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## Numerical modeling of graphene-coated fiber optic surface plasmon resonance biosensor for BRCA1 and BRCA2 genetic breast cancer detection

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Abstract. A numerical illustration of a hybrid design and numerical analysis of graphene-coated fiber-optic surface plasmon resonance (SPR) biosensor for BRCA-1 and BRCA-2 genetic breast cancer detection is provided. Two specific mutations named 916deITT in BRCA-1 gene and 6174deIT in BRCA-2 gene are being selected for detection of breast cancer numerically. This sensor is based on attenuated total reflection (ATR) method to detect individual point mutations in BRCA-1 and BRCA-2 genes. Based on the numerically obtained results, a momentous change is present in the SPR angle (minimum 35% more) and surface resonance frequency (SRF) (minimum 36% more) for probe DNA with various concentrations of target DNA corresponding to the mutation of the BRCA-1 and BRCA-2 genes. The variation of the SPR angle and SRF for mismatched DNA strands in BRCA-1 and BRCA-2 genes is quite negligible, whereas that for complementary DNA strands is considerable. This considerable change is essential for proper detection of genetic biomarkers (916delTT and 6174deIT) for early breast cancer. To the best of our knowledge, this is the first demonstration of such an adept biosensor for detecting BRCA1 and BRCA2 genetic breast cancer. Here, we used graphene as bimolecular acknowledgement element for improving sensor performance. At the end of the article, the performance in terms of sensitivity is analyzed. Therefore, the proposed biosensor opens a window toward detection of early detection of BRCA-1 and BRCA-2 genetic breast cancers. © 2019 Society of Photo-Optical Instrumentation Engineers (SPIE) [DOI: 10.1117/1 .OE.58.3.037104]

Keywords: attenuated total reflection; BRCA-1; BRCA-2; cancer; DNA hybridization; graphene; surface plasmon resonance; surface resonance frequency; 6174delT; 916delTT.

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#### 1 Introduction

The genetic tendency is responsible for 5% to 10% breast cancer and over 75% to 80% of inherited breast cancer cases are reported owing to mutations happening in BRCA1 and BRCA2 genes. 1,2 These genes are tumor suppressor associated with double strand DNA (dsDNA) repair accounts for abnormal uncontrolled tumor cell growth. Exceeding 400 different mutations in BRCA1 and BRCA2 genes leads to tumor growth and thus finally cancer.<sup>3</sup> Real-time detection of polymerase chain reaction (PCR) technologies has become the most traded and used for mutation detection with high specificity and sensibility.<sup>3</sup> Two prime factors, one is genetic and another is proteomic, are classified for breast cancer biomarkers. For example, BRCA1, BRCA2, and P53 are the genetic and ATM, CA125, CA153, CEA, CHEK2, HER-2, PTEN, and STK11 are the proteomic markers. These are responsible for breast cancer risk.<sup>4,5</sup> Every year above 350,000 deaths and 192,000 cases are reported due to breast cancer.<sup>4</sup> As yet, ultrasound, mammography, PCR screening, and other approaches have been applied to identify the diagnosis of breast cancer. This cancer cell is difficult to identify owing to tiny in size.<sup>4,6</sup> Thus, effective,

A great deal of attention on surface plasmon resonance (SPR)-based biosensors as a leading optical sensing technology has been achieved by the researchers because of its applications in numerous fields, including medical diagnostics, biomolecule detection, biochemical detection, environmental monitoring, proteins—DNA hybridization, and DNA—DNA hybridization. To DNA hybridization is regularly used biomolecular method in which the level of genetic relationship between pools of DNA sequences to obtain the genetic gap between two organisms has been estimated. This biochemical process of identifying DNA nucleotide bonding in proper orientation is very important for medical diagnostics. This technology is being used to obtain information on DNA molecular bonding and, from here, BRCA1 and BRCA2 breast cancer directly diagnosable.

Fundamentally, SPR is a charge density oscillation between metal—dielectric interface stimulated by TM polarized electromagnetic waves, typically known as surface plasmon waves (SPW). The SPW at the interface between a metal (e.g., Au, Ag, or Al) and a dielectric is used by the SPR biosensor to investigate the interactions between biomolecules. And these interactions are accomplished by practice of optical fiber-based SPR sensors. A fiber-optic

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precise approaches to breast cancer recognition and monitoring are immediately required.

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SPR sensor for wavelength interrogation mode has lots of advantages, such as light in weight, compactness, high sensitivity, ease of multiplexing and remote sensing, mechanical flexibility, and the ability to transmit optical signals over a long distance, which allows miniaturization and chemical or biological sensing in inaccessible locations. <sup>10</sup> As a fundamental part of fiber optic SPR biosensor, a thin metallic film (Au, Ag, or Al) is coated with a fiber to distinguish the sensing medium and the dielectric. <sup>12</sup> Here, the sensing layer is used to functionalize the gold film, which consequently improves the absorption of biomolecules. <sup>12–15</sup>

An essential constituent of the SPR sensing device is large surface area-based absorbing material; 13 recently discovered 2-D nanomaterial graphene has satisfied this requirement regarding this Ref. 16. Graphene and graphene oxide facilitate biomolecule adsorption because of their large surface area and rich  $\pi$  conjugation structure, <sup>17,18</sup> making them suitable as dielectric top layers for SPR sensing. 19 Graphene has great fields of applications in advanced electronics, optoelectronics, and plasmonic due to exclusive characteristics of high carrier mobility, high optical transparency, exceptional mechanical flexibility, mechanical strength, 17-22 low resistivity, tunable conductivity, and extreme mode confinement.<sup>23</sup> Recent studies have focused graphene-based diverse applications in optoelectronic devices, including touch panels, ultrafast lasers, solar cells, optical modulators, photodetectors, and polarizers.<sup>24,25</sup> Graphene supports surface plasmons at terahertz frequencies that show larger confinement in comparison with other noble materials and lower losses due to longer electron relaxation times.<sup>23</sup> An important feature of graphene plasmons is tunable via gating or chemical doping.<sup>26</sup> This property can be applied in transformation optic devices, plasmonic switches, and planer lenses.<sup>27</sup>

