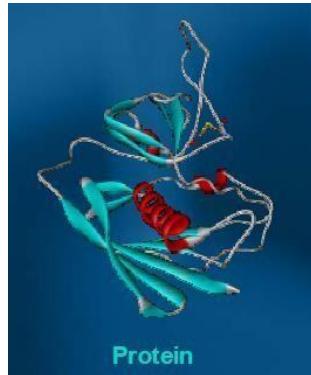
is a technique enabling purification of a biomolecule with respect to biological function or individual chemical structure. The substance to be purified is specifically and reversibly adsorbed to a ligand (binding substance), immobilized by a covalent bond to a chromatographic bed material (matrix). Samples are applied under favourable conditions for their specific binding to the ligand. Substances of interest are consequently bound to the ligand while unbound substances are washed away. Recovery of molecules of interest can be achieved by changing experimental conditions to favour desorption.

Aptamers are oligonucleotides (DNA or RNA molecules) that can bind with high affinity and specificity to a wide range of target molecules (proteins, peptides, drugs, vitamins and other organic or inorganic compounds).

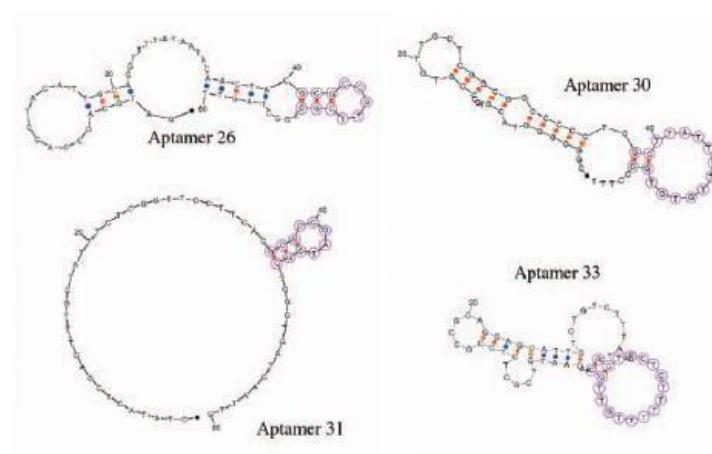
They were “discovered” in 1990 by the development of an in vitro selection and amplification technique, known as SELEX (Systematic Evolution of Ligands by Exponential enrichment).

(Ellington et al., **Nature** 346, 818; Tuerk and Gold, **Science** 249, 505)

Their name is derived from the Latin word “**aptus**” which means “to fit”.



Similar to proteins short oligonucleotides can adopt complex three-dimensional structures



*Starting point: Combinatorial oligonucleotide library*



A library containing a 25-nucleotide random region is represented by  $4^{25}$  ( $\sim 10^{15}$ ) individual sequences available for partitioning.

Normally, the starting round contains  **$10^{14}$ - $10^{15}$  individual sequences**.

**A, G, C, U(T)**

$$4^1 = 4$$

$$4^2 = 16$$

$$4^3 = 64$$

$$4^4 = 256$$

$$4^5 = 1024$$

.....

.....

.....

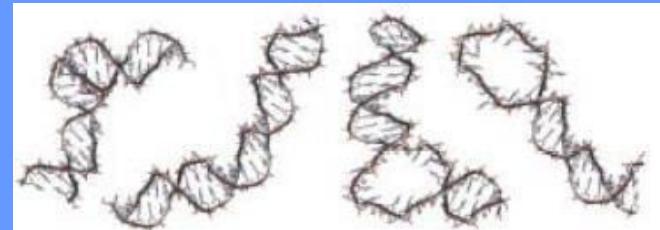
$$4^{25} = 1125899906842624$$



Pool of randomized DNA or RNA



**$10^{15}$  different sequences!!!!**





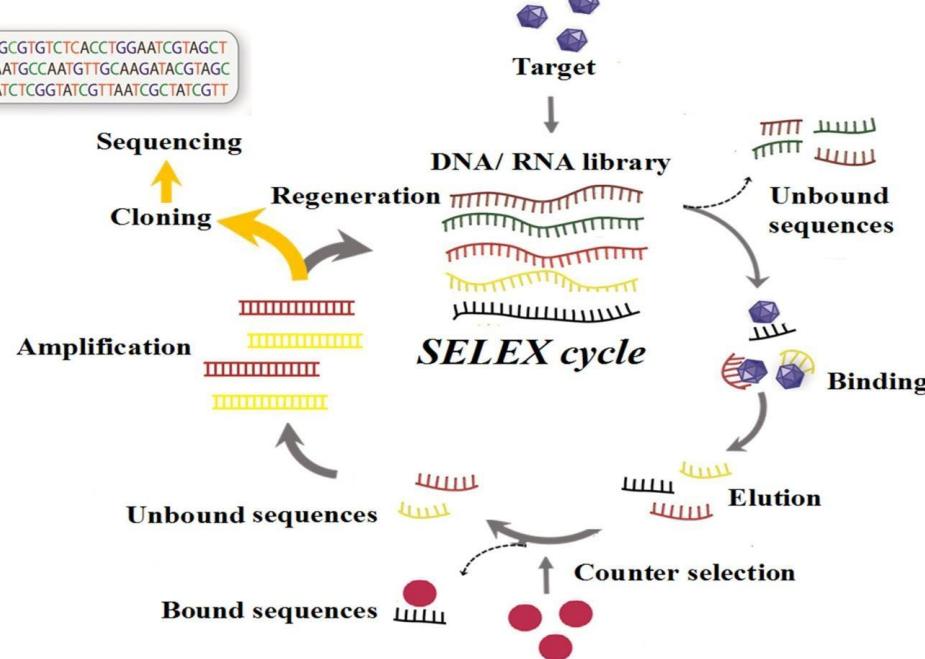
Review

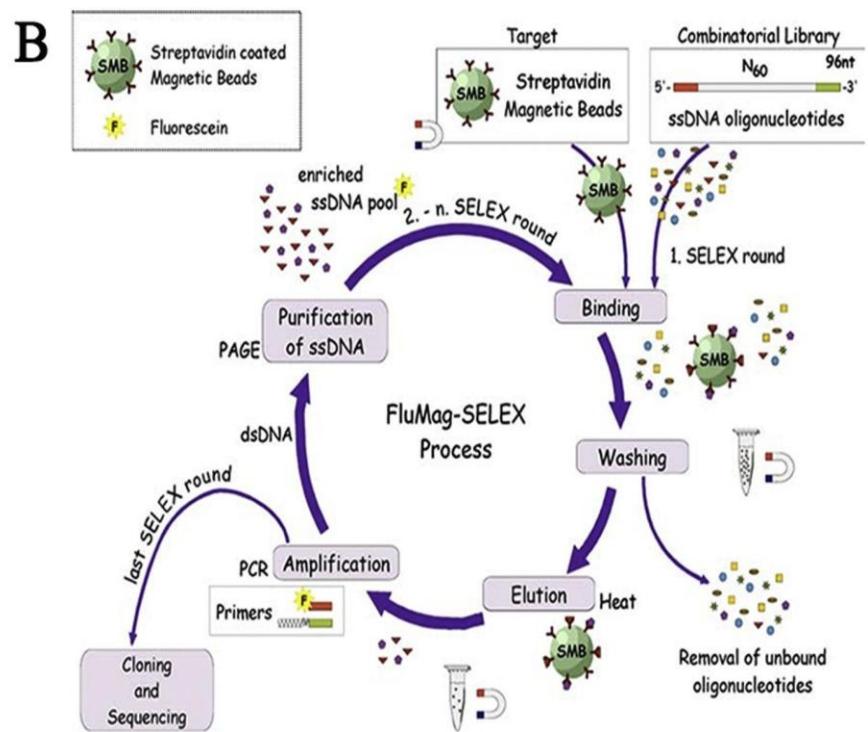
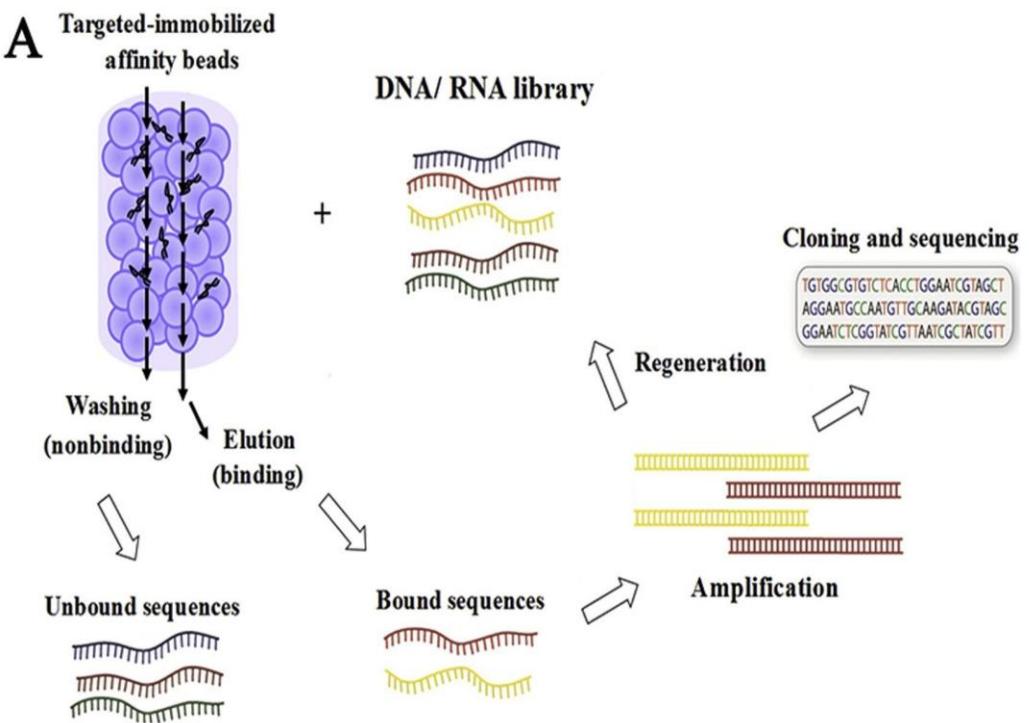
Aptamers used for biosensors and targeted therapy

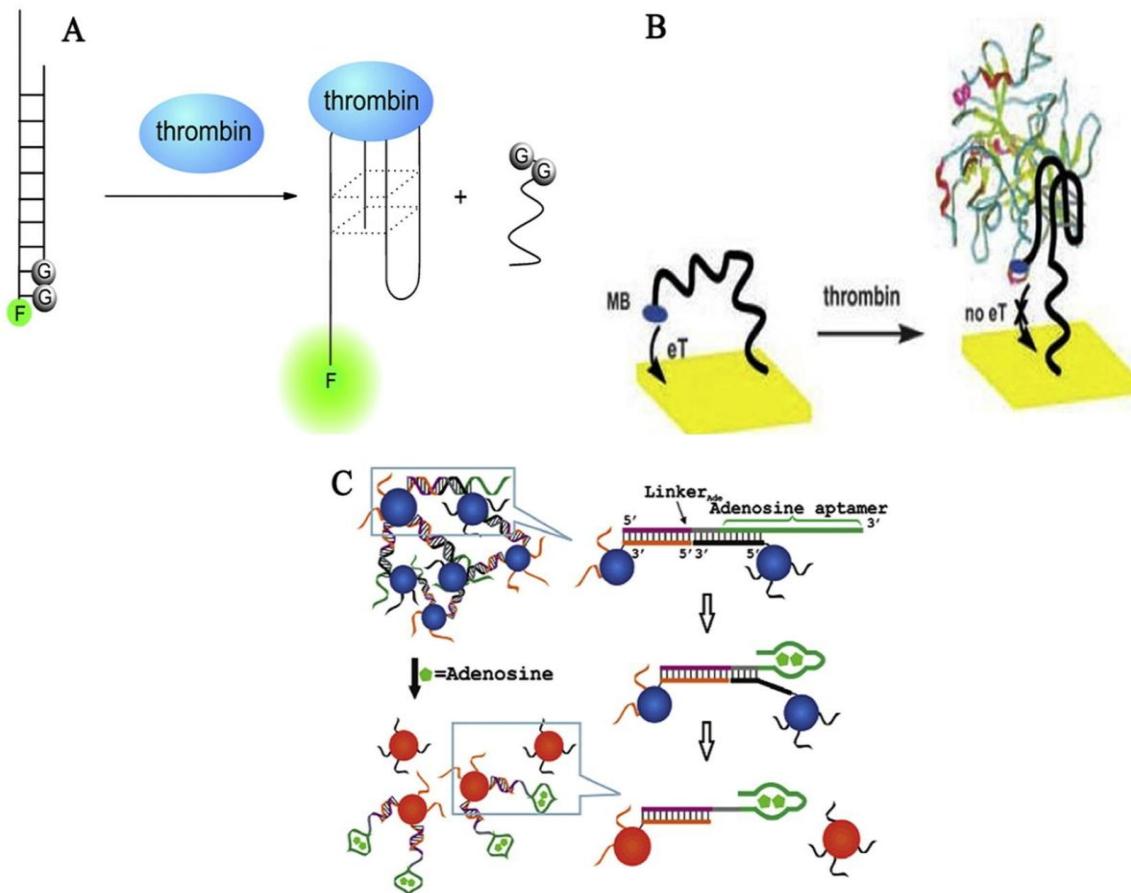


Yi Ning, Jue Hu, Fangguo Lu \*

Department of Microbiology, The Medicine School of Human University of Chinese Medicine, Chanosha, Human, 410208, PR China



