

Top-down trends in food analysis: towards nanotechnologies

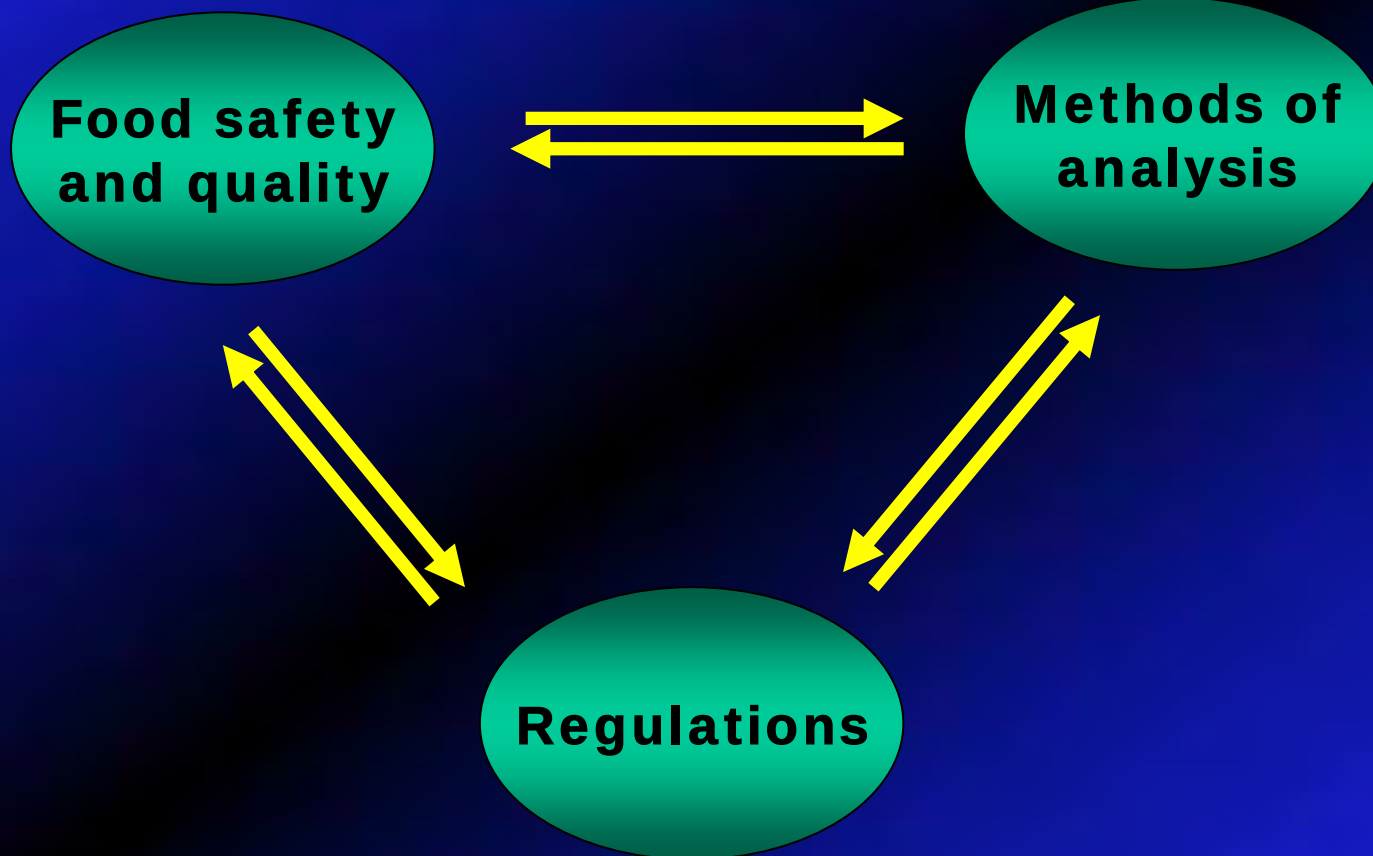
Rosangela Marchelli

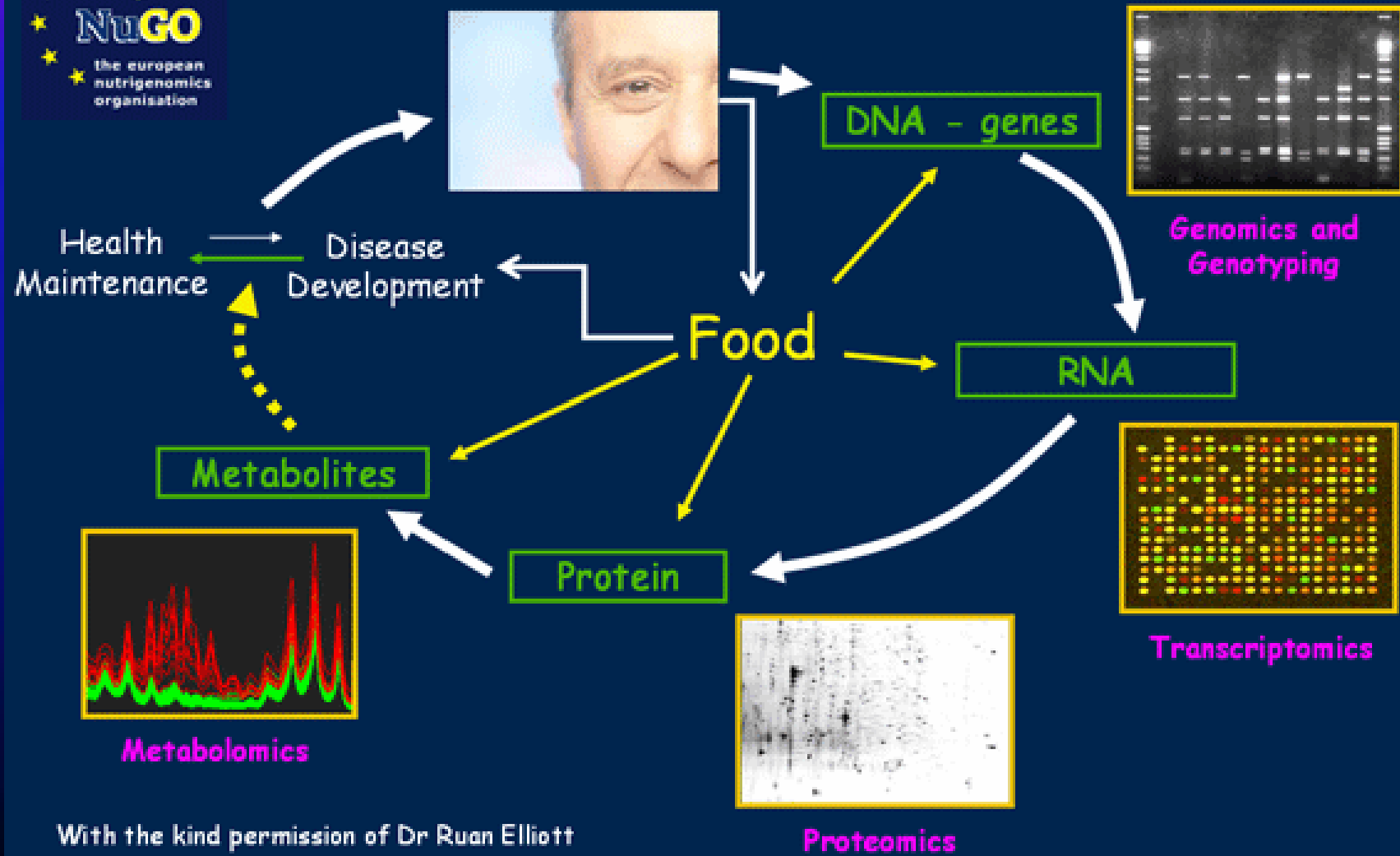
Department of Organic and Industrial Chemistry
Faculty of Agriculture
University of Parma



EU White Paper on Food Safety 2000

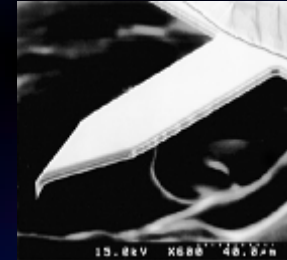
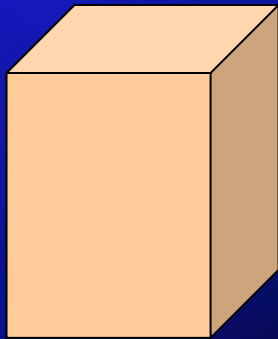
- “...there has been an enormous development, in the past decades, both in the methods of food production and processing and the controls required to ensure that acceptable safety standards are being met”





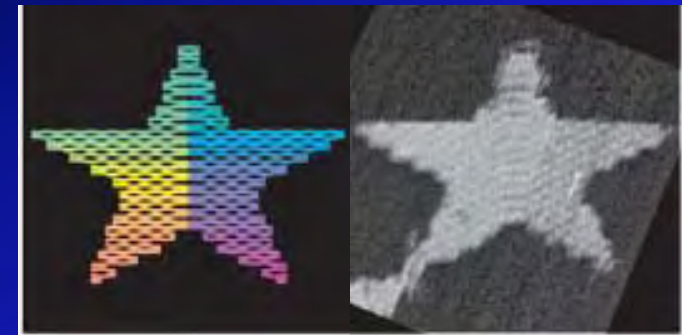
Top-down and bottom-up approaches

“Top-down” – building something by starting with a larger piece and carving away material (like a sculpture)



AFM tip, used to manipulate, image and measure atomic scale features.

“Bottom-up” – building something by putting together smaller pieces (like building a car engine).



DNA “origami”

Why is “micro and nano” good?

- Portable
- Lighter

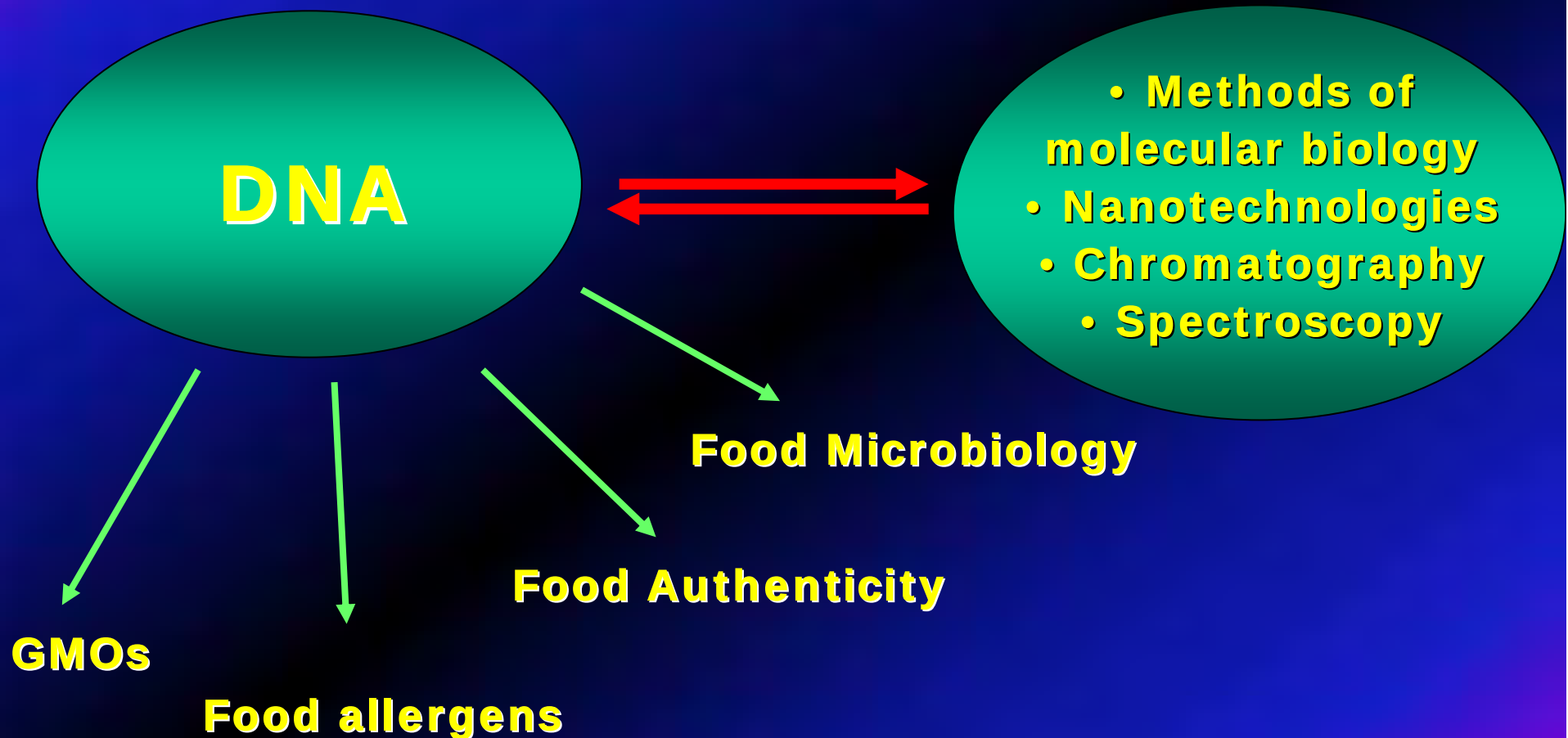


- Faster

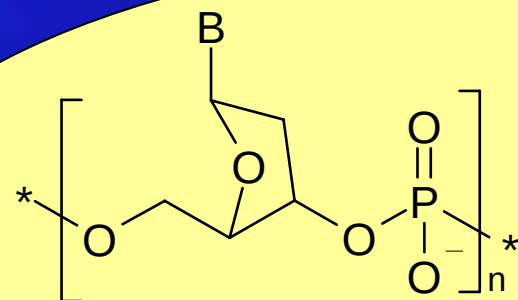


- Lower environmental impact (reagents/solvents)
- Cheaper
- More energy efficient
- Electronic reading allowing smart communication strategies
- Different properties at very small scale