In this paper, numerical modeling of a graphene-coated fiber optic SPR biosensor for breast cancer detection is reported. Since graphene has prominent characteristics, a graphene monolayer is sandwiched between gold film as an effective light absorption medium. We used  $R_{\rm min} \sim \theta_{\rm SPR}$  (minimum reflectance versus SPR angle) as well as  $T_{\rm max} \sim$  SRF (maximum transmittance versus SRF) as detecting attributors. This sensor is capable to recognize DNA hybridization event by checking the proper orientation of adenine (A), thymine (T), guanine (G), and cytosine (C) between immobilized mutant DNA sequence (sh-DNA) to mutated type complementary DNA (mr-DNA) in BRCA-1 and BRCA-2 gene shown in Figs. 2 and 3. The refractive index

(RI) changes nearby the graphene and gold (Au) owing to the perfectly matched hybridization event between mr-DNA sequences of 916delTT and 6174delT biomarkers with sh-DNA sequence. The RI change will cause to turn a change in the propagation constant of the SPR wave, which can be optically ascertained by ATR method. 9,10 If significant change is present in  $R_{\rm min} \sim \theta_{\rm SPR}$  as well as  $T_{\rm max} \sim {\rm SRF}$  curves, which indicates that perfectly matched complementary DNA hybridization has occurred between sh-DNA and mr-DNA sequences. And this significant change is the symptom for genetic breast cancer. If no significant change is found, it finally indicates no possibility of breast cancer. Phosphate-buffered saline (PBS) solution as bare sensing dielectric medium provides better adsorption of single-strand DNA biomolecules from mr-DNA sequences. 28,29 Our results demonstrated that SPR biosensors are able to detect mismatch sequences related to hereditary breast cancer with high sensitivity. We develop a pathological decision making table, whether there is breast cancer or not, on the basis of change minimum reflectance, SPR angle, maximum transmittance, and SRF. At the end of the article, we show the effect of adding graphene on the change of minimum reflectance, SPR angle, maximum transmittance, and SRF in terms of sensor sensitivity.

This paper is composed as follows: design methodology is given in Sec. 2. Section 2 consists of two subsections: one is mathematical modeling for proposed sensor and another is designated DNA sequences for detection of breast cancer. Section 3 discusses the details of the numerical results and discussions. Last, conclusion is given in Sec. 4.

#### 2 Design Methodology

The proposed graphene implicated fiber optic SPR biosensor design structure is given in Fig. 1. Fiber optic SPR sensor has a core diameter of  $50 \, \mu \text{m}^{10}$  and cladding diameter of  $50.25 \, \mu \text{m}$ . In sensor region, 5-mm cladding is detached and this slice of the cladding is considered to be implicated with a prism, gold (Au) thin film, and graphene with the thickness of dp =  $74.5 \, \text{nm}$ ,  $^{10} \, d_{\text{Au}} = 50 \, \text{nm}$ , and  $d_{\text{g}} = 0.34 \, \text{nm} * \pounds$  (where £ is the total of graphene coatings), respectively. This biosensor sensing region is involved with a total of four layers' Fresnel's structure. The first layer is a SF11 prism having high RI;  $n_p$  of 1.723, the second layer is Au, deposited on the base of the prism having complex RI of  $n_{Au} = 0.1726 + 3.4218i$ . The third layer is a graphene having high complex RI;  $n_g$  of 3 + 1.149106i and the fourth layer is a

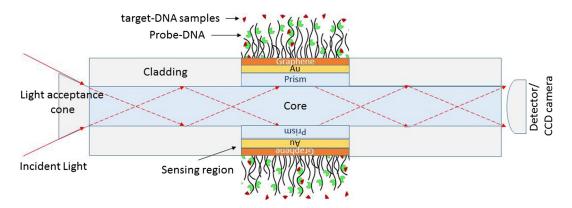


Fig. 1 Figural illustration of the four-layered model for fiber optic SPR biosensor: prism, Au (50 nm), graphene (0.34£ nm), and PBS contains p-DNA and complementary target mr-DNA or wt-DNA samples.

PBS solution as sensing dielectric medium for bare sensor cases. 9-14 The graphene-based biosensor as a adsorbing complementary molecule by the receptor produces a native increase in RI at the graphene to sensing medium intersurface. 30 Once there is molecular interaction, the immobilization surface can be renewed by applying an appropriate chemical with the intention of eliminating target DNA samples except denaturing the probe DNA (p-DNA). 31

In Fig. 1, light acceptance cone is the angle of an incident light hitting the fiber core and totally reflected by the cladding. In our proposed fiber optic SPR sensor, TM polarized He–Ne (wavelength,  $\lambda = 633$  nm) light is used as an incident light. The numerical results suggest that 633 nm allows the increased overall sensitivity of the sensor with the minimal possible Kerr effect.  $^{32}$ 

#### 2.1 Mathematical Modeling for Proposed Sensor

The light is launched into one end and the power available of the other end (dP<sub>T</sub>) at incident angle  $\theta$  can be expressed as follows:<sup>10,11</sup>

$$dP_T \propto \left[ \frac{n_c^2 \sin \theta \cos \theta}{(1 - n_c^2 \cos^2 \theta)^2} \right] d\theta, \tag{1}$$

where  $n_c$  (1.451<sup>10</sup>) is the RI of the fiber core. The normalized transmitted power of p polarized TM light is expressed at the core–prism–metal try interface as follows: <sup>10,11</sup>

$$P_{T} = \frac{\int_{\theta_{cr}}^{\Pi/2} R_{P}^{\left(\frac{L}{D \tan \theta}\right)} \left[\frac{n_{c}^{2} \sin \theta \cos \theta}{\left(1 - n_{c}^{2} \cos^{2} \theta\right)^{2}}\right] d(\theta)}{\int_{\theta_{cr}}^{\Pi/2} \left[\frac{n_{c}^{2} \sin \theta \cos \theta}{\left(1 - n_{c}^{2} \cos^{2} \theta\right)^{2}}\right] d(\theta)},$$
(2)

where L (5 mm) is the length of the sensing region,  $D(50~\mu\mathrm{m})$  is the diameter of the fiber core, and  $\theta_{cr}$  is the critical angle of the optical fiber [expressed as  $\theta_{cr} = a \sin(n_{cl}/n_c)^{10}$ ], where  $n_{cl}$  is the RI of the fiber cladding. The reflected power R can be obtained by four-layered (prism, gold, graphene, and sensing layer) Fresnel's model analysis as follows:  $^{9,12}$ 

$$R = \frac{A + \frac{B}{Z_f} - Z_i \left(C + \frac{D}{Z_f}\right)}{A + \frac{B}{Z_f} + Z_i \left(C + \frac{D}{Z_f}\right)},\tag{3}$$

where,  $Z_i$  and  $Z_f$  are initial (fiber core) and final layer (sensing layer) wave impedances, respectively, in the structure of Fig. 1. Specific layer wave impedance can be determined as follows:<sup>9,12</sup>

$$Z_f = \frac{k_s n_j \cos \theta}{\omega \varepsilon_j^2}.$$
(4)

