

1 **Naked Eye Y Amelogenin Gene Fragment Detection Using DNAzymes**
2 **on a Paper-based Device**

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13

14 **ABSTRACT**

15 Nowadays, there are strong efforts in developing new technology for rapid detection of specific DNA
16 sequences for environmental monitoring, forensic analysis and rapid biomedical diagnosis
17 applications. That is where microfluidic paper-based analytical devices are positioned as suitable
18 platforms for the development of point of care analytical devices, due to their simple fabrication
19 protocols, ease of use and low cost. Herein, a methodology for *in situ* single strand DNA detection
20 by using a colorimetric assay based on the formation of a DNAzyme within a paper substrate was
21 developed. A DNAzyme that could only be formed in the presence of a specific sequence of the Y
22 human amelogenin gene was designed. The performance of the DNAzyme was followed
23 colorimetrically first in solution and then in paper substrates. The reaction was found to be specific
24 to the Y fragment selected as analyte. The DNAzyme reaction on paper enabled the unequivocal
25 colorimetric identification of the Y single strand DNA fragment both qualitatively, with the naked
26 eye (143 ng), and quantitatively by image analysis (45.7 ng). As a proof of concept, a microfluidic
27 paper-based device, pre-loaded with all DNAzyme reagents, was characterized and implemented for
28 the simultaneous detection of X and Y single strand DNA fragments.

29 **Keywords:** DNAzyme, amelogenin, paper substrate, gene detection, colorimetric, paper based
30 microfluidics.

311. INTRODUCTION

32 There is an increasing interest in producing new technology for fast detection of specific DNA
33 sequences at the point of need for forensic analysis, environmental monitoring and rapid biomedical
34 diagnosis applications. Widely known methodologies such as qPCR or PCR-capillary electrophoresis
35 are mainly based on the amplification of DNA samples to achieve enough quantity of DNA to be
36 analyzed ¹. These methods require specialized procedures, equipment and personnel to carry out the
37 analysis.

38 On the other hand, point of care tests (POC) ² refer to devices that analyze, with minimum intervention
39 of the user, small volume of samples at the point of necessity in short times. Moreover, they are
40 usually designed to be mass-fabricated at low costs ³. Microfluidic paper-based analytical devices
41 (μ PADs) constitute suitable platforms for the development of POC systems since they fulfil all the
42 previously described characteristics ^{4,5}. Several paper-based types of materials, such as cellulose and
43 nitrocellulose, can be easily adapted to serve as μ PADs ⁶⁻⁸. Due to their fibers distribution, liquid can
44 flow inside of the material by imbibition, enabling the movement of liquids through its matrix without
45 the requirement of external pumps ⁹. Wax-printing, which involves the generation of hydrophobic
46 walls within the paper matrix by printing and melting of wax, is proposed as the best methodology
47 for the fabrication of μ PADs due to the easiness of the procedure and the capacity to be escalated to
48 mass-production ^{8,10}. Furthermore, μ PADs are disposable, becoming a more environmentally friendly
49 option than other microfluidic devices ¹¹. A number of μ PADs including, HIV Tests (Home Access,
50 USA), paper ELISA, and low-cost colorimetric diagnostic assays have been already described ^{12,13}
51 and commercialized.

52 Traditionally, available paper-based devices are centered on the detection of big molecules such as
53 proteins. Nevertheless, the development of novel μ PADs for nucleic acid detection, such as nucleic
54 acids capture on cellulose filter paper with *in situ* PCR ¹⁴, loop-mediated isothermal amplification of

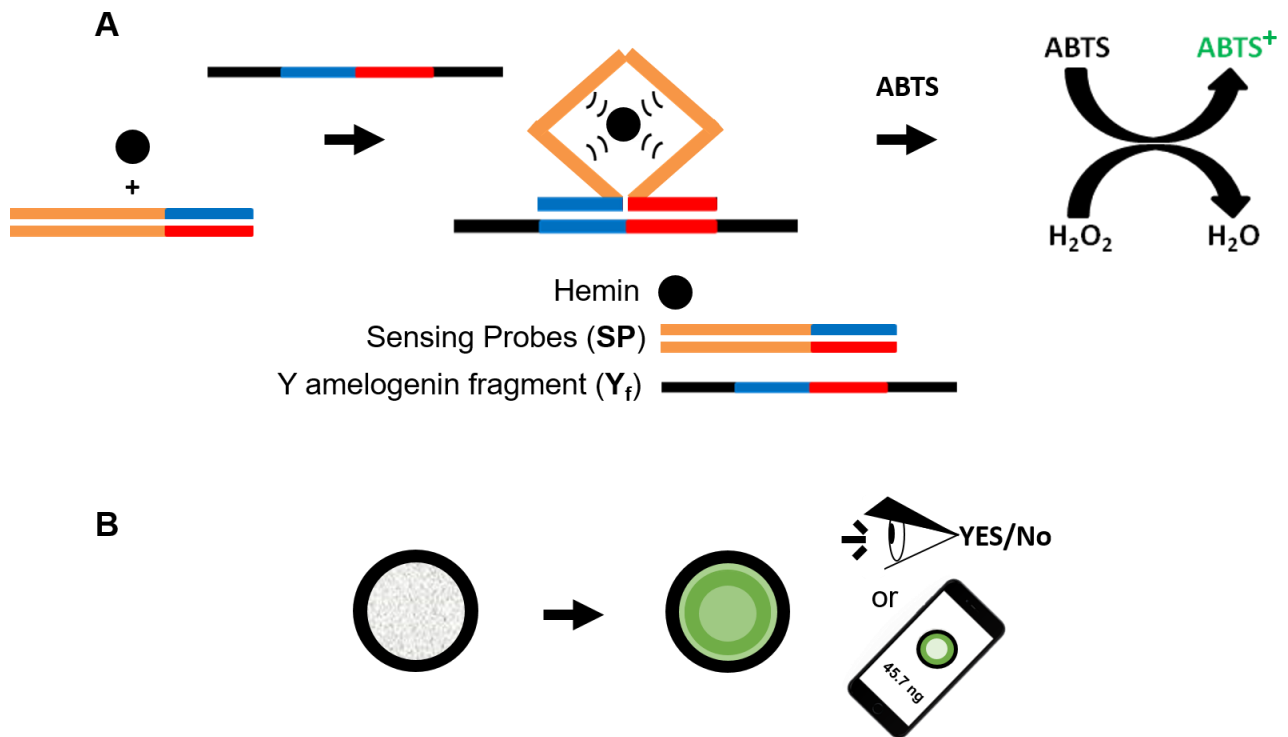
55 several DNA targets on paper device¹⁵ and colorimetric DNA detection through gold nanoparticles
56 capture and agglomeration in paper device¹⁶, among others, is a hot topic at the moment.

