

“COMPOUNDS AS STIMULI-RESPONSIVE PROBES, METHODS AND APPLICATIONS THEREOF”

The following specification particularly describes the invention and the manner in which it is to be performed.

TECHNICAL FIELD

The present disclosure relates to the field of synthetic pharmaceutical chemistry and biology. In particular, a compound of Formula I and a process of preparation thereof are disclosed. The disclosure further relates to methods and use of Formula I compounds as stimuli-responsive probes. Said Formula I compounds are employed for detecting ROS, and have related applications including but not limited to fluorescence spectroscopy, diagnostics, imaging and biomedical applications.

BACKGROUND OF THE DISCLOSURE

The regulation of redox homeostasis is essential for maintaining normal cellular functions such as signaling, growth, survival and death. Anomalous behavior of redox homeostasis adversely affects the normal physiological functions and is in turn responsible for numerous pathological conditions. Normally, cells in the diseased state exhibit high levels of aerobic glycolysis (Warburg effect), which results in oxidative stress. For example, the oxidative stress in cancer cells result in the accumulation of high levels of reactive oxygen species (ROS).

ROS constitute an important class of chemically reactive species that are essential for normal cellular functions including cell proliferation and differentiation. The optimum levels of ROS are controlled by various cellular redox homeostasis mechanisms, and an abrupt increase in their concentration levels is directly linked to oxidative stress-related disorders. Abnormally high levels of ROS are generated in response to adverse environmental and physiological stresses, exposure to ultraviolet (UV) light, and ionizing and heat radiations. It is crucial to monitor the levels of intracellular ROS for maintaining effective cellular homeostasis. Notably, different levels of ROS are responsible for different biological responses. Cell maintains different levels of ROS by activating the ROS-scavenging systems such as superoxide dismutases, glutathione peroxidase, redox enzymes (peroxiredoxins, glutaredoxin and thioredoxin) and catalase.

Mis-regulation in any of these ROS-scavenging processes leads to generation of excessive amounts of ROS. Accumulation of high levels of ROS causes oxidative damage of cellular components such as proteins, lipids and nucleic acids, which is responsible for ageing and many pathological conditions including cancer and cardiovascular, inflammatory and neurodegenerative diseases. It is known that cellular aging, also called cellular senescence, is a permanent cell cycle arrest state that results in increased production of ROS species. This increased ROS production is critical in maintaining the viability of the senescent cell. Therefore, it is necessary to develop molecular tools that are highly sensitive and can be activated by high levels of ROS to distinguish aged or diseased and normal cells.

ROS mainly comprises free radicals such as hydroxyl radical ($\text{OH}\cdot$) and superoxide ($\text{O}_2^{\cdot-}$) and reactive molecular species such as H_2O_2 . H_2O_2 is one of the most prominent and essential ROS in biological systems and its significantly higher levels are generated in aged and cancer cells than in normal cells. In fact, H_2O_2 is a small molecular metabolite and plays a vital role in the regulation of various physiological processes in living organisms. Most importantly, H_2O_2 serves as a messenger in normal cellular signal transduction and is also a known marker for oxidative damage in many disease-associated cells. In cells, H_2O_2 is generated through the receptor-mediated NADPH oxidase (Nox) activation, which affects the functioning of signaling proteins that control cell signaling, proliferation, senescence and death.

The biological significance of H_2O_2 in human physiology and pathology has generated great interest in understanding the mechanistic details of H_2O_2 generation, partition and its role in cellular function and signaling pathways. In comparison to other ROS, relatively higher stability and diffusion rates of H_2O_2 through the plasma membrane makes it an attractive candidate to study its signaling pathways in living cells. However, the role of H_2O_2 as an essential messenger, for cellular signal transduction and its chemical reactivity and chemical instability limits its spatiotemporal tracking in real-time, especially in living cells. Molecular imaging of H_2O_2 using fluorescence probes is a highly attractive tool for studying its generation, accumulation, and trafficking and its role in biological processes in a spatiotemporal manner in living cells.

In recent years, stimuli-responsive fluorescence probes are gaining momentum due to their flexibility in introducing diversity through chemical modification and liberation of biologically active probes at the site of target cellular organelles, in response to biological analytes of interest.

Moreover, targeting specific subcellular organelles (mitochondria) and biomolecules such as DNA and proteins using stimuli-responsive fluorescence probes is an emerging powerful imaging technique that presents enormous potential in biomedical applications related to diagnostics and therapeutics. Therefore, there is a need in the art to develop such stimuli responsive fluorescence probes in order to carry out imaging, biomedical research, diagnosis, treatment etc.

Accordingly, the present disclosure provides for DNA-binding fluorescence probes with a stimuli-responsive appendage, wherein in response to a specific stimulus (chemical or enzyme), the appendage functionality is cleaved to release an NIR-fluorescence ready probe, which upon binding the minor groove of DNA fluoresces strongly, thus, aiding the imaging and quantification of the stimulus.

BRIEF DESCRIPTION OF THE ACCOMPANYING FIGURES

In order that the disclosure may be readily understood and put into practical effect, reference will now be made to exemplary embodiments as illustrated with reference to the accompanying figures. The figures together with the detailed description below, are incorporated in and form part of the specification, and serve to further illustrate the embodiments and explain various principles and advantages, in accordance with the present disclosure where:

Figure 1 depicts H_2O_2 triggered release of NIR-fluorescence minor groove binder.

Figure 1(a) depicts the schematic representation of conversion of QCy-BA to *p*-quinone-methide and QCy-DT, a DNA minor groove binder, in the presence of H_2O_2 .

Figure 1(b) depicts the time-dependent 1H NMR spectral monitoring of slicing of phenyl boronic acid of QCy-BA in the presence of H_2O_2 . Red circles highlight the appearance of new signals for the newly-formed quinone system and QCy-DT. H_a and H_b represent the O-CH₂ (C-H_a) bearing phenyl boronic acid group and newly formed exocyclic (C-H_b) protons of *p*-quinone-methide, respectively.

Figure 1(c) depicts the change in solution color upon addition of H_2O_2 to QCy-BA, as visualized after 2 hours.

Figure 1(d) Fluorescence response of QCy-BA (5 μM) to various reactive oxygen species (ROS) at individual concentration of 100 μM .

Figure 2 depicts absorbance and fluorescence response of combination probe QCy-BA $\subset$ Drew-AT.

Figure 2(a) depicts the absorption spectra of combination probe QCy-BA $\subset$ Drew-AT (2 μ M) in the presence of H₂O₂ (100 μ M) in PBS-buffer solution as a function of time.

Figure 2(b) depicts the normalized fluorescence spectra of QCy-BA (2 μ M) in the presence of Drew-AT (2 μ M) upon excitation at 400 nm.

Figure 2(c) Fluorescence spectra of QCy-BA (2 μ M) in the presence of Drew-AT (2 μ M) upon excitation at 564 nm. All the spectra are acquired in the presence of H₂O₂ (100 μ M).

Figure 2(d) depicts the time-dependent fluorescence spectra of QCy-BA (2 μ M) in the presence of Drew-AT (2 μ M) after the addition of H₂O₂ (100 μ M) upon excitation at 400 nm.

Figure 2(e) depicts the schematic view of conversion of QCy-BA to *p*-quinone-methide and a DNA minor groove binder (QCy-DT) with turn-on NIR-fluorescence in the presence of H₂O₂.

Figure 3 depicts the glucose oxidase (GOx) assay.

Figure 3(a) depicts the schematic diagram showing the GOx-assay where Gox oxidizes glucose to gluconic acid, generating H₂O₂, followed by fluorescence reporting by the combination probe QCy-BA $\subset$ Drew-AT.

Figure 3(b) depicts the fluorescence spectra of combination probe QCy-BA $\subset$ Drew-AT in the presence of Gox(4 U/mL) and upon addition of glucose (1 mM).

Figure 3(c) depicts the time-dependent fluorescence of combination probe QCy-BA $\subset$ Drew-AT in the presence and absence of Gox (4 U/mL) upon addition of glucose (1 mM).

Figure 3(d) depicts the photographs of QCy-BA $\subset$ Drew-AT complex under UV-light in the presence of Gox(4 U/mL) with increasing glucose concentration 0.0 to 1.0 mM.

Figure 4 depicts cellular uptake of QCy-BA and fluorescence reporting of H₂O₂ in HeLa cells.

Figures 4(a) and 4b depict the fluorescence microscope and differential interference contrast (DIC) images of HeLa cells incubated with QCy-BA(5 μ M) in the absence of H₂O₂.

Figures 4(c) and 4(d) depict the fluorescence microscope and differential interference contrast (DIC) images of HeLa cells incubated with QCy-BA(5 μ M) in the presence of H₂O₂ (100 μ M).