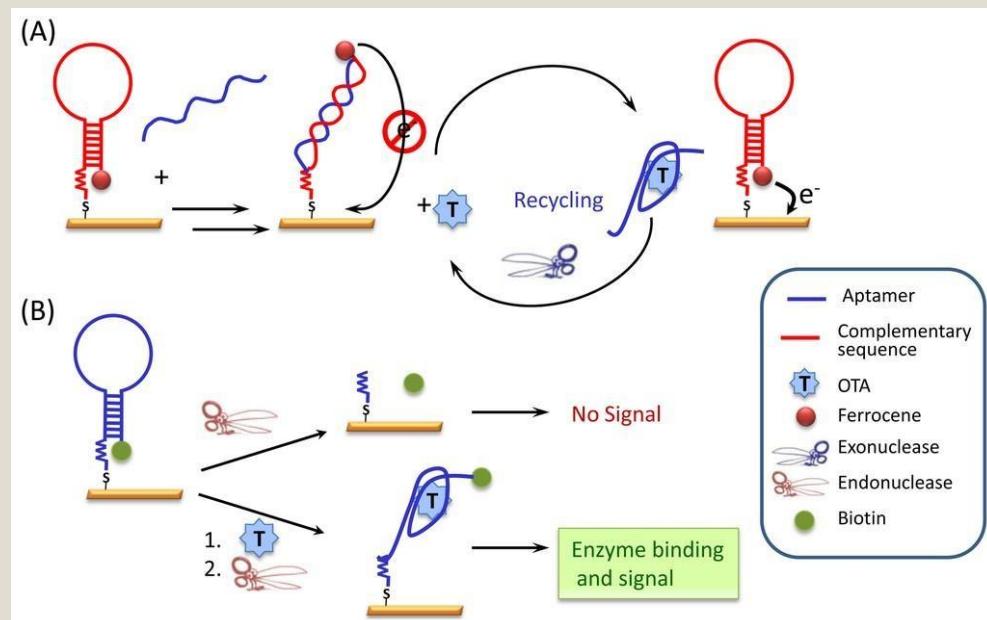
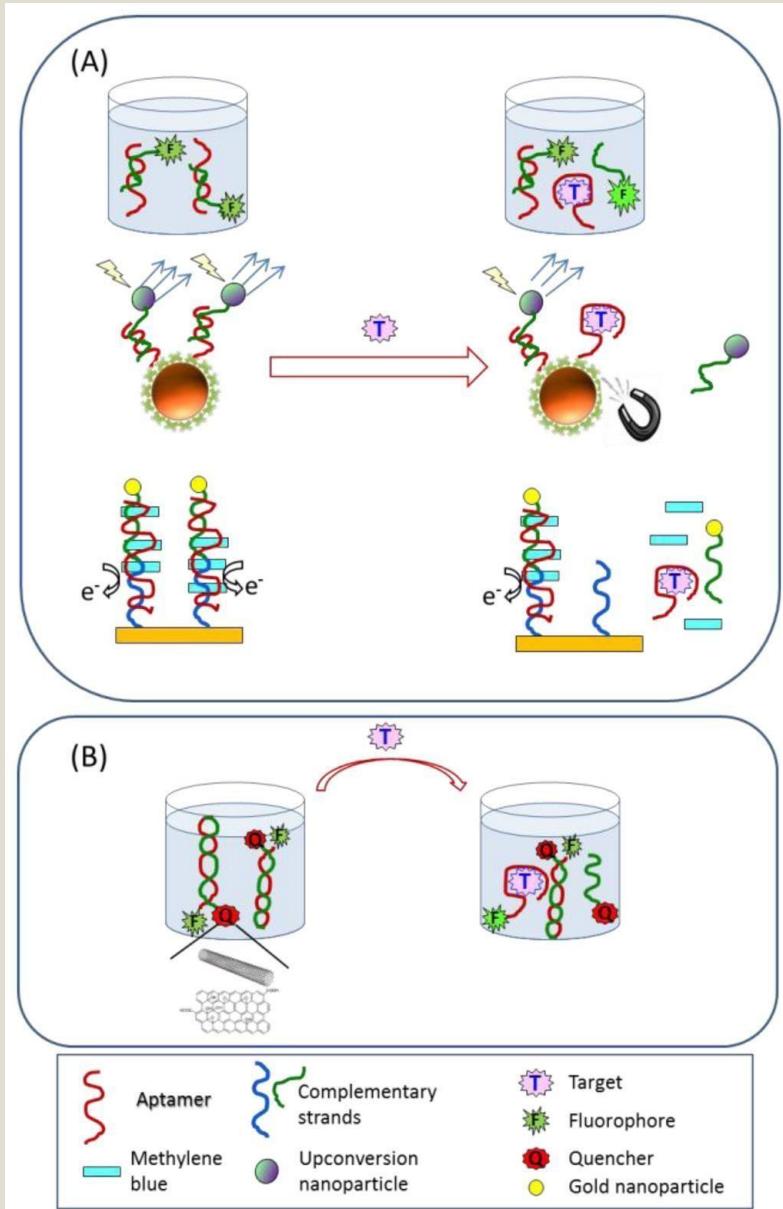




**Fig. 5.** Various signals generated by aptasensor based on structure-switching designs. (A) A schematic representation of the fluorescent aptasensor for thrombin assay. Thrombin-induced structure change of the aptamer from quenching-state into G-quartet structure could lead to fluorescence enhancement. **Fig. 5A** adapted from ref. [100]; (B) A schematic representation of the electrochemical aptasensor for thrombin assay. Before adding the thrombin, MB covalently labeled onto aptamer could transfer electron with the electrode surface due to the flexible conformation of the aptamer. Upon adding the thrombin A, G-quadruplex structure was formed and the MB moiety was far away from the electrode surface, resulting in the electrochemical signal-off. **Fig. 5B** adapted from ref. [105]; (C) A schematic representation of the colorimetric aptasensor for adenosine assay. Gold nanoparticles are functionalized with aptamer. Addition of the adenosine results in nanoparticles linking together and aggregating, thus causing the change in color. **Fig. 5C** adapted from ref. [107]. Copyright (2007) American Chemical Society.

## Aptamer-Based Analysis: A Promising Alternative for Food Safety Control

Sonia Amaya-González, Noemí de-los-Santos-Álvarez, Arturo J. Miranda-Ordieres and María Jesús Lobo-Castañón



[Check for updates](#)

Cite this: *Mater. Adv.*, 2020,  
1, 2663

## Detection and beyond: challenges and advances in aptamer-based biosensors

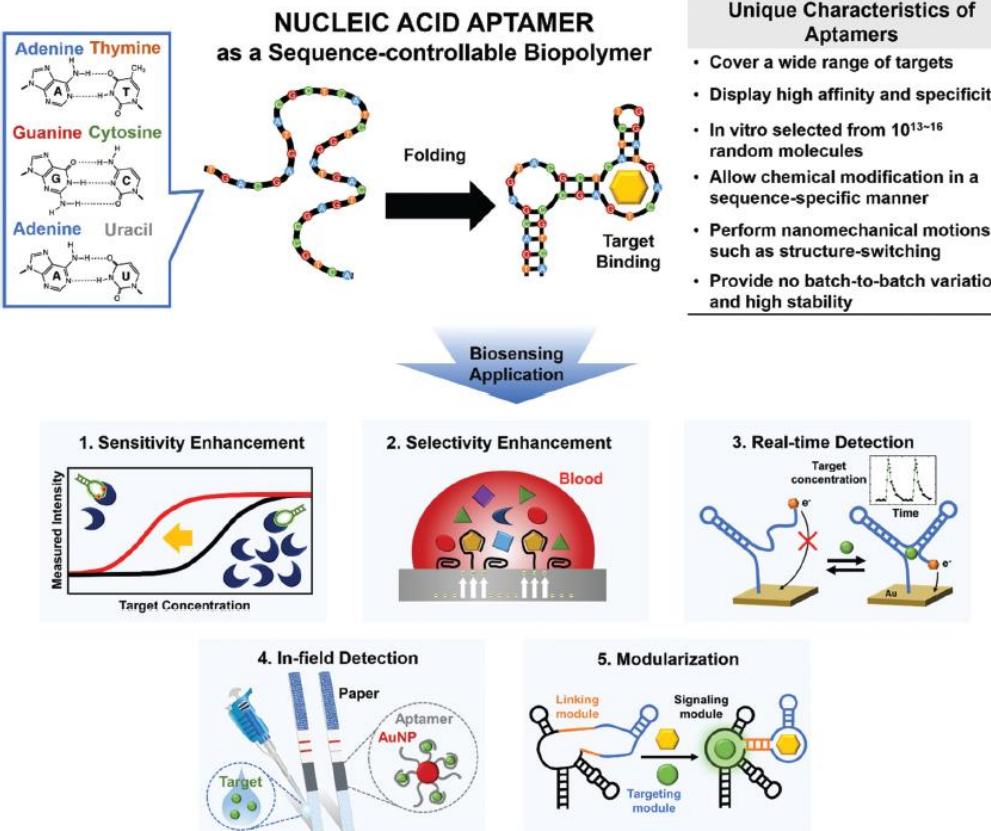
Hyebin Yoo,<sup>†<sup>a</sup></sup> Hyesung Jo<sup>†<sup>a</sup></sup> and Seung Soo Oh  <sup>\*,ab</sup>

Beyond traditional needs of biosensors such as high sensitivity and selectivity for analyte detection, newly emerging requirements including a real-time detection ability and in-field applicability have been gradually emphasized to address clinical and environmental availability. Highly programmable, synthetic aptamers that can specifically recognize a broad range of targets have the potential to fulfill these requirements; cooperative binding to target molecules achieves a significant increase in sensitivity, and binding-induced structure-switching enables target detection even in complex mixtures. Due to the availability of chemical synthesis and functional modifications, these artificial ligand materials are easily installed in many devices, and the amenability to modularization allows the aptamer-based biosensors to diversify detectable targets and signaling processes. In this review, we highlight current progress in the development of aptamer-based, next-generation biosensors including new types of field-effect transistors, electrochemical detectors, and microfluidic devices. As the nucleic acid aptamers have been rapidly generated by various *in vitro* selection techniques, the use of the versatile nanostructures is expected to expand further to include in-field and real-time biosensors.

Received 24th August 2020,  
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DOI: 10.1039/d0ma00639d

[rsc.li/materials-advances](http://rsc.li/materials-advances)



**Fig. 1** Main characteristics and functionalities of nucleic acid aptamers to overcome various limitations of conventional biosensors. Specific base pairing (top, left) folds the sequence-controllable biopolymers into thermodynamically-favored 3D nanostructures that enable molecular recognition (top, middle). The synthetic aptamers have unique features that can facilitate the development of next-generation biosensors (top, right). Here, we review technical advances in the development of aptamer-based biosensors, such as increases in sensitivity and selectivity, and actualization of newly emerging real-time and in-field detection applications, along with aptameric biosensors' interesting properties, such as amenability to modularization (bottom).

The sensitivity and selectivity can be improved using different strategies

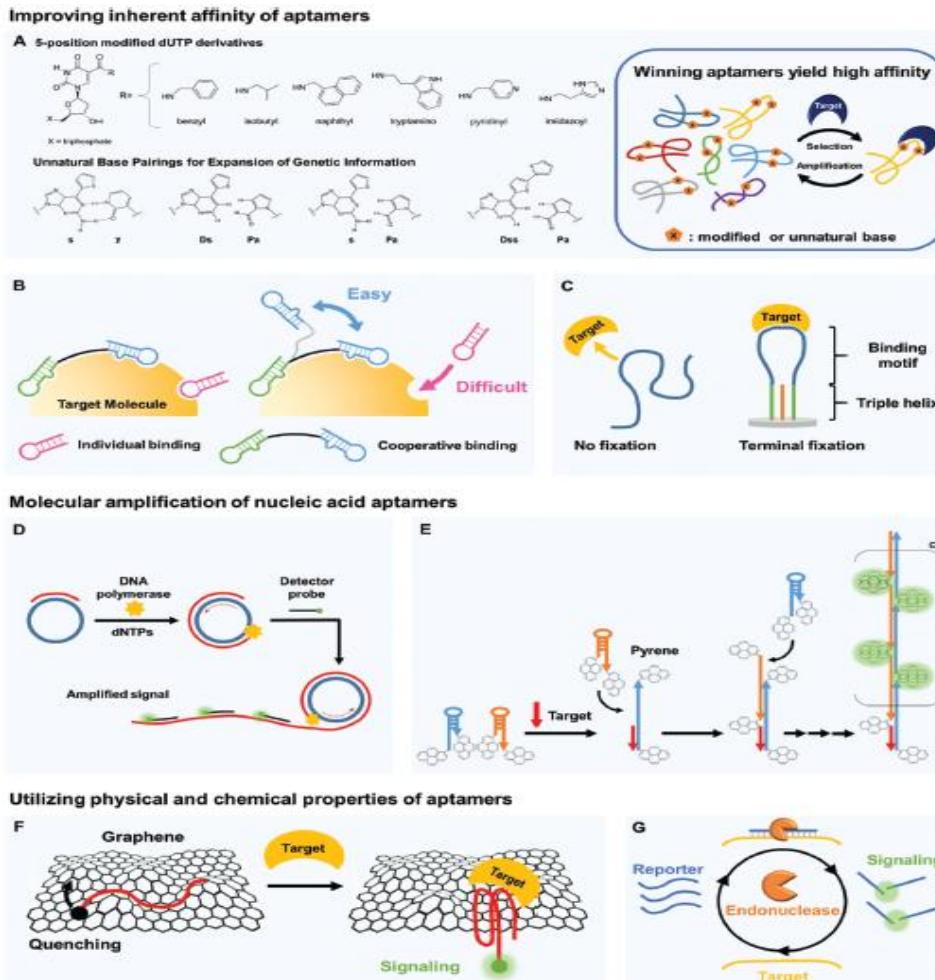
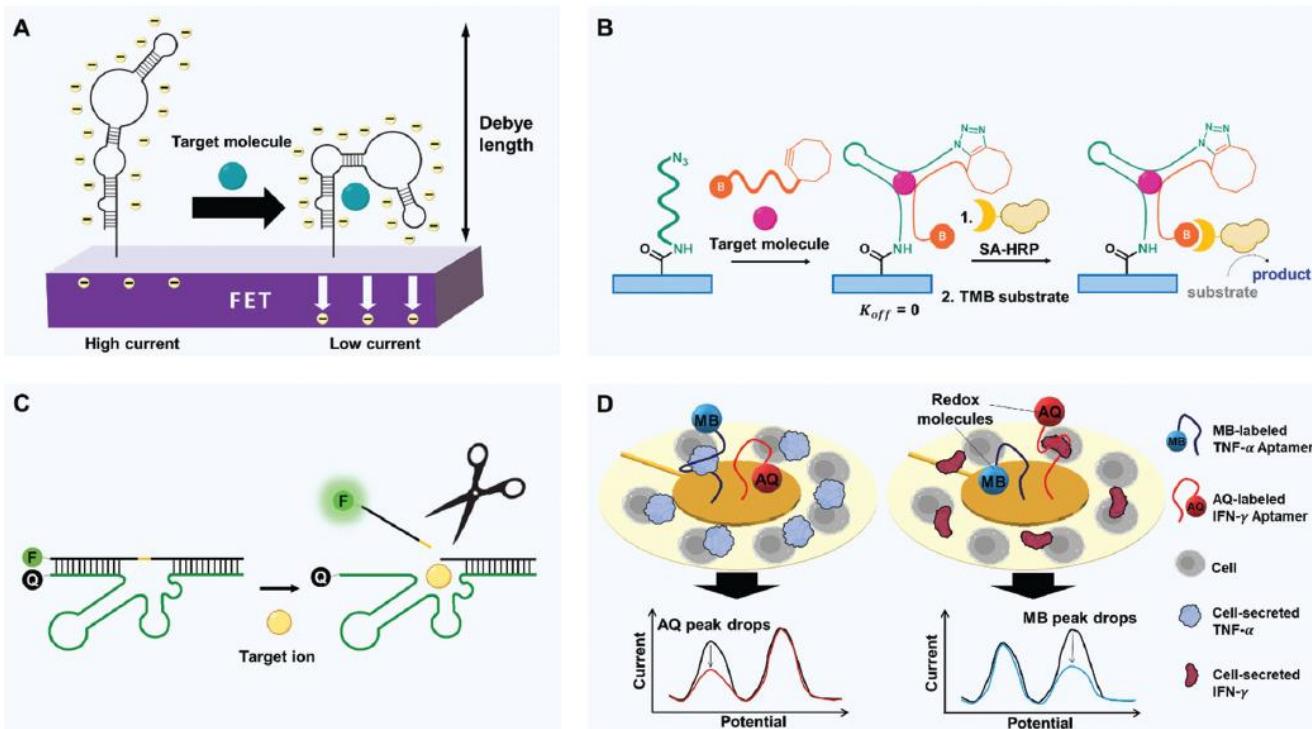
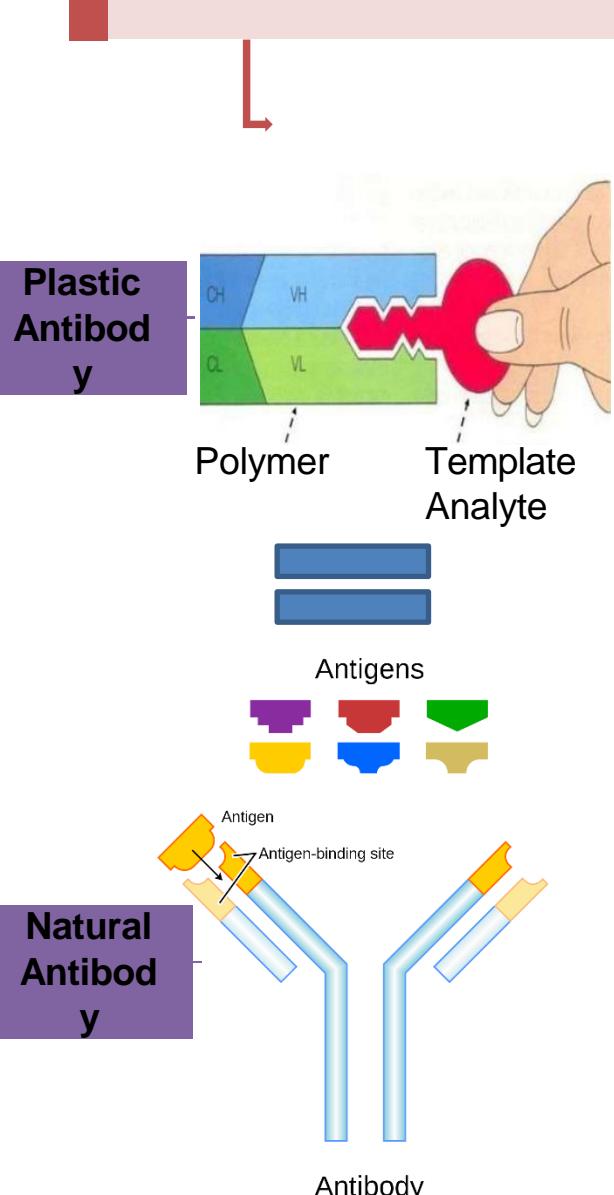