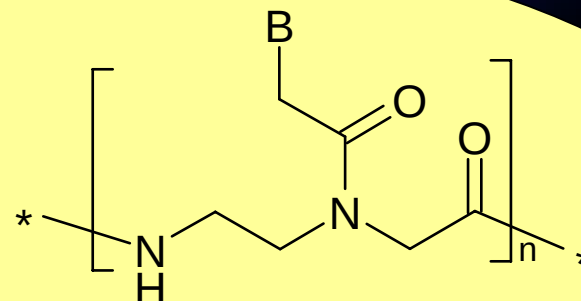
New methods based on genomics



PNA probes for food DNA detection



DNA



PNA

Microarrays

SPR

CD

CE

HPLC

**Light Up
Probes**

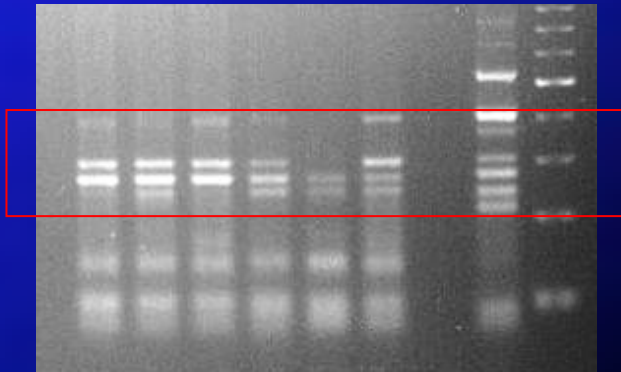
**Colorimetric
tests**

Beacons

High tech

Low tech

The microarray technology approach



**MULTIPLEX PCR
ANALYSIS**



Hybridization

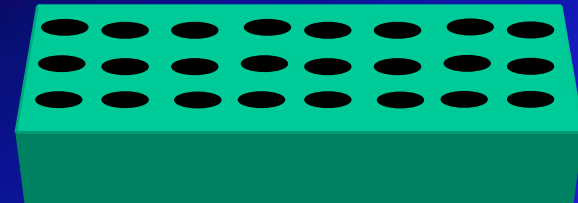


MICROARRAY

Deposition

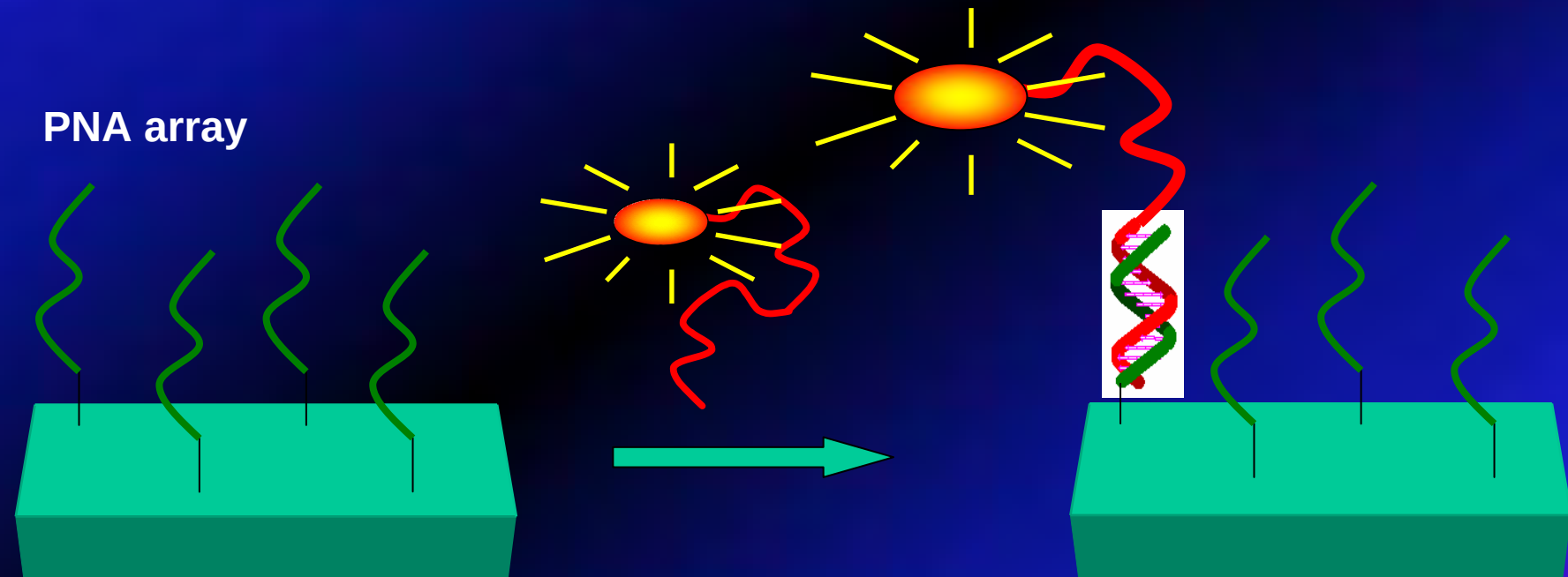


**PNA PROBE LIBRARY
DEDICATED TO FOOD**

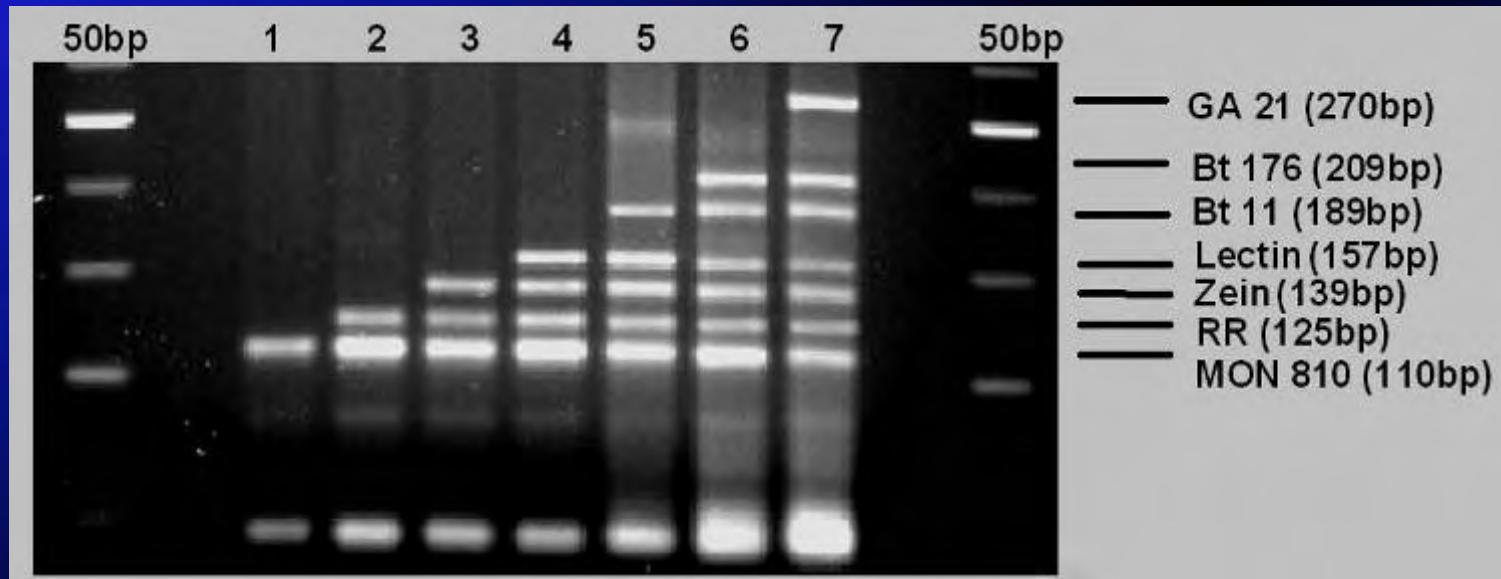


MICROARRAYS

1. Probe design and synthesis
2. Probe deposition
3. Hybridization with the target DNA

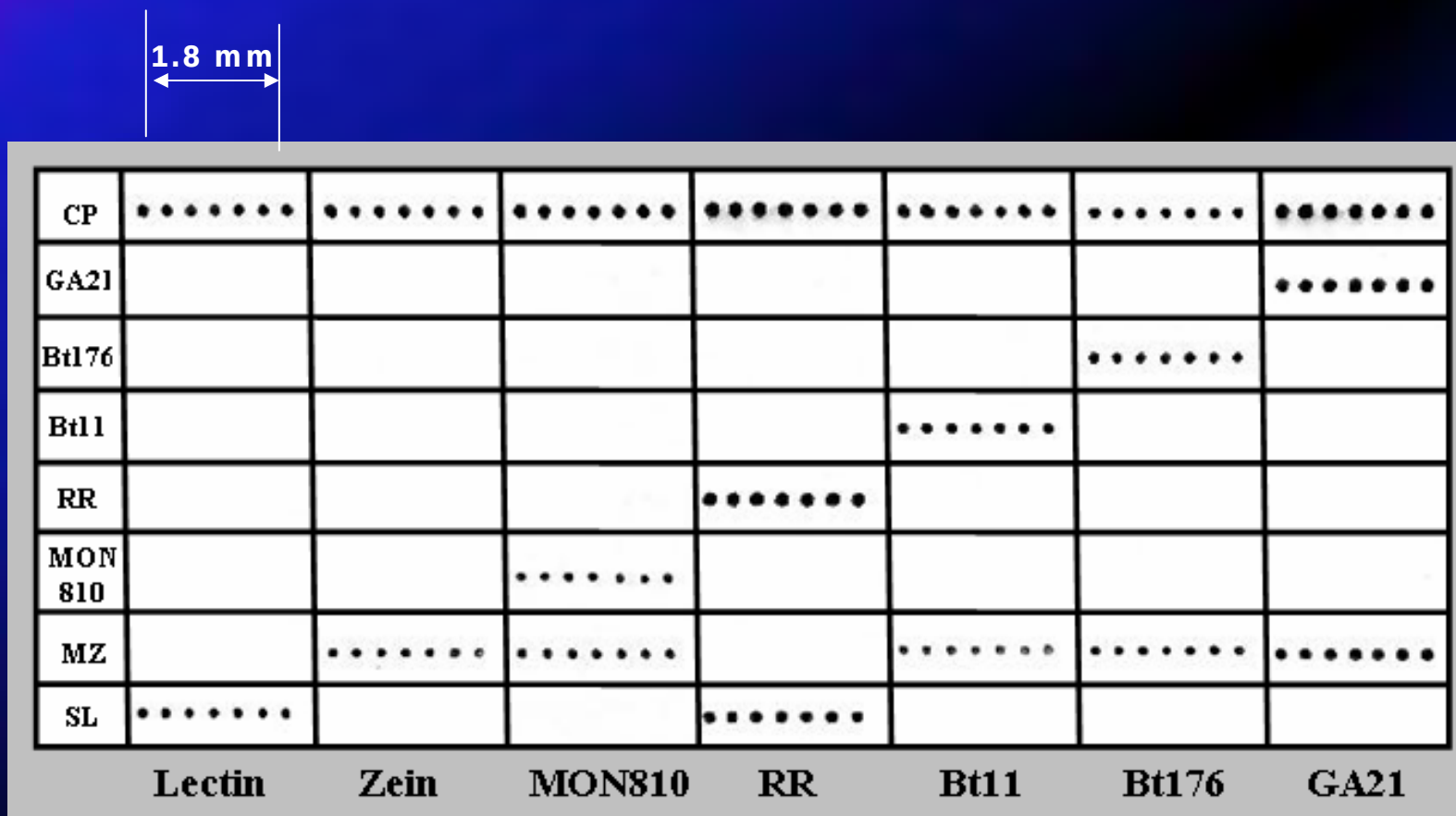


GMOs: development of a PCR multiplex



Germini A., Zanetti A., Salati C., Rossi S., Forre C., Schmid S., Fogher C., Marchelli, R.
Journal of Agricultural and Food Chemistry, 2004, 52, 3275-3280

Specificity hybridisation test

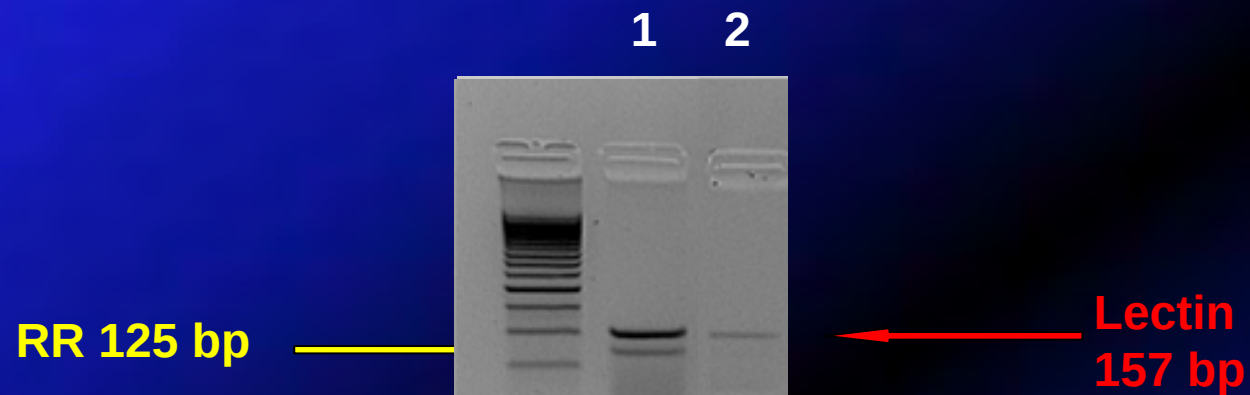


Germini A., Rossi S., Zanetti A., Corradini R., Fogher C., Marchelli R.
Journal of Agricultural and Food Chemistry, 2005, 53, 3958-3962.

In every sub-array: more than 1 thousand spots



PNA chip for RR-soy products



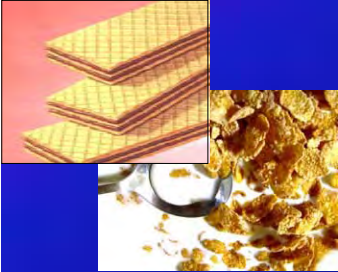
1) Soy seed

MZ	SL	RR	CP
			
			
			
			
			
			
			
			

2) Lecithin

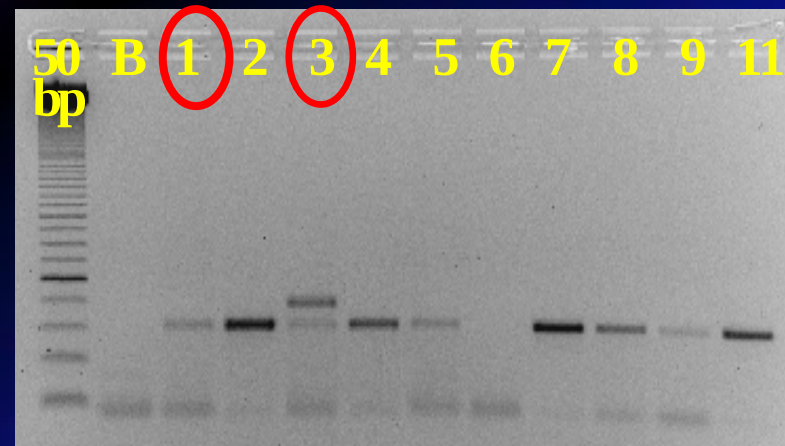
[illegible]

Detection of allergens in commercial products: hazelnut (Cor a1) and peanut (Ara h2)



PCR analyses performed on several foodstuff claiming to contain or to not contain hazelnut as ingredient or possible contaminants.

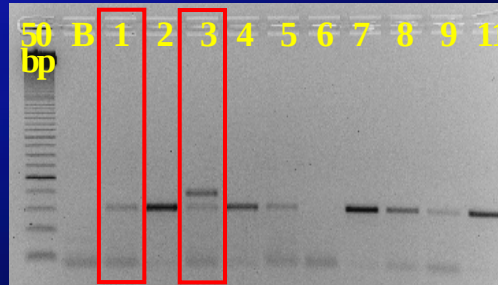
Sample	Product
1	Breakfast cereals
2	Breakfast cereals with chocolate and hazelnut
3	Müesli snack with milk chocolate
4	Müesli snack with cocoa
5	Snack with cereals and cocoa
6	Cocoa and vanilla wafer
7	Hazelnut wafer
8	Biscuit topped with cocoa cream and milk chocolate
9	Milk chocolate
10	Chocolate for topping
11	Chocolate cream with hazelnut



Hazelnut Cor a 1 (156bp) and Peanut Ara h 2 (201bp) amplicons.

Limit of detection : 50pg DNA.