57 DNAzyme is composed of a DNA oligonucleotide, capable of performing a chemical reaction, often
58 a catalytic reaction. In particular, DNAzymes sensing probes based on G-rich sequences, known as
59 G-quadruplex, are gathering special attention due to their simplicity, high sensitivity and selectivity
60 for the recognition of small molecules^{17,18}. The G-quadruplex creates a special hairpin on the DNA
61 sequences forming a complex with peroxidase activity, enabling the oxidation of substrates like 2,2'-
62 azino-bis(3-ethylbenzothiazoline-6-sulphonic acid) (ABTS) in the presence of hydrogen peroxide
63 (H₂O₂), producing an appreciable color change of the solution where dissolved¹⁹⁻²¹. By controlling
64 the design of the DNAzyme sequence, it is possible to specifically detect ribonucleic acid (RNA),
65 DNA, proteins and metal ions^{22,23}. In this regards, designing a DNAzyme able to hybridize to a single
66 strand DNA (ssDNA) sequence allows for the specific detection of the sequence of interest²⁴⁻²⁶.

67 The use of DNAzymes in paper substrates was recently presented for the design of novel analytical
68 devices. Zhang Y. *et al.*²⁷ demonstrated that hemin/G-quadruplex DNAzymes can be used on a paper
69 support for the detection of potassium cations. Monsur Ali M. *et al*²⁸ achieved the genetic detection
70 of *Escherichia coli* RNA on a paper-based filter support using DNAzyme catalytic reactions.
71 However, their detection method was based on fluorescence, which required of specific
72 instrumentation, thus increasing the complexity and cost of the methodology, whereas a colorimetric
73 detection could be a better choice to make²⁹.

74 Here, a colorimetric methodology for the detection of ssDNA sequences by specific DNAzymes in
75 solution, a Whatman filter paper 1 support and a paper-based device is presented, Figure 1. The
76 formation of the DNAzyme in the presence of the analyte allows the oxidation of ABTS producing a
77 green color, ABTS⁺, which is detected with the naked eye and quantitatively, using colorimetric
78 image analysis from a mobile camera. As a proof of concept, a synthetic fragment corresponding to

79 Y amelogenin gene, which differs from its X counterpart by 6 base pairs, was designed as a model to
80 test ssDNA detection³⁰⁻³².



81
82

83 **Figure 1.** Schematic representation of the DNAzyme formation and specific Y amelogenin fragment
84 (Y_f) identification of paper support. A) DNA sensing probes and hemin specifically bind to Y_f to
85 form a three dimensional structure with catalytic activity able to oxidize ABTS in the presence of
86 H₂O₂, producing green colored ABTS⁺. B) DNAzyme reaction on paper support produces an
87 appreciable green color in the presence of the Y_f thanks to the oxidation of ABTS. The reaction can
88 be monitored colorimetrically.

89

902. EXPERIMENTAL

91 2.1. Reagents and Equipment

92 **Oligonucleotides.** DNA sequences (DNA-active, DNA-sensing probes and Y and X amelogenin
93 fragments) were synthesized by Integrated DNA Technologies (IDT, Belgium).

94 **Reagents.** Hemin, ABTS, dimethyl sulfoxide (DMSO), HEPES buffer, sodium chloride, Triton X-
95 100 and dimethylsulfoxide were purchased from Sigma-Aldrich (Spain). H₂O₂ was obtained from E.
96 Merck (Germany). Potassium chloride was purchase from Panreac (Spain).