In Eq. (4),  $n_j$  and  $\varepsilon_j$  are RI and permittivity of j'th (specific) layer,  $\omega = 2\pi c/\lambda$  and  $k_s$  is the light wave vector defined by Eq. (8). The variables A, B, C, and D in Eq. (3) that can be calculated by solving the following matrix equation as follows:  $^{9,12}$ 

$$\begin{bmatrix} A & B \\ C & D \end{bmatrix} = \begin{bmatrix} A_2 & B_2 \\ C_2 & D_2 \end{bmatrix} \begin{bmatrix} A_3 & B_3 \\ C_3 & D_3 \end{bmatrix} \begin{bmatrix} A_4 & B_4 \\ C_4 & D_4 \end{bmatrix} \cdots \begin{bmatrix} A_{N-1} & B_{N-1} \\ C_{N-1} & D_{N-1} \end{bmatrix}.$$
(5)

Equation (5) shows n'th layer system constants. Here, the first layer represents reflectance, and consequently, it is misplaced in the matrix equation. In matrix, each element describes the reflection of the specific layer. These are the subscripted variables A, B, C, and D of each matrix and can be found:  $^{9,12}$ 

$$A_{j} = \cos(d_{j}\omega\varepsilon_{j}^{2}\cos\theta_{j})$$

$$B_{j} = Z_{j}\cos(d_{j}\omega\varepsilon_{j}^{2}\cos\theta_{j})$$

$$C_{j} = \frac{\sin(d_{j}\omega\varepsilon_{j}^{2}\cos\theta_{j})}{Z_{j}}$$

$$D_{j} = \cos(d_{j}\omega\varepsilon_{j}^{2}\cos\theta_{j})$$
(6)

where  $d_j$ ,  $\theta_j$ , and  $Z_j$  are thickness, angle of incidence, and wave impedance of the j'th layer, respectively. The angle of incidence of j'th layer is unknown that can be found as a function of the RI of the initial and j'th layer as follows:  $^{12}$ 

$$\theta_j = \cos^{-1} \left[ \sqrt{1 - \frac{n_i}{n_i} \sin \theta^2 \theta_j} \right]. \tag{7}$$

The light generated an evanescent wave normally known as SPW, when travels through prism and is reflected at the prism–gold–graphene interface. The SPW propagation constant ( $k_{\text{SPW}}$ ) that propagates along the horizontal direction is described in Eq. (8), where  $n_{\text{Au}}$  is the RI of the gold and  $n_{\text{S}}$  is the RI of the sample: 12

$$k_{\text{SPW}} = \frac{2\pi}{\lambda_{633}} \sqrt{\frac{n_{\text{Au}}^2 n_S^2}{n_{\text{Au}}^2 + n_S^2}}.$$
 (8)

The SPR applications concerned of real RI changes owing to biomolecule interaction and enterprise to the change in SPW. 11-14 SPW propagates indifferent propagation constant of incident light, which is known as surface plasmon resonance point (SPRP). 9,12,33 The frequency at SPRP is called SRF. If SPW changes, it makes the SRF change, which is given in the following: 12

$$SRF = \frac{k_{SPW}C}{2\pi\sqrt[3]{n_{An}n_Sn_G}},\tag{9}$$

where *c* is light speed. Transmittance versus SRF curve is normally known as surface plasmon resonance frequency (SPRF) response curvature. Here, four layered Fresnel model is applied to create an SPRF curve for the proposed fiber optic SPR sensor to determine transmitted light intensity. The reflectance versus angle of incidence or reflectance versus wavelength characteristics curve is known as the SPR curve. The SPR angle that can be represented as follows:<sup>12</sup>

$$\theta_{\rm SPR} = a \, \sin\left(\frac{1}{n_P} \sqrt{\frac{n_{\rm Au}^2 n_s^2}{(n_{\rm Au}^2 + n_s^2)}}\right).$$
 (10)

The shifting characteristics of the SPR angle and SRF owing to the change of RI are established as the following equation: 9-11

$$n_s^2 = n_s^1 + c_a \frac{dn}{dc}. ag{11}$$

Equation (11) shows how RI of the sensing medium is changed after the hybridization between sh-DNA and mr-DNA sequences. The SPR angle and SRF are RI-dependent parameter of sensing medium. Here,  $n_s^1$  is the RI of the sensing dielectric before adsorption of sh-DNA molecule. When dielectric sample (probe or target) is absent inside the sensing medium, then  $n_s^1$  is the RI of PBS saline ( $n_4 = 1.34$ ), which is available in bare sensor.  $C_a$  is the concentration of adsorbed bio molecules, for example, if 1000 nM concentrated mr-DNA has been added into the sensing medium, then the value of  $C_a = 1000$  nM. The  $\frac{dn}{dc}$  is the RI increment parameter, suppose, after adding 1000 nM concentrated molecules, the RI of the sensing layer has been changed because the sensing layer now consists not only PBS but also 1000 nM concentrated molecules. The changed value of RI from PBS is  $\frac{dn}{dc}$  (=0.181 cm<sup>3</sup>/gm for PBS as bare case<sup>12</sup>). And  $n_s^2$  is the RI of the sensing dielectric after adsorption of mr-DNA. PBS is made by containing 0.0003 mol of NaH<sub>2</sub>PO<sub>4</sub> (monosodium phosphate), 0.0016 mol of Na<sub>2</sub>HPO<sub>4</sub> (disodium phosphate), 0.022 mol of NaCl, and 0.00054 mol of KCl mixing in Milli-Q water.<sup>29</sup> PBS buffer solution allows easy discrimination of perfectly matched DNA from mismatch DNA sequences.<sup>29</sup> The RI increment parameter is a (dn/dc) value of  $0.18 \pm 0.03$  cm<sup>3</sup>/g, irrespective of the identity of the protein and the buffer ions in the occasion of utilizing PBS buffer solution.<sup>28</sup> The sensitivity of the optical SPR sensor is defined as the ratio of the change of output parameters (SRF or  $\theta_{\rm SPR}$ ) to the change in the concentration of biomolecules,  $\Delta C_a$ .