PNA - microarray detection of hazelnut and peanut DNA in commercial products



Sample 1. Breakfast cereals

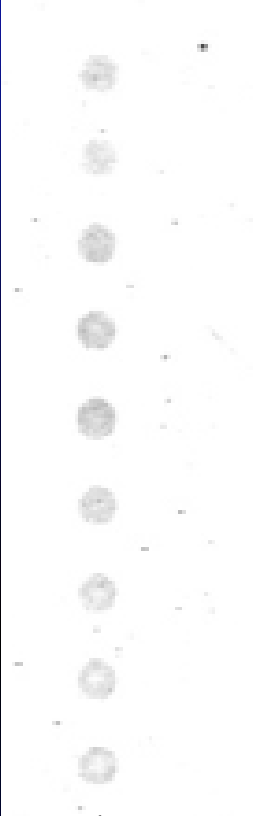
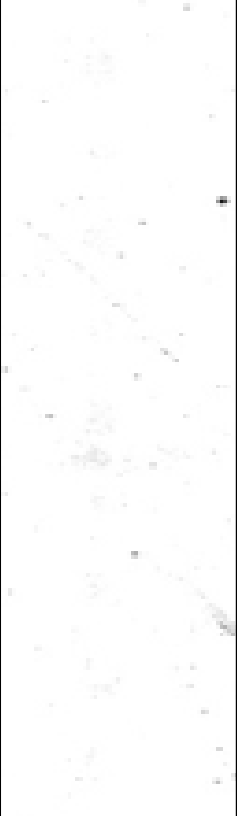
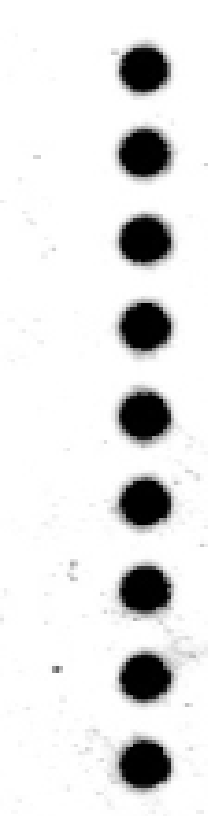
H	P	A	CP
•			•
•			•
•			•
•			•
•			•
•			•
•			•
•			•
•			•
•			•

Sample 3. Muesli snack

H	P	A	CP
•	•		•
•	•		•
•	•		•
•	•		•
•	•		•
•	•		•
•	•		•
•	•		•
•	•		•
•	•		•

Rossi S., Scaravelli E., Germini A., Corradini R., Fogher C., Marchelli R.
European Food Research Technology, 2006, 223, 1-6

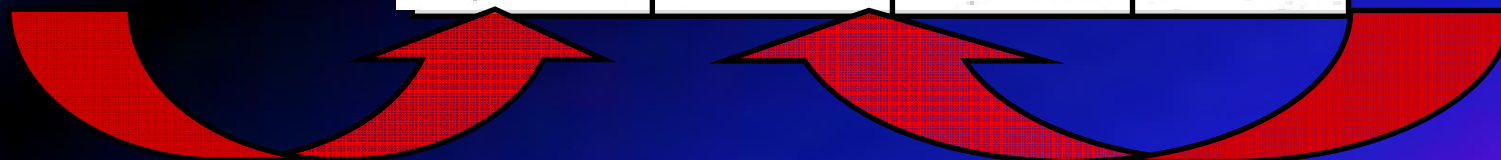
Detection of hazelnut oil (5%) in extra-virgin olive oil

PNA Cor	PNA Ara	PNA Pru	Control
			

- Cor a 1
Hazelnut
- Ara h 2
Peanut
- Pru du 2.1
Almond

**Detection limit
50fmol**

**PNA probes
specificity**



Authenticity: Olive oil cultivars

Cultivar	60	120	183	198	345
Biancolilla	A	A	G	C	G
Canino	A	R	R	S	G
Carolea	A	A	G	C	G
Coratina	A	A	G	C	G
Frantoio	A	A	G	C	G
Leccino	A	A	G	C	R
Nocellara belice	A	A	G	C	R
Ogliarola leccese	T	A	G	C	G
Moraiolo	A	R	R	C	G
Bosana	A	R	R	C	G
Nocellara etnea	A	A	G	C	R
Arbequina	A	A	G	C	G



**SNPs in the
Actin gene**

R = T o C
S = G o C

Olive variety identification by PNA microarrays

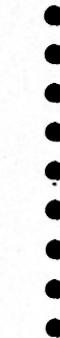
"Ogliarola leccese"

PNA	A 60	T 60	C198	G198	CP
DNA	T	A	G	C	CP

"Canino"

PNA	A 60	T 60	C198	G198	CP
DNA	T	A	G	C	CP

PNA

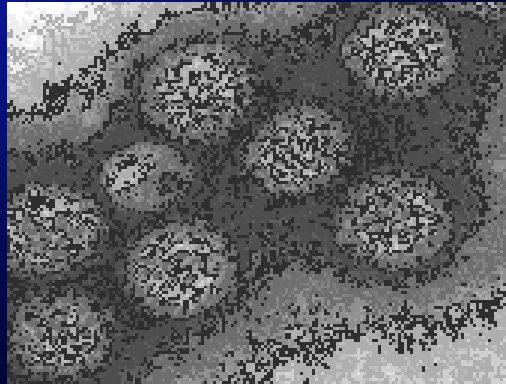
A 60	T 60	C198	G198	CP
				
T	A	G	C	CP

DNA

"Frantoio"

NOROVIRUS

Caliciviridae family: single strand **RNA virus**



Main responsible for non bacterial gastroenteritis

23 milion cases per year estimated in USA

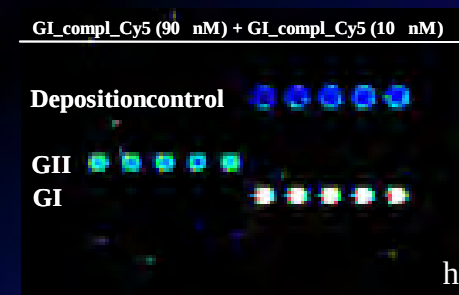
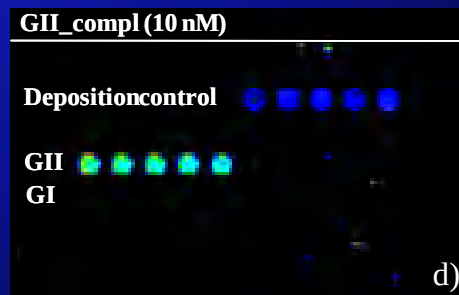
40% food-transmitted

water, mytilidae, fruits, vegetables

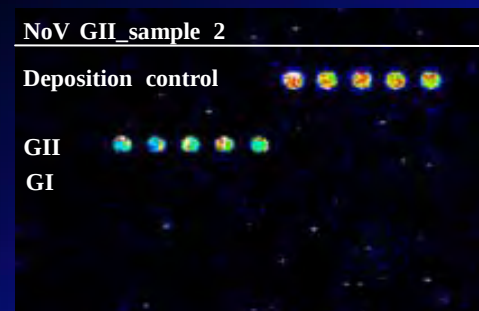
Not cultivable in vitro

Microarrays for Norovirus

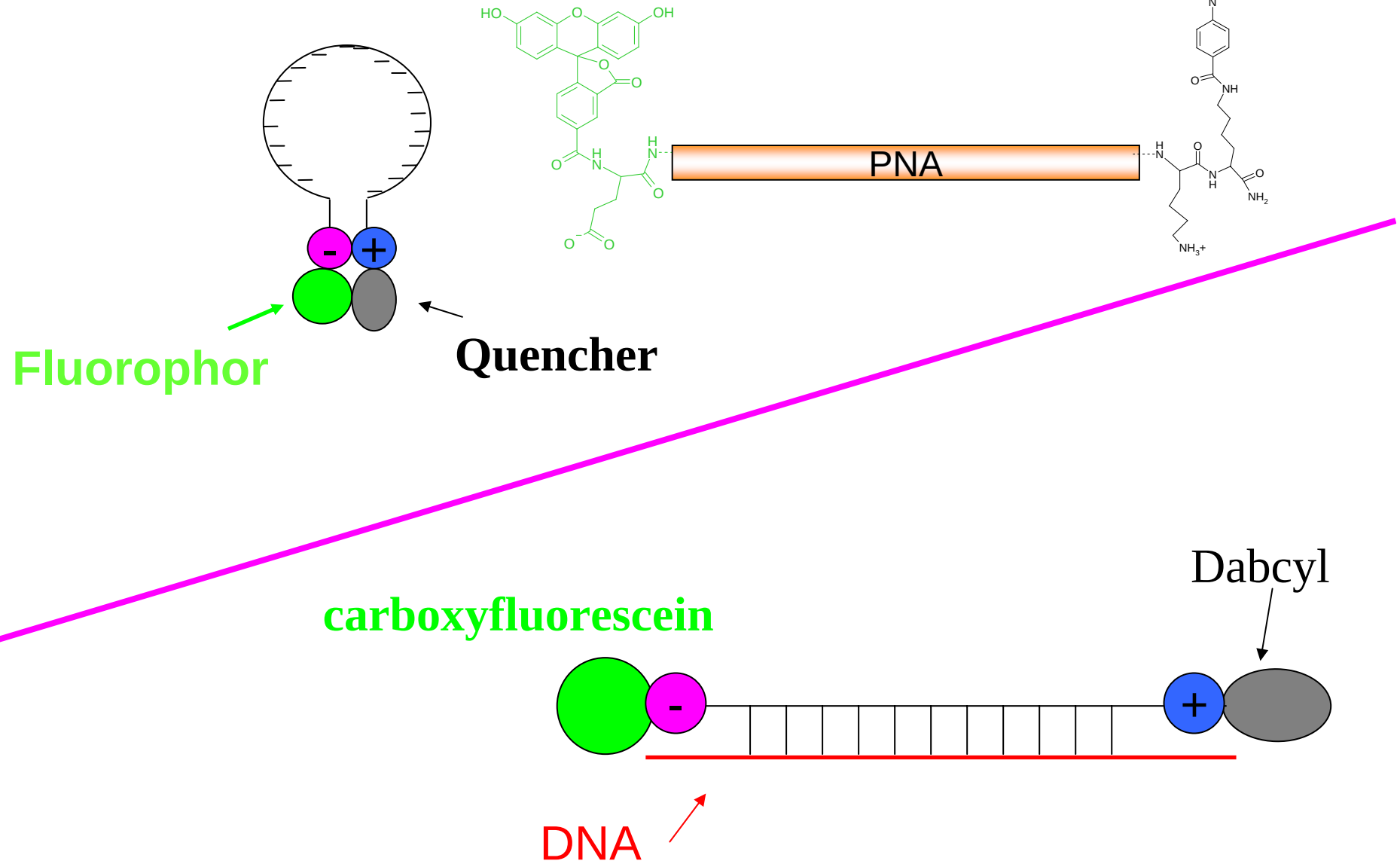
Oligonucleotides



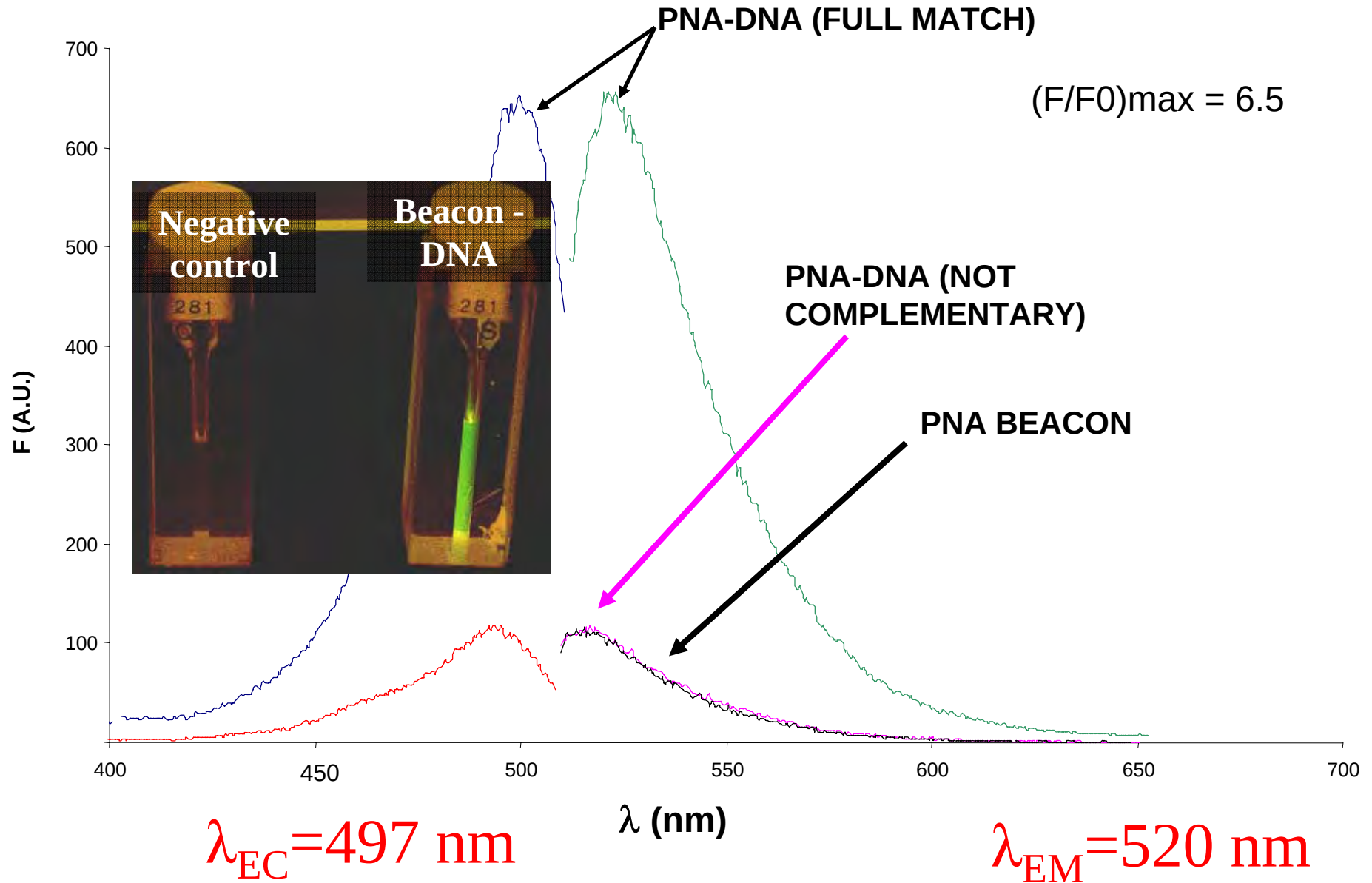
PCR products



PNA BEACON (STEMLESS)

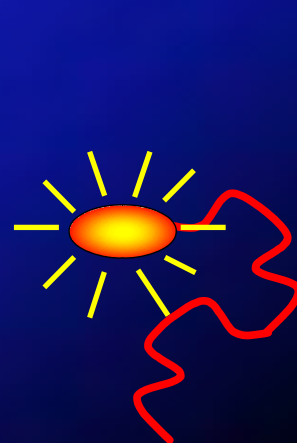
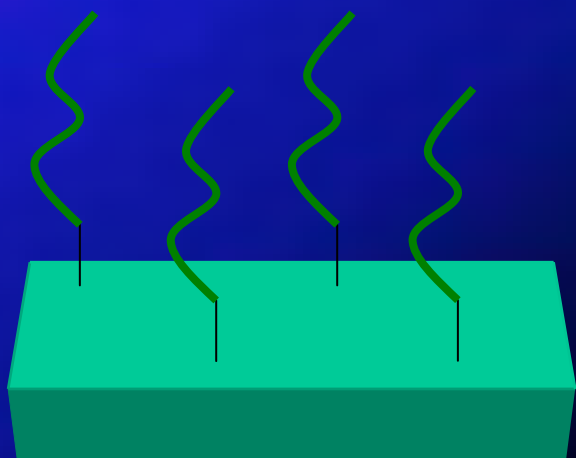