97 **Equipment.** *Solution assays* were performed in a microtiter plate CORNING 96 wells and
98 colorimetric values were recorded with a Beckman Coulter DTX 880 Multimode Detector.

99 *Paper assays* were performed on Whatman filter paper 1(Sigma Aldrich, Spain) and Hi-Flow Plus
100 HFC 12004 nitrocellulose (EMD Millipore, Ireland). The paper assays and the paper-based device
101 were printed by a Xerox ColorQube 8570 wax printer and the wax barriers were generated with an
102 oven, set at 125 °C for 5 min.

103 Photographs were taken by a Sony Xperia Z3 D6603 mobile phone camera (20.7 MP, f/2.0, 25 mm
104 (wide), 1/2.3", 1.12 μm, AF) or a LaserJet Pro 400 MFP scanner (Scanning Method: Flatbed; Scanner
105 Type Flatbed: DF; Bit Depth: 30-bit).

106
107 **Buffer solution:** HEPES 2.5 10⁻² M (Sigma-Aldrich, Spain), Potassium Chloride 2 10⁻² M (Panreac,
108 Spain), sodium chloride 0.2 M (Sigma-Aldrich, Spain), Triton X-100 0.05 % and dimethylsulfoxide
109 1 % (Sigma-Aldrich, Spain), at pH 7.4 with sodium hidroxide (Sigma-Aldrich, Spain) in water, was
110 freshly prepared in the laboratory.

111 112 **2.2. DNAzyme performance in solution**

113 For specific Y_f detection, 1 μL of a mix of DNA-sensing probes 1 and 2 (SP1 and SP2, Scheme 1)
114 (100 μM each probe) and 47 μL of buffer solution were pipetted in the microtiter plate and incubated
115 for 1 h. Then, 1 μL of hemin solution (100 μM) was added and incubated for 1 h. After that, 1 μL of
116 either Y_f or X amelogenin fragment (X_f) solutions ranging from 50 to 150 μM were added and
117 incubated for 30 min (n = 3). Finally, 25 μL ABTS solution (28.8 mM) and 25 μL H₂O₂ solution (1.76
118 mM) were pipetted. The concentrations were previously optimized for this reaction, see supporting
119 information SI-1, as well as the type of probe used for the assay, see supporting information SI-2.

120 The absorbance was measured at 415 nm every 5 min for 30 min using a Beckman Coulter DTX 880
121 Multimode Detector.

Y amelogenin fragment (Y_f) CCCTGGGCTCTGTAAAGAATAGTGGGTGGATTCTTCATCCCAAATAAAGTGGTTCTCAAGTGGTCCCAATTTT
X amelogenin fragment (X_f) CCCTGGGCTCTGTAAAGAATAGTGTGTTGATTCTTTATCCCAGATGTTTCTCAAGTGGTCCCTGATTTACAGTTCC
DNA-active TTTGGGTAGGGCGGGTAGGG
DNA-sensing probe 1 (SP1) CGGGATGGGTTT TGGGATGAAGATCCA
DNA-sensing probe 2 (SP2) GAAACCACTTTATAAAGGGTAGGG

122

123 **Scheme 1** DNA sequences: Y_f, X_f, DNA-active and sensing probes 1 and 2 (SP1 and SP2) used in
124 this study.

125

126 2.3. Y amelogenin fragment discrimination and analysis in paper substrate

127 The Y_f detection through DNAzyme was carried out in a paper substrate. Therefore, circle-shaped
128 structures (1 cm, inner diameter after heating) were wax-printed to set the sensing areas in the two
129 paper supports. After printing, the circles were heated in the oven for 5 min at 125 °C. The heating
130 step is necessary in order to melt the wax trough the paper and generate the hydrophobic barriers^{8,10}.

131

132 *Determination of the most adequate paper substrate: Whatman filter paper 1 or nitrocellulose*
133 *membrane*

134 In order to test the most suitable type of paper for Y_f detection, the oxidation of ABTS with a
135 DNAzyme-active solution, serving as a control, was carried out in Whatman filter paper 1 and
136 nitrocellulose membrane. First, 1 cm wax-circles were printed on both type of substrates. Then, 0.5
137 μL of DNA-active (100 μM), 0.5 μL of hemin (100 μM) and 24 μL of buffer solutions were pipetted
138 and incubated for 1 h, at room temperature (RT). The concentrations of DNA-active and hemin were
139 previously optimized for this reaction in solution, see supporting information SI-1. Finally, 12.5 μL
140 of ABTS solution (28.8 mM) and 12.5 μL of H₂O₂ solution (1.76 mM) were added in order to generate
141 the color assay. The concentrations of ABTS/H₂O₂ were previously optimized for this reaction in

142 solution, see Supporting Information SI-1. Photos were taken with a mobile phone camera before the
143 addition of ABTS and H₂O₂ solutions and 5 and 60 min after addition.