### **2.2** Designated DNA Sequences for Detection of Breast Cancer

In this study, we selected 916delTT and 6174delT specific mutations in BRCA-1 and BRCA-2 genes, respectively, for the detection of breast cancer. Mutated type (mr) oligonucleotide, wild-type (wt) oligonucleotide, immobilization mutant (sh), and control target (mf) oligonucleotide represent complementary DNA sequence, mismatch DNA sequence, immobilization p-DNA sequence, and control target DNA sequence, respectively. These sequence orientation specification data in 916delTT and 6174delT are tabulated in Tables 1 and 2, respectively. <sup>2,31,34</sup>

In Tables 1 and 2, p-DNA sequence (sh-DNA), target fully complementary sequence (mr-DNA), and target mismatch sequence (wt-DNA) are shown in bold, italic, and bold italic, respectively, whereas control target sequence (mf-DNA) is revealed in black.

Perfectly matched hybridization event between sh-DNA sequence and mr-DNA sequence in 916delTT and 6174delT specific mutations in BRCA-1 and BRCA-2 genes

**Table 1** Sequence orientation of adenine (A), thymine (T), guanine (G), and cytosine (C) sequence in of BRCA1 (916deITT).

Immobilized mutant (sh) sequence	$5'\text{-SH-(CH}_2)_6\text{-TTT TTT TTT TTT TTT}\\ \textbf{GTT CTG TCA AACT-}3'$
Mutated type (mr) target sequence	5'-TGC CAC ATG GCT CCA CAT GCA AGT TTG ACA GAA CTA CCC TGA TAC TTT TCT GGA TGCC-3'
Wild type (wt) target sequence	5'-TGC CAC ATG GCT CCA CAT GCA AGT TTG AAA CAG AAC TAC CCT GAT ACT TTT CTG GAT GCC-3'
Control target (mf) sequence	5'-GGC ATC CAG AAA AGT ATC AGG GTA GTT CTG TTT CAA ACT TGC ATG TGG AGC CAT GTG GCA-3'

**Table 2** Sequence orientation of adenine (A), thymine (T), guanine (G), and cytosine (C) sequence in BRCA2 (6174deIT).

Immobilized mutant (sh) sequence	5'-AGC TGG TCT <b>GAC GTT TAT GAA</b> TGT TCG TTACT-3'
Mutated type (mr) target sequence	5'-GGTA CGA CAC GAT TTT TAG GGAC TTC ATA AAC GTC TAC TCT GAC-3'
Wild type (wt) target sequence	5'-GGTA CGA CAC GAT TTT TAG GGAC TTC ATC GAC ATC TAC TCT GAC-3'
Control target (mf) sequence	5'-GTC AGA GTA GAT GTC GAT GAA GTC CCT AAA AAT CGT GTC GTACC-3'

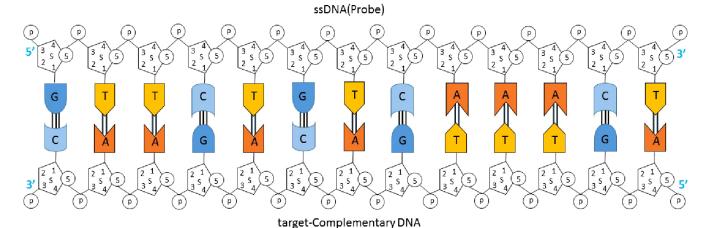
revealed the biomarker of genetic breast cancer, shown in Figs. 2(a) and 3(a), respectively.

On the other hand, Figs. 2(b) and 3(b) are the figural representation of mismatched hybridization event between wt-DNA sequence and mr-DNA sequence in 916delTT and 6174delT specific mutations in BRCA-1 and BRCA-2 genes, which revealed the biomarker of no possibility of genetic breast cancer. This mismatched and perfectly matched hybridization between sh-DNA sequence to mr-DNA sequence and sh-DNA sequence to wt-DNA sequence are numerically performed at concentrations ranging from 1 to100 nM in PBS solution. Mf-DNA, HCL, and the appropriate chemical have flowed with PBS solution with eliminating mr-DNA and wt-DNA samples and generating sh-DNA sample.<sup>23</sup>

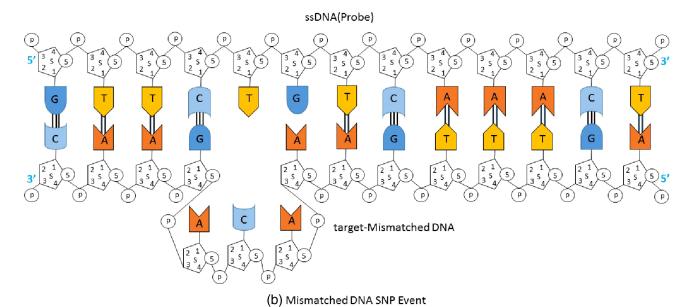
#### 3 Numerical Results and Discussion

#### 3.1 Sensor Detection Consideration

The addition of sh-DNA sequence as p-DNA on the sensing layer, reflectance versus SPR angle ( $R_{\rm min} \sim \theta_{\rm SPR}$ ), and transmittance versus SRF ( $T_{\rm max} \sim {\rm SRF}$ ) curve shift rightward. This is due to the addition of a probe changes the RI of the sensing dielectric medium (PBS in bare). By insertion of electron-rich DNA molecules, the number of carriers of graphene layer changes, which leads to modification of the propagation constant of SPW described in Eq. (8). Finally, a change in SRF is found due to the modification of the propagation constant of SPW described in Eq. (9). Therefore, the proposed SPR sensor with high performance is successfully



#### (a) Perfectly Matched (dsDNA) Hybridization Event



**Fig. 2** (a) Figural orientation between sh-DNA and mr-DNA sequence of BRCA-1 gene (916deITT): perfectly matched hybridization happened (A–T bond by dihydrogen bonding, G–C bond by trihydrogen bonding). (b) Figural orientation between sh-DNA and wt-DNA sequence: mismatch hybridization happened.