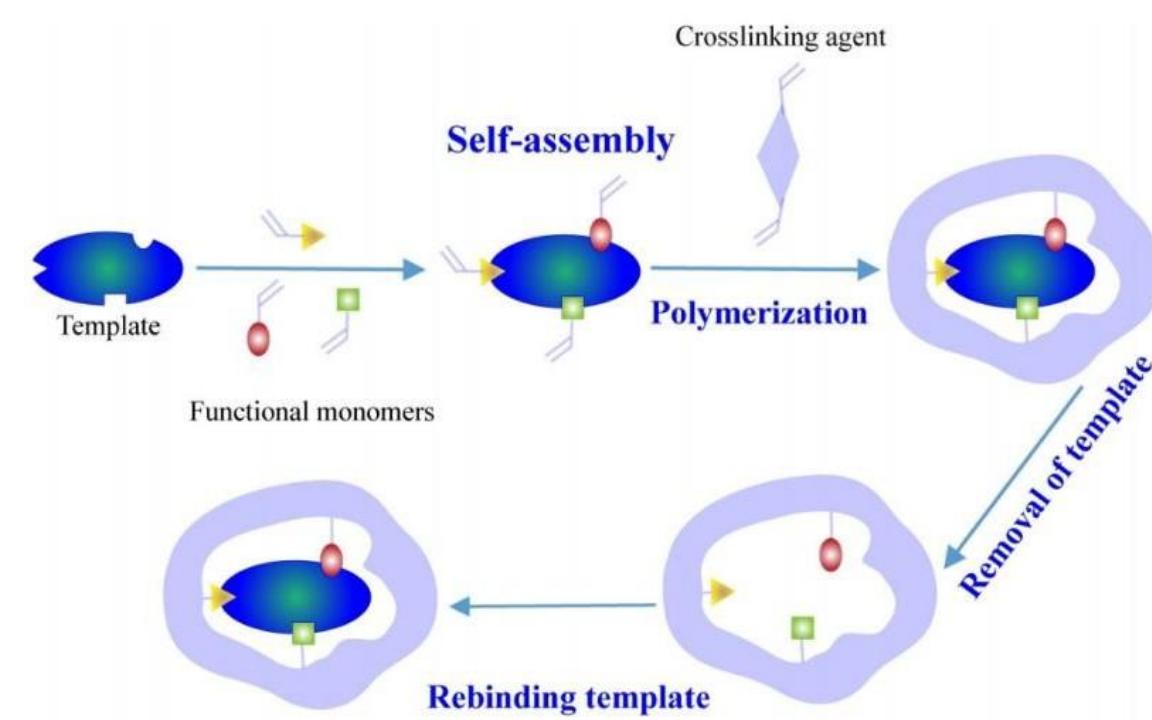
Fig. 2 Aptamers can be key components of biosensors to increase sensitivity in target detection. (A) The inherent binding affinity of aptamers can be strengthened by chemical modification of nucleic acids, e.g., by attaching hydrophobic moieties to nucleotides and by adding non-canonical pairing bases. (B) Multivalency by cooperative binding of multiple aptamers and (C) terminal fixation of folding structures can be also effective to improve the target binding capability of aptamers. (D) Molecular amplification techniques such as rolling circle amplification (RCA) can be useful to develop ultra-sensitive biosensors by significantly increasing detectable signals. In RCA, primers are bound to circular templates; polymerases extend the primers to yield long single-stranded concatemers with tandem repeat structures, and the repeated hybridization of dye-labelled strands with tandem repeats produces amplified fluorescent signals. (E) Hybridization chain reaction (HCR) can also be used to increase sensitivity. In HCR, introduction of DNA targets can trigger a hybridization cascade of signaling probes such as pyrene-conjugated hairpin probes and thereby facilitate ultra-sensitive target detection. (F) Unique physical properties of nucleic acid aptamers can contribute to highly sensitive target detection. By  $\pi-\pi$  stacking, single-stranded nucleic acids bind well to graphene surfaces, whereas the target-bound aptamers are released due to folding in tertiary structures. This folding change of aptamers yields changes in fluorescent or electrical signals, which can be easily detected. (G) Target-bound aptamers are less vulnerable to nuclease digestion than their target-free forms, and this feature can be applied to signal accumulation; the exonuclease-based, enzyme-assisted target recycling (EATR) technique can significantly decrease the limit of detection by summing fluorescence signals.



**Fig. 3** Diverse strategies with aptamers to improve selectivity of biosensors. (A) Binding-induced self-conformational change of aptamers. FETs modified with target-specific structure-switching aptamers enable selective electronic target detection. Within or near the Debye length, target-induced reorientations of stem-loop aptamers near semiconductor channels deplete the channels electrostatically and thereby decrease transconductance. (B) Binding-induced hetero-conformational change of split aptamers. Upon target binding, split aptamers can be covalently linked to each other by click chemistry. When biotinylated aptamer fragments recruit streptavidin-horseradish peroxidase (SA-HRP), chromogenic substrates such as TMB can be oxidized to emit detectable signals. (C) Binding-activated catalytic reaction. By target binding-induced cleavage of aptazymes, the release of fluorophore-linked fragments can be activated to emit highly target-specific fluorescence by reducing physisorption-derived signaling. (D) High selectivity-driven multiplexing. A multiplex analysis can be conducted by aptamers that are linked to redox molecules. Surrounded by cells, Au electrodes can be modified with different aptamers-redox reporter constructs. Binding to cytokines (TNF- $\alpha$  and IFN- $\gamma$ ) causes target-dependent conformation changes that decrease electron-transfer efficiency and thereby decrease the current. Redox molecules with different potential enable the simultaneous detection of multiple targets.


**Plastic Antibody**  
**Polymer**  
**Template Analyte**  
**Antigens**  
**Natural Antibody**  
**Antibody**

**Molecularly imprinted polymers (MIPs) are synthetic receptors for a targeted molecule. As such, they are analogues of the natural antibody–antigen systems**  
*DOI: 10.1021/acs.chemrev.8b00171 Chem. Rev. 2019, 119, 94–119*

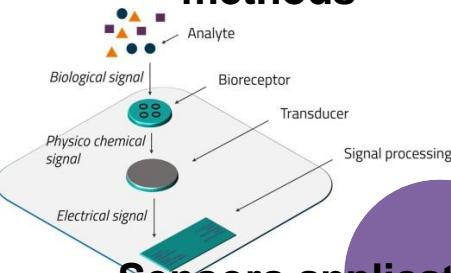

**Crosslinking agent**  
**Self-assembly**  
**Polymerization**  
**Functional monomers**  
**Removal of template**  
**Rebinding template**

**Scheme 1. Schematic representation of the synthesis of molecularly imprinted Polymers (MIPs).**  
*Abdellatif Ait Lahcen<sup>[a]</sup> and Aziz Amine<sup>\*[a]</sup>, 2018*

# MIP-State of the art

## MIPs Applications

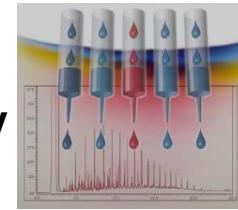
### Sample preparation in bio analytical methods



### Sensors applications

MIPs are excellent materials with high selectivity and are widely used for:

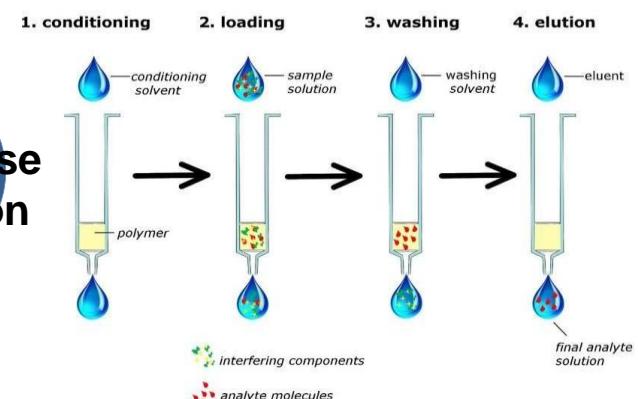
### Chromatography



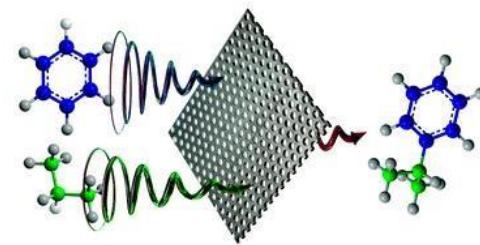
### Drug delivery



### Solid phase extraction



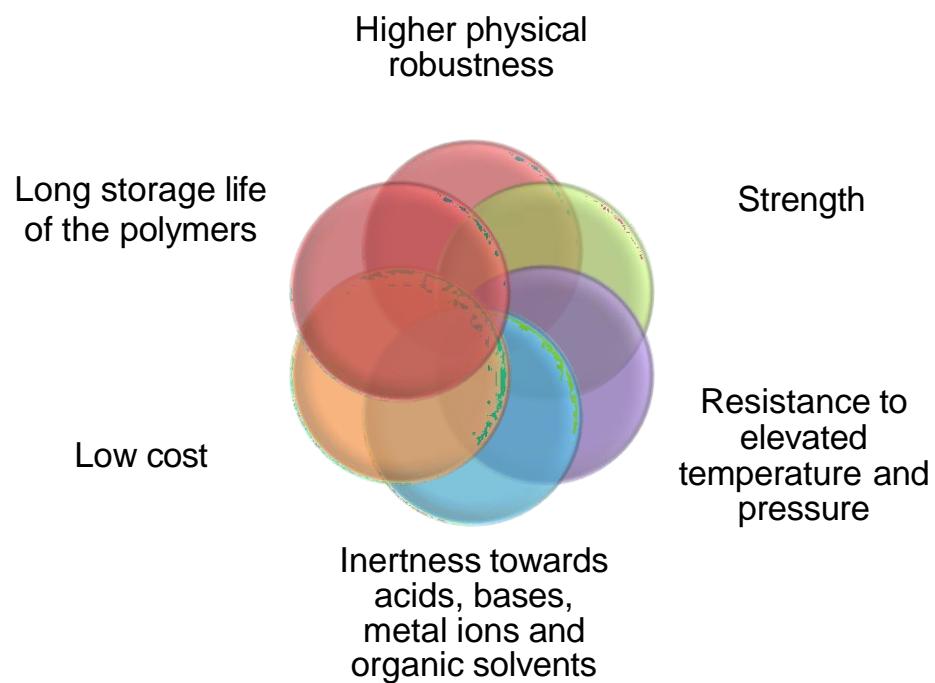
### Catalysis



## *Advantages of MIPs*

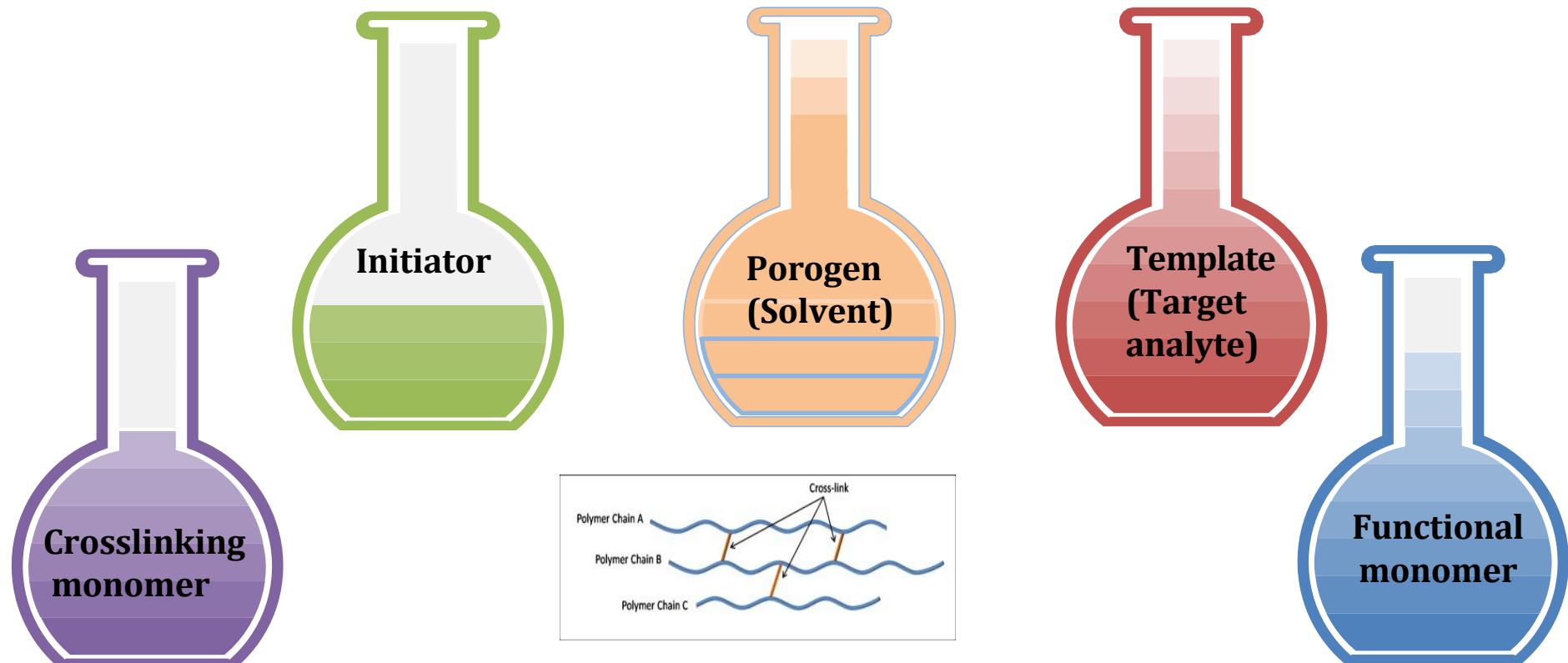
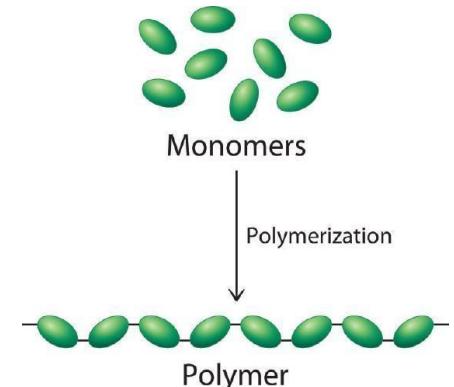
- ❖ High **selectivity** and **affinity** for the **target molecule** used in the imprinting procedure.