PNA BEACON FOR ROUNDUP READY-SOY

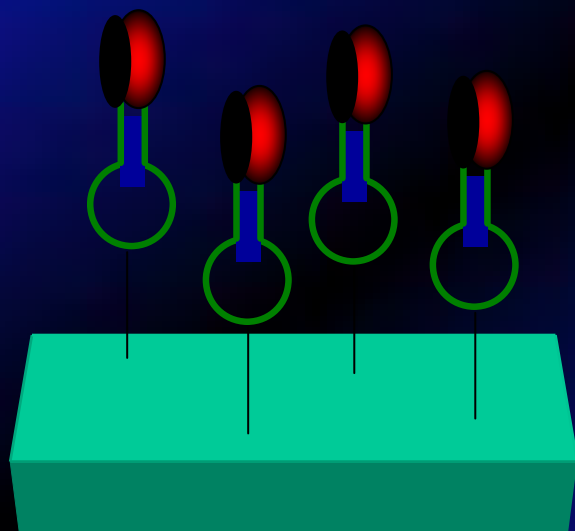
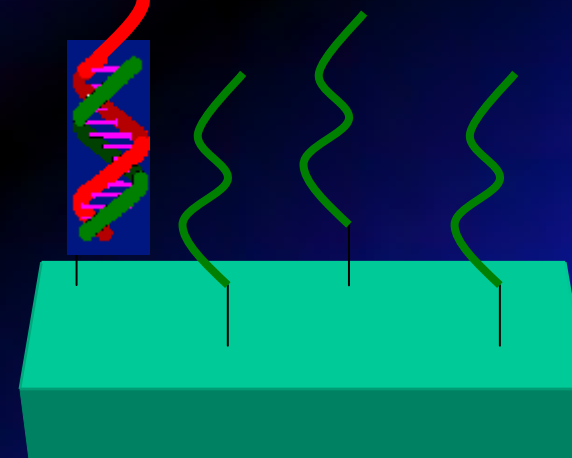
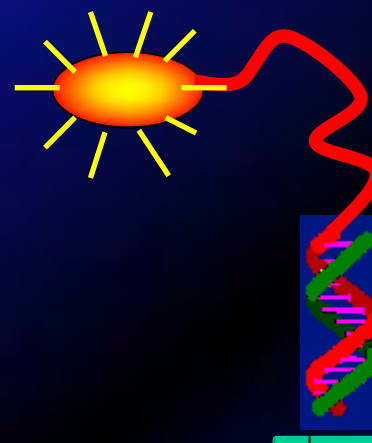


Towards a complete "lab on a chip" system

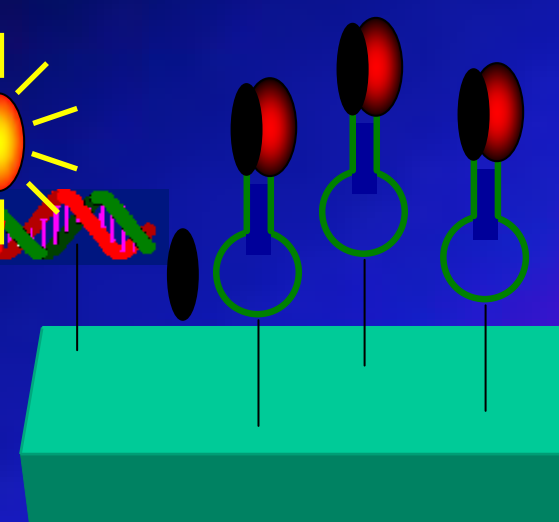
PNA array



Labelled
DNA



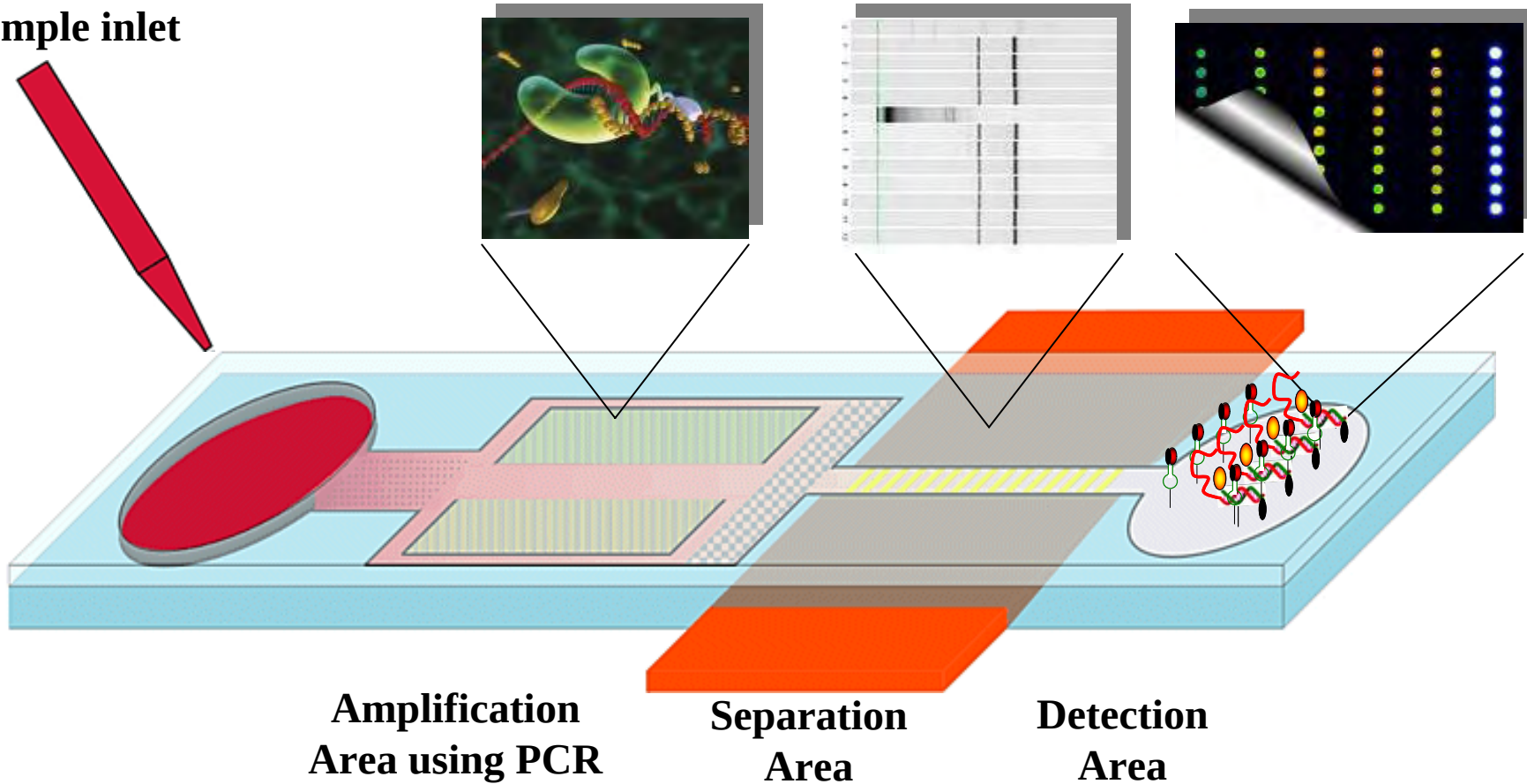
Unlabelled DNA



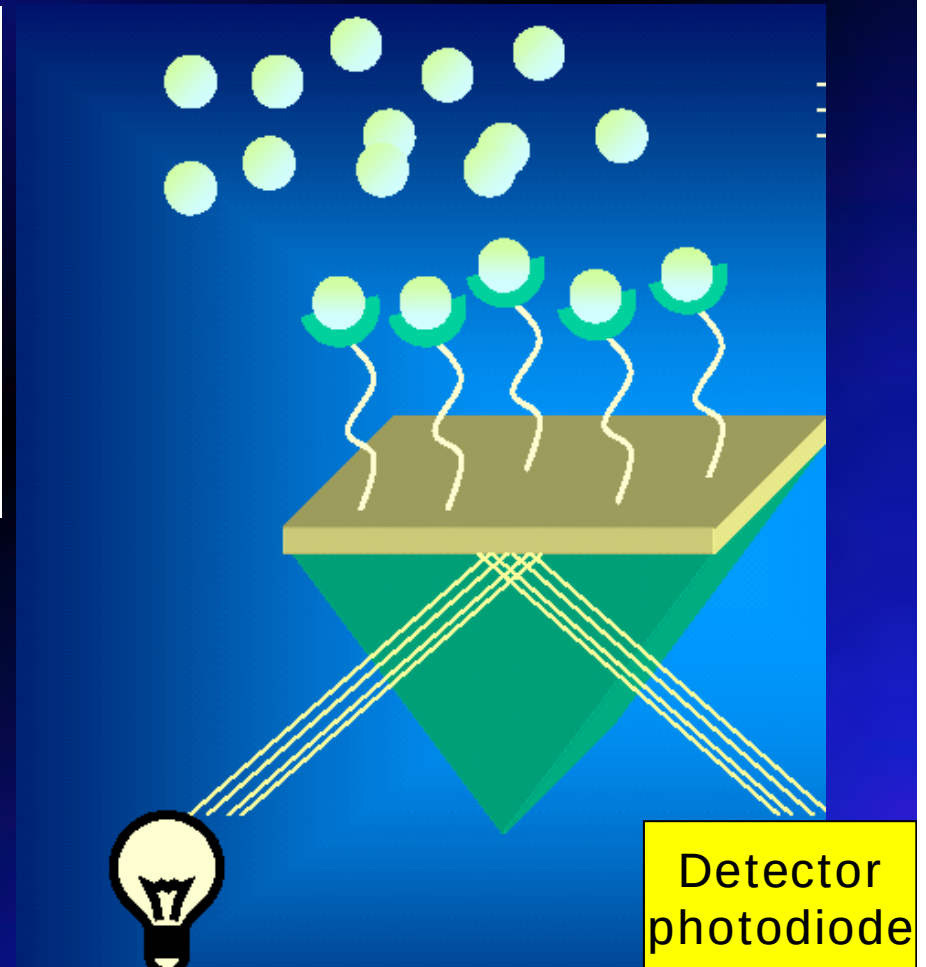
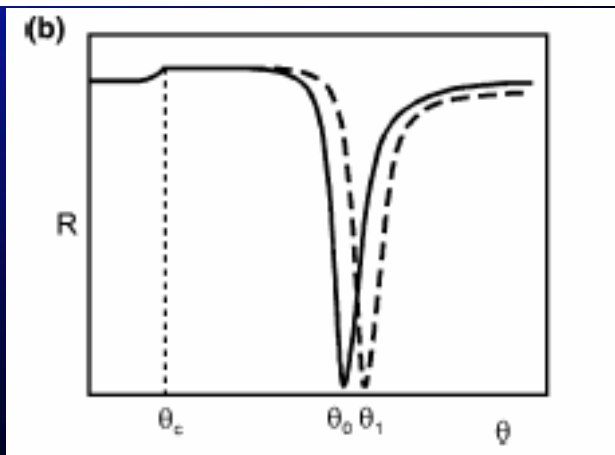
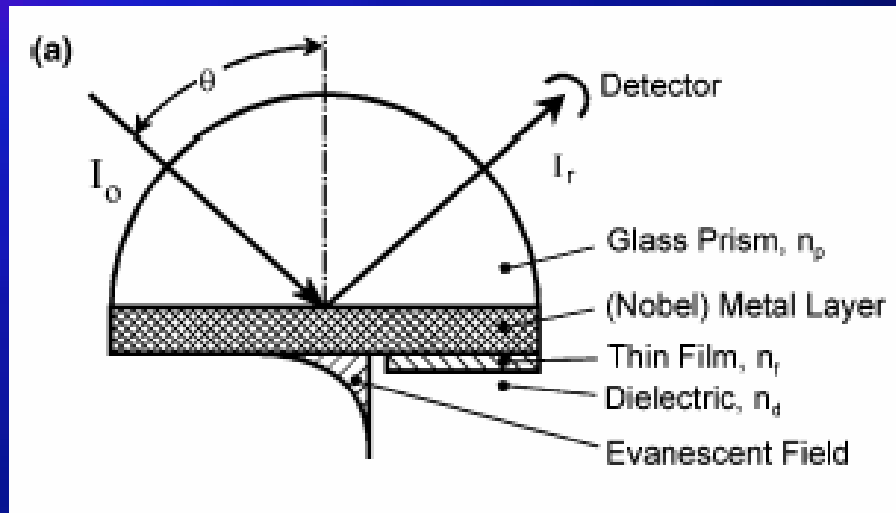
Array with PNA beacon

The “Lab on a chip” system

Sample inlet

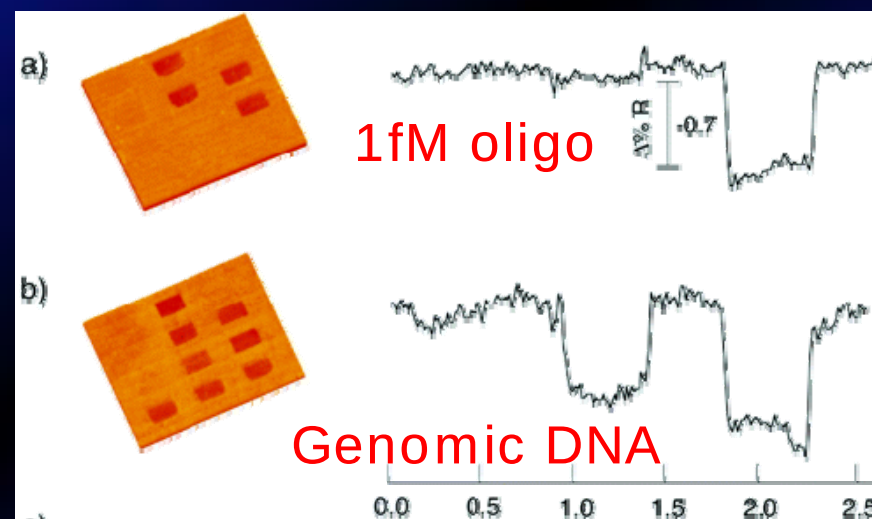
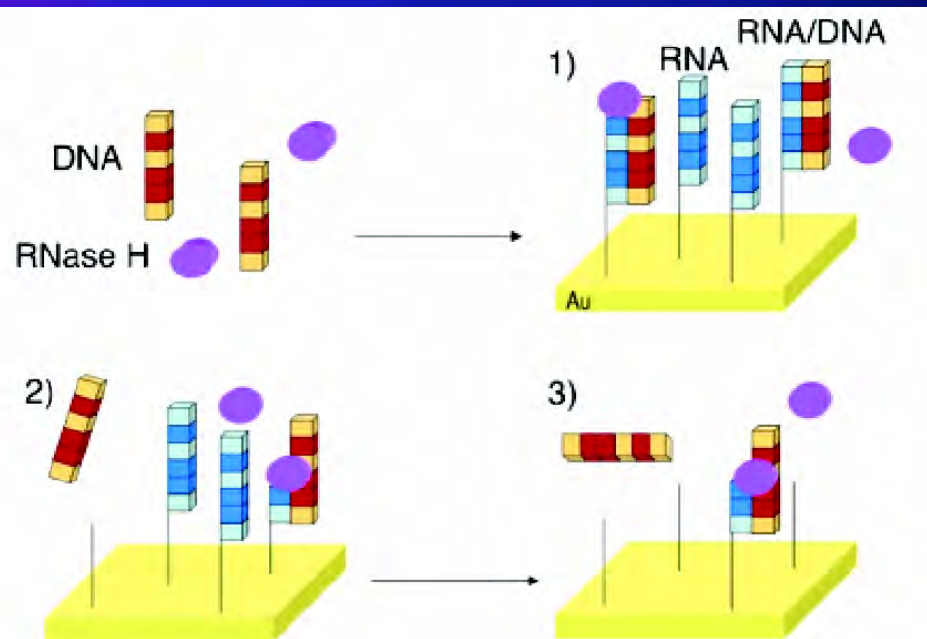


Surface Plasmon Resonance



Typical reflectivity scan (R versus incident angle θ) before (full curve) and after the deposition of a thin film (broken curves).