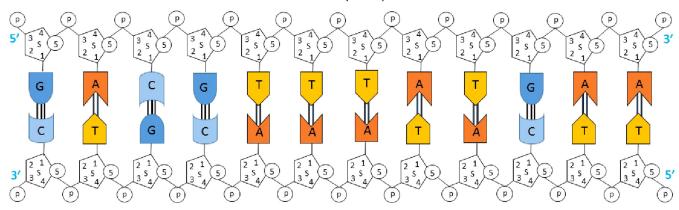
used to detect early breast cancer by observing the hybridization nature (perfectly matched or mismatched) based on the SPR angle and SRF variations. It is observed from Figs. 4(a) and 4(b) by adding sh-DNA as p-DNA (denoted by dip-blue color line), on the graphene surface, the SPR angle and SRF are considerably right shifted by 1.10<sup>0</sup> THz (from 57.75<sup>0</sup> to 58.80<sup>0</sup>) and 1.04 THz (from 92.34 THz to 93.38 THz), respectively. This phenomenon can illustrate the dependency of SPR angle, an SRF on the immobilization of the p-DNA, and hybridization of the complementary target DNA (mr-DNA in this work).

In Figs. 4(a) and 4(b), light green color line shows the change of SPR angle and SRF, respectively, with the submersion of the wt-DNA as mismatch target-DNA to the immobilized probe (sh-DNA) on SPR device. The results reveal that the SPR angle and SRF are 58.84<sup>0</sup> and 93.94 THz, respectively, after submersion of mismatched DNA (wt-DNA in this work). These values indicate no significant

change in the SPR angle (the change of the SPR angle while emerging mismatched target is  $\Delta SPR=0.04^0$ ) as well as no significant change in the SRF (the change of SRF while emerging mismatched target is  $\Delta SRF=0.56$  THz). When the sh-DNA sequence is contacted with wt-DNA sequence, there is no hydrogen bonding between the sh-DNA and wt-DNA strands according to Figs. 2(b) and 3(b) in 916delTT and 6174delT specific mutations in BRCA-1 and BRCA-2 genes, respectively, because of the presence of a mismatched pair. Therefore, there is no change of charges in the sensing medium due to its nonbonding reaction between two pairs of DNA strands since they cannot be hybrid. This information reveals that there is no possibility of genetic breast cancer.

Figures 4(a) and 4(b) also show the  $R_{\rm min} \sim {\rm SPR}$  angle and  $T_{\rm max} \sim {\rm SRF}$  characteristics, respectively, of the proposed model for different concentrated target DNA (mr-DNA), where each color line represents individual concentration of mr-DNA molecules. Here, black line shows  $R_{\rm min} \sim {\rm SPR}$ 

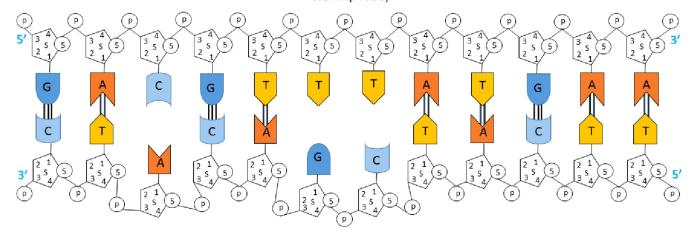
#### ssDNA(Probe)



target-Complementary DNA

#### (a) Perfectly Matched (dsDNA) Hybridization Event

#### ssDNA(Probe)



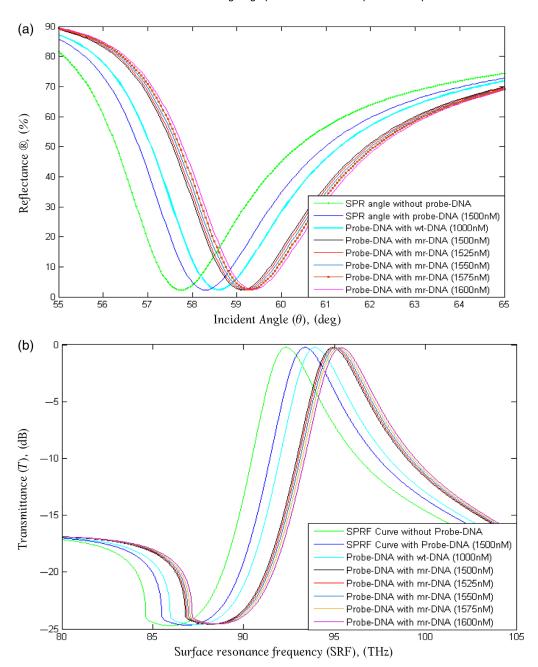
#### target-Mismatched DNA

#### (b) Mismatched DNA SNP Event

**Fig. 3** (a) Figural orientation between sh-DNA and mr-DNA sequence of BRCA-2 gene (6174delT): perfectly matched hybridization happened (A–T bond by dihydrogen bonding, G–C bond by trihydrogen bonding). (b) Figural orientation between sh-DNA and wt-DNA sequence: mismatch hybridization happened.

angle and  $T_{\text{max}} \sim \text{SRF}$  characteristics curve after addition of 1500 nM perfectly matched mr-DNA, red line shows  $R_{\min} \sim \text{SPR}$  angle and  $T_{\max} \sim \text{SRF}$  characteristics curve after addition of 1525 nM perfectly matched mr-DNA, light blue line shows  $R_{\min} \sim \text{SPR}$  angle and  $T_{\max} \sim \text{SRF}$  characteristics curve after addition of 1550 nM perfectly matched mr-DNA, yellow line shows  $R_{\min} \sim \text{SPR}$  angle and  $T_{\max} \sim \text{SRF}$  characteristics curve after addition of 1575 nM perfectly matched mr-DNA, and magenta color line shows  $R_{\min} \sim SPR$  angle and  $T_{\text{max}} \sim \text{SRF}$  characteristics curve after addition of 1600 nM perfectly matched mr-DNA, respectively. The results reveal that the SPR angles are  $59.10^{\circ}$ ,  $59.15^{\circ}$ ,  $59.20^{0}$ ,  $59.30^{0}$ , and  $59.35^{0}$  as well as the SRF are 94.90 THz, 94.99 THz, 95.10 THz, 95.22 THz and 95.36 THz respectively, for the five cases. According to these analytical data, four important factors—the SPR angle, the minimum reflectance, the SRF, and the maximum transmittance-play a significant role in detecting early breast cancer by observing DNA hybridization events. The sensing parameter  $R_{\rm min}$ ,  $\theta_{\rm SPR}$ , SRF, and  $T_{\rm max}$  changes with different concentrated mr-DNA and wt-DNAs (ranging 1 to 100 nM) is figured in the Figs. 4(a) and 4(b), furthermore, table data have been extracted in Table 3. It is evidently decided that the significant right shift of  $\theta_{\rm SPR}$  and SRF curve is a sign of perfectly matched hybridization happened in 916delTT and 6174delT,<sup>34</sup> which reveals mutations in 916delTT and 6174delT biomarkers in BRCA1 and BRCA2, respectively.