Compared to biological systems such as proteins and nucleic acids MIP has:



## MIPs Synthesis

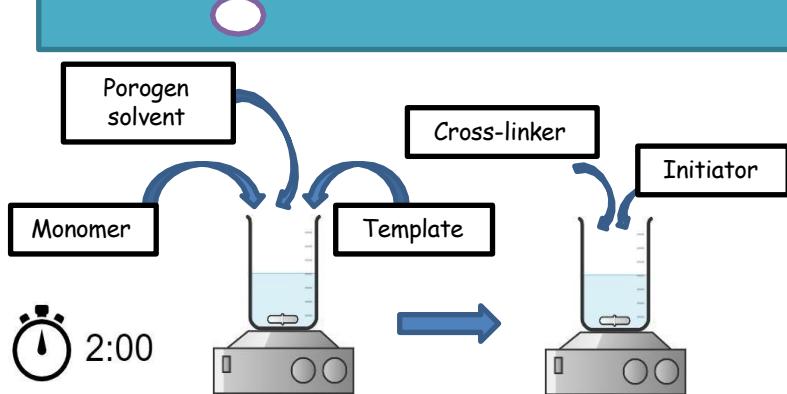
### Components of MIP Mixture



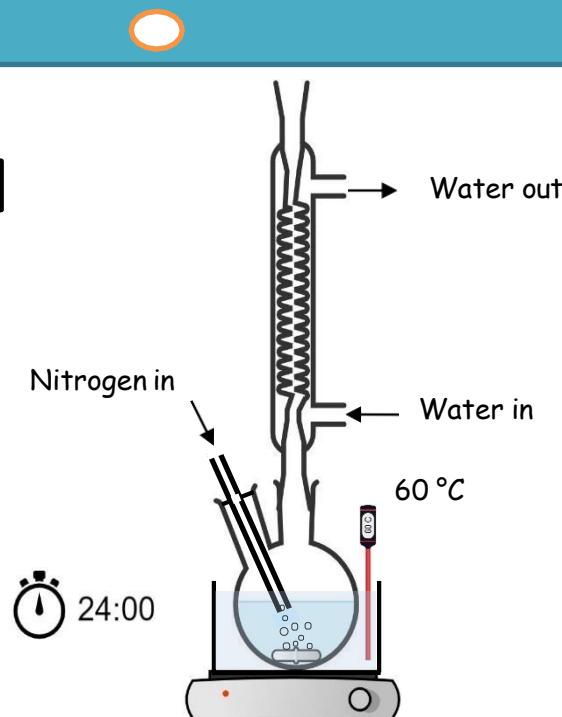
## MIPs Synthesis

### General procedure

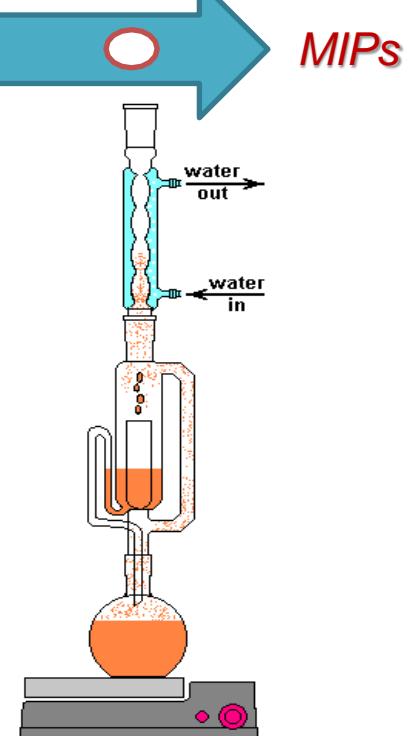
#### Self-assembly step



#### Polymerization

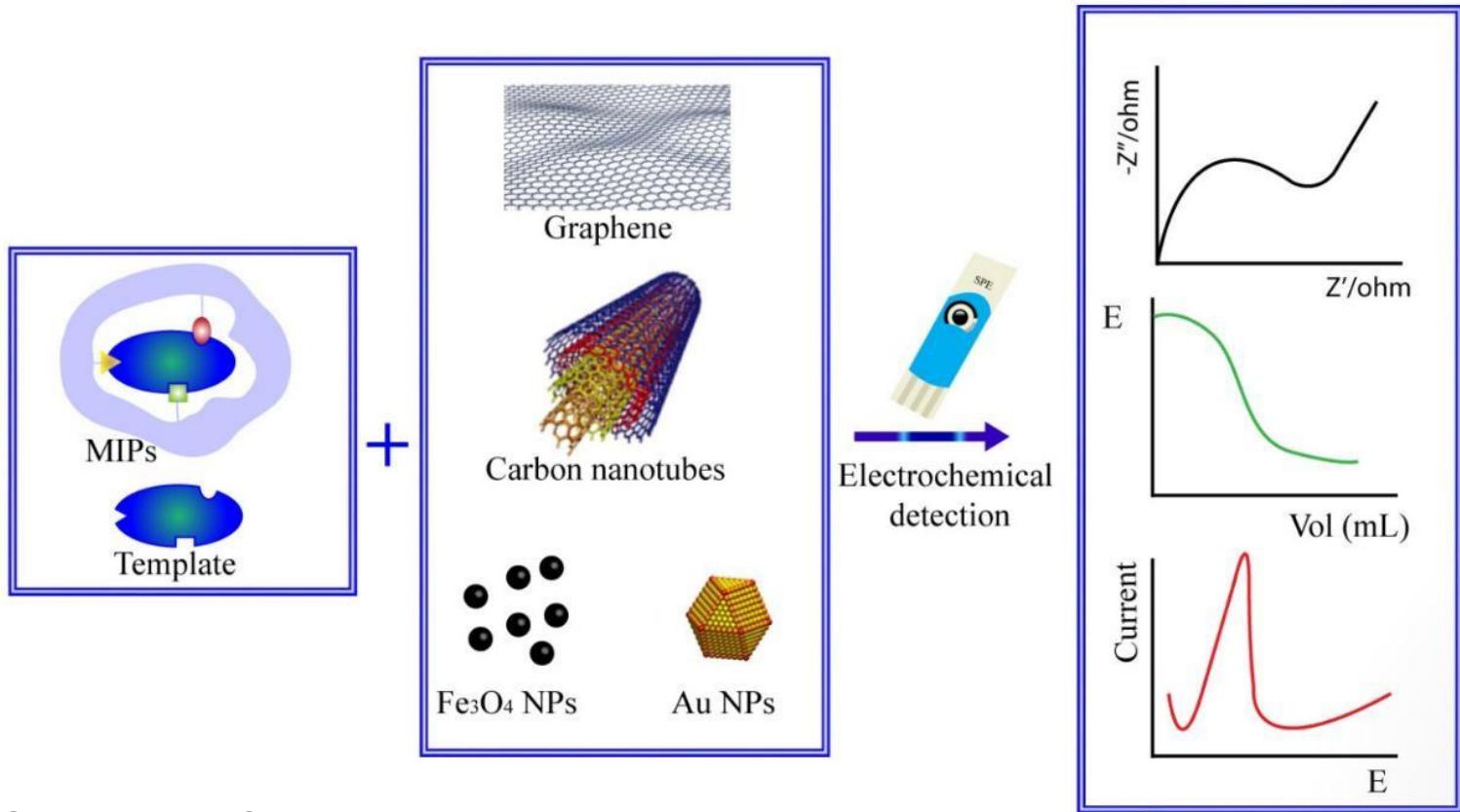


#### Extraction



Soxhlet extractor

## *MIP based electrochemical sensors and nanomaterials*

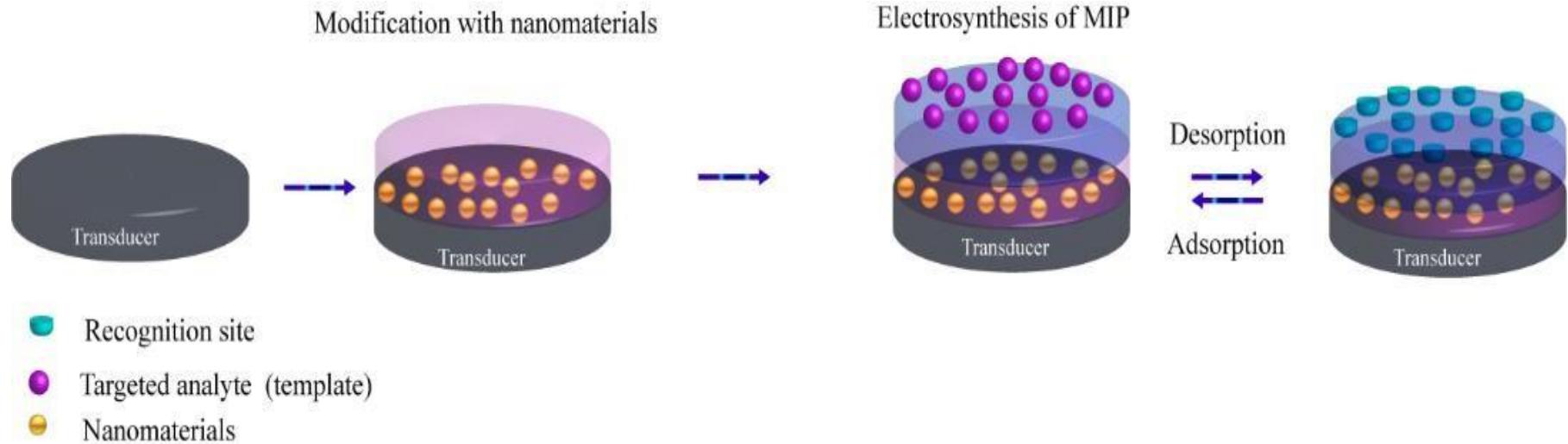


**Scheme 2.** Schematic illustration of MIP based electrochemical sensors and nanomaterials.

*Abdellatif Ait Lahcen<sup>[a]</sup> and Aziz Amine<sup>\*[a]</sup>, 2018*

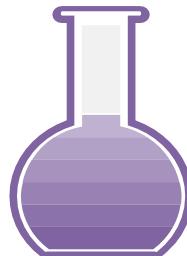
# *MIP based electrochemical sensors and nanomaterials*

## *Electrosynthesis of MIPs*

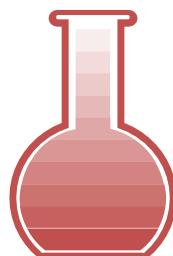


~~Initiator~~  
~~Crosslinking monomer~~

**Template  
(Target  
analyte)**



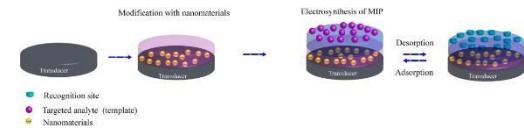
**Porogen  
(Solvent)  
Buffer**



**Functional  
monomer**



# MIP based electrochemical sensors and nanomaterials



Food  
contaminant

Template

Electrochemical  
behaviour

Non-electroactive

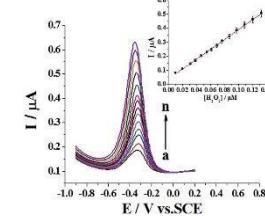
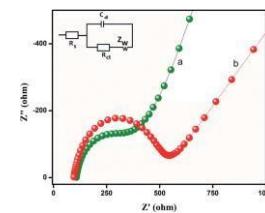
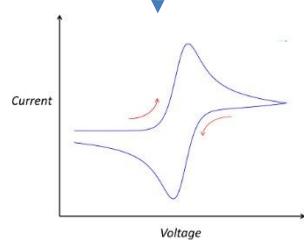
Electroactive

Detection

Voltammetric  
measurement of  
redox marker  
ferricyanide

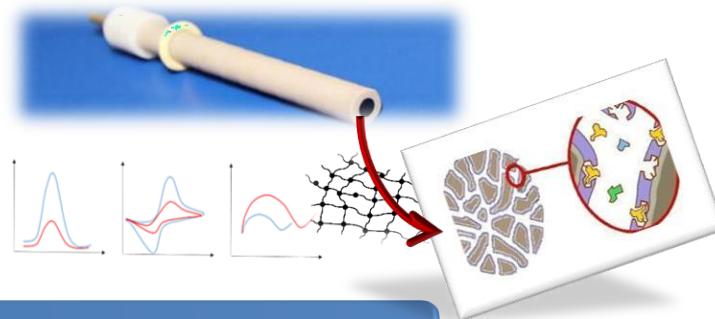
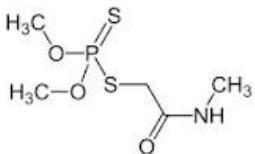
Impedance  
spectroscopy