144

145 *Detection and quantification of Y_f on the paper substrate: Whatman filter paper 1.*

146 0.5 cm wax circles (inner circle, after melting) were wax-printed. 0.25 μL of a 1:1 mix of SP1 and
147 SP2 (100 μM each), 0.25 μL of hemin-stock solution (100 μM), 6 μL of ABTS-stock solution (40
148 mM) and 7 μL of buffer solution were mixed in an Eppendorf. The resulting mixture was pipetted on
149 the paper wax-circle. After overnight storage at 4 °C, either 0.25 μL of Y_f (25, 50, 100 or 200 μM),
150 X_f (25, 50, 100 or 200 μM) or buffer solution were pipetted in the wax-circles (n = 3) and 30 min
151 incubated at RT. Then, 6 μL of H₂O₂ (2.4 mM) was pipetted in all wax-circles. Images were taken by
152 a mobile phone camera after 5 min.

153

154 *Detection and quantification of Y_f on the paper substrate in the paper-based device*

155 For the specific detection of Y_f fragment in the paper-based device (See Supporting Information SI-
156 3, for device specifications). An optimized mixture of the reagents, ABTS and H₂O₂ were used for
157 these experiments, see supporting information SI-1 and SI-2.

158 0.25 μL of a 1:1 mix of SP1 and SP2 (100 μM), 0.25 μL of hemin-stock solution (100 μM), 6 μL of
159 ABTS-stock solution (21.6 mM) and 7 μL of buffer solution were pipetted in the device reservoirs
160 (Y and X wax-circles,), in three different devices. The mix of this reagents forms the Y_f-DNAzyme
161 cocktail. After overnight storage at 4 °C, 0.25 μL of Y_f and 0.25 μL of X_f (0.5 μM) were pipetted on
162 their respective detection zones and incubated for 3 min at RT. Finally, 12 μL of H₂O₂ solution (1.32
163 mM) was pipetted on the inlet (center of the device) and the solution was left to run until it reached
164 both detection zones. Photographs were taken by the mobile phone camera 5 min after H₂O₂ solution
165 reached the detection zones.

166

167 **2.4. Image and data analysis**

168 All images were transformed to 8-bit and grey intensities (corresponding to the total color intensity
169 ^{12,33}). They were analyzed using Image-J software. Paper background intensity was subtracted from
170 all data. Data plotting and statistical analysis were carried out in Excel and Origin Pro 2018.

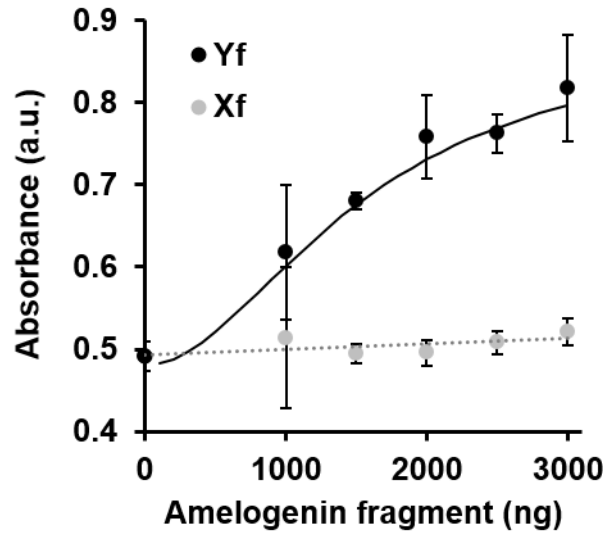
171

1723. **RESULTS AND DISCUSSION**

173 In order to investigate the use of DNazymes on paper substrates for naked eye detection of ssDNA
174 fragments and quantification by image analysis with a mobile camera, a DNzyme for the detection
175 of a specific fragment, Y_f , of the Y amelogenin gene was designed. Then, a reaction cocktail
176 comprised of hemin, ABTS and two sensing probes (SP1 and SP2, which bind specifically to Y_f) was
177 developed; see supportive information SI-1 and SI-2 for detailed information of the characterization
178 of the different components of the cocktail.

179 To evaluate the sensitivity and selectivity of the designed DNzyme assay for the detection of Y_f
180 *versus* X_f , the DNzyme cocktail was incubated with different concentrations of Y_f and X_f ranging
181 from 0.5 to 1.5 μ M (corresponding to a range between 1000 and 3000 ng of ssDNA per sample), and
182 the absorbance of the final solutions were measured in a conventional microtiter plate reader after 5
183 min of incubation. An increase in absorbance was appreciated over time and when increasing the
184 concentration of Y_f . On the other hand, the absorbance values for the reaction performed with the X_f
185 remained constant over time and for all concentrations investigated, Figure 2. This results
186 demonstrate the specific reaction of the DNzyme with the Y_f fragment, while the same DNzyme
187 was not able to detect the X_f fragment, in solution.

188



189

190 **Figure 2. Graphical representation of Y_f detection in solution.** Y axis indicates the absorbance
 191 obtained from the oxidation of ABTS with different amounts of Y_f and X_f at 5 min. Fitting curve of
 192 Y_f was done through the equation $y = 0.89 - (0.41 / (1 + (x/1.58)^{1.89}))$, $R^2 = 0.9909$. Error bars mean \pm SD
 193 (n = 3 samples per experimental condition).