Figures 4(a) and 4(b) realistically explain the  $R_{\rm min} \sim \theta_{\rm SPR}$  (SPR curve) characteristics and  $T_{\rm max} \sim {\rm SRF}$  (SPRF curve) characteristics, respectively, without p-DNA, with p-DNA, and flowing the specific concentrated complementary target wt-DNA or mr-DNAs of BRCA-1 and BRCA-2 genes. The RI is changed owing to the immobilization of different concentrated breast cancer biomarker (916delTT and 6174delT)



**Fig. 4** (a) The SPR curves characteristics. (b) The SPRF curves characteristics of SPR sensor without p-DNA, with p-DNA, and flowing the different concentrated complementary target wt-DNA or mr-DNA of BRCA-1 and BRCA-2 genes in PBS.

DNA molecules. The RI of detecting medium has a significant impact on the SPRF and the SPR curve. The significant right shift of  $\theta_{\rm SPR}$  and SRF is a sign of recognition of specific mutations 916delTT and 6174delT interaction with p-DNAs, as well as that clarifies the detection of hereditary breast cancer. The detecting parameter changes with differently concentrated mr-DNAs have been tabulated in Table 3, accordingly, a decision will be made based on the change of detecting parameter.

In Table 3, the breast cancer recognizing attributors  $(\delta R_{\min}, \delta \theta_{\rm SPR}, \delta T_{\max}, \text{ and } \delta {\rm SRF})$  are calculated by choosing the change of  $R_{\min}, \theta_{\rm SPR}, T_{\max}$ , and SRF value after adding p-DNA as standard. And after adding p-DNA, the interaction between p-DNA to mismatched (wt sequence) DNA and

p-DNA to different concentrated perfectly matched target (mr sequence) DNAs has been determined and tabulated in Table 4, in order to make a decision according to Table 5. The following Eq. (12) is used to determine standard attributor's value, in which value is essential for proper detection of genetic biomarkers (916delTT and 6174delT) for early breast cancer recognized specific mutations of BRCA-1 and BRCA-2 genes as follows:

$$\delta R_{\min}^{\text{Cancer}} = |R_{\min}^{\text{probe}} - R_{\min}^{mr=1500}| = 0.09$$

$$\delta \theta_{\text{SPR}}^{\text{Cancer}} = |\theta_{\text{SPR}}^{\text{probe}} - \theta_{\text{SPR}}^{mr=1500}| = 0.80 \text{ deg.}$$

$$\delta T_{\max}^{\text{Cancer}} = |T_{\max}^{\text{probe}} - T_{\max}^{mr=1500}| = 0.0083 \text{ dB}$$

$$\delta \text{SRF}^{\text{Cancer}} = |\text{SRF}^{\text{probe}} - \text{SRF}^{\text{mr}=1500}| = 1.52 \text{ THz}$$

**Table 3** Data of change of detecting attributors ( $\delta R_{\min}$ ,  $\delta \theta_{SPR}$ ,  $\delta T_{\max}$ , and  $\delta SRF$ ) for introducing different concentrated target WT-DNA or MR-DNA of BRCA-1 and BRCA-2 gene.

Concentration $(C_a)$ (nM)	R <sub>min</sub> (%)	$ heta_{SPR}$ (deg)	T <sub>max</sub> (dB)	SRF (THz)	$\delta R_{ ext{min}}^{P-T} = \  R_{ ext{min}}^{ ext{probe}} - R_{ ext{min}}^{ ext{target}} $	$\delta  heta_{ extsf{SPR}}^{P-T}( extsf{deg}) = \   heta_{ extsf{SPR}}^{ extsf{probe}} -  heta_{ extsf{SPR}}^{ ext{target}} $	$\delta T_{max}^{P-T}(dB) = \  T_{max}^{probe} - T_{max}^{target} $	$\delta SRF^{P-T}(THz) =  SRF^{probe} - SRF^{target} $
0 (no-sh-DNA)	2.26	57.75	0.2288	92.34	<del></del>	<del></del>	_	_
1500 (sh-DNA)	2.31	58.80	0.2339	93.38	0.00	0.00	0.0000	0.00
1000 (wt-DNA)	2.34	58.84	0.2368	93.94	0.03	0.04	0.0029	0.56
1500 (mr-DNA)	2.40	59.10	0.2422	94.90	0.09	0.80	0.0083	1.52
1525(mr-DNA)	2.40	59.15	0.2427	94.99	0.09	0.85	0.0088	1.61
1550 (mr-DNA)	2.41	59.20	0.2433	95.10	0.10	0.90	0.0094	1.72
1575 (mr-DNA)	2.42	59.30	0.2441	95.22	0.11	1.00	0.0102	1.84
1600 (mr-DNA)	2.42	59.35	0.2449	95.36	0.11	1.05	0.0110	1.98

Table 4 Decision making based upon different situations that happened to detecting attributors.