Direct  
electrochemical  
detection



	Thermal heating	Ultrasound		Electro-polymerization
		Bath	Probe	
Complexity	medium	medium	easy	Medium
Time of synthesis	24 h	2-4 h	10 min	10 min
Temperature	60-70 °C	60-70 °C	60-70 °C	Room temperature
Synthesis of high amount	high	medium	medium	Low
Template	All templates except those sensible to high temperature such as proteins, bacterial cells,etc.	All templates except those sensible to high temperature and ultrasonic waves such as proteins, bacterial cells,etc.		It is preferred that templates are soluble in water

# MIP-MEPS based sensing strategy for the selective assay of dimethoate. Application to wheat flour samples



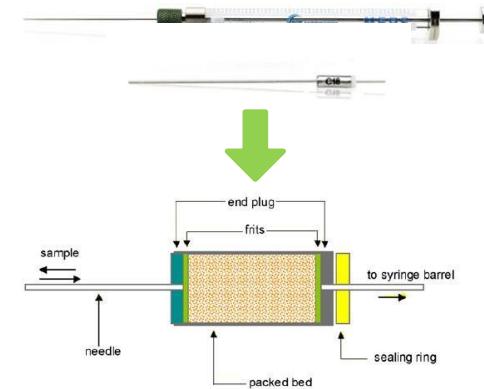
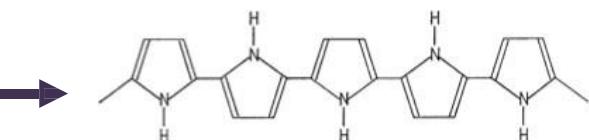
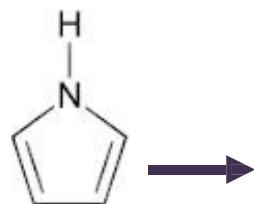
## DIMETHOATE MONITORING IN WHEAT FLOUR

### SAMPLE PREPARATION

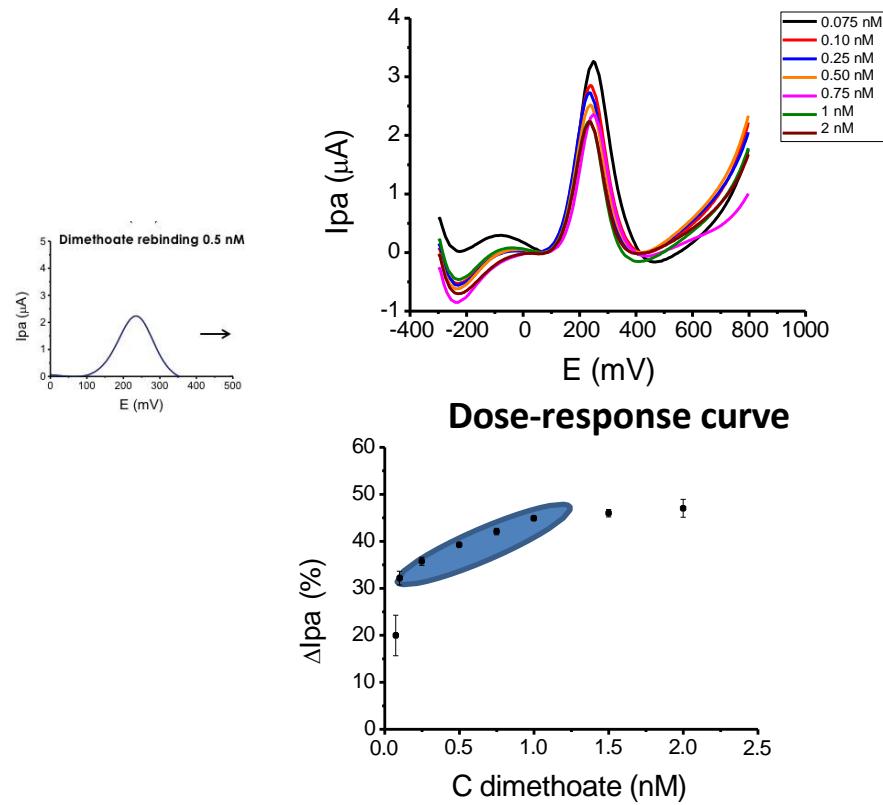
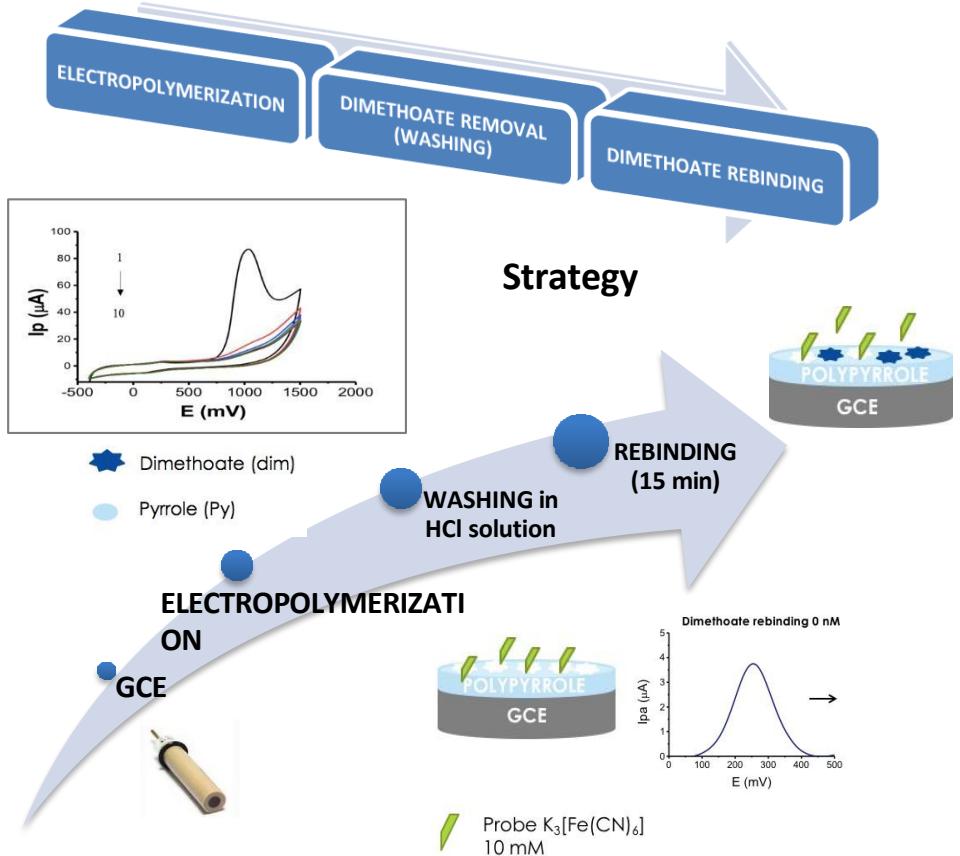
MICROEXTRACTION  
BY PACKED SORBENT  
(MEPS)

### ANALYTE DETECTION

MIP-GLASSY  
CARBON ELECTRODE

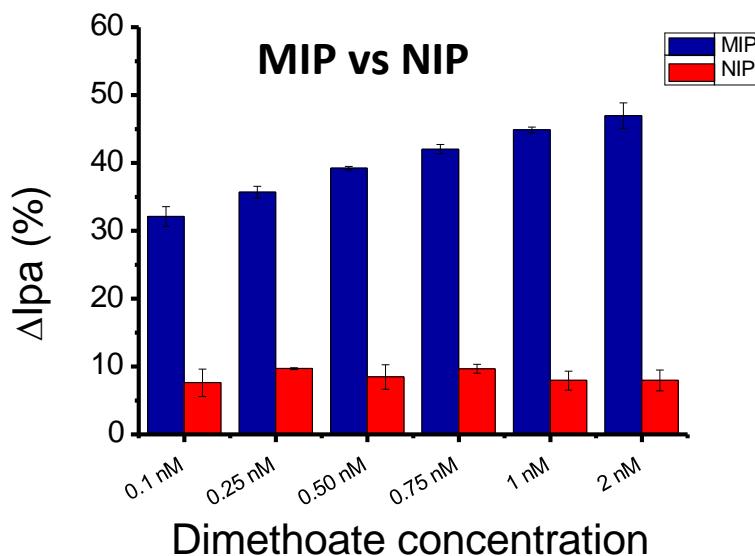


# MIP-MEPS based sensing strategy for the selective assay of dimethoate. Application to wheat flour samples

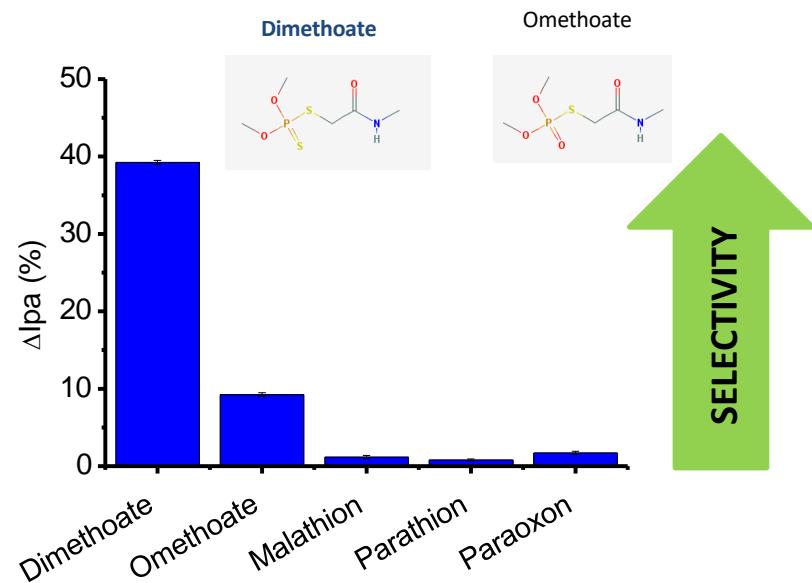


## MIP-MEPS based sensing strategy for the selective assay of dimethoate. Application to wheat flour samples

$\Delta\alpha$ (%)	Repeatability (RSD %)	Reproducibility (RSD %)
0.5 nM dimethoate (n=3)	0.68	2.72
1 nM dimethoate (n=3)	0.95	5.51



$\Delta\alpha$  (%) for malathion, parathion and paraoxon after the rebinding step was negligible; **omethoate** gave a response of **23%**.



## MIP-MEPS based sensing strategy for the selective assay of dimethoate. Application to wheat flour samples

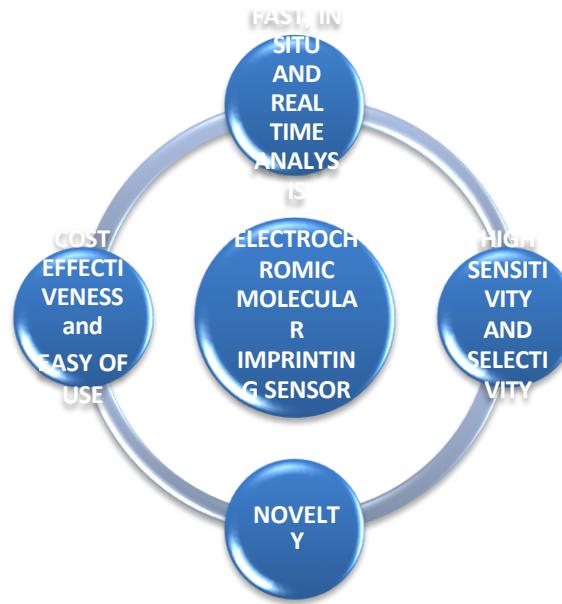
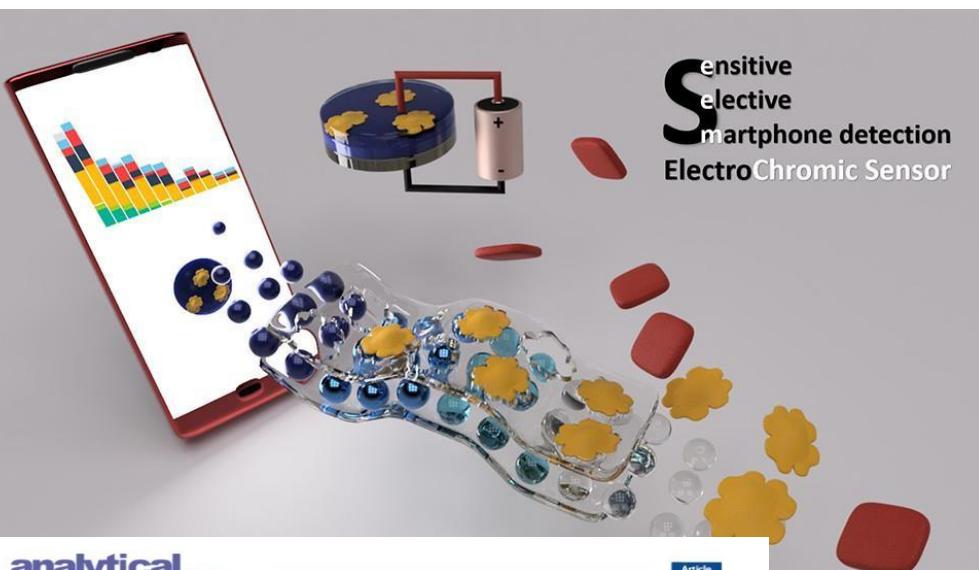
Wheat flour samples: MIP vs. UHPLC-MS/MS



Samples1	MIP-GCE	MIP-GCE
	RELATIVE ERROR (%) of dimethoate concentration ( $\mu\text{g kg}^{-1}$ )	SD of dimethoate concentration ( $\mu\text{g kg}^{-1}$ )
Wheat flour spiked with dimethoate 0.5 MRL	+13.5	0.52
Wheat flour spiked with dimethoate 0.5 MRL + mix	+4.6	2.37
Wheat flour spiked with dimethoate MRL	-21.1	1.24
Wheat flour spiked with dimethoate MRL + mix	-21.2	1.36
Wheat flour spiked with dimethoate 1.5 MRL	+16.7	0.74
Wheat flour spiked with dimethoate 1.5 MRL + mix	-0.4	1.69
Wheat flour spiked with dimethoate MRL + omethoate (1:1)	+3.5	2.70
Wheat flour spiked with dimethoate MRL + omethoate (1:10)	-15.5	0.86