194

195 Y_f absorbance values were fitted to a 4 parametric logistic (4PL) curve, which was used before in
 196 immunoassays and binding ligand assays³⁴. A limit of detection (LoD) of 655 ng of Y_f was calculated
 197 following the equation 1,

$$LoD = mean_{blank} + 3 SD_{blank} \quad (eq. 1)$$

199 where $mean_{blank}$ is the absorbance value of the reaction mix without Y_f and X_f, and SD is the standard
 200 deviation (n = 3) of the mean value of the blank. And a limit of quantification (LoQ) of 1462 ng of
 201 Y_f was calculated following the equation 2,

$$LoQ = mean_{blank} + 10 SD_{blank} \quad (eq. 2)$$

203 where $mean_{blank}$ is the absorbance value of the reaction mix without Y_f and X_f, and SD is the standard
 204 deviation (n = 3) of the mean value of the blank. High SD values were observed at low amounts of
 205 Y_f and X_f, (1000 ng). This higher SD corresponds to the experimental error, which comes from the
 206 number of dilutions needed to reach this 1000 ng solution from the X_f and X_f main stock solution.

207 This error could be reduced by increasing the number of experiments per concentration. Therefore,
208 the calculated LoQ value was taken as more certain value in the analysis of the dynamic range of the
209 reaction. The dynamic detection range (1462 – 3000 ng) is limited by the increasing signal of the
210 background over time, provided by the spontaneous ABTS oxidation in the reaction mix, as
211 previously described in literature ^{35,36}.

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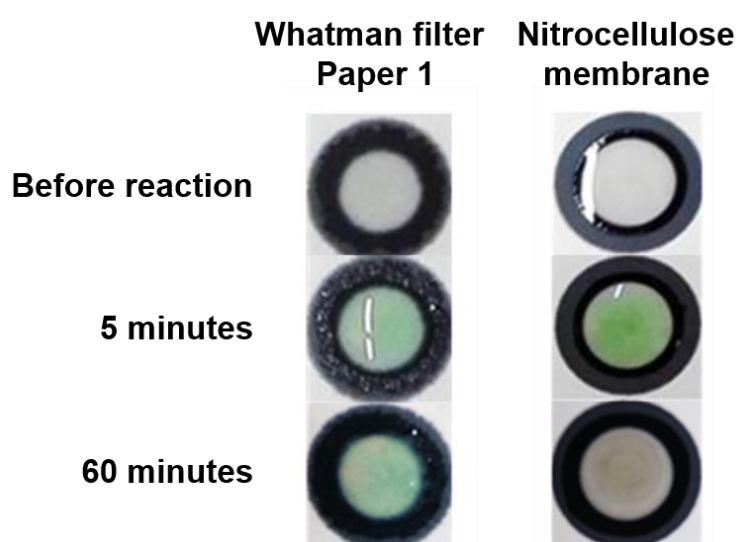
213 **3.2. Selectivity and Sensitivity of the assay in the Whatman filter paper substrate**

214 In order to get an assay that is suitable for a paper-based device configuration, it is necessary that all
215 the reagents of the assay are paper compatible, can be preloaded and, if possible, stored for long
216 periods of time and at different ambient conditions (*e.g.* light and temperature). Therefore, the
217 stability of the assay over time, was investigated for two light conditions (daylight and dark at RT) as
218 well as at 4 °C in the dark for 24 h storage; see Supporting Information SI-4. The absorbance values
219 of the assay in solution were lower than those carried out without storing for 24 h, but they clearly
220 allow the detection of the Y_f in the same range of concentrations and times than when no storage was
221 carried out. Therefore, these protocols can be implemented during the fabrication of a μ PAD, with
222 all the reagents preloaded and stored in paper.

223 The assay solutions stored at daylight and at RT for 24 h did not significantly have an increase in
224 color due to ABTS oxidation, see section Figure SI-4 B and C. Moreover, the assay kept at 4 °C in
225 the dark presented higher absorbance values due to a better preservation of the reagents at low
226 temperature, see Figure SI-4 D, being this protocol (4 °C in the dark) the one used in experiments
227 hereafter.

228 After confirming the possibility to detect Y_f using the designed DNAzyme in solution, the conditions
229 to perform the detection of the assay on a paper substrate were investigated. Two different papers,
230 nitrocellulose membrane and Whatman filter paper 1, which are commonly used for the fabrication
231 of lateral flow assays and μ PADs, were investigated. Using a wax printer, 1 cm diameter wax-circles

232 were printed on both types of substrates. Initial experiments on paper were performed using a positive
233 control mix comprised of DNA-active sequence and hemin group to form a DNAzyme without the
234 presence of the target analyte (DNAzyme-control mix). First, 0.5 μL of DNAzyme-control mix and
235 24 μL of buffer solution were pipetted and incubated for 1 h at RT in the paper substrates. Then, 12.5
236 μL of ABTS solution (28.8 mM) and 12.5 μL H_2O_2 (1.76 mM) solution were added. Photos of the
237 substrates were taken with a mobile phone camera 0, 5 and 60 min after addition of ABTS and H_2O_2
238 (Figure 3).



239

240 **Figure 3. Pictures of DNAzyme control mix in Whatman filter paper 1 and nitrocellulose**
241 **membrane.** Photographs were taken at times 0, 5 and 60 min after ABTS/ H_2O_2 solutions addition.
242 Green color indicates the presence of ABTS^+ .