Figures 4(e) and 4(f) depict the FACS/flow cytometry analysis showing the PerCP mean fluorescence intensity in HeLa cells. Figure 4(e) depicts the fluorescence intensity of QCy-BA (5 μ M) in HeLa cells upon addition of H₂O₂ (100 μ M) and N-acetyl-L-cysteine (NAC) (8 mM). Figure 4(f) depicts the fluorescence intensity of QCy-BA (5 μ M) in HeLa cells upon addition of epithelial growth factor (EGF) (500 ng/mL) and N-acetyl-L-cysteine (NAC) (8 mM). Error bars represent $\pm$ standard deviation.

Figure 5 depicts cellular uptake of QCy-BA in primary MRC5 cells and genotoxic stress-induced H₂O₂ detection in primary and cancer cells.

Figures 5(a) and 5(b) depict the fluorescence microscope and differential interference contrast (DIC) images of MRC5 cells incubated with QCy-BA(5 μ M) in the absence of H₂O₂.

Figures 5(c) and 5(d) depict the fluorescence microscope and differential interference contrast (DIC) images of MRC5 cells incubated with QCy-BA(5 μ M) in the presence of H₂O₂ (100 μ M).

Figure 5(e) depicts H₂O₂ detection in attached live HeLa cells using QCy-BA (5 μ M) after treatment with BrdU from 0 to 200 μ M or doxorubicin (0.1 μ M) for 48 hours. Fold change of fluorescence per cell is normalized to 1 for control cells (n=3).

Figure 5(f) depicts MRC5 cells treated with BrdU from 0 to 200 μ M or doxorubicin (0.1 μ M) for 72 hours. H₂O₂ levels estimated using QCy-BA (5 μ M) dye and fold change of fluorescence per cell is normalized to 1 for control cells (n=3).

Figure 6 depicts absorption and emission spectra of QCy-BA, wherein,

Figure 6(a) depicts the absorption spectra of QCy-BA (5 μ M) in presence of H₂O₂ (1 mM).

Figure 6(b) depicts the emission spectra of QCy-BA (5 μ M) in presence of H₂O₂ (1 mM) in PBS-buffer solution as a function of time.

Figure 7 depicts the plot of the fluorescence intensity at 550 nm against the concentration of [H₂O₂] PBS-buffer solution. Each data point is acquired after addition of H₂O₂ at 25 °C. The detection limit (5.3 μ M) is calculated with $3\sigma/k$; where σ is the standard deviation of blank measurement, k is the slop (-0.30).

Figure 8 depicts the absorption and emission spectra of QCy-BA in presence of PBS.

Figure 8(a) depicts the absorption spectra of QCy-BA (2 μ M) in presence of Drew-AT (2 μ M). Figure 8(b) depicts the emission spectra of QCy-BA (2 μ M) in presence of Drew-AT (2 μ M) in PBS-buffer solution.

Figure 9 depicts the time-dependent fluorescence spectra of QCy-BA (2 μ M) in presence of Drew-AT (2 μ M) after the addition of H₂O₂ (100 μ M) upon excitation at 564 nm.

Figure 10 depicts the plot of $\ln(F_{\infty}-F)$ of combination probe QCy-BA $\subset$ Drew-AT (2 μ M) as a function of time at 650 nm upon addition of H₂O₂ (1 mM), where F_{∞} and F are the fluorescence intensities at 650 nm at time t_{∞} (= 60 min) and t , respectively. The k_{obs} calculated from the slope of this plot is $1.04 \times 10^{-3} \text{ s}^{-1}$.

Figure 11 depicts the fluorescence spectra of combination probe QCy-BA $\subset$ Drew-AT.

Figure 11(a) depicts the fluorescence spectra of combination probe QCy-BA $\subset$ Drew-AT in presence of Gox (4 U/mL) and glucose (1 mM) with time, upon excitation at 400 nm.

Figure 11(b) depicts the plot of fluorescence intensity of combination probe QCy-BA $\subset$ Drew-AT at 650 nm as function of time, in presence of Gox (4 U/mL) with increasing concentration of glucose from 0 to 1 mM upon excitation at 564 nm.

Figure 12 depicts the plot of fluorescence intensity of combination probe QCy-BA $\subset$ Drew-AT in the presence of GOx (4 U/mL) against the concentration of glucose at 650 nm. Each data point is acquired 1 hour after addition of glucose and GOx at 25°C. The detection limit (6.11 μ M) is calculated by $3\sigma/k$, where σ is the standard deviation of blank measurement and k is the slope (0.12).

Figure 13 depicts the plot of $\ln(F_{\infty}-F)$ of combination probe QCy-BA $\subset$ Drew-AT in presence of Gox (4 U/mL) as a function of time at 650 nm upon addition of glucose (1 mM), where F_{∞} and F are the fluorescence intensities at 650 nm at time t_{∞} (= 60 min) and t , respectively. The k_{obs} calculated from the slope of this plot is $6.87 \times 10^{-4} \text{ s}^{-1}$.

Figure 14 depicts the study of the combination probe QCy-BA $\subset$ Drew-AT under catalase activity.

Figure 14(a) depicts schematic diagram representing the catalase activity in presence of H₂O₂ and combination probe **QCy-BA $\subset$ Drew-AT**.

Figure 14(b) depicts fluorescence intensity of QCy-BA at 565 nm. Where Q: QCy-BA, QH: QCy-BA+H₂O₂, QHD: QCy-BA+H₂O₂+Drew-AT, QCH: QCy-BA+Catalase+H₂O₂, QCHD: QCy-BA+Catalase+H₂O₂+Drew-AT.

Figure 14(c) depicts time dependent fluorescence of combination probe QCy-BA $\subset$ Drew-AT in presence and absence of catalase upon addition of H₂O₂ (1 mM) upon excitation at 564 nm.

Figure 15 depicts dose dependent cell viability of HeLa cells by taking 0.0-25 μ M of probe QCy-BA for 24 h. Error bars represent $\pm$ standard deviation.

Figure 16 depicts Flow activated cell sorter (FACS) analysis showing the PerCP mean fluorescence intensity in HeLa cells in presence of QCy-BA (5 μ M).

Figure 17 depicts H₂O₂ detection in attached live HeLa cells using DCFDA after treatment with BrdU (100 μ M) and doxorubicin (0.1 μ M) for 48 h. Fold change of fluorescence per cell is normalized to 1 for control cells (n=3).

Figure 18 depicts ¹HNMR spectrum of compound 2 i.e. 2-(4-(iodomethyl)phenyl)-4,4,5,5-tetramethyl-1,3,2-dioxaborolane.

Figure 19 depicts ¹³CNMR spectrum of Compound 2.

Figure 20 depicts ¹HNMR spectrum of Compound 3 i.e. 4-(4-(4,4,5,5-tetramethyl-1,3,2-dioxaborolan-2-yl)benzyloxy)isophthalaldehyde.

Figure 21 depicts ¹³CNMR spectrum of Compound 3.

Figure 22 depicts ¹HNMR spectrum of QCy-BA.

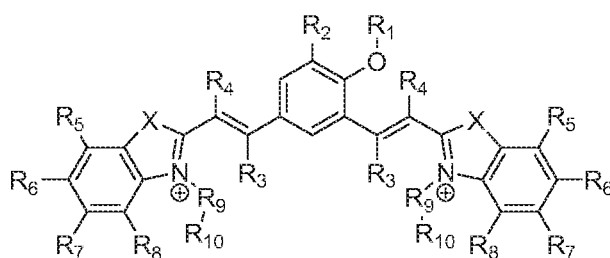
Figure 23 depicts ^{13}C NMR spectrum of QCy-BA.

Figure 24 depicts HRMS mass data of QCy-BA.

Figure 25 depicts HPLC trace of QCy-BA.