# Chlorpyriphos

## Electrochromic Molecular Imprinting Sensor for Visual and Smartphone-Based Detections



### Electrochromic Molecular Imprinting Sensor for Visual and Smartphone-Based Detections

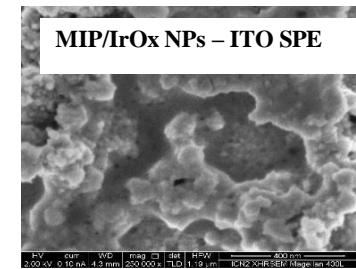
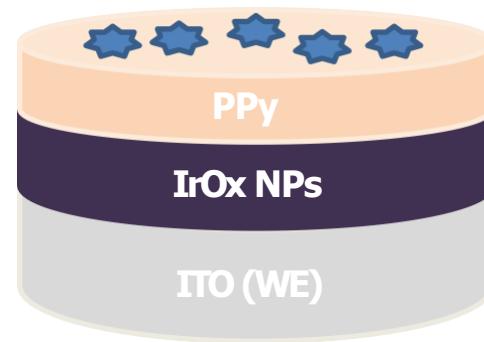
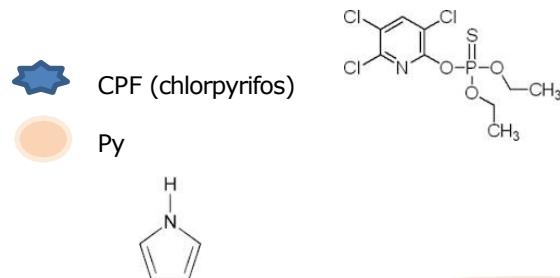
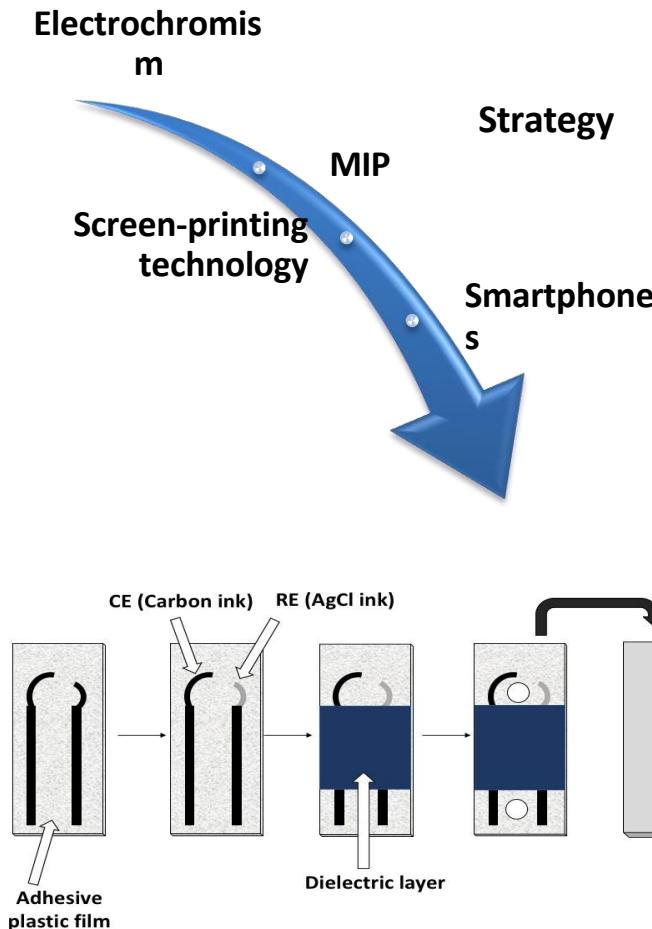
Denise Capoferrini,<sup>†,‡,§</sup> Ruslan Álvarez-Diduk,<sup>†,§</sup> Michele Del Carlo,<sup>‡</sup> Dario Compagnone,<sup>‡</sup> and Arben Merkoci<sup>\*,†,||</sup>

<sup>†</sup>Nanobioelectronics and Biosensor Group, Catalan Institute of Nanoscience and Nanotechnology (ICN2), CSIC, The Barcelona Institute of Science and Technology, Campus UAB, Bellaterra, 08193, Barcelona, Spain

<sup>‡</sup>Faculty of Biosciences and Technologies for Food, Agriculture and Environment, University of Teramo, via R. Balzarini 1, 64100 Teramo, Italy

<sup>§</sup>Catalan Institution for Research and Advanced Studies (ICREA), Pg. Lluís Companys 23, 08010 Barcelona, Spain

# Electrochromic Molecular Imprinting Sensor for Visual and Smartphone-Based Detections



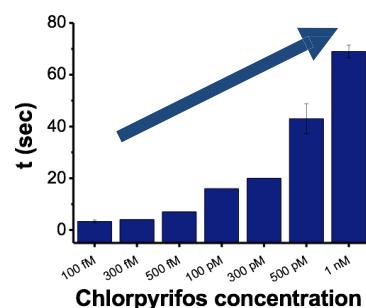
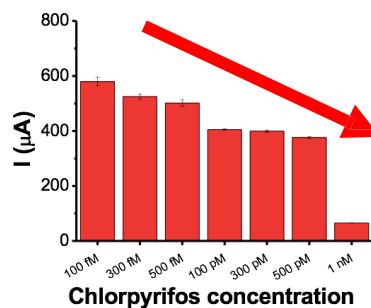
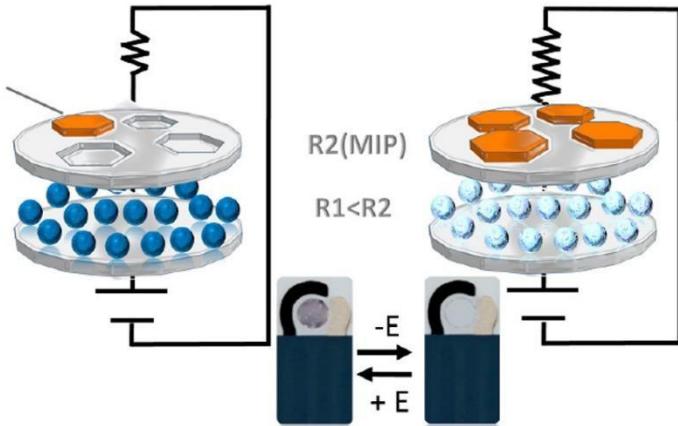
ELECTROCHROMIC MOLECULAR IMPRINTING  
SENSOR for an  
ORGANOPHOSPHATE  
PESTICIDE

# Electrochromic Molecular Imprinting Sensor for Visual and Smartphone-Based Detections

## WORKING PRINCIPLE

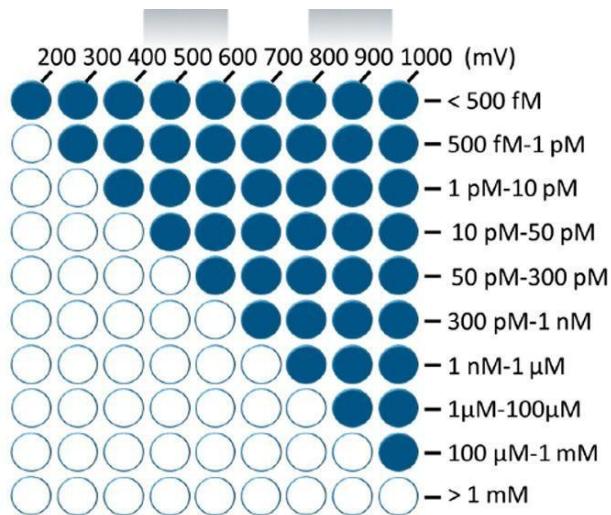
**A**

Analyte  
R1(MIP)  
IrOxNp  
ITO W.E.

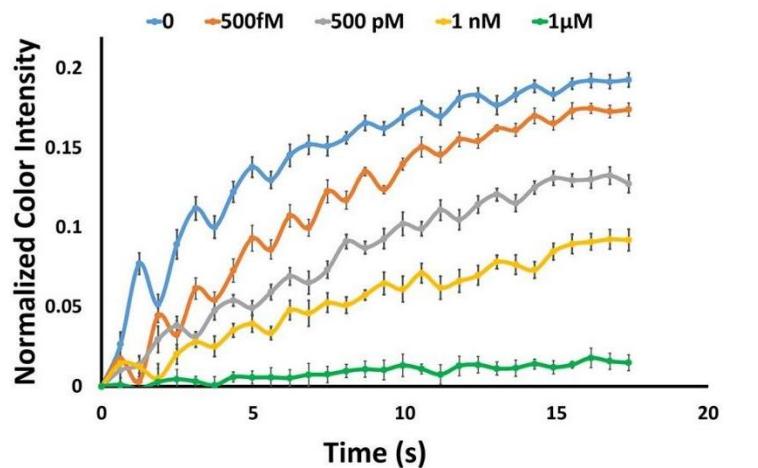


# Electrochromic Molecular Imprinting Sensor for Visual and Smartphone-Based Detections

## VISUAL APPROACH

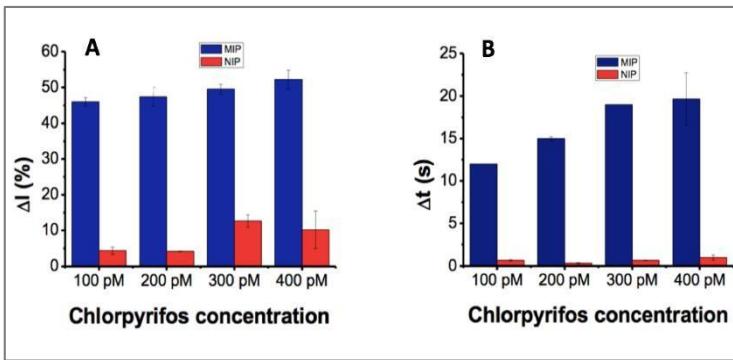


## SMARTPHONE APPROACH

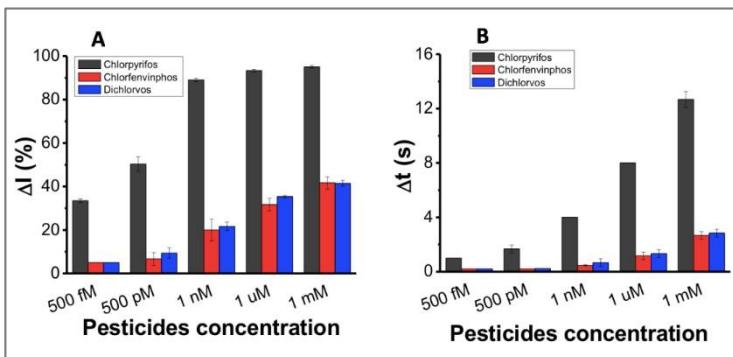


# Electrochromic Molecular Imprinting Sensor for Visual and Smartphone-Based Detections

## MIP vs NIP



## SELECTIVITY (500 mV-1000 mV)

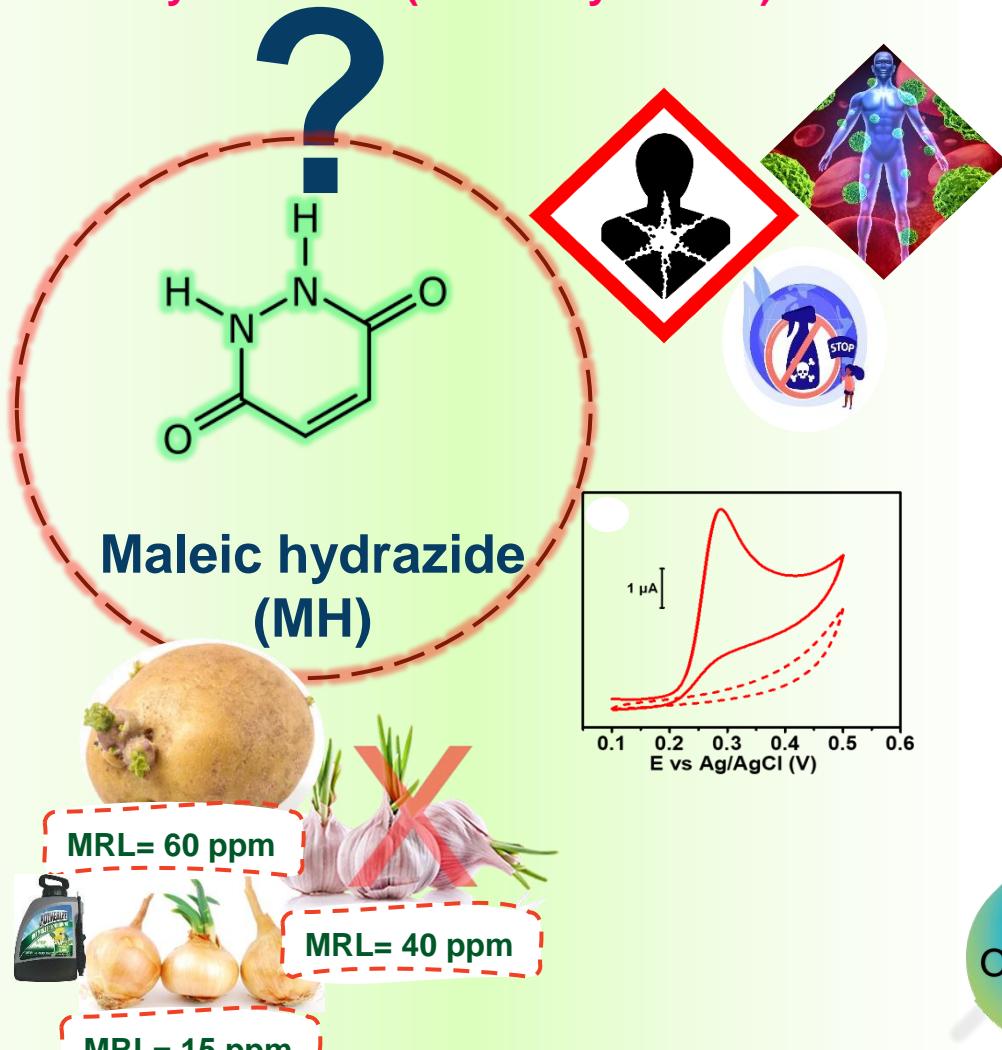


**Recovery values of chlorpyrifos in spiked drinking water samples ( $n = 3$ ) using the current response**

Added (Spiked)	Found	Recovery (%)	RSD (%)
500 fM	517.19 fM	<b><math>103.44 \pm 16.14</math></b>	15.60
500 pM	471.45 pM	<b><math>94.29 \pm 17.92</math></b>	19.00
1 nM	0.99 nM	<b><math>99.50 \pm 19.90</math></b>	20.00
1 μM	0.98 μM	<b><math>97.55 \pm 25.87</math></b>	26.52
1 mM	1.07 mM	<b><math>106.57 \pm 15.30</math></b>	14.36

# 02 MIP-Pesticides : Maleic hydrazide

## ► Why herbicide (Maleic hydrazide)?



## ► Sparking idea

**Biosensors and Bioelectronics** 24 (2009) 2323–2327

Contents lists available at [ScienceDirect](http://www.ScienceDirect.com)

**Biosensors and Bioelectronics**

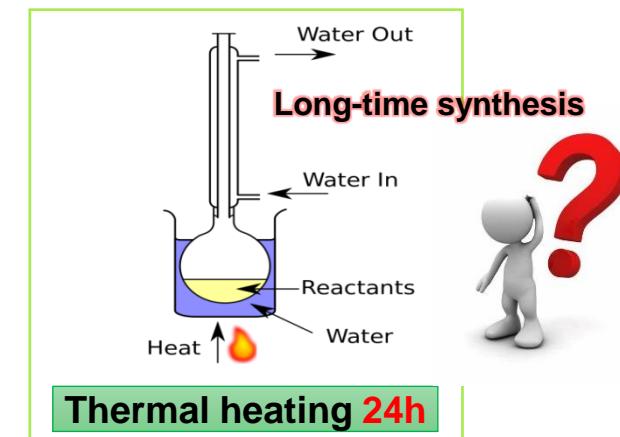
journal homepage: [www.elsevier.com/locate/bios](http://www.elsevier.com/locate/bios)