243 The Whatman filter paper 1 presented a better performance than the nitrocellulose. The Whatman
244 paper 1 had high ability to absorb the reagents and integrate them within the paper fibers, with the
245 reaction taking place in the paper substrate. After the assay was carried out (addition of ABTS/ H_2O_2
246 solutions), the paper substrate kept the green color, which is the result of the enzymatic reaction, even
247 after 60 min (dry paper). Considering this result it can be speculated that the fibers of the Whatman
248 paper 1 were able to stabilize the ABTS^+ over time, up to 60 min.

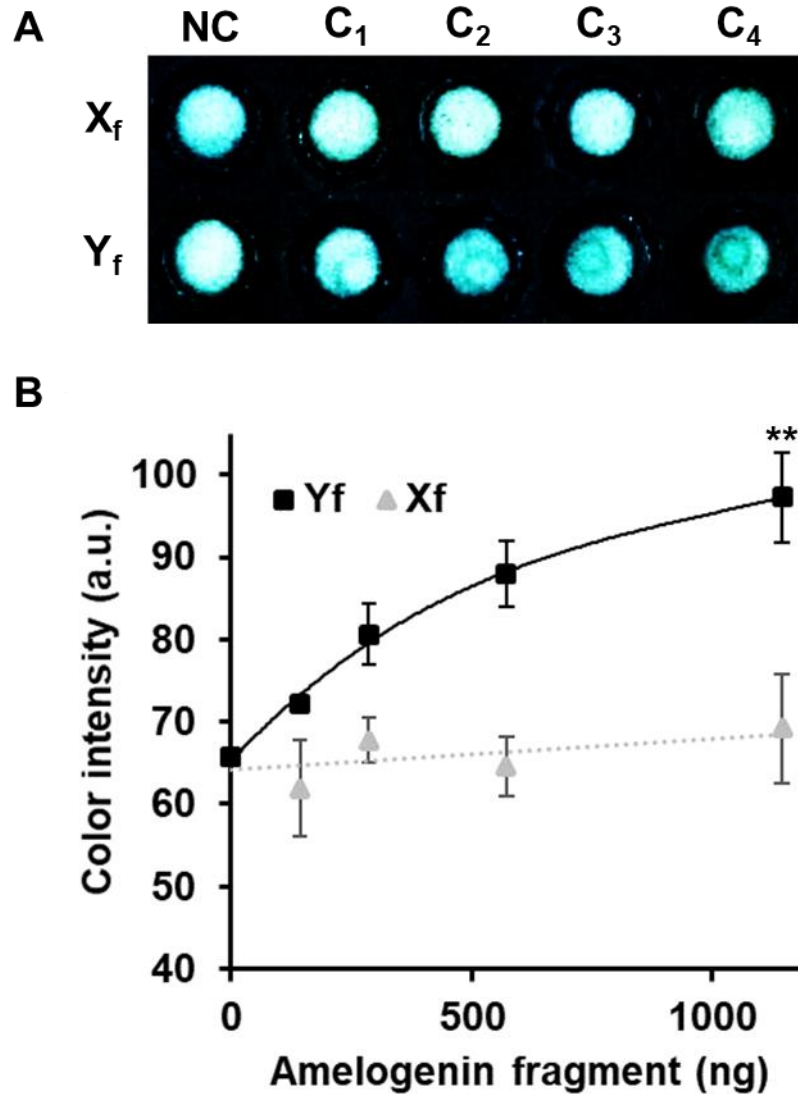
249 On the other hand, the nitrocellulose paper substrate did not absorb the reagents, remaining in solution
250 in the drop, held by the wax-printed-circle. After the assay was carried out (addition of ABTS/ H₂O₂
251 solutions), the oxidation reaction took place in solution and not in the fibers of the nitrocellulose
252 paper. The oxidized ABTS⁺ molecules (green color) were not stabilized by the paper substrate and
253 continued the oxidation process. Under prolonged oxidative conditions, ABTS⁺ suffers over-
254 oxidation into ABTS²⁺, which is not stable in aqueous solution and decompose into several uncolored
255 by-products, losing its characteristic green color ³⁷⁻³⁹. In view of these results, the Whatman filter
256 paper was selected as the right support for the DNAzyme reaction.

257

258 The detection of Y_f on Whatman paper 1 was evaluated using the DNAzyme on a 2 by 5 array of 0.5
259 cm diameter (internal diameter, after heating) wax-printed circles. 12 μL of a solution containing 0.25
260 μL of the sequences SP1 and SP2 (1 μM) with 0.25 μL of hemin solution (1 μM), 6 μL of ABTS (40
261 mM) and 7 μL of buffer were pipetted and subsequently dry over night at 4 °C in the dark in each
262 circle. Then, 0.25 μL of solutions with increasing concentrations of Y_f or X_f ranging from 0 (negative
263 control) to 200 μM were loaded in each reaction circle and let dry for 30 min at RT. Finally, 6 μL of
264 H₂O₂ (2.4 mM) were added to each reaction circle. After 5 min a green color was appreciated in all
265 the reaction circles, by naked eye. A pronounced “coffee ring” effect was identified in all the circles
266 containing Y_f, while this coffee ring was not appreciated in the circles containing X_f. The shape of
267 the coffee ring corresponded with the shape of the drop of X_f previously loaded on the substrate. The
268 reason for the formation of this ring can be explained by the concentration of the reagents during the
269 drying process of the sample drop pipetted in the paper support ⁴⁰. Additionally, a darker green color
270 intensity was appreciated in those circles containing higher concentrations of Y_f. Differently, the
271 negative control and the circles containing X_f exhibited the same light green color coming from the
272 uncontrolled residual oxidation of ABTS, Figure 4A.