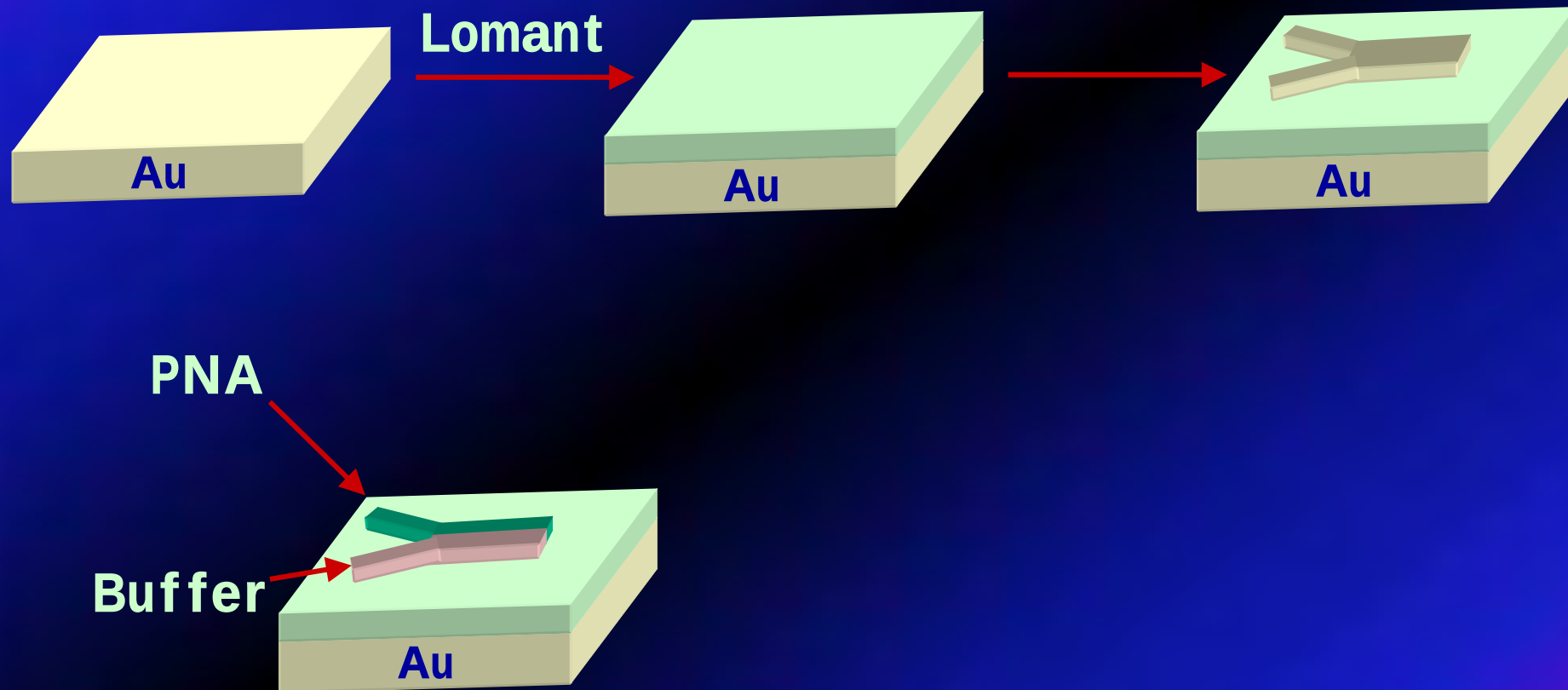
PCR-free DNA Detection



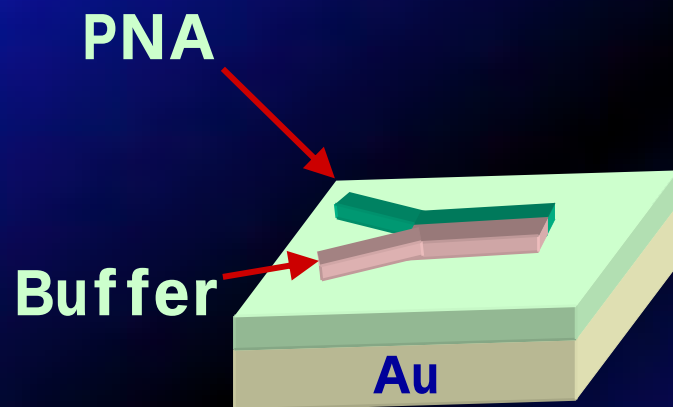
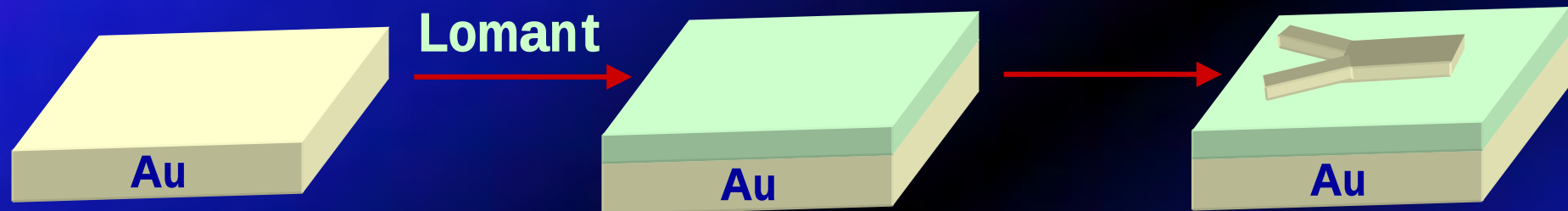
Scheme of PCR-free detection of genomic DNA using a RNaseH amplification of the SPR signal depletion. Detection limit of 10^{-15}M was attained

T.T. Goodrich, H.J. Lee, R.M. Corn *J. AM. CHEM. SOC.* 2004, 126, 4086-4087

SPR microfluidic platform based on Peptide nucleic acids (PNAs)

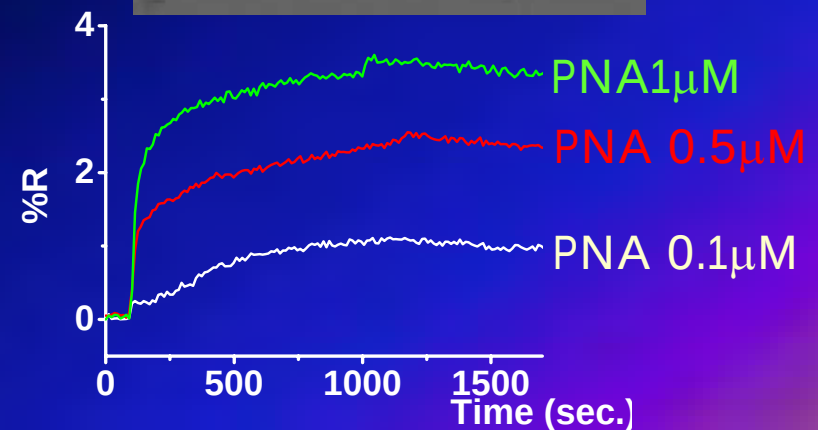
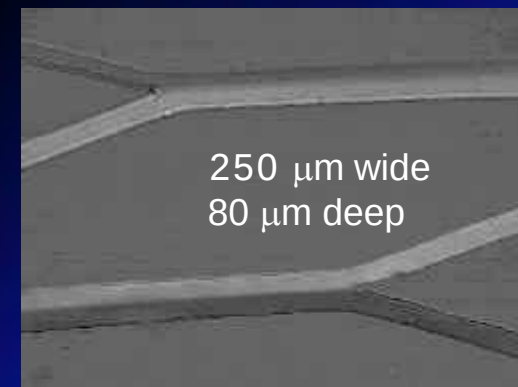


SPR microfluidic platform based on Peptide nucleic acids (PNAs)

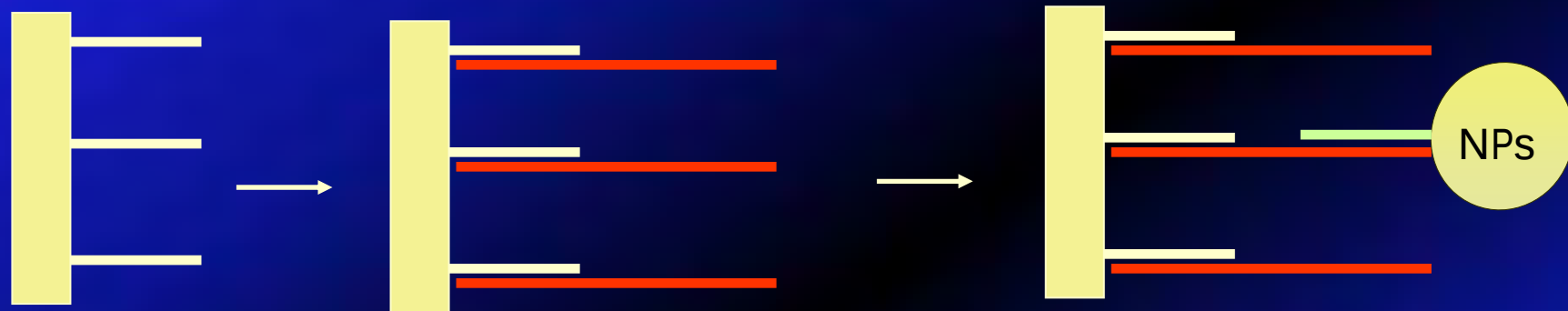


Coverage: 5×10^{12} molecule cm^{-2}

In collaboration with G. Spoto
(University of Catania)



SPR microfluidic platform based on Peptide nucleic acids (PNAs)

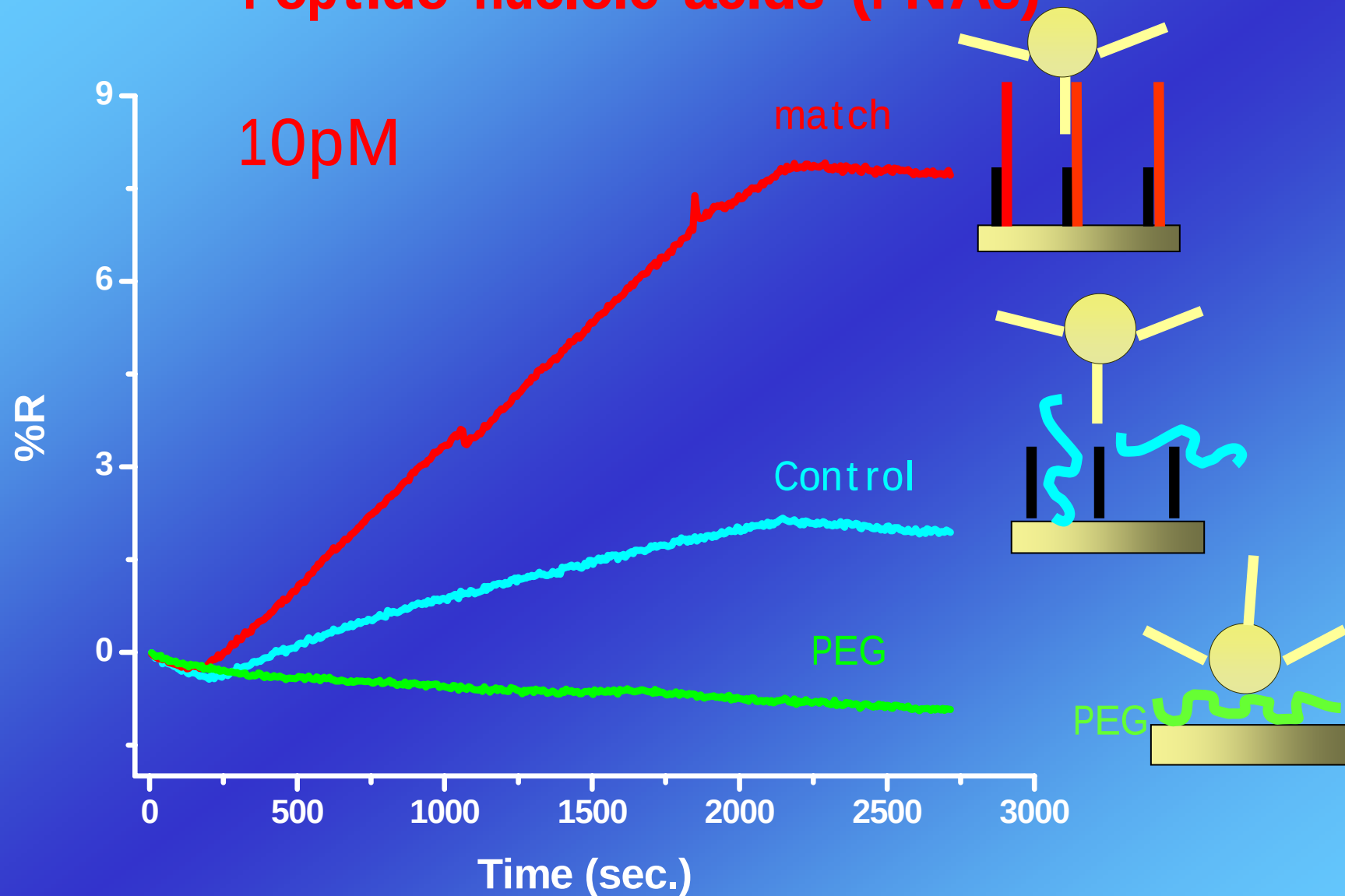


— 5'-LL-AAACCCTTAATCCCA-3' PNA-15mer

— 3'-TTTGGGAATTAGGGTTTTTTTTTTTCGTCGAATAGCA-5'
ssDNA-36mer-match

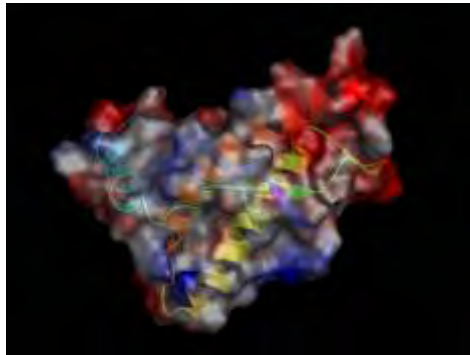
5'-GCAGCTTATCGT-3'-Biotin — NPs
ssDNA-12merC₆Biotin

SPR microfluidic platform based on Peptide nucleic acids (PNAs)



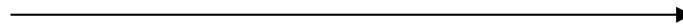
Prion protein detection in attomole amounts