STATEMENT OF THE DISCLOSURE

The present disclosure relates to a compound of Formula I:



Formula I

wherein, 'X' is selected from a group comprising oxygen, sulphur, and selenium;

'R₁' is selected from a group comprising alkyl chain, allyl group, aryl group, benzyl group, aryl and alkyl group, acid, amine, ester, methyl phenyl boronic acid, boronic ester, carbonate, phosphate, silane, quaternary ammonium, amide, imine and any other moiety which can be triggered by chemical or enzymatic tools;

'R₂' is selected from a group comprising H, OH, halogen, alkyl and substituted alkyl, and wherein, the halogen is selected from a group comprising bromide, chloride and iodide;

'R₃' or 'R₄' is selected from a group comprising H, alkyl, aryl, nitrile, acid and halogen, and wherein the halogen is selected from a group comprising, chloride, fluoride, bromide and iodide;

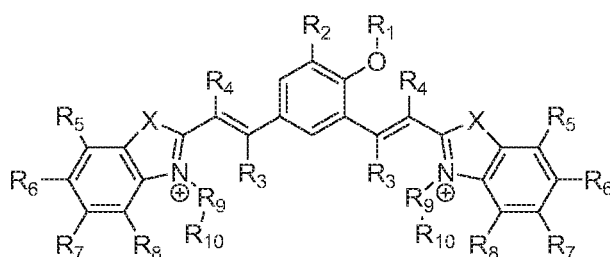
'R₅', 'R₆', 'R₇' or 'R₈' is selected from a group comprising H, OH, alkyl, aryl, halogen, nitro, sulfonates (SO₃⁻) and nitrile group;

'R₉' is selected from a group comprising H and -(CH₂)_n-, wherein 'n' is 1-6;

'R₁₀' is selected from a group comprising hydrogen, -OH, methyl, amine, terminal alkyne, alkene, alkyl acid, amine acid and sulfonates (SO₃⁻);

or its salt, derivative, tautomer, isomer, polymorph, analog, solvate or intermediate thereof;

a process for preparation of compound of Formula I:



Formula I

wherein, 'X' is selected from a group comprising oxygen, sulphur, and selenium;

'R₁' is selected from a group comprising alkyl chain, allyl group, aryl group, benzyl group, aryl and alky group, acid, amine, ester, methyl phenyl boronic acid, boronic ester, carbonate, phosphate, silane, quaternary ammonium, amide, imine and any other moiety which can be triggered by chemical and enzymatic tools;

'R₂' is selected from a group comprising H, OH, halogen, alkyl and substituted alkyl, and wherein, the halogen is selected from a group comprising bromide, chloride and iodide;

'R₃' or 'R₄' is selected from a group comprising H, alkyl, aryl, nitrile, acid and halogen, and wherein the halogen is selected from a group comprising, chloride, fluoride, bromide and iodide;

'R₅', 'R₆', 'R₇' or 'R₈' is selected from a group comprising H, OH, alkyl, aryl, halogen, nitro, sulfonates (SO₃⁻) and nitrile group;

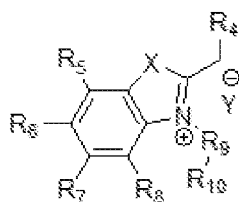
'R₉' is selected from a group comprising H and -(CH₂)_n-, wherein 'n' is 1-6;

'R₁₀' is selected from a group comprising hydrogen, -OH, methyl, amine, terminal alkyne, alkene, alkyl acid, amine acid and sulfonates (SO₃⁻);

or its salt, derivative, tautomeric form, isomer, polymorph, analog, solvate or intermediates thereof;

said process comprising:

- a. reacting compound of Formula II with compound of Formula III to obtain compound of Formula IV



Formula IV

wherein,

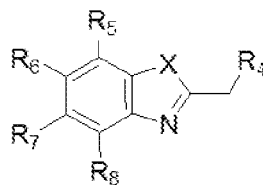
'R₄' is selected from a group comprising H, alkyl, aryl, nitrile, acid and halogen, and wherein the halogen is selected from a group comprising, chloride, fluoride, bromide and iodide;

'R₅', 'R₆', 'R₇' or 'R₈' is selected from a group comprising H, OH, alkyl, aryl, halogen, nitro, sulfonates (SO₃⁻) and nitrile group;

'R₉' is selected from a group comprising H and -(CH₂)_n-, wherein 'n' is 1-6; and

'R₁₀' is selected from a group comprising hydrogen, -OH, methyl, amine, terminal alkyne, alkene, alkyl acid, amine acid and sulfonates (SO₃⁻);

'Y' is either Br or I;

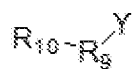


Formula II

wherein,

'R₄' is selected from a group comprising H, alkyl, aryl, nitrile, acid and halogen, and wherein the halogen is selected from a group comprising, chloride, fluoride, bromide and iodide; and

'R₅', 'R₆', 'R₇' or 'R₈' is selected from a group comprising H, OH, alkyl, aryl, halogen, nitro, sulfonates (SO₃⁻) and nitrile group;



Y = Br or I

Formula III

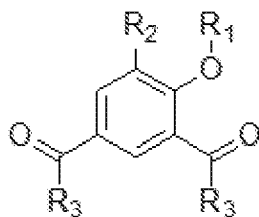
wherein,

'R₉' is selected from a group comprising H and -(CH₂)_n-, wherein 'n' is 1-6;

'R₁₀' is selected from a group comprising hydrogen, -OH, methyl, amine, terminal alkyne, alkene, alkyl acid, amine acid and sulfonates (SO₃⁻);

and,

- b. reacting the compound of Formula IV with compound of Formula V in presence of piperidine and alcohol.



Formula V

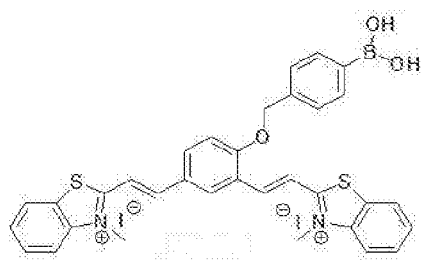
wherein,

‘R₁’ is hydrogen;

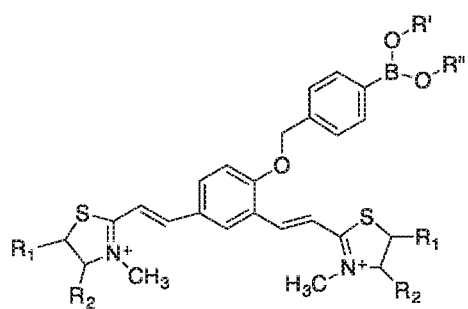
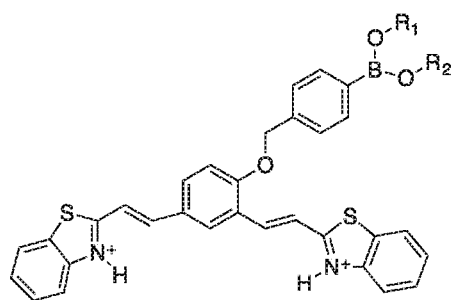
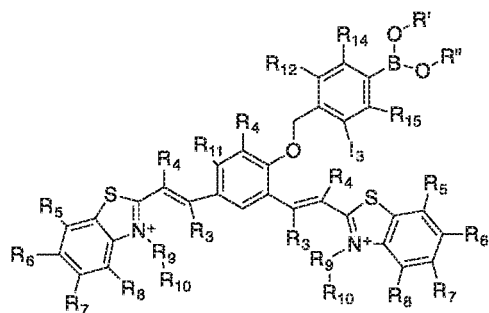
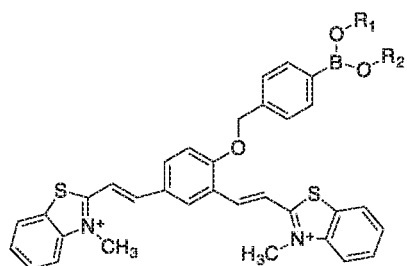
‘R₂’ is selected from a group comprising H, OH, halogen, alkyl and substituted alkyl, and wherein, the halogen is selected from a group comprising bromide, chloride and iodide; and

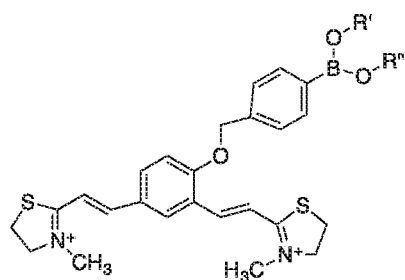
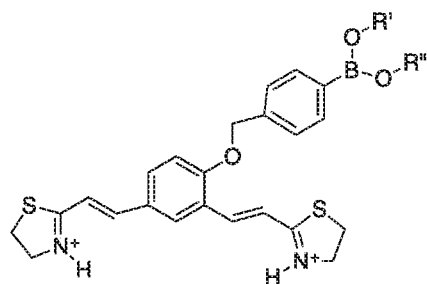
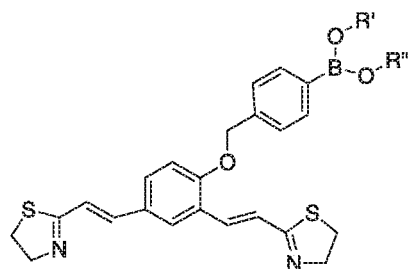
‘R₃’ is selected from a group comprising H, alkyl, aryl, nitrile, acid and halogen, and wherein the halogen is selected from a group comprising, chloride, fluoride, bromide and iodide;

a compound of Formula I as described above selected from



QCy-BA





R_1 to R_{14} or R' and R'' are independently selected from a group comprising alkyl, aryl, alicyclic, heterocyclic, hetero atom selected from a group comprising O, N, P, S, halogens, cyclic, acyclic ring system;

a pharmaceutical composition comprising the compound of Formula I its salt, derivative, tautomeric form, isomer, polymorph, analog, solvate and intermediates thereof, optionally along with at least one pharmaceutically acceptable excipient;

a method of detecting or quantifying the presence of reactive oxygen species (ROS) in a biological sample, said method comprising the act of contacting the compound of Formula I or its salt, derivative, tautomer, isomer, polymorph, analog, solvate or intermediates thereof, or the

composition of claim 8 with the sample and detecting the fluorescence indicative of the presence of ROS in the biological sample;

a method for detecting or quantifying a ROS compound in vivo in a subject, said method comprising:

- a. administering to a subject, a compound of Formula 1;
- b. allowing said compound of Formula 1 to react with a ROS; and
- c. detecting or quantifying the fluorescence, indicative of the presence of ROS compound in vivo;

a method of diagnosing a disease condition in a subject, said method comprising contacting the compound of Formula I or its salt, derivative, tautomer, isomer, polymorph, analog, solvate or intermediates thereof, or the composition of comprising compound of formula I with a sample obtained from the subject;

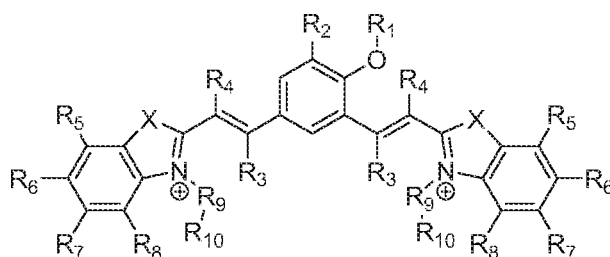
a method of inhibiting growth of a cell, said method comprising contacting the compound of Formula I or its salt, derivative, tautomer, isomer, polymorph, analog, solvate or intermediates thereof, the composition comprising compound of formula I with the cell;

a method of treating a disease characterized by abnormal levels of ROS in a subject, said method comprising step of administering the compound of Formula I or its salt, derivative, tautomer, isomer, polymorph, analog, solvate or its intermediates thereof, or the composition comprising compound of formula I in said subject to treat the disease; and

a kit for detecting reactive oxygen species (ROS) in a sample, wherein the kit comprising the compound of formula 1 or its salt, derivative, tautomer, isomer, polymorph, analog, solvate or intermediates thereof; or a composition comprising compound of formula I, wherein the said compound is present in an amount effective to detect the presence of ROS.

DETAILED DESCRIPTION OF THE DISCLOSURE

The present disclosure relates to a compound of Formula I:



Formula I

wherein,

‘X’ is selected from a group comprising oxygen, sulphur, and selenium;

‘R₁’ is selected from a group comprising alkyl chain, allyl group, aryl group, benzyl group, aryl and alkyl group, acid, amine, ester, methyl phenyl boronic acid, boronic ester, carbonate, phosphate, silane, quaternary ammonium, amide, imine and any other moiety which can be triggered by chemical or enzymatic tools;

‘R₂’ is selected from a group comprising H, OH, halogen, alkyl and substituted alkyl, and wherein, the halogen is selected from a group comprising bromide, chloride and iodide;

‘R₃’ or ‘R₄’ is selected from a group comprising H, alkyl, aryl, nitrile, acid and halogen, and wherein the halogen is selected from a group comprising, chloride, fluoride, bromide and iodide;

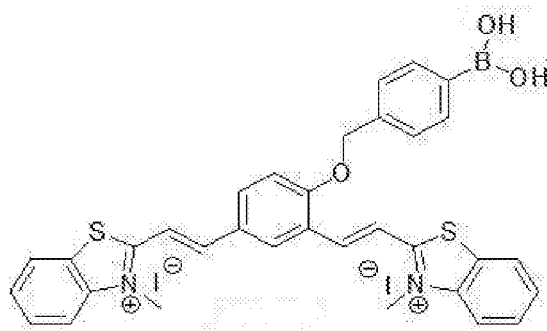
‘R₅’, ‘R₆’, ‘R₇’ or ‘R₈’ is selected from a group comprising H, OH, alkyl, aryl, halogen, nitro, sulfonates (SO₃⁻) and nitrile group;

‘R₉’ is selected from a group comprising H and -(CH₂)_n-, wherein ‘n’ is 1-6;

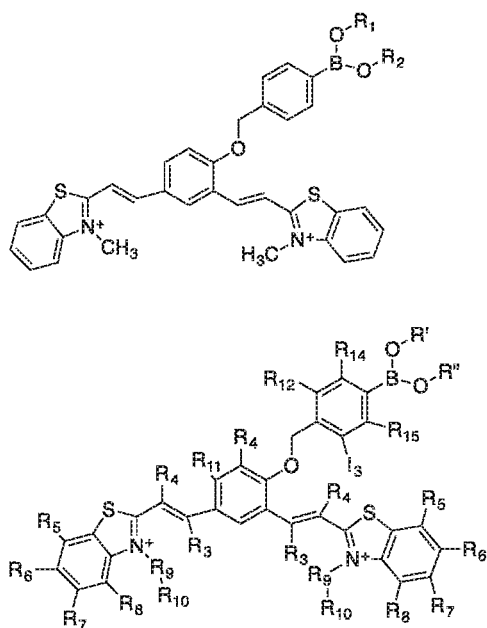
‘R₁₀’ is selected from a group comprising hydrogen, -OH, methyl, amine, terminal alkyne, alkene, alkyl acid, amine acid and sulfonates (SO₃⁻).

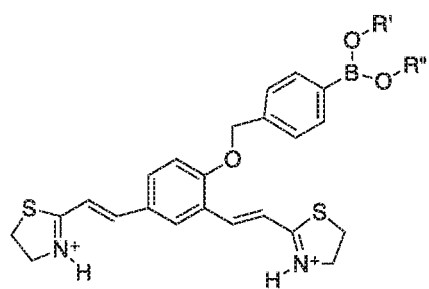
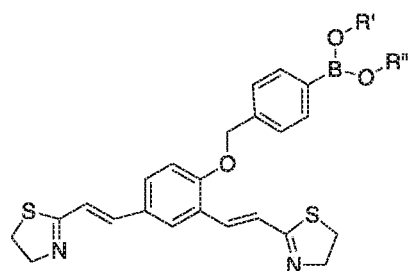
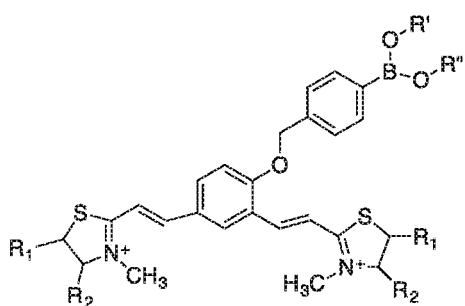
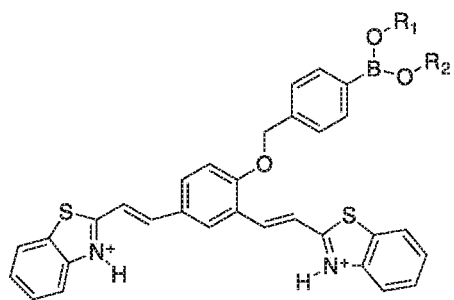
In an embodiment of the present disclosure, the compound of Formula I includes but is not limited to 4-((N, N-dimethyl, 2,4-bis((E)-2-(benzo[d]thiazol-2-ylinium) vinyl) phenoxy) methyl) phenylboronic acid.

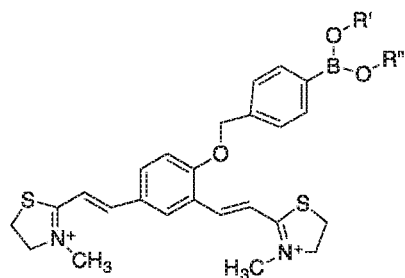
In another embodiment, the structure of 4-((N, N-dimethyl, 2,4-bis((E)-2-(benzo[d]thiazol-2-ylinium)vinyl)phenoxy)methyl)phenylboronic acid (QCy-BA) is provided below:



In yet another embodiment of the present disclosure, compound of Formula I is selected from







R_1 to R_{14} or R' and R'' are independently selected from a group comprising alkyl, aryl, alicyclic, heterocyclic, hetero atom selected from a group comprising O, N, P, S, halogens, cyclic, acyclic ring system;

In an embodiment of the present disclosure, tautomers, isomers, analogs, derivatives and salts of Formula I compounds are also provided.

The present disclosure relates to compound of Formula I which is a probe and is specific or selective to reactive oxygen species. In an exemplary embodiment, the Formula I compound QCy-BA is specific or selective to H_2O_2 .

In another embodiment of the present disclosure, Formula I compounds, particularly compounds with aryl boronates (R_1) are unique chemical moieties with selective reactivity towards the amphiphilic H_2O_2 , a desirable property to achieve specificity and selectivity over other biologically relevant reactive oxidative species (ROS).