273 Mixtures of Y_f and X_f were also investigated in the 2 by 6 array of wax-printed circles. The same
274 green color intensity than samples containing only Y_f were obtained, indicating that the presence of
275 the X_f does not inhibit the formation of the Y_f - DNAzyme (supporting information Figure SI-5).

276 The quantification of the DNAzyme reaction was done analyzing the color intensity of the circles
277 (green color intensity) by just taking pictures of the array with a mobile phone color camera. The
278 whole area of the reaction circle was selected, without differentiating the presence or absence of the
279 coffee ring. Figure 4 B shows the plot of the color intensity *versus* Y_f and X_f concertation. As
280 qualitatively observed by eye before, the green color intensity increased with the concentration of Y_f ,
281 while it remained constant for the assay containing X_f . There was a significant difference in the signal
282 obtained with and without Y_f in all the cases, indicating that this assay could be used in paper for
283 quantification of specific fragments of ssDNA. Y_f values were fitted to a 4PL curve. A LoD of 45.7
284 ng and a LoQ of 172 ng of Y_f were calculated using eq. 1 and eq. 2, respectively, obtaining a dynamic
285 range of 172 - 1200 ng, which is much lower than in solution. Therefore, transfer the reaction to a
286 paper support greatly improved the LoD, reducing 14 times the minimum amount of Y_f that can be
287 detected. This improvement can be attributed to the higher concentrations of reagents with respect to
288 the detection zone available in the paper substrate. Moreover, after quantification it can be concluded
289 that at least 143 ng of Y amelogenin fragment could be detected by naked eye.



290

291 **Figure 4. Pictures and graphical representation of Y_f detection on Whatman filter paper 1. A)**

292 Picture of the 2 by 6 array printed in paper after 5 min DNAzyme reaction. C₁, C₂, C₃ and C₄ refers to

293 the different concentrations of Y_f or X_f solutions added to the paper substrate (25 (143 ng), 50 (286

294 ng), 100 (572 ng) and 200 (1144 ng) μM, respectively) and NC is the negative control. B) Graphical

295 representation of the color intensity obtained from the picture taken by the mobile camera 5 min after

296 reaction. Fitting curve of Y_f was done through the equation $y = 97 - (31.44 / (1 + (x/51.7)^{1.23}))$, $R^2 =$

297 0.9917. Error bars mean ± SD (n = 3 samples per experimental condition). Statistical significance;

298 paired two-tailed t-test (** p < 0.01).

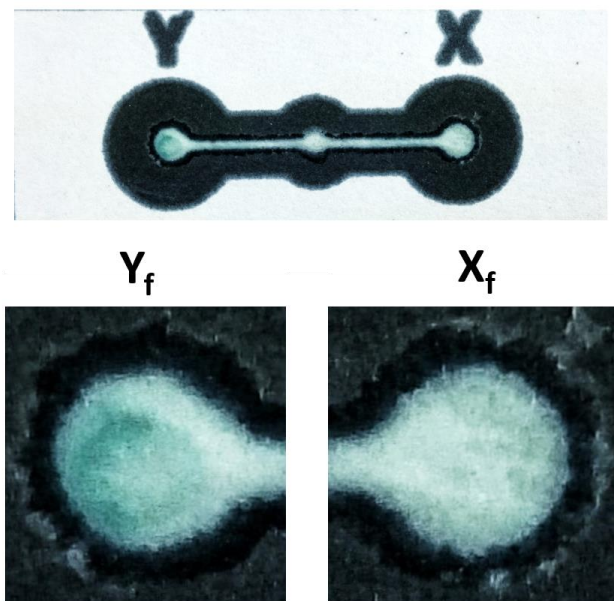
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300 In order to fabricate a paper-based device useful in real applications, it is desirable that the reagents
301 are stable for long period of time embedded in the device, preferably in a controlled environment.
302 Therefore, the stability of the reagents (SP1, SP2, hemin and ABTS) forming the assay was tested by
303 storing them at 4 °C from 24 h to 48 h and 7 days, see Supporting Information Figure SI-6. For the
304 three times investigated, the “coffee ring” was clearly visible in all samples containing Y_f while the
305 green color was imperceptible with the naked eye for the X_f samples, See Figure SI-6A. Pictures of
306 the wax-circles were taken and analyzed, see Figure SI-6B, allowing for the quantification of the
307 ABTS oxidation process. The values obtained for each experiment are comparable within the error
308 ($n = 3$ experiments), indicating that the reagents are well preserved in the paper substrate maintaining
309 their chemical properties at least for one week, under refrigeration.