Flow injection chemiluminescence sensor using molecularly imprinted polymers as recognition element for determination of maleic hydrazide

Yanjun Fang<sup>a</sup>, Shoulei Yan<sup>b</sup>, Baoan Ning<sup>a</sup>, Nan Liu<sup>a</sup>, Zhixian Gao<sup>a,\*</sup>, Fuhuan Chao<sup>a</sup>

<sup>a</sup> Institute of Hygienic and Environmental Medicinal Science, Tianjin 300050, PR China

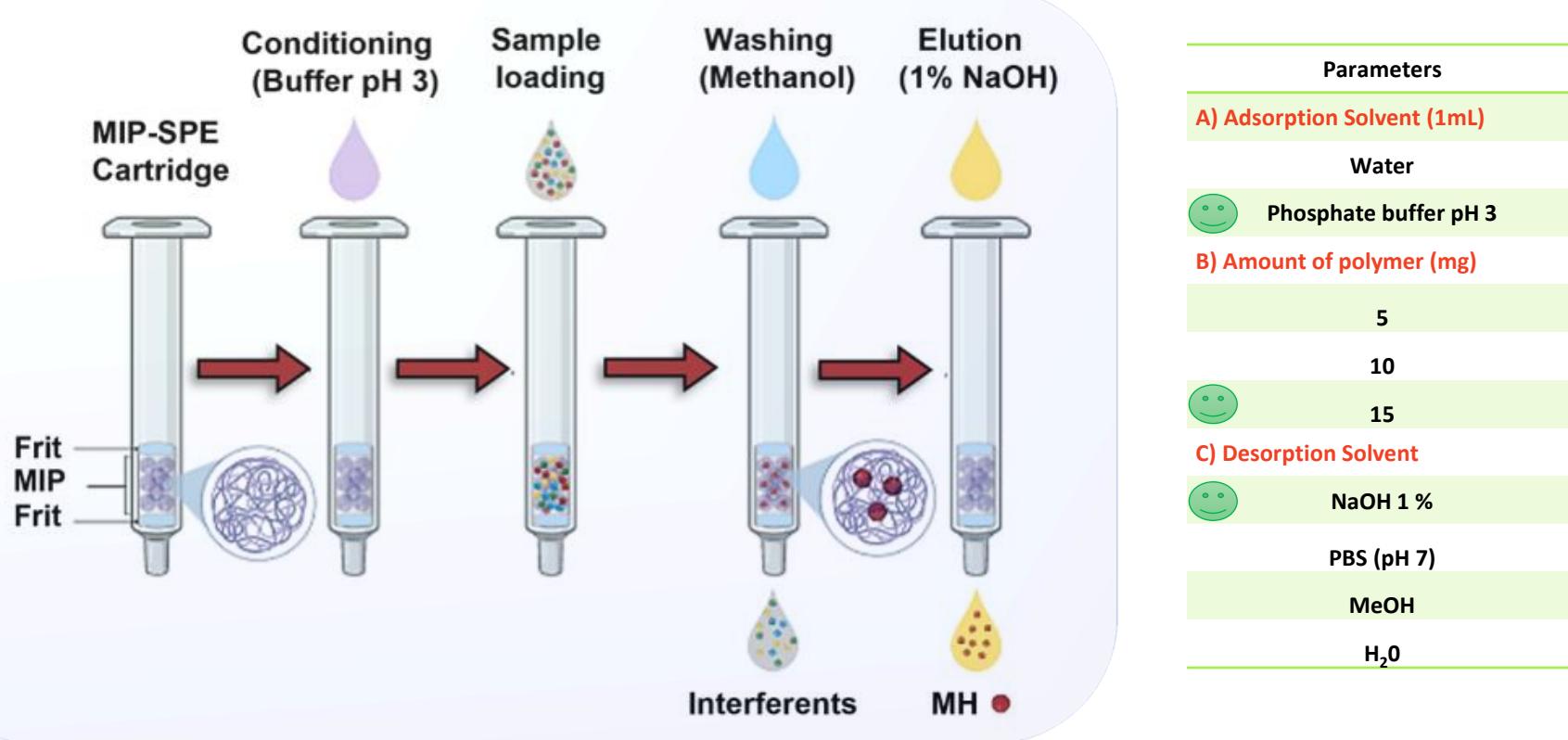
<sup>b</sup> College of Food Science and Technology Huazhong Agricultural University, Wuhan 430070, PR China



Development of a fast synthesis strategy based on a high-power ultrasound probe to synthesize effective MIPs for maleic hydrazide

## 02 MIP-MH based solid phase extraction

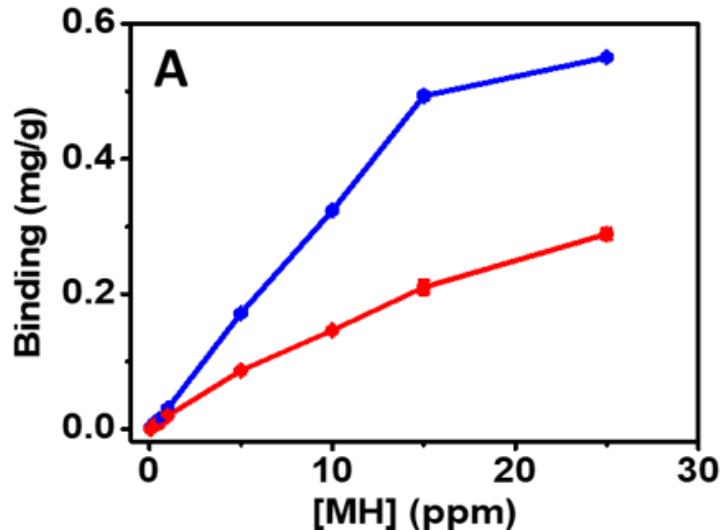
### ► Solid phase extraction



**Scheme .** Graphical scheme of the solid-phase extraction procedure based on the cartridge containing the MIP.

## 02 MIP-MH based solid phase extraction

### ► Binding capacity



$$Q = \frac{(C_i - C_f)}{m} * V$$

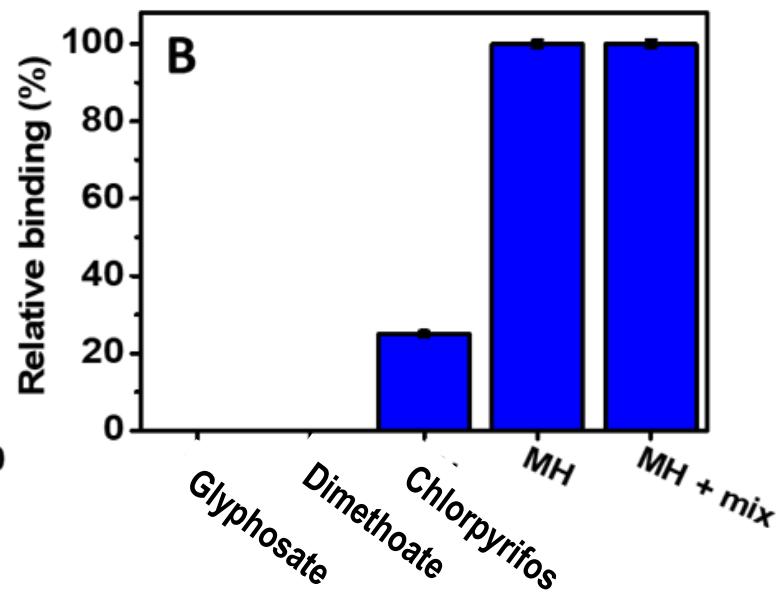
$$IF = \frac{Q(\text{MIP})}{Q(\text{NIP})}$$

$$IF = 2.30 \pm 0.02 \ (n=3)$$

-Imprinting factor (IF)

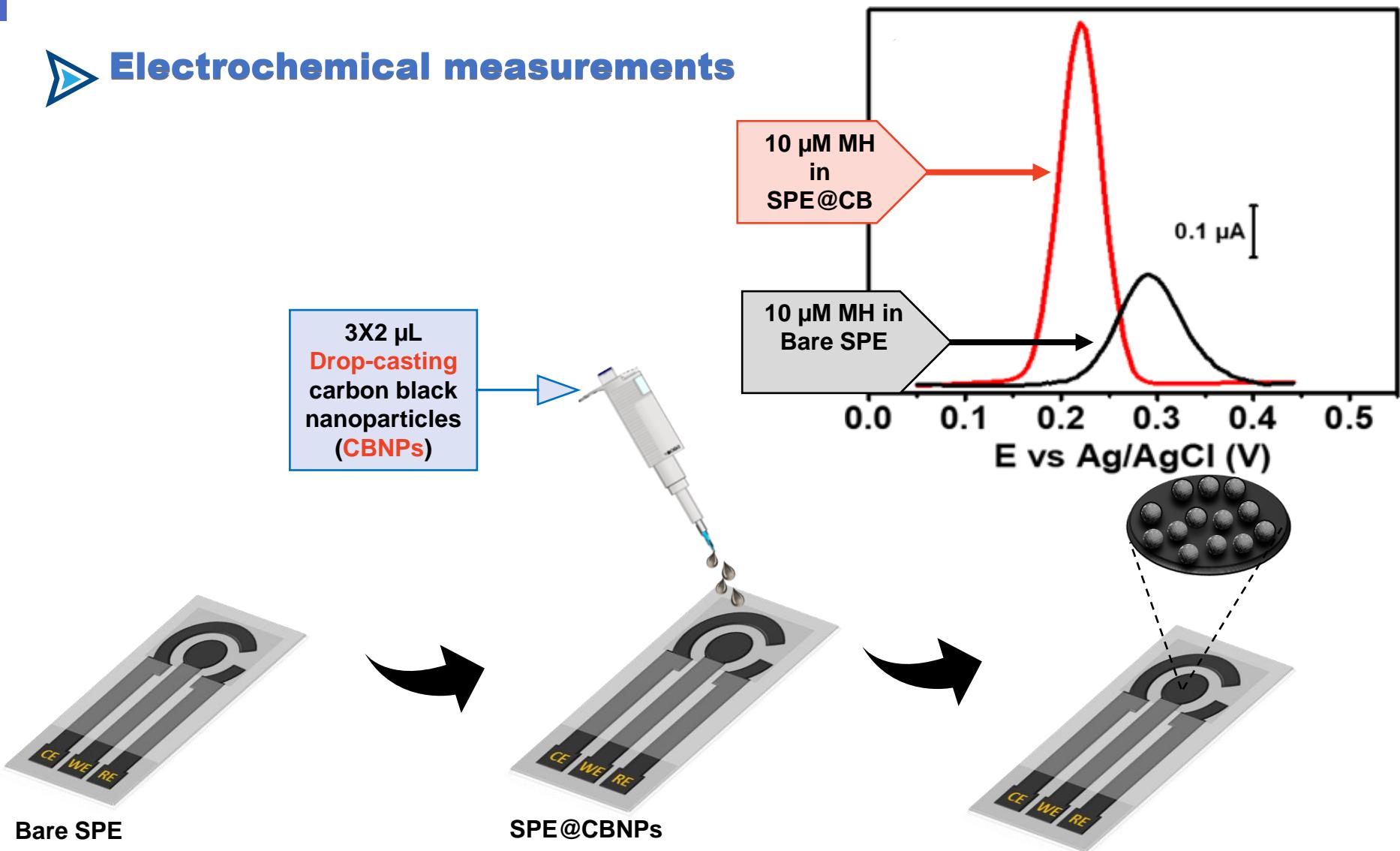
-Non-Imprinted Polymer (NIP)

### ► Selectivity



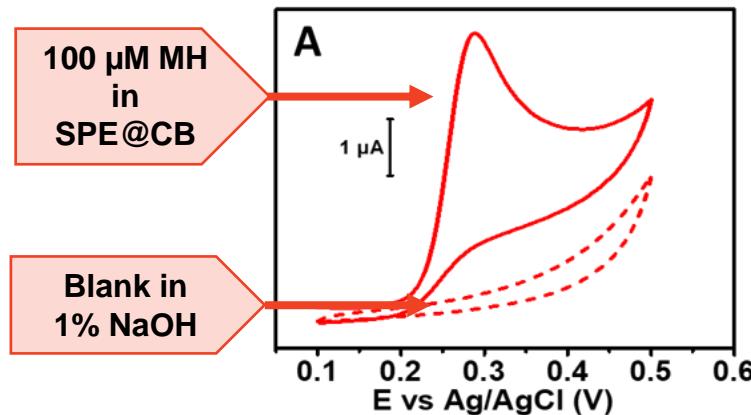
## 03 MIP-MH combined EC sensor

### ► Electrochemical measurements

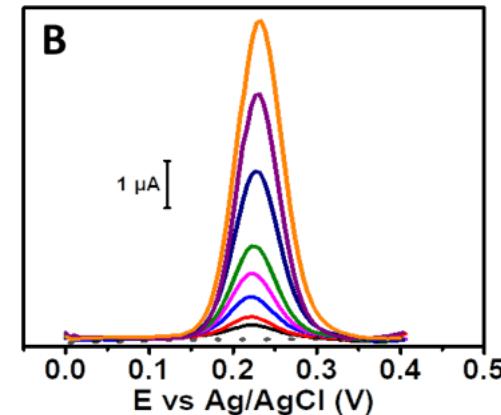


## ► Electrochemical measurements

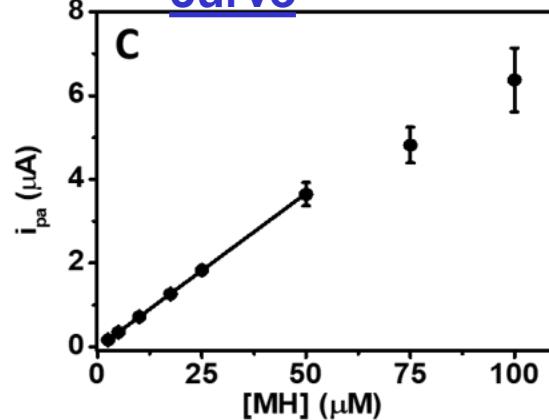
### ► Cyclic voltammetry



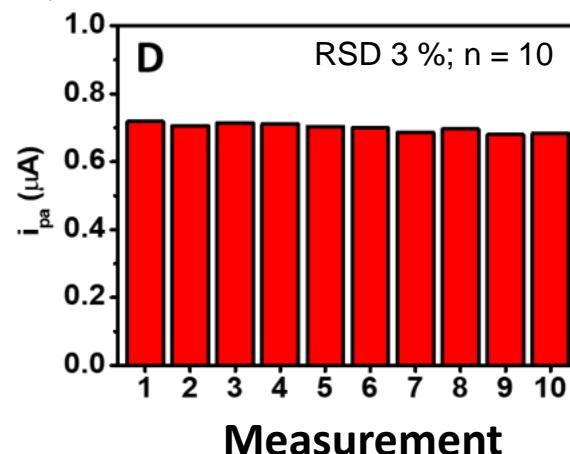
### ► DPV, 2.5-100 $\mu\text{M}$ [MH]