Pathology no.	Decision making condition (with r	Decision		
01	$\delta R_{\min}^{P-T} \ge \delta R_{\min}^{\text{Cancer}}$ && $\delta \theta_{\text{SPR}}^{P-T} \ge \delta \theta_{\text{SPR}}^{\text{Cancer}}$		$\delta T_{\text{max}}^{P-T} \ge \delta T_{\text{max}}^{\text{Cancer}} \text{ && } \delta \text{SRF}^{P-T} \ge \delta \text{SRF}^{\text{Cancer}}$	Cancer detect
02	$\delta R_{\rm min}^{P-T} < \delta R_{\rm min}^{\rm Cancer} \ \&\& \ \delta \theta_{\rm SPR}^{P-T} < \delta \theta_{\rm SPR}^{\rm Cancer}$		$\delta T_{\max}^{P-T} < \delta T_{\max}^{\text{Cancer}} \&\& \delta \text{SRF}^{P-T} < \delta \text{SRF}^{\text{Cancer}}$	No cancer detect
03	$\delta R_{\min}^{P-T} \ge \delta R_{\min}^{\text{Cancer}}$ && $\delta \theta_{\text{SPR}}^{P-T} \le \delta \theta_{\text{SPR}}^{\text{Cancer}}$	OR	$\delta T_{\max}^{P-T} \le \delta T_{\max}^{\text{Cancer}} \text{ \&\& } \delta SRF^{P-T} \ge \delta SRF^{\text{Cancer}}$	Further try again
04	$\delta R_{\min}^{P-T} \le \delta R_{\min}^{\text{Cancer}}$ && $\delta \theta_{\text{SPR}}^{P-T} \ge \delta \theta_{\text{SPR}}^{\text{Cancer}}$		$\delta T_{\max}^{P-T} \ge \delta T_{\max}^{\text{Cancer}} \&\& \ \delta \text{SRF}^{P-T} \le \delta \text{SRF}^{\text{Cancer}}$	Further try again

**Table 5** Data of  $R_{\min}$ ,  $\theta_{SPR}$ ,  $T_{\max}$ , and SRF without graphene layer and with the total number of increasing graphene layer in fiber optic SPR biosensor.

Graphene layer	R <sub>min</sub> (%)	$\theta_{\rm SPR}$ (deg)	$T_{\rm max}$ (dB)	SRF (THz)	$\delta R_{\rm min}$ [%]	$\delta  heta_{\mathrm{SPR}}$ [deg]	$\delta T_{\rm max}$ [dB]	$\delta$ SRF [THz]
Without layer (£ = 0 nm)	0.25	57.35	0.0252	91.92	0.00	0.00	0.0000	0.00
First layer (£ = 1 $\times$ 0.34 nm)	2.26	57.70	0.2285	92.28	2.01	0.35	0.2033	0.36
Second layer (£ = $2 \times 0.34$ nm)	5.40	58.10	0.5550	92.68	5.15	0.75	0.5298	0.76
Third layer (£ = $3 \times 0.34$ nm)	9.05	58.50	0.9481	93.08	8.80	1.15	0.9229	1.16
Fourth layer (£ = $4 \times 0.34$ nm)	12.87	58.95	1.3770	93.51	12.62	1.60	1.3518	1.59
Ninth layer (£ = $9 \times 0.34$ nm)	30.51	61.50	3.6405	95.93	30.26	4.15	3.6153	4.01
Tenth layer (£ = $10 \times 0.34$ nm)	33.55	62.10	4.0867	96.47	33.30	4.75	4.0615	4.55

Equation (12) describes that breast cancer recognizing attributors minimum value is stated in such way that is equal to the right shift of SPR and SRF curvatures after submersing mutated type target (1500 nm concentrated mr sequence) DNA into p-DNA. Concededly, Table 4 is the decision-making table, which can be applied for breast cancer detection, taking advantage of the detecting attributor's values. The minimum attributor values are  $\delta R_{\rm min}^{\rm Cancer} = 0.09\%$ ,  $\delta \theta_{\rm SPR}^{\rm Cancer} = 0.80$  deg,  $\delta T_{\rm max}^{\rm Cancer} = 0.0083$  dB, and  $\delta {\rm SRF}^{\rm Cancer} = 1.52$  THz for detection of specific mutations in PBS solutions.

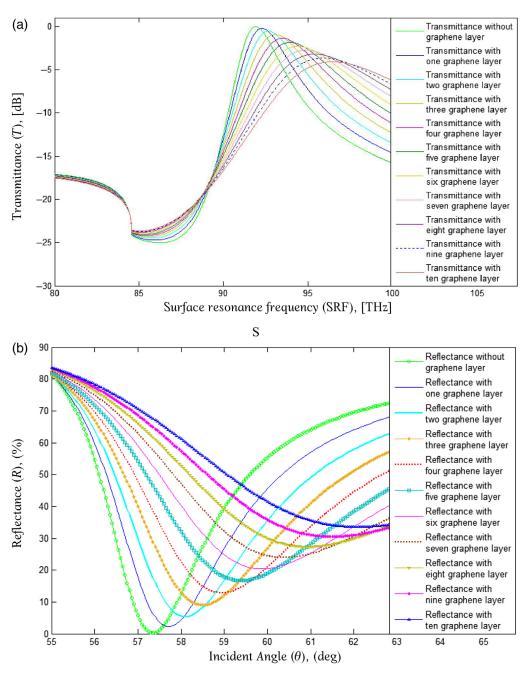
Table 4 represents the pathological decision for different cases. After submersing an arbitrary molecular sample on the sensing region of SPR device, either perfectly matched sequence (mutated type target sequence, mr-DNA) or mismatched sequence (wild-type target sequence, wt-DNA) in 916delTT and 6174delT with two specific mutations in BRCA-1 and BRCA-2 genes, the detecting attributors—such as  $R_{\min}$  (%),  $\theta_{\rm SPR}$  (deg),  $T_{\rm max}$  (dB), and SRF (THz)—are changed according to the nature of hybridization. When the change of  $\delta R_{\min}^{P-T}$ ,  $\delta \theta_{\rm SPR}^{P-T}$ ,  $\delta T_{\max}^{P-T}$ , and  $\delta {\rm SRF}^{P-T}$  are greater than or equal to  $\delta R_{\min}^{\rm Cancer}$ ,  $\delta T_{\rm SPR}^{\rm Cancer}$ , and  $\delta {\rm SRF}^{\rm Cancer}$ ,

then specific mutations (916delTT and 6174delT) of BRCA-1 and BRCA-2 genes are sensed (breast cancer detect) by SPR sensor (described this decision by pathology No. 01 in Table 4) and if the change of  $\delta R_{\min}^{P-T}$ ,  $\delta \theta_{\rm SPR}^{P-T}$ ,  $\delta T_{\max}^{P-T}$ , and  $\delta {\rm SRF}^{P-T}$  is smaller than  $\delta R_{\min}^{\rm Cancer}$ ,  $\delta \theta_{\rm SPR}^{\rm Cancer}$ ,  $\delta T_{\max}^{\rm Cancer}$ , and  $\delta {\rm SRF}^{\rm Cancer}$ , then it confirms that no breast cancer mutations are sensed (described this decision by pathology No. 02 in Table 4); apart from both of these decisions, no consistencies consequence will be recognized. Hence, the change of  $\delta R_{\min}^{P-T}$ ,  $\delta \theta_{\rm SPR}^{P-T}$ ,  $\delta T_{\max}^{P-T}$ , and  $\delta {\rm SRF}^{P-T}$  is due to the variability arising from other interferents in the biomolecules and the recognition process should be further tried again (described this decision by pathology Nos. 03 and 04 in Table 4). In such way, a graphene-implicated SPR sensor is used to make a pathological decision from the achieved results,

respectively, Table 4 and its pathology nos. indicated the detection and can be utilized for confirming detection of 916delTT and 6174delT mutation.