Onisko et al., J.Am.Soc.Mass Spec., 2007, 18, 1070-1079



Prion protein

Reduction, alkylation, digestion

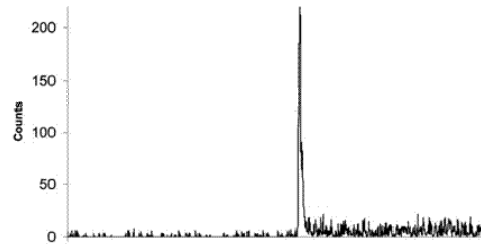


nanoLC/MS/MS

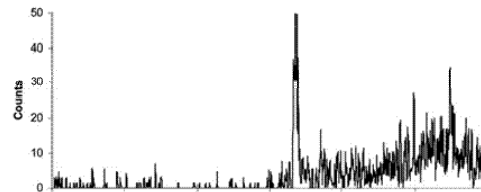


Detection of the specific peptide
VVEQMCTTQYQK

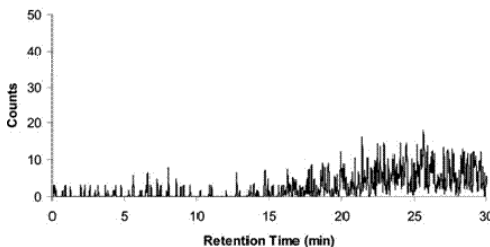
900 attomol



300 attomol

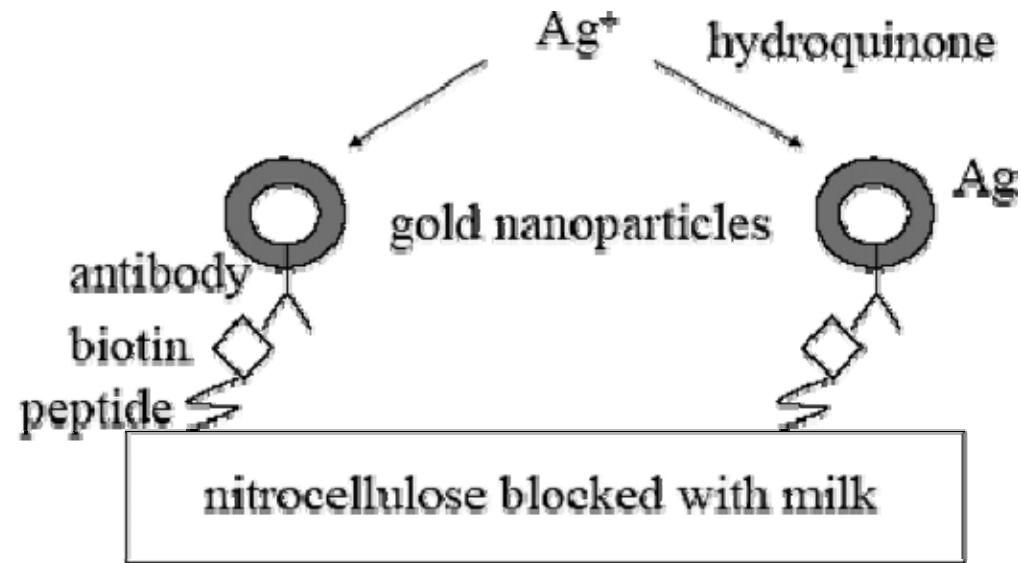


0 attomol

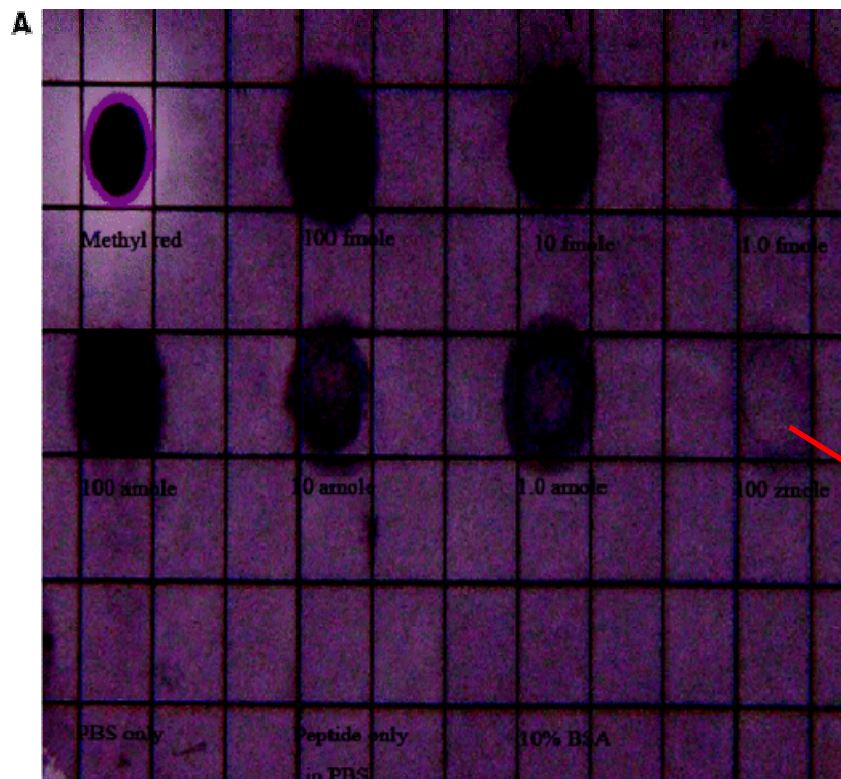


Zeptomole and attomole detection of biotin-peptides by a dot-blot gold nanoparticle immunoassay

Hou S.Y. Et al., Analytical Chemistry, 2007, 79, 980-985



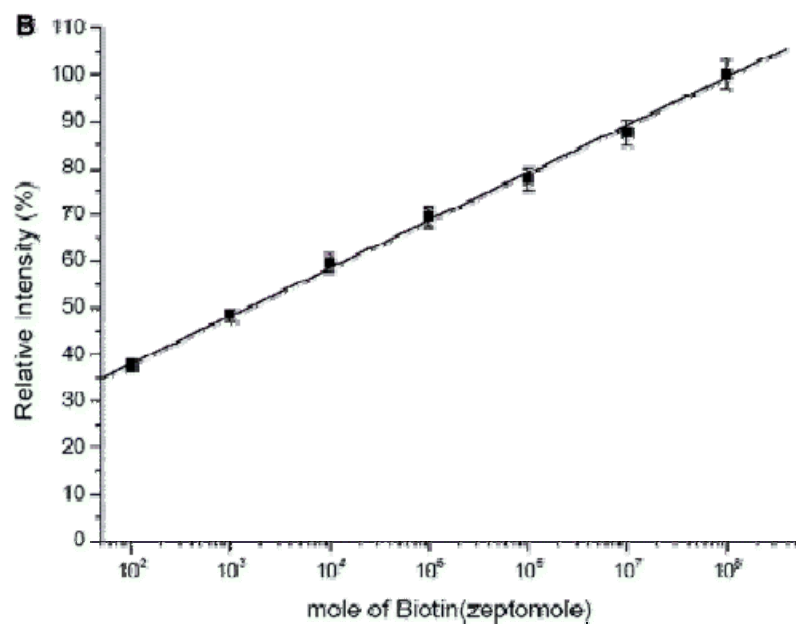
peptide sequence: SGQSWRPQGRFG



Detection of biotin-peptide with silver enhancement.

(A) Dot images for different levels of the biotin-peptide after silver enhancement.

100 zeptomoles



B) Detection curve for the biotin-peptide

Conclusions

- **The trend in food analysis is to devise micro(nano) systems, which can provide affordable, portable, fast and user friendly equipments**
- **Nanotechnologies allow to push down the detection limits almost to “nothing”**
- **How low should the legal limits be set in food analysis ?**

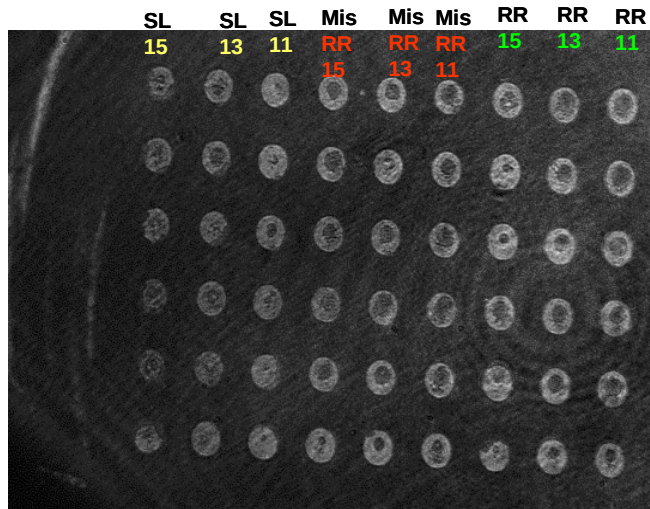
An aerial photograph of a city square. In the center is a large, light-colored octagonal tower with a tiled roof and a small spire. The square is paved and surrounded by various buildings, including a large one on the left and several smaller ones on the right. The image has a slightly grainy, vintage quality.

Acknowledgments

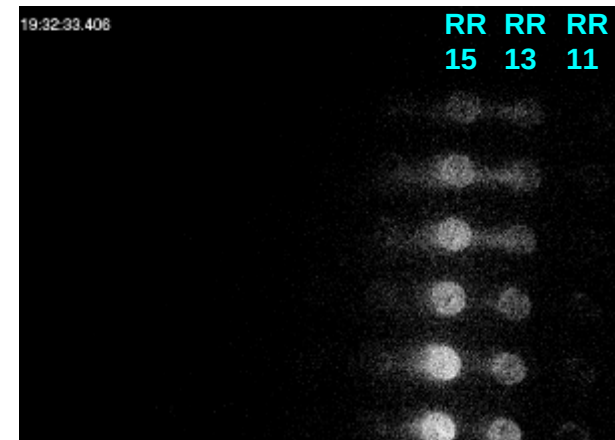
**Roberto Corradini
Stefano Sforza
Gianni Galaverna
Arnaldo Dossena
Tullia Tedeschi
Andrea Germini
Chiara Dall'Asta**

**Stefano Rossi
Elena Scaravelli
Andrea Faccini
Filbert Totsingan
Alessandro Accetta
Alessandro Calabretta
Alessandro Tonelli
Valeria Cavatorta
Mattia Mangia**

Surface Plasmon Microscopy



SPM image of Array B against air. All Parma PNAs ($1\mu\text{M}$ probe concentration)



SPFM image of the array after hybridisation with 1% RR125 GMO target solution

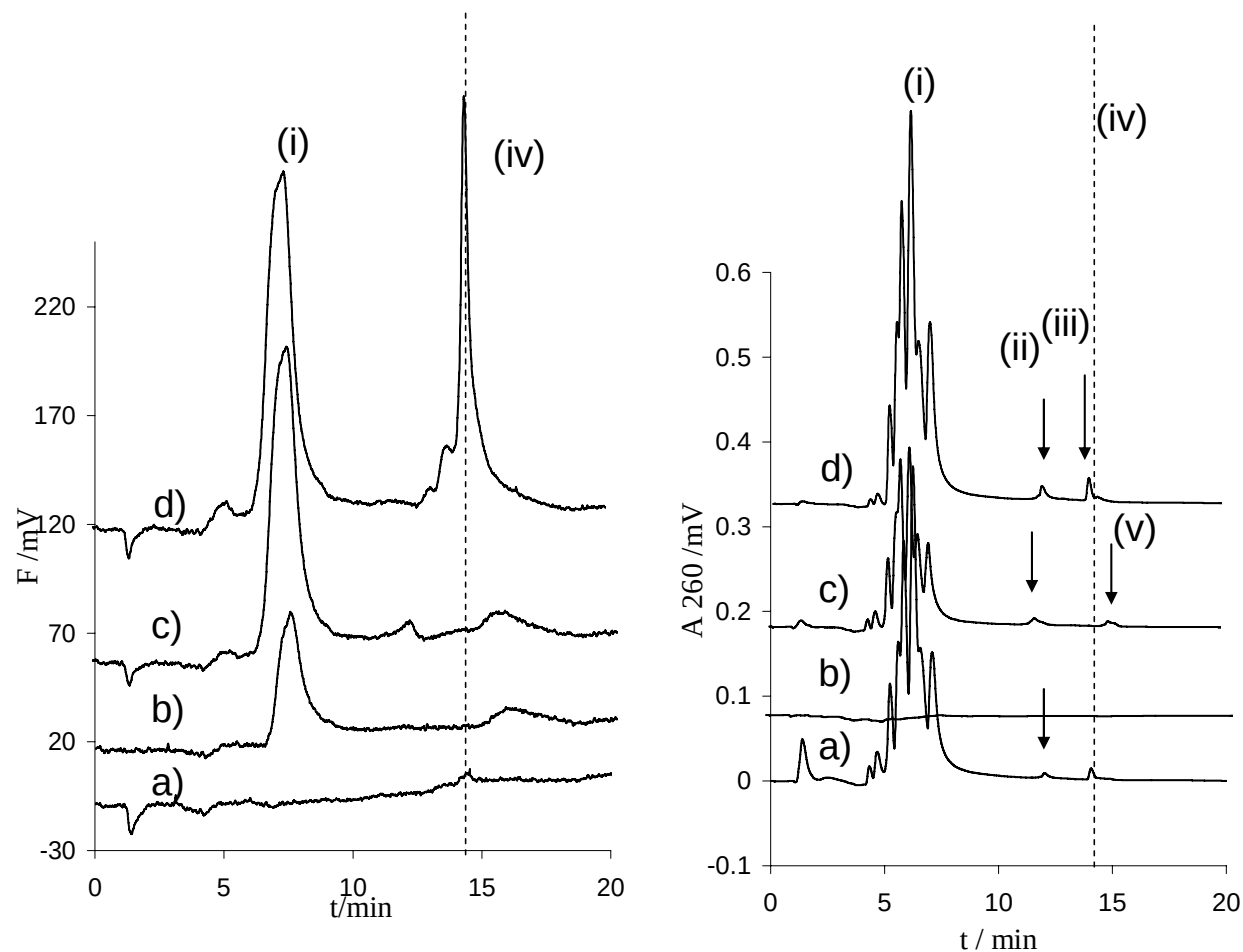
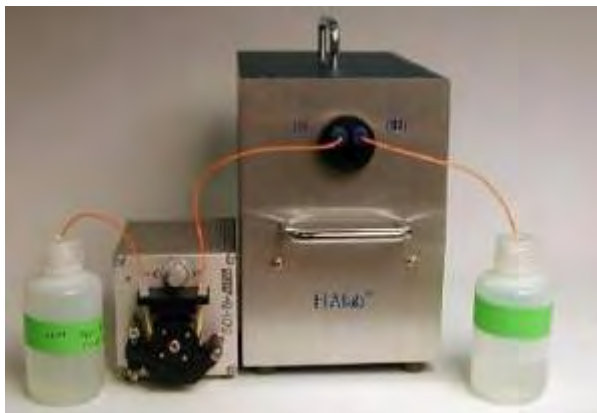
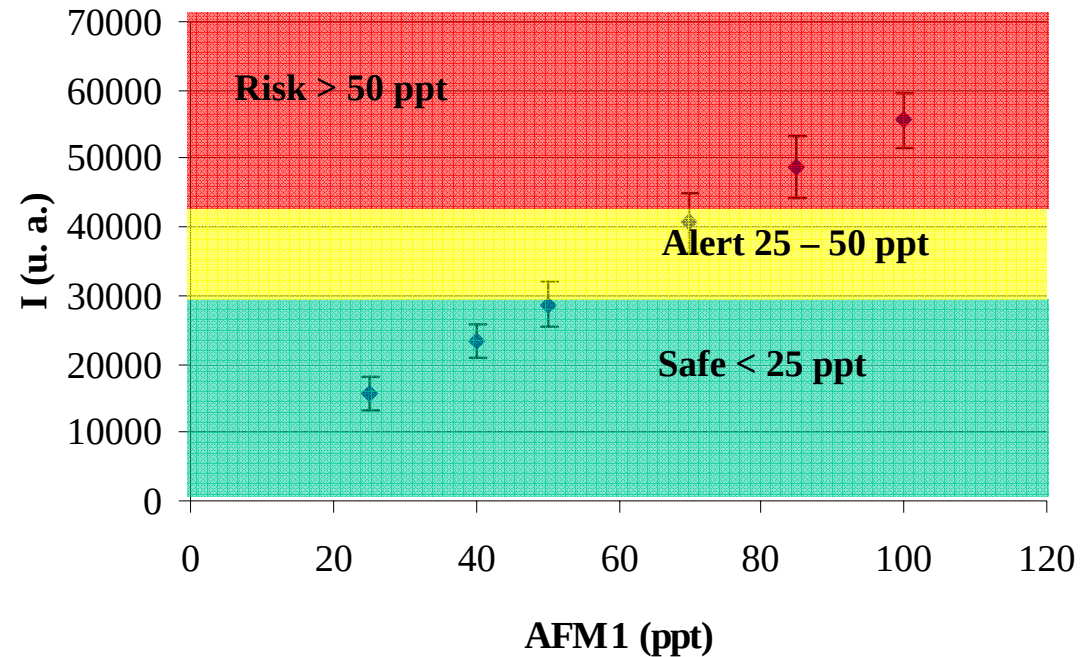
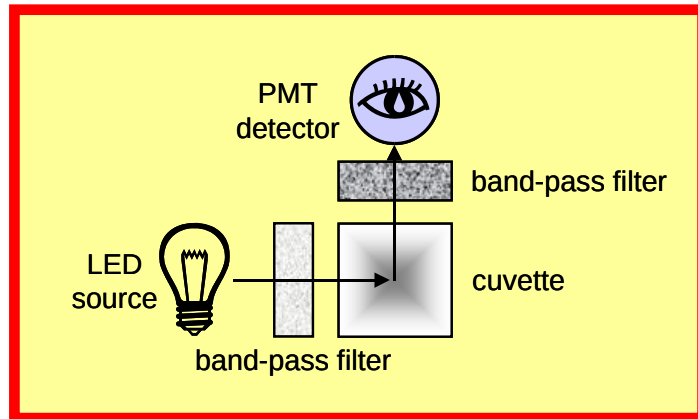


Figura 14. Tracciato IE-HPLC con detector a fluorescenza ($\lambda_{\text{ex}} = 497 \text{ nm}$, $\lambda_{\text{em}} = 520 \text{ nm}$, sinistra) e UV (260 nm, destra) relativo a : a) amplificato di PCR da solo; b) beacon RR-soia (1 mM) solo; c) beacon RR-soia (1 mM) + amplificato di PCR aspecifico; d) beacon RR-soia (1 mM) + amplificato di PCR specifico. I vari picchi sono stati identificati come: i) beacon e componenti della reazione di PCR; ii) prodotti di amplificazione aspecifici (primer-dimers); iii) prodotto di PCR 79-mer (soia RR) iv) ibrido PNA beacon-DNA; v) prodotto di PCR aspecifico da 201bp (controllo negativo).). Colonna: TSK-gel DNA NPR; eluenti: A = TRIS 0.02M, pH = 9.0, B = NaCl 1M in eluent A. Gradiente lineare: from 100%A to 100% B in 20 min; flusso: 0.5 mL/min.