In yet another embodiment, initially, the boronate functionality in Formula I acts as an electrophilic center and reacts with the nucleophile to generate the tetrahedral-boronate complex. Subsequently, the carbon-boron (C-B) bond becomes labile and acts as a nucleophile towards the electrophilic oxygen center of H_2O_2 . In particular, the aryl boronate functionality becomes a specific reorganization center for H_2O_2 among all other biological oxygen metabolites and ROS, which operate through one electron transfer or electrophilic oxidation pathways.

In an embodiment, the Formula I compound QCy-BA is QCy-DT compound (a DNA minor groove binder) functionalized with methyl phenyl boronic acid. QCy-BA possesses two positively charged nitrogen atom-containing benzothiazoles, with distinct conjugation patterns around the central phenolic moiety derivatized with phenylboronic acid functionality. Furthermore, the electron delocalization in QCy-BA is disrupted as a consequence of masking

the central phenolic hydroxyl with phenyl boronic acid functionality. Upon slicing of the phenyl boronic acid functionality in response to the H_2O_2 stimulus, QCy-BA transforms into the negatively charged phenolate of QCy-DT. The generation of phenolate restores the electron transfer towards one of the positively charged nitrogen atoms of the benzothiazole acceptor. This restores internal charge transfer (ICT) to generate a highly electron delocalized π -system with NIR-fluorescence in the presence of DNA (as shown in Figure 1(a)).

In another embodiment, Formula I compounds, particularly, phenyl boronic acid-functionalized quinone-cyanine (QCy-BA) in combination with AT-rich DNA (Drew-AT), *i.e.*, QCy-BA $\subset$ Drew-AT as a stimuli-responsive NIR fluorescence probe for *in vitro* levels of H_2O_2 is presented. In response to cellular H_2O_2 stimulus, QCy-BA converts into QCy-DT, a one-donor-two-acceptor (D2A) system that exhibits switch-on NIR fluorescence upon binding to DNA minor groove. Fluorescence studies on combination probe QCy-BA $\subset$ Drew-AT show strong NIR fluorescence selectively in the presence of H_2O_2 . Further, glucose oxidase (Gox) assay confirms that the combination probe QCy-BA $\subset$ Drew-AT is highly efficient for probing H_2O_2 generated *in situ* through Gox-mediated glucose oxidation. Quantitative analysis through fluorescence plate reader, flow cytometry and live imaging approaches show that the Formula I compounds including QCy-BA is a promising probe to detect normal as well as elevated levels of H_2O_2 produced by EGF/Nox pathways and post-genotoxic stress in primary cells as well as senescent cells. Thus, the Formula I compounds including QCy-BA, in combination with exogenous or cellular DNA are versatile probes to quantify and image H_2O_2 in normal and disease-related cells.

In an embodiment, the phrase AT-rich DNA refers to DNA which consists more percentage of AT-base pairs than GC-base pairs.

In another embodiment, functionalization of the compounds of Formula I (such as QCy-BA) with a stimuli-responsive appendage (H_2O_2) is a promising and efficient method to *in situ* generate the active probe (QCy-DT) in response to H_2O_2 . Such a probe with signal capturing and amplification assisted by an additional recognition event, DNA-binding in the present disclosure, is highly advantageous for probing specific biological activity in a cellular environment.

Structurally, QCy-DT has a free hydroxyl group readily available for functionalization with a large number of chemically or enzymatically cleavable appendages to make it a versatile and promising stimuli-responsive probe.

In another embodiment, the probe QCy-BA in response to specific stimulus (chemical or enzyme), is cleaved to release an NIR-fluorescence ready QCy-DT probe, which upon binding the minor groove of DNA fluoresces strongly, thus, aiding the imaging and quantification of the stimulus.

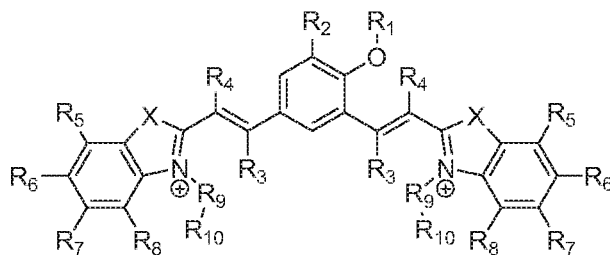
In another embodiment, the probe QCy-BA acts as a stimulus receiver or protectant and upon sensing the stimulus gets converted to the activated dye/probe QCy-DT in order to bind to the minor groove of the DNA.

In another embodiment, functionalizing the DNA binding fluorescence probe QCy-DT with aryl boronates is an attractive strategy for the development of a stimuli-responsive fluorescence probe (Formula I compounds) for H_2O_2 . Accordingly, QCy-DT hydroxyl group is functionalized to obtain methyl phenyl boronic acid (QCy-BA, Figure 1(a)), which reacts selectively with H_2O_2 to release the parent DNA binding dye. In another embodiment, phenyl boronate is the preferred appendage owing to the fact that the reaction between H_2O_2 and boronic acid or ester is highly chemospecific, bioorthogonal and biocompatible while the byproducts are non-toxic to living cells.

In an exemplary embodiment, methyl phenyl boronic acid conjugated one-donor-two-acceptor (D2A) π -electron-based quinone-cyanine (QCy-BA) moiety is provided, which in combination with AT-rich DNA (such as Drew-AT) acts as a stimuli-responsive NIR fluorescence probe for H_2O_2 (Figure 1(a)). The major advantage with NIR fluorescence combination probe is that it circumvents the false positive results by means of a double check on the signal outcome and background fluorescence from *in vitro* and *in vivo* studies. In the presence of H_2O_2 , QCy-BA releases non-fluorescent QCy-DT, which upon binding AT-rich DNA shows enhanced fluorescence in the NIR region. Nuclear magnetic resonance (NMR) and UV-vis absorption studies reveal the efficient conversion of QCy-BA to QCy-DT and *p*-quinonemethide through selective 1,6-elimination rearrangement in the presence of H_2O_2 over other ROS (Figure 1(a)). Fluorescence studies of QCy-BA show strong NIR fluorescence enhancement in the presence of

H₂O₂ and AT-rich DNA (Drew-AT), which clearly endorses the use of combination probes such as QCy-BA⊂Drew-AT for the fluorometric detection of H₂O₂ ratiometrically. Glucose oxidase assay confirm that the combination probe QCy-BA⊂Drew-AT is capable of monitoring *in situ* generated H₂O₂ by the oxidation of glucose using glucose oxidase (GOx). Live cell imaging studies show the staining of cell nucleus by the *in situ* generated NIR fluorescence-ready probe in the presence of H₂O₂. Fluorescence activated cell sorting (FACS) analysis show that probe QCy-BA is equally efficient in detecting the normal and *in situ* generated H₂O₂ in response to Nox-mediated growth factor signaling pathways. Further, probe QCy-BA is capable of detecting elevated levels of H₂O₂ generated either through insult by doxorubicin (Dox) or 5-bromo-2'-deoxyuridine (BrdU) in primary or cancer cells, which results in the induction of cellular senescence. Therefore, Formula I compounds such as probe QCy-BA in combination with exogenous or cellular DNA is a promising stimuli-responsive NIR fluorescence combination probe for investigating H₂O₂ production and concentration levels in living cells, which can further assist the imaging and diagnosis of disease-related cells.

The present disclosure also relates to a process for preparation of compound of Formula I:



Formula I

wherein, 'X' is selected from a group comprising oxygen, sulphur, and selenium;

'R₁' is selected from a group comprising alkyl chain, allyl group, aryl group, benzyl group, aryl and alkyl group, acid, amine, ester, methyl phenyl boronic acid, boronic ester, carbonate, phosphate, silane, quaternary ammonium, amide, imine and any other moiety which can be triggered by chemical and enzymatic tools;

'R₂' is selected from a group comprising H, OH, halogen, alkyl and substituted alkyl, and wherein, the halogen is selected from a group comprising bromide, chloride and iodide;

'R₃' or 'R₄' is selected from a group comprising H, alkyl, aryl, nitrile, acid and halogen, and wherein the halogen is selected from a group comprising, chloride, fluoride, bromide and iodide;

'R₅', 'R₆', 'R₇' or 'R₈' is selected from a group comprising H, OH, alkyl, aryl, halogen, nitro, sulfonates (SO₃⁻) and nitrile group;

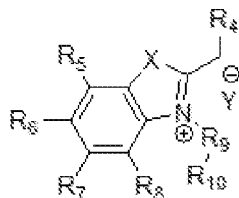
'R₉' is selected from a group comprising H and -(CH₂)_n-, wherein 'n' is 1-6;

