

## Characterization of genomic SNP via colorimetric detection using a single gold nanoprobe

Fábio Ferreira Carlos<sup>1,2</sup>, Orfeu Flores<sup>2</sup>, Gonçalo Doria<sup>1</sup>, Pedro Viana Baptista<sup>1</sup>

<sup>1</sup>Nanomedicine@FCT, UCIBIO, Departamento de Ciências da Vida, Faculdade de Ciências e Tecnologia, Universidade Nova de Lisboa. Campus da Caparica, 2829-516 Caparica,

<sup>2</sup>STABVIDA, Investigação e Serviços em Ciências Biológicas, Lda. Madan Parque, Caparica, Portugal  
[fa.carlos@campus.fct.unl.pt](mailto:fa.carlos@campus.fct.unl.pt)

### Abstract

Identification of specific nucleic acid sequences mediated by gold nanoparticles derivatized thiol-modified oligonucleotides (Au-nanoprobes) has been proven a useful tool in molecular diagnostics [1,3-5]. Here, we demonstrate that, upon molecular optimization, detection may be simplified and results attained using a single Au-nanoprobe to detect SNP in homo- or heterozygous condition (Figure 1). Au-nanoprobes are becoming extremely useful tools for routine molecular diagnostics involving single-base mismatch detection that can be a fast, cheap and reliable alternative to standard techniques [4].

We demonstrate the robustness of this approach through validation using clinical samples and screening for the SNP rs9939609 in the FTO gene locus [2]. For the first time, we demonstrate that, upon comprehensive optimization, a single Au-nanoprobe may be used alone to detect SNP by presenting distinct threshold for each genetic status (wild type, heterozygous and mutant) with high degree of sensitivity (87.50%) specificity (91.67%) [5].

Results were validated using Sanger sequencing as gold standard. Sensitivity, specificity and limit of detection (LOD) were determined and statistical differences calculated by one-way analysis of variance (ANOVA) and a post hoc Tukey's test to ascertain whether there were any differences between Au-nanoprobe genotyped groups. From the 20 samples genotyped via Sanger sequencing, 8 samples were wild type (T/T), 7 samples heterozygous (T/A) and 5 samples mutated (A/A) [4]. Genotyping using the Au-nanoprobe determined 9 wild type (T/T) samples, 7 heterozygous (T/A) and 4 mutated (A/A). The LOD for Au-nanoprobe FTOWt20 was set at 20  $\mu\text{g ml}^{-1}$ . This simple and fast approach requires low complexity apparatus (UV/Vis spectroscopy) but may also be evaluated by the naked eye.

### References

- [1] Baptista P, Pereira E, Eaton P, Doria G, Miranda A, Gomes I, et al, *Anal. Bioanal. Chem.*, **391** 2008 943-50
- [2] Carlos FF, Silva-Nunes J, Flores O, Brito M, Doria G, Veiga L et al, *Diabetes Metab. Syndr. Obes.*, **11(6)** 2013 241-5.
- [3] Doria G, Franco R, Baptista P, *IET Nanobiotechnol.*, **1(4)** 2007 53-7.
- [4] Doria G, Conde J, Veigas B, Giestas L, Almeida C, Assunção M, Rosa J, Baptista PV, *Sensors*, **12(2)** 2012 1657-1687.
- [5] Carlos FF, Flores O, Doria G, Baptista PV, *Anal. Biochem.*, **465C** 2014 1-5.

### Figures

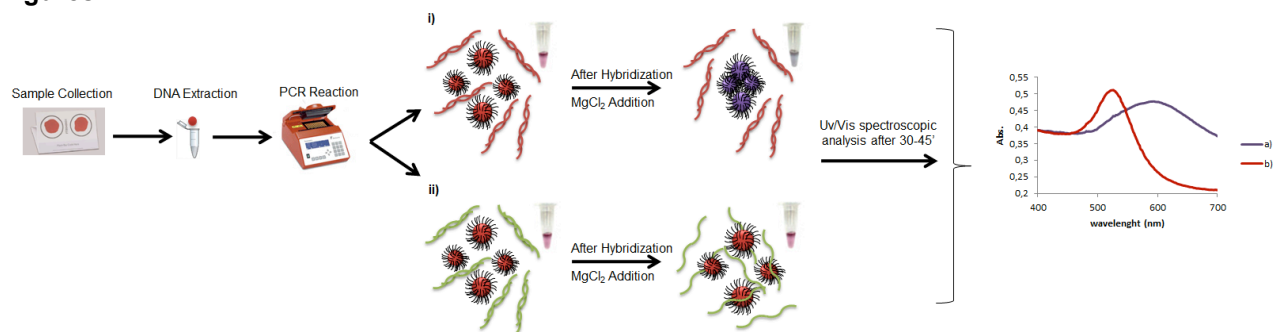


Figure 1 – Blood sample was collected from finger prick using a lancet and stored in FTA™ Indicated Micro Card (Whatman, UK). DNA extraction was carried out for subsequently PCR reactions. After PCR reaction, Au-nanoprobe was mixed with PCR products and a hybridization step was carried out i) after salt addition in the presence of a non-complementary or unrelated PCR a colorimetric change is visible

from red to purple after a certain time, ii) after salt addition in the presence of a fully complementary PCR product no colorimetric change is visible and the solution remains red. This color observation can also be observed by UV/Vis spectroscopy.