#### 3.2 Sensor Performance Consideration

In this section, a numerical study is presented on the effect of including the graphene layer in terms of SPR curve ( $R_{\rm min}$  versus  $\theta_{\rm SPR}$ ) and SRF curve ( $T_{\rm max}$  versus SRF) characteristics. Here, we used PBS solution ( $n_4=1.34$ ) before adsorbing probe sh-DNA molecules. The  $\theta_{\rm SPR}$  and SRF without the graphene layer are 57.35 deg and 91.92 THz, furthermore, with a single layer of graphene is 57.70 deg and 92.28 THz, respectively, shown in Figs. 5(a) and 5(b). The SPR angle (change of SPR angle,  $\delta\theta_{\rm SPR}=0.35\,$  deg) and SRF (change



**Fig. 5** (a) The SPR curves. (b) The SPRF curves for the fiber optic SPR biosensor without graphene and with graphene implicated sensor (for  $\pounds = 1,2,3,\ldots,10$ ) before the adsorption of target BRCA DNA molecules.

of SRF,  $\delta$ SRF = 0.36 THz) have been increased owing to adding graphene film, which is enhanced the sensitivity of this sensor. This enhanced sensitivity is due to the electron loss of graphene, high carrier mobility, high optical transparency, exceptional mechanical flexibility, strength, <sup>35</sup> low resistivity, tunable conductivity, and extreme mode confinement, <sup>25–27</sup> which is accompanying the imaginary dielectric constant. This increased SPR angle will lead to increased sensitivity of the sensor as sensitivity is directly related to the variation of SPR angle discussed in Refs. 8–15. And thus, the detecting attributors  $\delta\theta_{\rm SPR}$  and  $\delta$ SRF are shifting rightward because of adding the graphene layer. These values have been tabulated in Table 5, as well as table data depicted in Figs. 5(a) and 5(b), respectively.

In Table 3, the breast cancer recognizing attributors  $R_{\min}$ ,  $\theta_{SPR}$ ,  $T_{\max}$ , and SRF are increasing with the number of the graphene layer (£ = n \* 0.34 nm;  $1 \le n \le 10$ ) included on the Au thin film, which causes enhanced sensitivity of the proposed fiber optic SPR sensor.

The sensor performance is determined in terms of sensor sensitivity (S). Sensor sensitivity is defined as function to the change of output (change of  $\Delta\theta_{SPR}$  or  $\Delta SRF$ ) with respect to the change of input (concentration of the sensing layer,  $\Delta C_a$ ).

For  $R_{\min} \sim \text{SPR}$  angle attributor, sensitivity (S) is defined as follows: 12

$$S = \frac{\Delta\theta_{\text{SPR}}}{\Delta c_a}$$

$$= \frac{\Delta\theta_{\text{SPR}}^0 + \Delta\theta_{\text{SPR}}^L}{\Delta c_a}$$

$$= \frac{\Delta\theta_{\text{SPR}}^0 + 0.35\Delta\theta_{\text{SPR}}^0}{\Delta c_a}$$

$$= \frac{(1+0.35L)\Delta\theta_{\text{SPR}}^0}{\Delta c_a}$$

$$= (1+0.35L)s^O$$
(13)

where  $\Delta\theta_{\rm SPR}$  is the change of SPR angle due to the presence of mr-DNA sequence and  $\Delta c_a$  is the change of concentration of the sensing dielectric after adsorption of perfectly matched mr-DNA molecule. Here,  $S^0$  is the sensitivity without graphene, and  $S^L$  is the sensitivity with graphene. Equation (13) is derived from Table 5 and is seen if we use a single graphene layer (£ = 1,0.34 nm), the sensitivity has been increased 35% more with respect to conventional sensor.

For  $T_{\text{max}} \sim \text{SRF}$  attributor, sensitivity (S) is defined as follows:

$$S = \frac{\Delta SRF}{\Delta c_a}$$

$$= \frac{\Delta SRF^0 + \Delta SRF^L}{\Delta c_a}$$

$$= \frac{\Delta SRF^0 + 0.36\Delta SRF^0}{\Delta c_a}$$

$$= \frac{(1 + 0.35L)\Delta SRF^0}{\Delta c_a}$$

$$= (1 + 0.36L)s^O$$
(14)

where  $\Delta$ SRF is the SRF change of sensor summing with and without graphene sublayers, S0 is the sensitivity of the devices without graphene film, and SG is the sensitivity for different number of graphene sublayers (L=1,2,3,...). Equation (14) is derived from Table 5 and is seen if we use a single graphene layer (£ = 1,0.34 nm), the sensitivity has

been increased 36% more with respect to conventional sensor.

#### 4 Conclusion

We present a numerical modeling of graphene-coated fiber optic SPR biosensor for proper detection of genetic biomarkers (916delTT and 6174delT) for early breast cancer by means of DNA hybridization. From the variation of the SPR angle and SRF, the proposed sensor can differentiate between perfectly matched and mismatched DNA interaction between p-DNA (sh-DNA mutant) and target DNA (mutation type mr-DNA). Numerical results show that the use of graphene can be more sensitive compared with the usual SPR biosensors. This increased sensitivity is due to the absorption ability of graphene. This sensor provided the easiest way of monitoring cancer tumor cell and confirming detection of specific mutations of BRCA-1 and BRCA-2 genes. The thought offered in this article is expected to be understood easily, and because of enhanced sensitivity, the proposed hybrid SPR biosensor has great prospective in future industrial applications of genetic biomarkers (916delTT and 6174delT) for early breast cancer detection.

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