'R₁₀' is selected from a group comprising hydrogen, -OH, methyl, amine, terminal alkyne, alkene, alkyl acid, amine acid and sulfonates (SO₃⁻);

or its salt, derivative, tautomeric form, isomer, polymorph, analog, solvate and intermediates thereof;

said process comprising:

- b. reacting compound of Formula II with compound of Formula III to obtain compound of Formula IV



Formula IV

wherein,

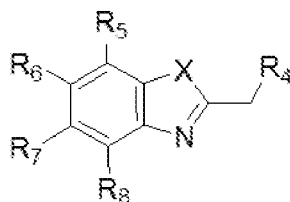
'R₄' is selected from a group comprising H, alkyl, aryl, nitrile, acid and halogen, and wherein the halogen is selected from a group comprising, chloride, fluoride, bromide and iodide;

'R₅', 'R₆', 'R₇' or 'R₈' is selected from a group comprising H, OH, alkyl, aryl, halogen, nitro, sulfonates (SO₃⁻) and nitrile group;

'R₉' is selected from a group comprising H and -(CH₂)_n-, wherein 'n' is 1-6; and

'R₁₀' is selected from a group comprising hydrogen, -OH, methyl, amine, terminal alkyne, alkene, alkyl acid, amine acid and sulfonates (SO₃⁻);

'Y' is either Br or I;



Formula II

wherein,

'R₄' is selected from a group comprising H, alkyl, aryl, nitrile, acid and halogen, and wherein the halogen is selected from a group comprising, chloride, fluoride, bromide and iodide; and

'R₅', 'R₆', 'R₇' or 'R₈' is selected from a group comprising H, OH, alkyl, aryl, halogen, nitro, sulfonates (SO₃⁻) and nitrile group;



Y = Br or I

Formula III

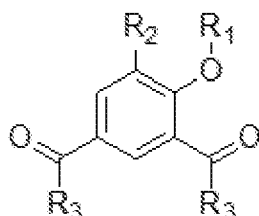
wherein,

'R₉' is selected from a group comprising H and -(CH₂)_n-, wherein 'n' is 1-6;

'R₁₀' is selected from a group comprising hydrogen, -OH, methyl, amine, terminal alkyne, alkene, alkyl acid, amine acid and sulfonates (SO₃⁻);

and,

- b. reacting the compound of Formula IV with compound of Formula V in presence of piperidine and alcohol.



Formula V

wherein,

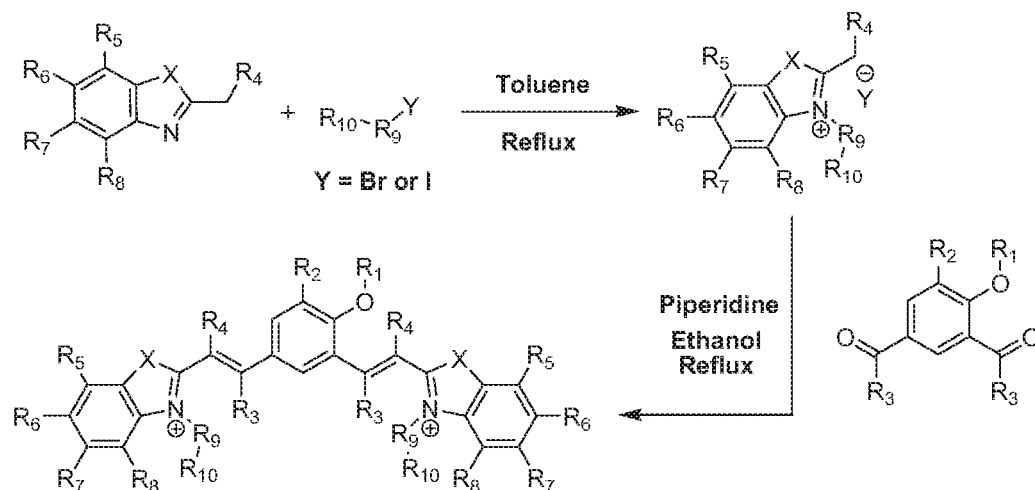
'R₁' is hydrogen;

'R₂' is selected from a group comprising H, OH, halogen, alkyl and substituted alkyl, and wherein, the halogen is selected from a group comprising bromide, chloride and iodide; and

'R₃' is selected from a group comprising H, alkyl, aryl, nitrile, acid and halogen, and wherein the halogen is selected from a group comprising, chloride, fluoride, bromide and iodide;

Thus, the present disclosure relates to a method for preparing the compounds of Formula I, wherein the method comprises the act of reacting N-alkylted benzothiazole derivatives with 4-hydroxy isophthalaldehyde derivatives (Scheme 1).

Scheme 1: General scheme for the synthesis of compounds under Formula 1



In an embodiment, the Formula I compound 4-((N,N-dimethyl,2,4-bis((E)-2-(benzo[d]thiazol-2-ylinium)vinyl)phenoxy)methyl)phenylboronic acid [QCy-BA] is prepared by

- a. reacting 2-methyl benzothiazole with methyl iodide to obtain N-methyl-2-methylbenzothiazole; and
- b. reacting the N-methyl-2-methylbenzothiazole with 4-(4-(4,4,5,5-tetramethyl-1,3,2-dioxaborolan-2-yl)benzyloxy)isophthalaldehyde in presence of piperidine and ethanol.

The processes as described above, wherein said process is carried out at a temperature ranging from about 30°C to 80°C, and for a time period ranging from about 2 hours to 24 hours.

The process as described above, wherein the steps a) and b) further comprise isolation, purification or a combination thereof of the corresponding product; wherein said isolation and purification is carried out by acts selected from a group comprising addition of solvent, washing with solvent, quenching, filtration, extraction, chromatography and combinations thereof.

The present disclosure also relates to pharmaceutical compositions or formulations comprising one or more compounds of Formula I, optionally along with pharmaceutically acceptable excipients. In an embodiment, the pharmaceutically acceptable excipient is selected from a group comprising adjuvant, diluent, carrier, granulating agents, binding agents, lubricating agents,

disintegrating agent, sweetening agents, glidant, anti-adherent, anti-static agent, surfactant, anti-oxidant, gum, coating agent, coloring agent, flavouring agent, coating agent, plasticizer, preservative, suspending agent, emulsifying agent, plant cellulosic material, spheronization agent, other conventionally known pharmaceutically acceptable excipient or any combination of excipients thereof. In another embodiment, the pharmaceutical composition of the present disclosure is administered to a subject through modes selected from a group comprising intravenous administration, intramuscular administration, intraperitoneal administration, hepatoporal administration, intra articular administration and pancreatic duodenal artery administration, or any combination thereof.

The present disclosure also provides for a method of protecting a fluorescent probe, particularly QCy-DT, wherein the method comprises the step of reacting QCyDT with a protecting agent. In an embodiment, the protecting agent is selected from group comprising phenyl boronate, esters, phosphates, ethers and carbonates. In another embodiment, the method provides a protected fluorescent probe QCy-BA.

The present disclosure provides a method of detecting or quantifying the presence of reactive oxygen species (ROS) in a biological sample, said method comprising the act of contacting the compound of Formula I or its salt, derivative, tautomer, isomer, polymorph, analog, solvate or intermediates thereof, or the composition comprising a compound of formula I with the sample and detecting the fluorescence indicative of the presence of ROS in the biological sample. In an embodiment of the present disclosure, ROS compound is selected from a group comprising hydrogen peroxide, tertbutyl hydroperoxide, superoxide, hydroxyl radical, tert-butoxy radical, hypochlorite and ONOO^- . In another embodiment of the present disclosure, the biological sample is cells, tissue, biological fluids, or combinations thereof.

The present disclosure provides a method for detecting or quantifying a ROS compound in vivo in a subject, said method comprising:

- a. administering to a subject, a compound of Formula 1;
- b. allowing said compound of Formula 1 to react with a ROS; and
- c. detecting or quantifying the fluorescence, indicative of the presence of ROS compound in vivo.

In an embodiment of the present disclosure, detecting or quantifying further comprises detecting the location of ROS compound in said subject. In another embodiment of the present disclosure, the detection or quantifying is by technique selected from fluorescence microscopy, fluorescence spectroscopy, confocal laser scanning microscopy, total internal reflection fluorescence microscopy, Near infra-red fluorescence and combinations thereof. In yet another embodiment of the present disclosure, the compound of Formula I is provided as a combination probe with DNA sequences for detecting and quantifying the ROS, and wherein the DNA sequence is exogenous DNA or endogenous nuclear DNA, or a combination thereof.