310

311 **3.3 DNA samples analysis on a paper-based device.**

312 A final test was performed to evaluate the applicability of Y_f detection on a paper –based device for
313 the simultaneous analysis of Y_f and X_f samples. As a proof of concept a microfluidic paper-based
314 device was designed and wax-printed. It has a channel with two analysis zones, one at each edge of
315 the channel and an inlet for reagent loading in the middle of the channel (Figure 5). The Y_f -
316 DNAzyme cocktail was pre-loaded in both analysis zones at kept for 24 h at 4 °C (dark conditions).
317 Then, two samples containing 570 ng of Y_f or X_f were loaded in each analysis zone and let dry for 30
318 min. Finally, 12 μ L of H_2O_2 (1.32 mM) were pipetted in the inlet of the device. The H_2O_2 flowed
319 equally in both detections reaching the analysis zones in less than 10 s. The ABTS oxidation reaction
320 took place and a clear green coffee ring appeared only in the analysis zone containing Y_f after 5 min
321 (supporting information video in SI-7) while the intensity of the green color in the analysis zone
322 containing the X_f sample was very low.



323

324 **Figure 5.** Paper-based device for Y_f detection. Pictures were taken 5 min after the H_2O_2 solution

325 reached the analysis zones. Detection circles correspond to sample loading of Y_f (left) and X_f (right).

326 Increased green intensity relates to increasing presence of $ABTS^+$.

327

3284. CONCLUSIONS

329 Rapid tests for DNA detection have application in forensic practice, biomedical diagnostics, and

330 environmental monitoring among others. Current ways to analyze DNA sequences often require long

331 procedures at centralized labs, and there is a need to developed technologies that enable in-field and

332 rapid sample analysis and detection of the DNA of interest.

333 An optimized methodology for easy and in-field ssDNA detection in a paper-based device was

334 presented. By using a DNAzyme specific to the desired DNA sequence, a label-free, easy to use and

335 colorimetric based detection method for ssDNA was developed. This reaction was optimized in

336 solution in order to obtain a distinguishable specific and positive signal in a short time with a LOD

337 of 655 ng.

338 The assay developed in solution was then transferred to a paper support and to a paper-based device.
339 Two types of paper were tested, Whatman filter paper 1 and nitrocellulose membrane, both highly
340 used in paper microfluidics and/or DNA research⁶⁻⁸. It was found out that Whatman filter paper 1
341 was more adequate for the DNAzyme reaction and thus was chosen as the desired substrate. The
342 DNAzyme reaction on paper enabled the unequivocal colorimetric identification of the Y_f ssDNA
343 fragment by qualitative detection, with the naked eye, and quantitative determination by image
344 analysis. At least 143 ng of Y amelogenin fragment could be detected by naked eye, while 45.7 ng
345 could be detected by the quantification of the color intensity values from a picture of the paper
346 substrate. The transfer of the reaction to a paper support significantly reduced the LoD, improving
347 the sensitivity of the reaction.

348 Multisampling analysis on a paper-based device pre-loaded with the DNAzyme cocktail was carried
349 out. Although the configuration of the device is not optimized, two analytes, Y_f and X_f, were analysed.
350 The analysis of different genes from a single individual would need of further validation and
351 optimization of the process, including device flow control and prevention of cross-contamination of
352 samples. Furthermore, the devices were stored for up to 7 days at 4 °C before use, indicating that they
353 could be stored and used at the point of care

354 This work constitutes the first steps into what could become an in-field paper-based device for several
355 applications. In the case of fast typifying of human samples for forensic analysis it should be noted
356 that human samples are more complex than the proposed 74 bases synthetic sequence, usually
357 presenting the genome in a double strand structure with high packing, safeguarded inside the cell's
358 nucleus. Also, while the detection limits achieved are optimal for our synthetic sequence, the total
359 genetic material found per human cell can be very low (between 3 and 6 picograms per cell) and the
360 amount of cells that can be found in human samples can vary significantly, depending on the sample
361 origins⁴¹, being this scenario a limitation of this technology in its current configuration. Therefore,
362 for the incorporation of real human samples, research on the integration and the optimization of

363 extraction methods (Chelex 100⁴²) and DNA denaturalization methods (chemical denaturalization
364⁴³) inside the paper supports would be necessary. Conventional techniques for human genome
365 analysis, such as qPCR, only require nanograms of whole genomes to produce data with high
366 specificity. Our proposed methodology would extremely benefits from complementary amplification
367 methods of the genetic material, such as isothermal amplification, that can be incorporated in the
368 future to an in-field device for the successful detection of human DNA^{44,45}. The technology presented
369 in this manuscript could also be applied in the field of biomedicine for the detection and diagnosis of
370 bacterial and viral infections, which present shorter genomes and more variety of DNA structures,
371 including ssDNA^{46,47}.

372

373 **ACKNOWLEDGEMENTS**

374 This work was supported by Gobierno de España, Ministerio de Economía y Competitividad, with
375 Grant No. BIO2016-80417-P; funding from Basque Government (Grant: IT1271-19) and European
376 Union funds: DNASURF (H2020-MSCA-RISE-778001). E.A.-H. acknowledges funding from the
377 Basque Government, Department of Education, for predoctoral fellowship 2016.

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