Aflatoxins: FIAlab PMT-FL Detector (Fluorescence In-flow Assay)



LED: 360 nm
Band-Pass Filter maximum: 450 nm

A portable fluorometer for the rapid screening of M1 aflatoxin
C. Cucci, A.G. Mignani, C. Dall'Asta, R. Pela, A. Dossena
Sensors and Actuators B, 2007, in press

Future implementation: fiber optics.

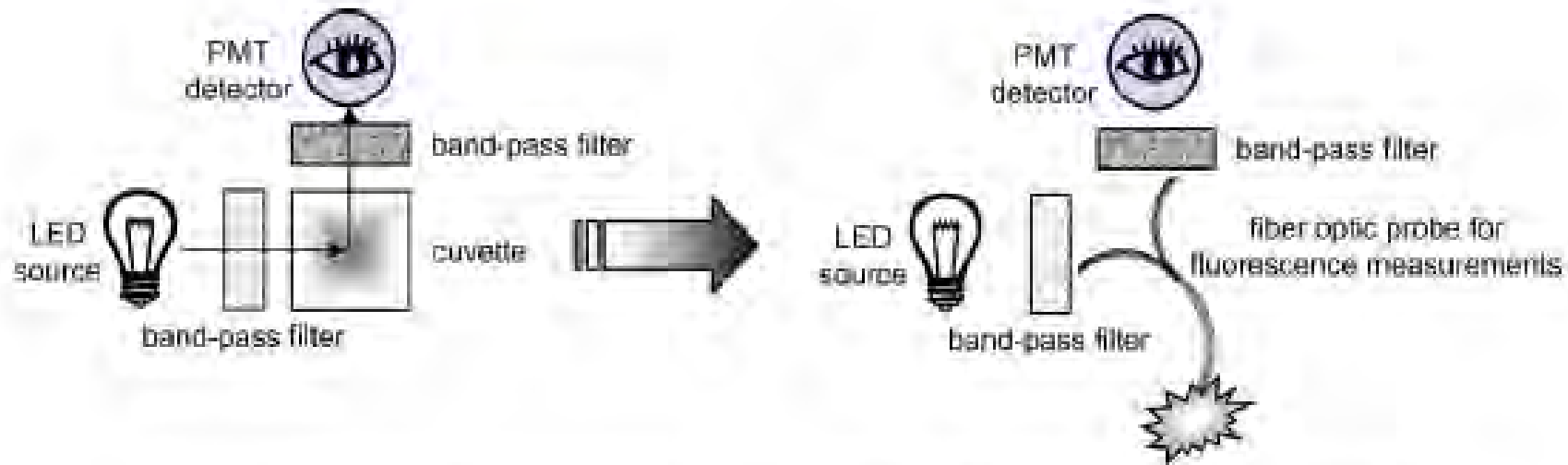
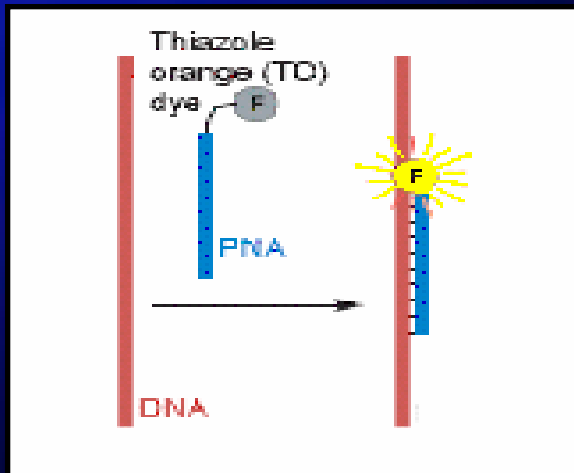
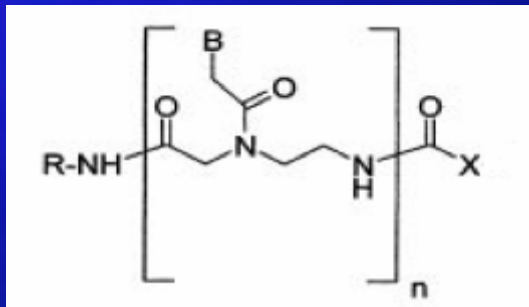


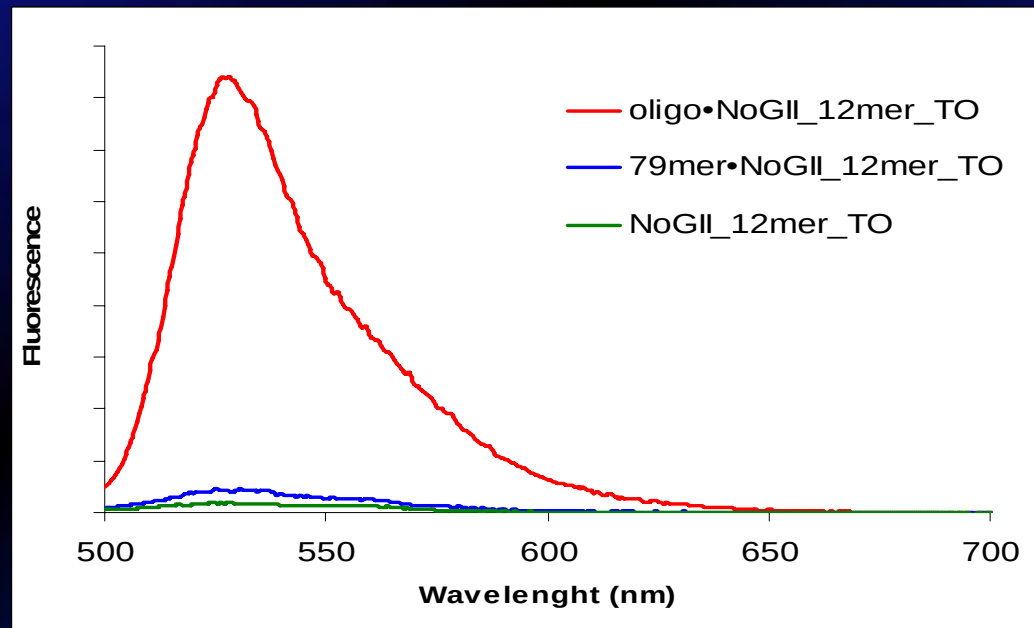
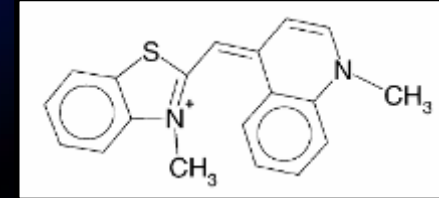
Fig. 1. Schematic diagram of the fluorometer (left) and its possible implementation using fiber optics (right).

LightUp® Probes

PNA

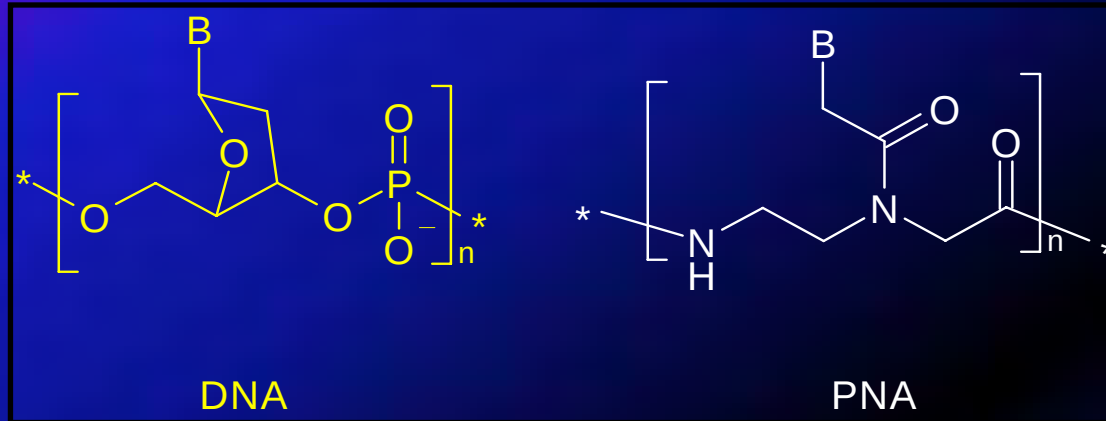


Thiazole Orange (TO)



Fluorescence emission spectra of free NoGII_12mer_TO (green line) and PNA_12mer_TO in the presence of non-complementary (blue line) and complementary (red line) DNA oligonucleotide. Samples were analyzed at a concentration of 1 μ M. Excitation wavelength: 470 nm. All samples were heated at 95 °C for 3 min and spectra were acquired after 5 min at 45 °C

Identification of specific DNA sequences by Peptide Nucleic Acids (PNAs)



DUPLEX STABILITY:

PNA-DNA >> DNA-DNA

10-mer: $T_m = 50^\circ\text{C}$ $T_m = 35^\circ\text{C}$

POINT-MUTATION RECOGNITION:

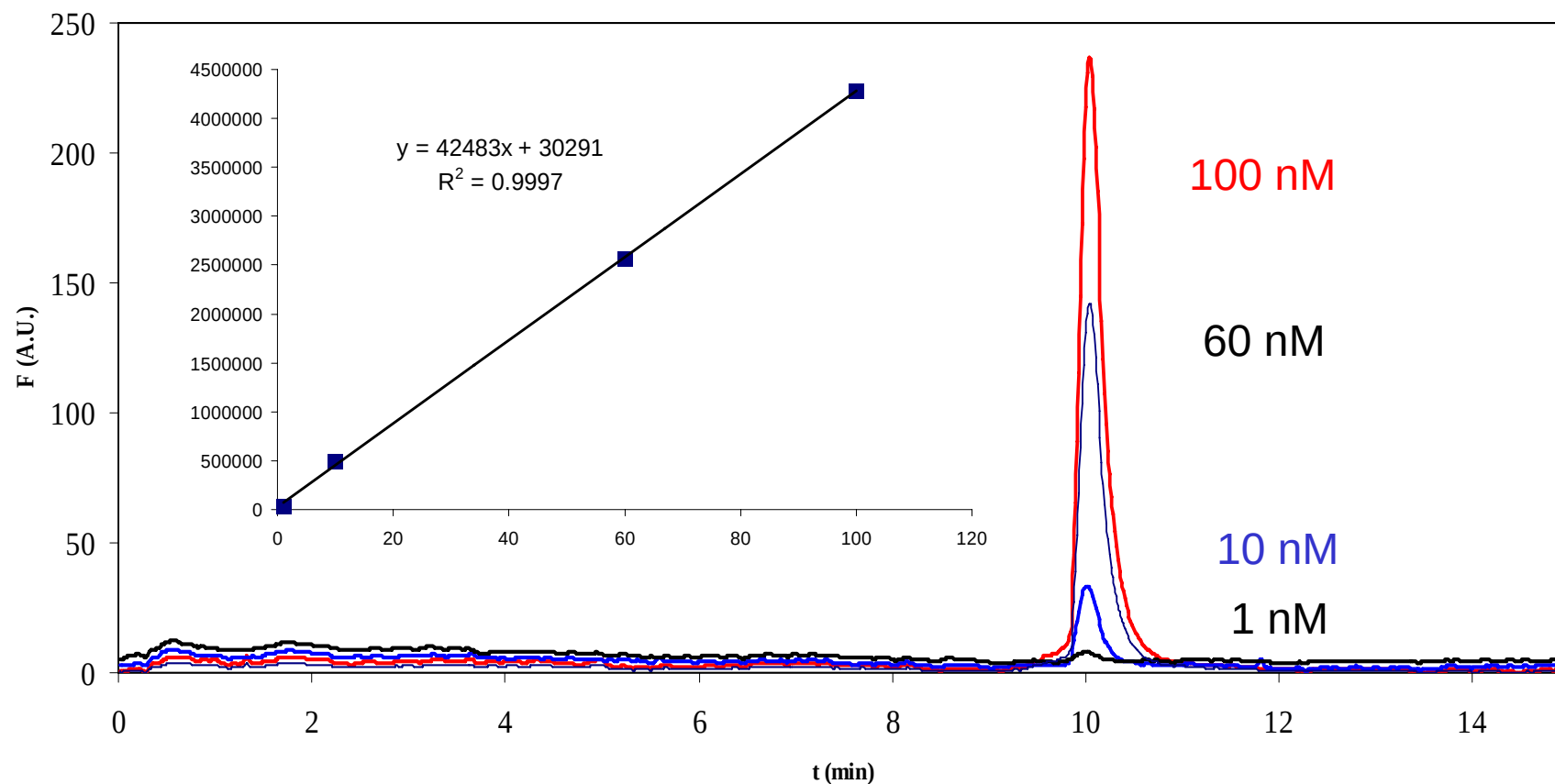
PNA-DNA > DNA-DNA

**15-mer: T_m drop
with a mismatch**

$T_m: 69^\circ\text{C} \rightarrow 56^\circ\text{C}$ $T_m: 53^\circ\text{C} \rightarrow 49^\circ\text{C}$