In an embodiment of the present disclosure, the compound in the presence of ROS is cleaved to release a fluorescent probe quinone cyanine-dithiazole (QCy-DT), which is capable of binding to the AT-rich DNA for fluorometric detection and quantification of the ROS

The present disclosure provides for a method of diagnosing a disease condition in a subject, wherein the method comprises the step of contacting the compound of Formula I the composition comprising a compound of formula I with sample obtained from the subject. In an embodiment, the subject is a mammal or a plant, and wherein the sample from mammal is selected from group comprising blood, serum, in-vitro sample, synthetic sample, any bodily fluid and combinations thereof. In an embodiment, the sample is selected from group comprising blood, serum, *in-vitro* sample, synthetic sample, any bodily fluid and combinations thereof. In an embodiment, the disease is caused by the excessive presence of ROS leading to oxidative stress. In another embodiment, the ROS includes but is not limited to H_2O_2 . In another embodiment, the disease is selected from a group comprising cancer, cardiovascular dysfunction, neurodegenerative diseases, gastroduodenal pathogenesis, inflammatory disorders, metabolic dysfunction of organs, premature aging and combinations thereof. In an embodiment of the present disclosure, the disease is diagnosed by detecting and optionally quantifying reactive oxygen species (ROS) in the sample. In an exemplary embodiment, the Formula I compound detects ROS, to diagnose the disease condition.

The present disclosure also relates to the use of Formula I compounds for detecting ROS in a sample.

The present disclosure also relates to a method of inhibiting growth of a cell, said method comprising contacting the compound of Formula I or its salt, derivative, tautomer, isomer, polymorph, analog, solvate or intermediates thereof, the composition comprising compound of formula I with the cell. In an embodiment, the cell is unicellular an eukaryotic cell selected from a group comprising cancerous cells, cells infected with microorganisms, parasite or unicellular protozoan and other cells characterized by abnormal levels of ROS, and wherein the parasite is *Plasmodium*.

The present disclosure also relates to a method of treating a disease characterized by abnormal levels of ROS in a subject, said method comprising step of administering the compound of Formula I or its salt, derivative, tautomer, isomer, polymorph, analog, solvate or its intermediates thereof, or the composition comprising compound of formula I in said subject to treat the disease.

The present disclosure also relates to the use of compound of Formula I or its salt, derivative, tautomer, isomer, polymorph, analog, solvate or intermediates thereof, or the composition of claim 8 as a probe for detecting and optionally quantifying ROS, diagnosing a disease caused by abnormal ROS levels, inhibiting growth of a cell, treating a disease characterized by abnormal levels of ROS.

The present disclosure also relates to a kit for detecting reactive oxygen species (ROS) in a sample, wherein the kit comprising the compound of formula I or its salt, derivative, tautomer, isomer, polymorph, analog, solvate or intermediates thereof; or a composition comprising compound of formula I, wherein the said compound is present in an amount effective to detect the presence of ROS.

The present disclosure is further described with reference to the following examples, which are only illustrative in nature and should not be construed to limit the scope of the present disclosure in any manner.

EXAMPLES

Standard Parameters:

All the chemicals, reagents, self-complementary Drew-AT, Hoechst 33258, Phosphate buffer saline (PBS), 2',7'-dichlorofluorescein (DCFDA), 5-bromo-2'-deoxyuridine (BrdU) and doxorubicin (dox) and N-acetyl-L-cysteine (NAC) are purchased from Sigma-Aldrich. All synthesized compounds are purified by column chromatography using Rankem silica gel (60-120 mesh). ^1H and ^{13}C NMR spectra are recorded on a Bruker AV-400 MHz spectrometer with chemical shifts reported as parts per million (*ppm*) (in CDCl_3 , $\text{DMSO}-d_6$, tetramethylsilane as an internal standard) at 20°C. High resolution mass spectra (HRMS) are obtained on Agilent Technologies 6538 UHD Accurate-Mass Q-TOF LC/MS spectrometer. The UV-vis absorption and emission spectra are recorded on Agilent Technologies Cary series UV-vis-NIR absorbance and Cary Eclipse fluorescence spectrophotometers respectively. UV-vis absorption and emission spectra are measured in quartz cuvettes of 1 cm path length. HeLa cells and MRC 5 PDL 23 cells used in the biological studies are obtained from "Molecular reproduction, development and genetics lab Indian institute of science Bangalore, India".

Sample preparation for UV-vis and Fluorescence measurements

All biophysical studies (UV-vis and fluorescence) are carried out at the concentration 0-5 μM of probe, volume = 500 μL , temperature = 25 °C and time of incubation = 2 min.

Stock solution of probe QCy-BA is prepared in milli molar concentration in milli Q-water (MQ-water) and stored at -10 °C. DNA stock solutions are prepared by dissolving oligos in double distilled water in the order of 10^{-4} M. Double stranded DNA samples are prepared in PBS (10 mM, pH = 7.4) buffer solution and subjected to annealing by heating up to about 85 °C for 15 minutes and subsequently cooled to room temperature for 7 h and stored in refrigerator for about 4 hours.

HeLa cells maintenance: Human cervix carcinoma cell line (HeLa) is cultured in DMEM (Dulbecco's Modified Eagle's Medium) with 10 % FBS (Fetal Bovine Serum). The antibiotics penicillin and streptomycin (1 %) is mixed with 10 % FBS medium. The cells are incubated at 37 °C temperature and 5 % CO_2 humidified chamber. All cell culture work is carried out under laminar flow hood and auspicious conditions.

Cytotoxicity studies on HeLa cells (MTT Assay): MTT [(3-(4,5-dimethylthiazol-2yl)-2,5-diphenyltetrazolium bromide] assay is carried out with probe QCy-BA on HeLa cells to determine the cytotoxicity effect. In a tissue cultured 96-well plate, 10,000 cells per well are plated and incubated for 24 h, for cells to attach. After completion of 24 h incubation, the attached and healthy cells are treated with various concentrations (25 μ M, 12.5 μ M, 6.25 μ M, 3.125 μ M and 0 μ M) of probe QCy-BA and further incubated for 24 h. All the treatments are carried out in triplicates. The required concentrations of QCy-BA are made from stock solution in 0.2% DMEM. Stock solution of probe QCy-BA (1 mg/mL) is made in water. Four hours before stipulated time of experiment, MTT-solution (5 mg/mL of 20 μ L) is added in each well and incubated to form formazan crystals. The culture medium is completely removed by 1 mL pipette and 200 μ L of DMSO is added to dissolve formazan crystals. The purple colored formazan is estimated by determining absorbance at 590 nm with the help of spectrophotometer (Bio-RAD model 1680, Microplate reader). The results are shown in bar graphs (concentration of QCy-BA *vs* % cell viability).

Exogenous and endogenous detection of H₂O₂ in HeLa cells by QCy-BA: In each well, 3×10^6 cells are plated in 12-well tissue culture plates and incubated for 24 hours. These cells are serum deprived for 1 h. In addition, the serum deprived cells of 6-well are treated with N-acetyl-L-cysteine (NAC) (8 mM) solution and incubated for 1 hour and cells are treated with probe QCy-BA (5 μ M) and incubated for 30 minutes. After 30 minutes incubation, the cells are washed with DPBS (Dulbecco's Phosphate buffer saline) to remove the excesses of QCy-BA. These cells are harvested after trypsinization. Exogenously, H₂O₂ (100 μ M) is added to QCy-BA and NAC + QCy-BA treated cells and incubated for 15 min. These samples are subjected to FACS analysis.

Epidermal growth factor (EGF) produced H₂O₂ detection by QCy-BA in HeLa cells: To determine the EGF produced ROS in cells by QCy-BA. In a 12-well plate, 3×10^6 HeLa cells are plated in each well and incubated for 24 hour. These cells are serum deprived for 1h. Cells are incubated with EGF (500 ng/mL) for 40 minutes and further treated with NAC (8 mM) for 1 hour. Then cells are treated with QCy-BA (5 μ M) for 30 minutes. After 30 minutes incubation of cells with QCy-BA, cells are washed with DPBS (Dulbecco's Phosphate Buffer Saline) to remove the excesses of QCy-BA. These cells are harvested after trypsinization. The samples are subjected to FACS.

Immunofluorescence studies with QCy-BA for detection of H_2O_2 in HeLa cells: An Immunofluorescence studies are carried out on HeLa cells to validate exogenous and endogenous detection of H_2O_2 by QCy-BA. The HeLa cells (10,000 cells) are grown on cover slips. These cells are treated with 5 μM concentration of QCy-BA for 30 minutes. The cells are washed with DPBS for several times to remove the excesses of QCy-BA. The cells are treated with H_2O_2 (100 μM) for 30 minutes. These samples are subjected to confocal microscopy for immunofluorescence images.

Detection limit of H_2O_2 in presence of QCy-BA: Concentration dependent studies are performed using microplate reader. In the well-plates, first QCy-BA (5 μM) in buffer solution is taken, then increasing concentration of H_2O_2 from 0 to 100 μM is added. Upon excitation at 400 nm, the emission at 550 nm as function time after addition of H_2O_2 is collected. The fluorescence intensity at 550 nm is plotted as a function of concentration of hydrogen peroxide and each experiment is done in triplicates.