### ► Dose-response curve



### ► Repeatability

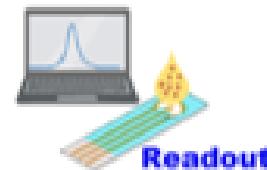


## 02 MIP-Maleic hydrazide

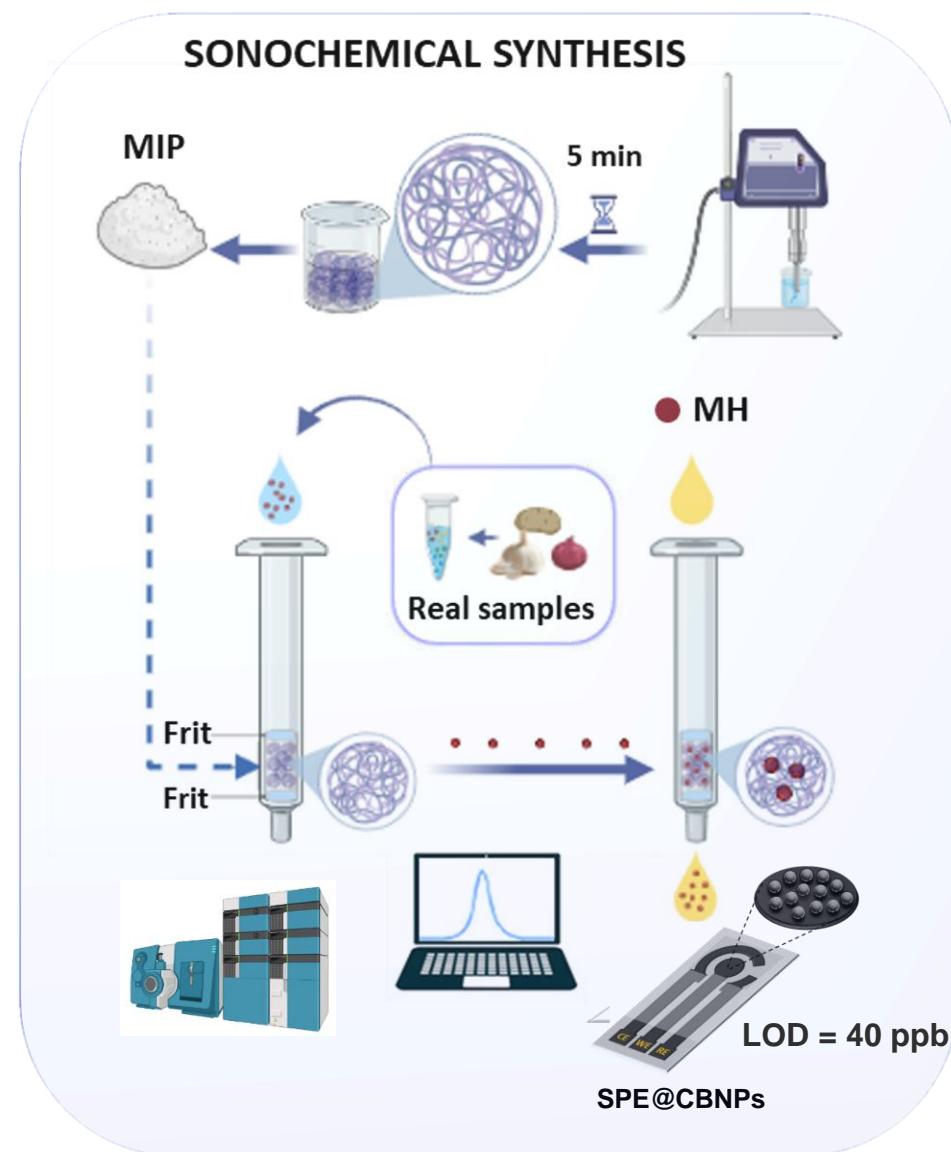
### MIP-SPE and sample analysis

**Table .** Results of MH MIPs-SPE combined with CB-based electrochemical determination applied in food samples

MH added (ppm)	EC found (ppm)	Recovery (%)	LC-MS/MS found (ppm)	Relative Error (%)
<b>Onion</b>				
5	4.6±0.2	92	4.8±0.02	-4.2
10	8.9±0.5	88.5	8.5±0.9	4.3
15	14.2±0.7	94.5	14.2±0.8	0.2
<b>Garlic</b>				
5	4.1±0.1	82.2	4.5±0.4	-8.0
10	10.5±0.4	105.1	9.6±0.3	9.4
15	15.3±0.6	102.1	14.6±0.2	5.0
<b>Potato</b>				
5	5.3±0.3	106	5.5±0.3	-3.6
10	9.7±0.3	97	10.0±0.3	-3.6
15	13±0.2	80.0	14.2±0.6	-8.5

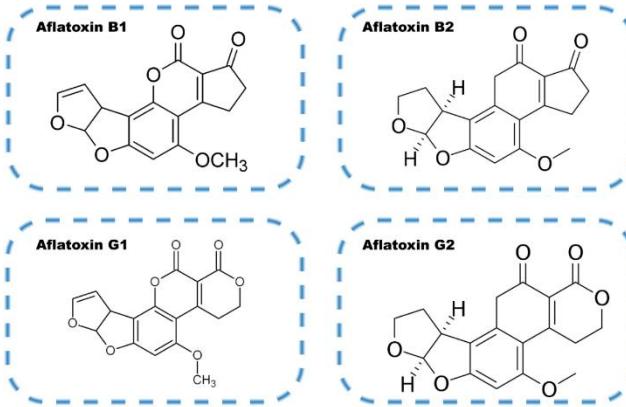


In Europe, the maximum residue limits (MRLs) for MH in **potatoes**, **garlic**, and **onions** are **60**, **40**, and **15 ppm**, respectively.



# 03 MIP-Mycotoxins : Aflatoxins

## Commercial immunoaffinity column for Aflatoxins



VS

Aflatoxins

MRL of AFG1, AFG2, AFB1, and AFB2 for processed food at **2 µg/kg**



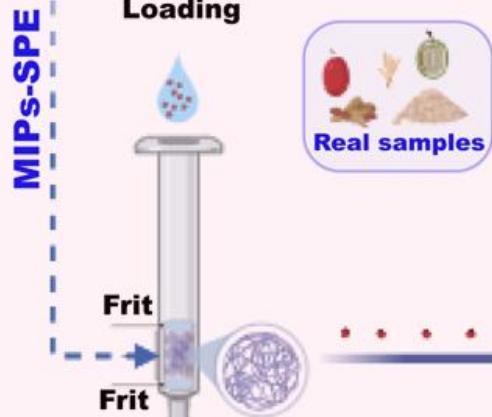
## MIPs-Aflatoxins synthesis

Dry polymer



Ultrasound Probe

Loading



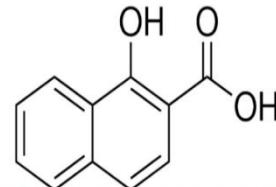
Frit

Frit

Elution

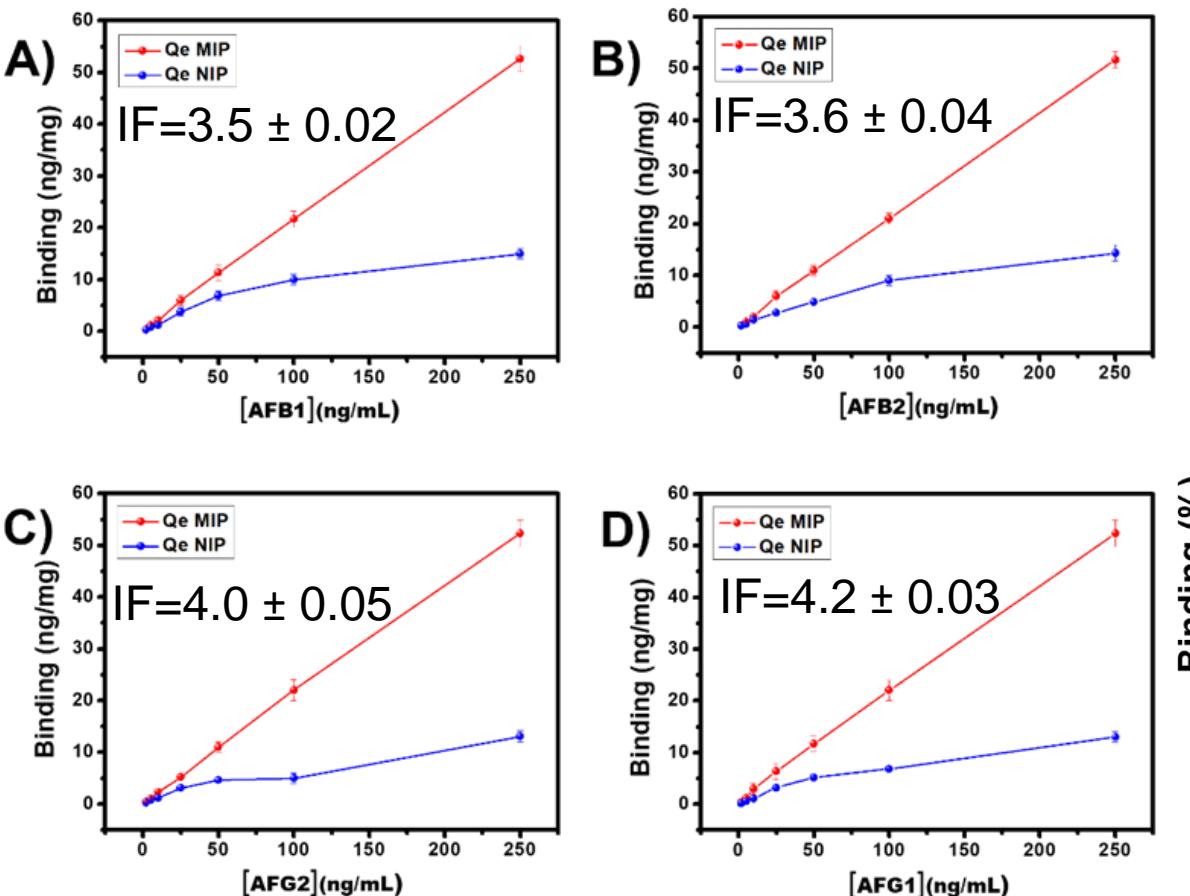
Readout

1-Hydroxy-2-naphthoic acid (Dummy Template)



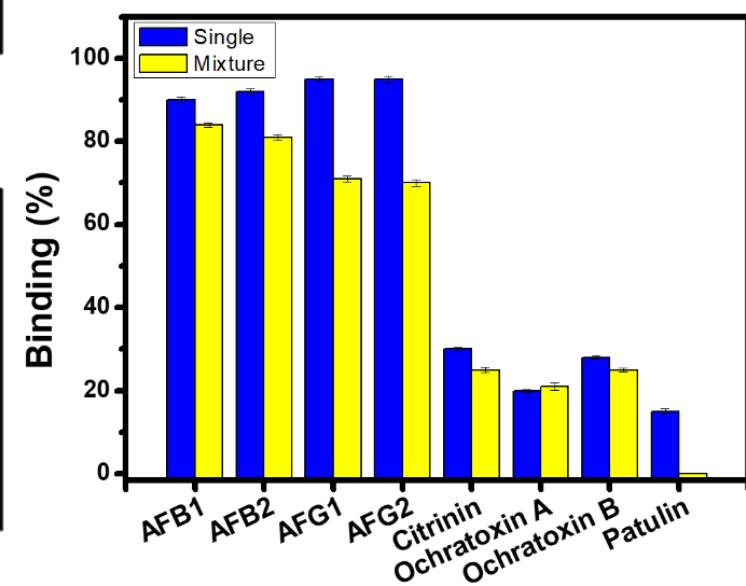
## 03 MIP-Mycotoxins : Aflatoxins

### ► Adsorption performance



**Figure.** Adsorption capacity (%) of the MIPs (red) and NIPs (blue) obtained analyzing different aflatoxin amounts (from 2 to 250 ng/mL). Graphs (A–D) correspond to aflatoxins B1, B2, G1, and G2, respectively.

### ► Selectivity



**Figure.** Selective adsorption of MIPs at a concentration of 5 ng/mL.

## 03 MIP-Mycotoxins : Aflatoxins

**Table.** Data obtained analyzing food samples using the MIPs-based and IAC-based procedures.

Sample	AFG1				AFG2				AFB1				AFB2			
	MIPs-SPE (%)		IAC (%)		MIPs-SPE (%)		IAC (%)		MIPs-SPE (%)		IAC (%)		MIPs-SPE (%)		IAC (%)	
	RC	ME	RC	ME												
Ginger	60	+9	50	-30	63	-6	48	+28	64	-1	50	+25	81	-14	52	+30
Echinacea purpurea	83	+9	73	+15	66	+10	75	+15	90	-1	80	+10	60	-8	72	+12
Ginseng	75	+16	66	-26	60	+1	46	-23	90	+7	67	-28	78	+15	61	-29
Hypericum	50	+15	50	27	60	+15	38	+30	53	+11	52	+25	72	+1	50	+30
Red elm	77	+10	64	+5	70	+7	63	+15	90	-4	80	+10	74	-6	80	+8
Saffron	61	+16	60	+12	69	+11	62	+20	60	+15	60	+17	68	+9	64	+15
Mango	67	+10	65	+15	65	+7	60	+15	65	+10	70	+15	65	+10	68	+12
Red rice	76	+11	50	+25	65	+15	55	+20	89	+3	65	+20	70	+9	66	+20
Parsley	60	+9	50	+30	60	+10	40	+30	76	+1	45	+28	60	+4	43	+19
Red fruits	60	+7	55	+20	60	+10	56	+21	79	+6	60	+18	60	+10	58	+22
Grapefruit	61	+11	65	+15	69	+11	70	+12	68	+12	70	+15	71	+4	68	+16
Magnolia	61	-7	50	+30	64	+12	50	+20	64	+15	50	+18	77	+8	54	+20
Tilia cordata	60	+1	50	+20	62	+12	45	+18	62%	+11	50	+19	72	+2	50	+20
Salsapariglia root	62	+3	55	+20	67	+6	52	+18	72	+15	50	+20	69	+14	70	+17
Hop	60	+2	50	+20	72	+12	51	+20	74	+15	60	+17	57	+12	67	+18
Verbena officinalis	72	+5	56	+18	60	+15	55	+21	69	-8	67	+18	65	+13	66	+16
Galega officinalis	78	+4	64	+15	77	+17	65	+20	73	+14	65	+20	75	+3	74	+20

**Appreciable recoveries  
(65–90%; RSD < 6%,  
n = 3)**

**low matrix effect  
(ME < 15%)**



**RC:** Recovery  
**ME:** Matrix effect  
**IAC:** Commercial immunoaffinity column