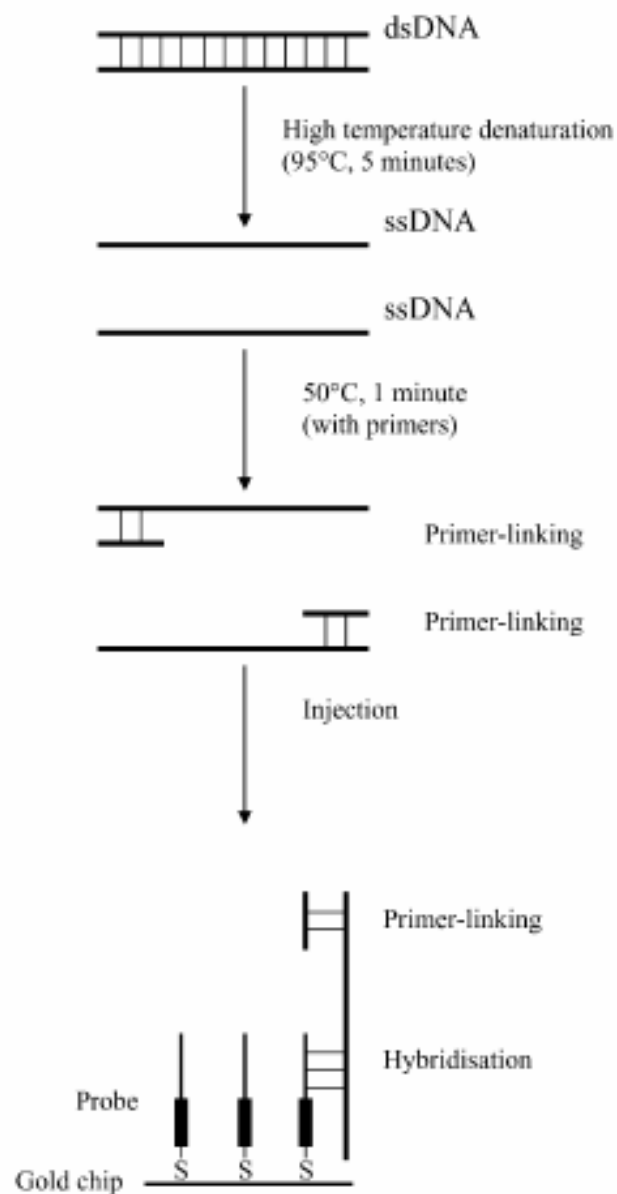
GMOs

Quantitative analysis of unlabelled DNA by RR-beacon/IE HPLC



Conditions: column TSK-gel DEAE-NPR 4.6mm x 7.5cm. Gradient elution from 100%A (tris, pH = 9.0) to 100% B (1M NaCl in eluent A) in 20 min. Fluorescence detector ($\lambda_{ex} = 497$ nm $\lambda_{em} = 520$ nm). Flow rate: 0.5 mL/min

PCR-FREE DNA analysis with DNA probes



35S genomic DNA detection from GM *Nicotiana glauca*, by piezoelectric sensing

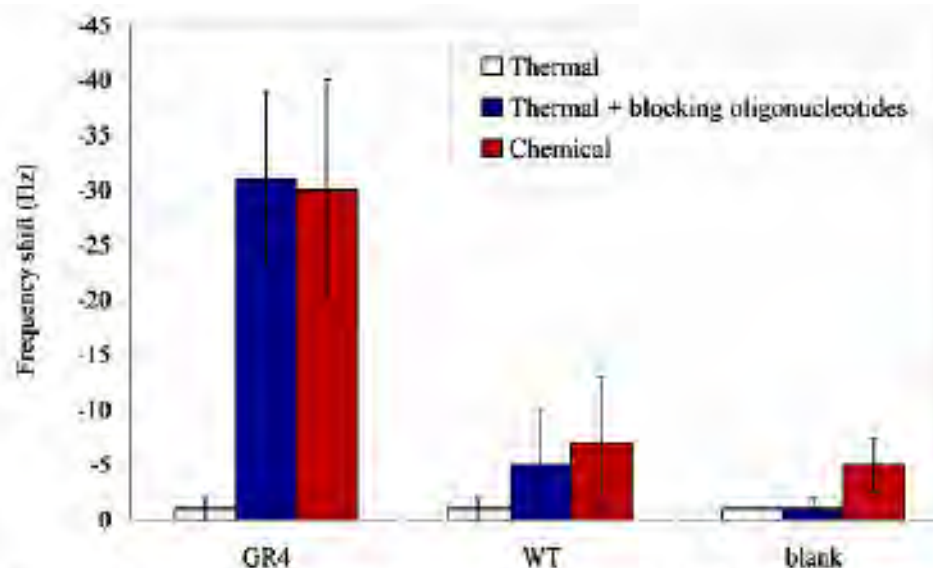


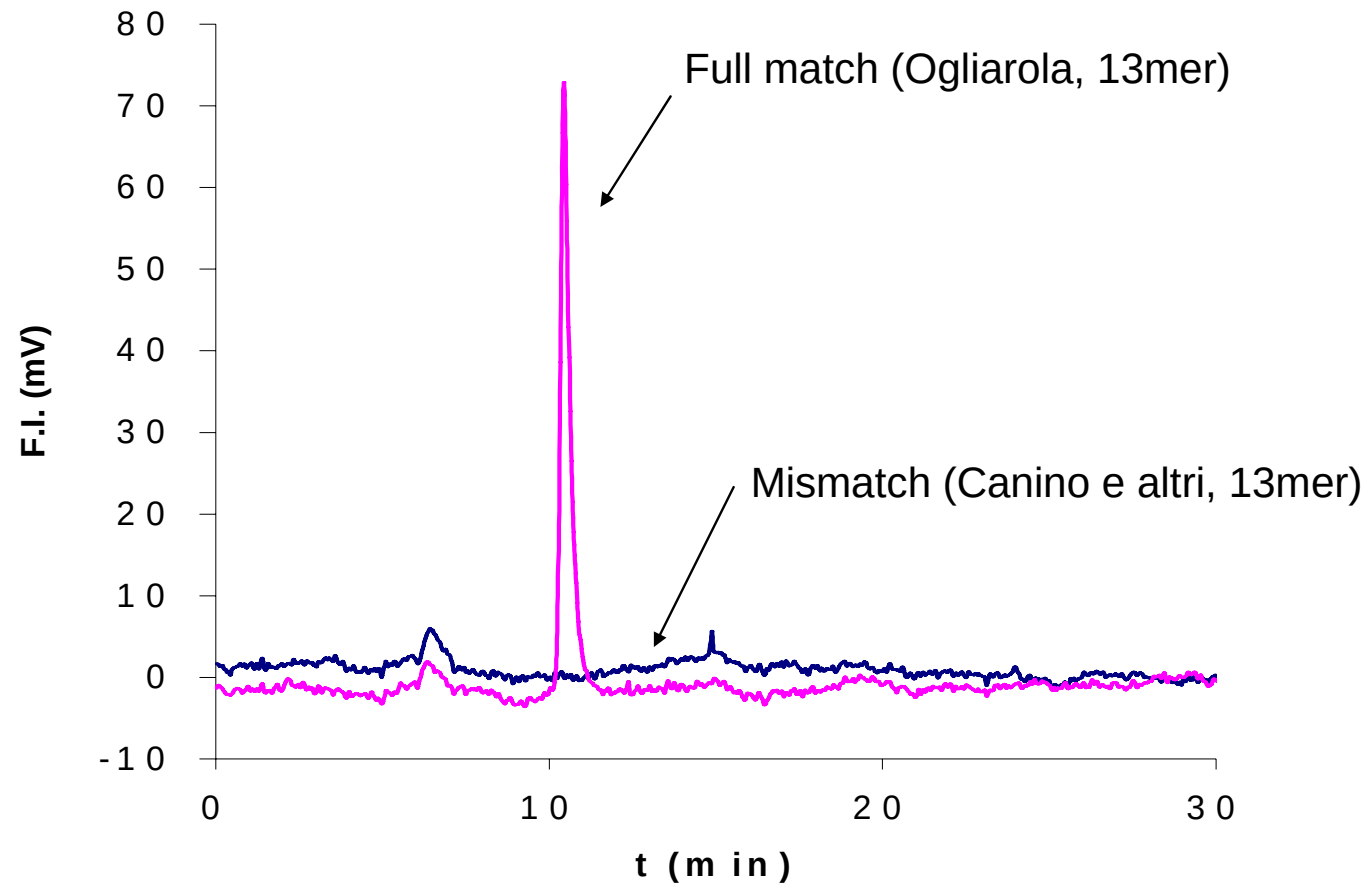
Figure 2. Experiments with digested genomic DNA (10 ppm), transgenic GR4 DNA ($n = 6$), nontransgenic WT DNA ($n = 6$), and blanks ($n = 3$).

Minunni M., Tombelli S., Fonti J, Spiriti M.M., Mascini M., Bogani P., Buiatti M.

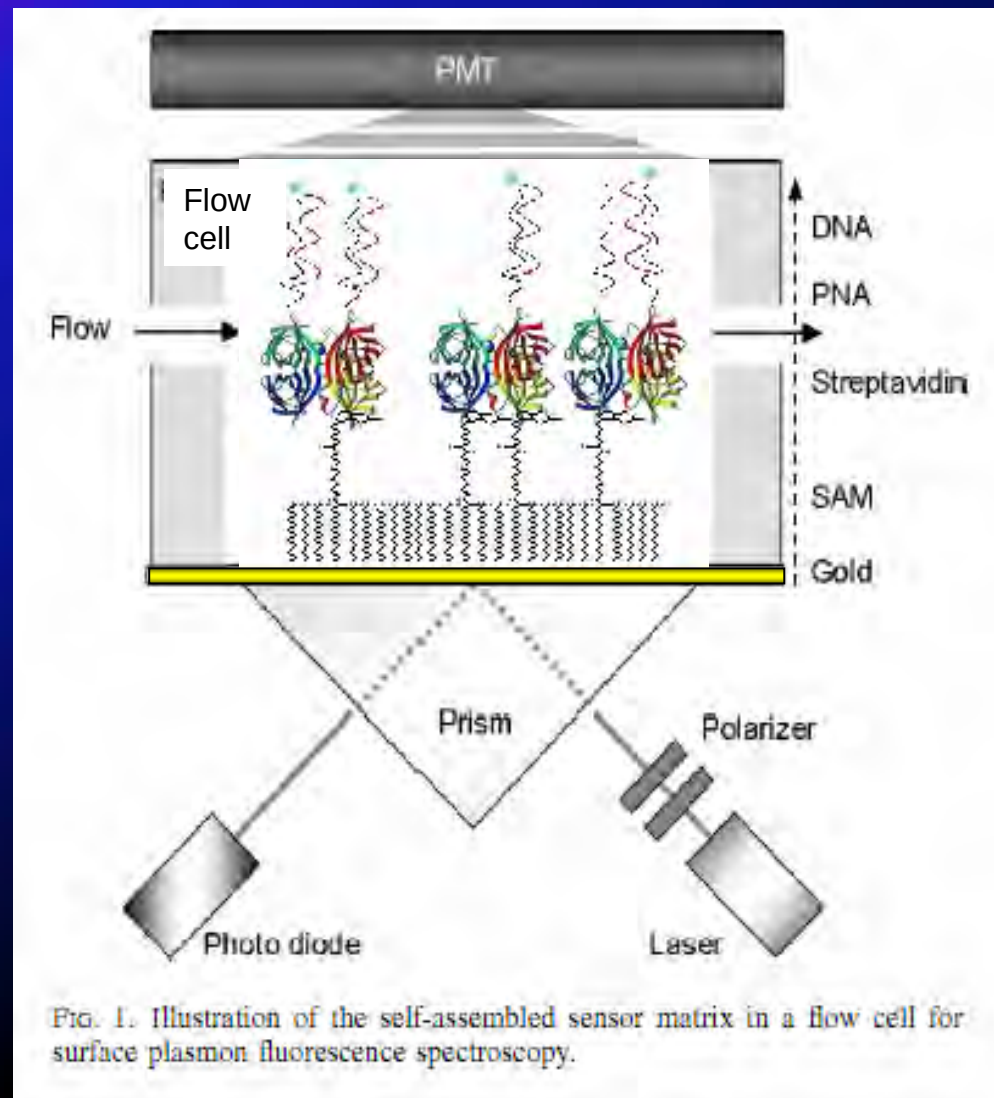
Journal of the American Chemical Society, 2005, 127, 7966-7967

PNA beacon

A 60 (11mer) Fluo-Glu-x xxx A xxx xxx-Lys-Lys(Dabcyl)-NH₂



Bottom-up: building sensors on self-assembled monolayers (SAM)



Surface-Plasmon Fluorescence Spectroscopy (SPFS)

Measures:

Affinity
Kinetics
Concentration
Arraying/Imaging is
possible

Selective detection of RR-soy PCR

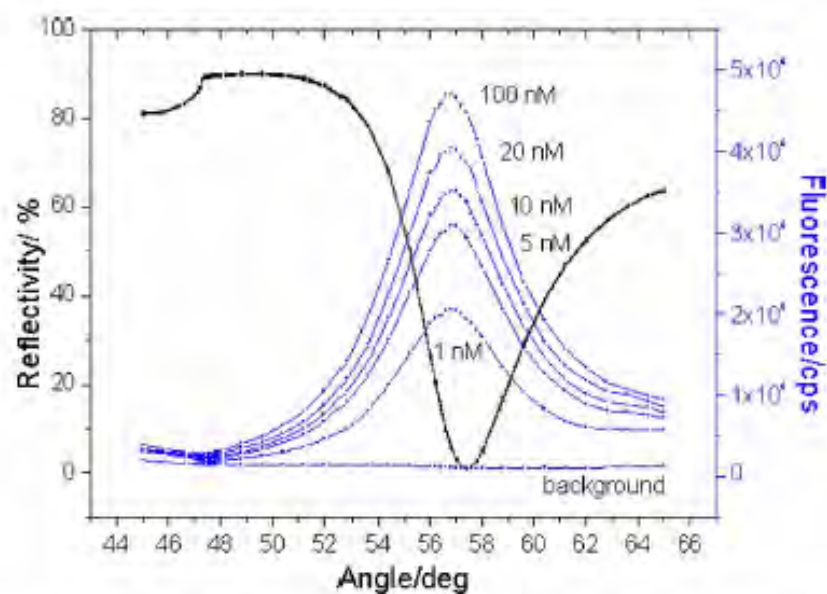


FIG. 7. Angular scans for PNA/DNA RR-169 hybridization taken in 10 mM phosphate buffer after saturation of the fluorescence was reached for 1, 5, 10, 20, and 100 nM target solutions, respectively, together with background fluorescence intensity recorded after rinsing. Typical R^2 after fitting is 0.999.

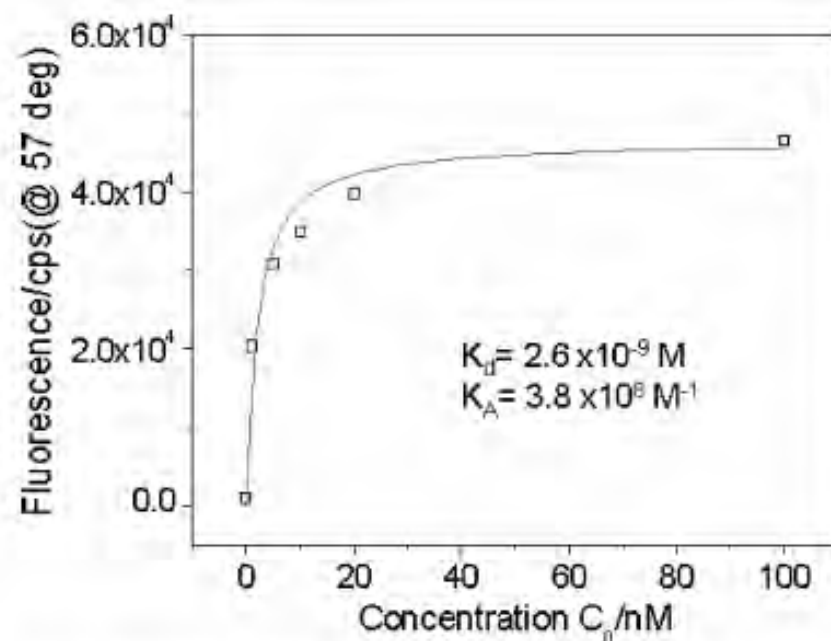


FIG. 8. Maximum fluorescence intensity [taken from Fig. 7(b) at $\theta=57^\circ$] vs target concentration. The full curve corresponds to a fit according to the Langmuir isotherm with $K_A=3.8 \times 10^8 M^{-1}$. Typical R^2 after fitting is 0.999.