Detection of ROS using fluorescence plate reader: The cells are incubated with QCy-BA (5 μM) for 30 minutes in dark, washed with PBS and analysed to detect QCy-BA dye fluorescence using Infinite M1000 Pro, Tecan, Austria). Wavelengths used for excitation and emission for QCy-BA dye is 400nm/650nm. In all ROS measurement assays are done using plate reader and after fluorescence measurements, cells are washed, trypsinized and counted to estimate fluorescence per cell recordings.

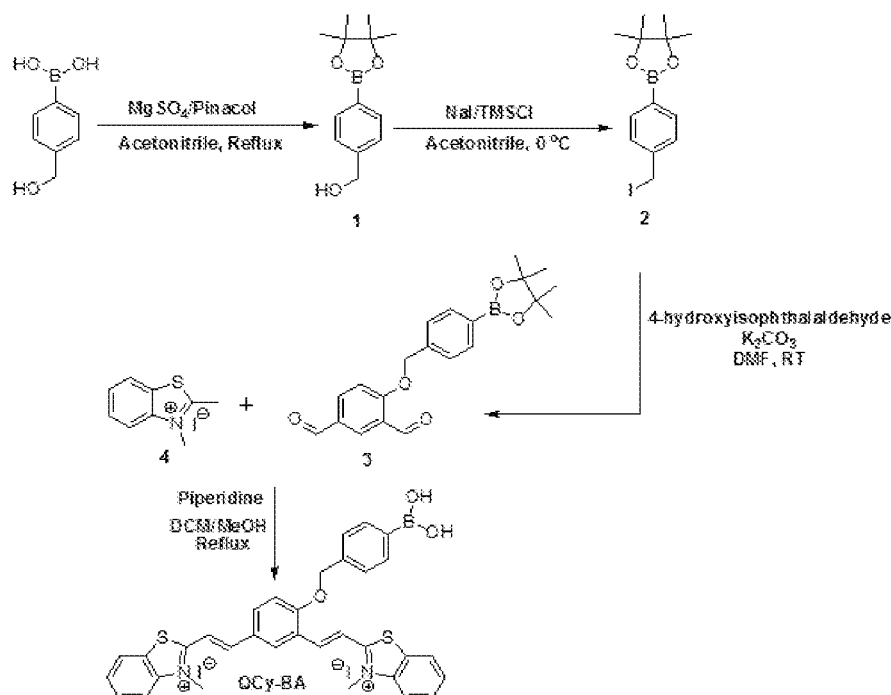
Live cell imaging of MRC5 cells: MRC5 PDL 23 cells are seeded overnight and treated with Hydrogen Peroxide to do live cell imaging after addition of QCy-BA (5 μM) for 30 minutes. Images are acquired using Olympus IX 83 inverted epifluorescence microscope using a 20X objective.

Absorption and emission spectra: The UV-vis absorption spectra are recorded on a Perkin Elmer Model Lambda 900 spectrophotometer. Emission spectra are recorded on Perkin Elmer Model LS 55 spectrophotometer. Temperature dependent absorption measurements (UV-Vis melting studies) are carried out on Cary 5000 UV-vis-NIR spectrophotometer equipped with Cary temperature controller in the range of 10 $^{\circ}C$ to 90 $^{\circ}C$ with ramp rate of 1 $^{\circ}C$ /min.

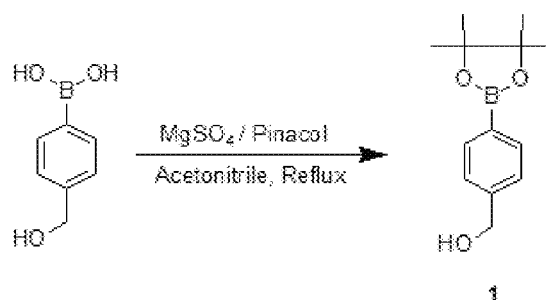
Example 1: Preparation of QCy-BA

Synthesis of QCy-BA is achieved by treating 4-(hydroxymethyl)phenyl boronic acid with a pinacol in the presence of magnesium sulfate in acetonitrile to obtain 4-(hydroxymethyl)phenylboronic ester (**1**). The phenylboronic ester **1** is treated with NaI and trimethylsilyl chloride in acetonitrile at 4°C to give 4-(iodomethyl)phenylboronic ester (**2**). The 4-(iodomethyl)phenyl boronic ester (**2**) is coupled to 4-hydroxy isophthalaldehyde using potassium carbonate as a base in dimethylformamide (DMF) at room temperature to obtain phenyl boronic ester dialdehyde (**3**) in good yield. Finally, the dialdehyde (**3**) is coupled with *N*-methylated benzothiazolein the presence of piperidine to yield the probe QCy-BA. All the intermediates and probe QCy-BA are characterized by NMR and high-resolution mass spectroscopy (HRMS).

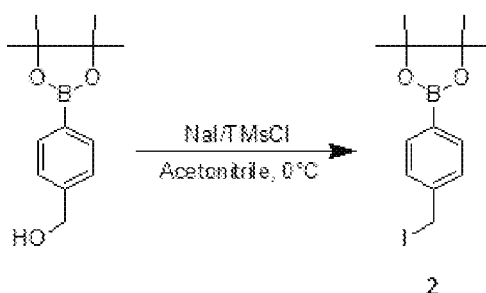
The overall synthesis of QCy-BA has been described below in a schematic fashion. Also, the preparation of the intermediates has been described below individually which leads to the preparation of QCy-BA.



Scheme 2: Synthesis of QCy-BA

a) Synthesis of Compound 1 (4-(4,4,5,5-tetramethyl-1,3,2-dioxaborolan-2-yl)phenyl) methanol

To a stirred solution of 4-(hydroxymethyl) phenylboronic acid (0.4 g, 2.63 mmol) in acetonitrile (15 mL), MgSO_4 (3 g) and pinacol (0.37 g, 3.15 mmol) are added. The reaction mixture is heated up to about 80°C and allowed to reflux for about 24 hours. After completion of the reaction, solvent is evaporated under vacuum. The crude mixture is dissolved in dichloromethane and filtered. The obtained product (4-(4,4,5,5-tetramethyl-1,3,2-dioxaborolan-2-yl)phenyl) methanol (i.e. compound **1**) is used for further reaction without purification.

b) Synthesis of compound 2 (2-(4-(iodomethyl)phenyl)-4,4,5,5-tetramethyl-1,3,2-dioxaborolane)

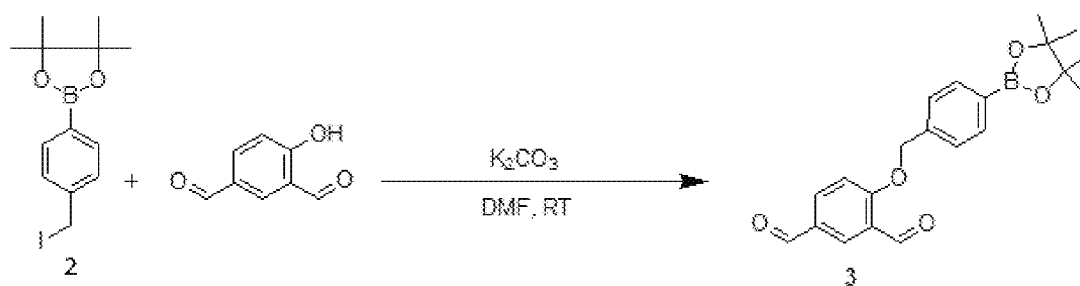
To a stirred solution of 4-(4,4,5,5-tetramethyl-1,3,2-dioxaborolan-2-yl)phenyl) methanol (**1**) (0.55 g, 2.35 mmol) in acetonitrile (20 mL), sodium iodide (1.1 g, 7.05 mmol) and Trimethylsilyl chloride (0.65 mL, 7.05 mmol) are added at about 0°C . The reaction mixture is allowed to stir at room temperature for about 1 hour. After completion of the reaction, solvent is evaporated under vacuum. The crude product is dissolved in saturated solution of $\text{Na}_2\text{S}_2\text{O}_3$ to quench the unreacted iodide and the product is extracted with dichloromethane. The crude product is purified by

column chromatography on silica gel using ethyl acetate/hexane (5:95) as an eluent to give product 2-(4-(iodomethyl)phenyl)-4,4,5,5-tetramethyl-1,3,2-dioxaborolane (**2**) in excellent yield (90%).

^1H NMR (400 MHz, CDCl_3) δ_{ppm} 7.73 (d, $J = 8$ Hz, 2H), 7.37 (d, $J = 8$ Hz, 2H), 4.45 (s, 2H), 1.34 (s, 12H) (Figure 18).

^{13}C NMR (100 MHz, CDCl_3) δ_{ppm} 142.3, 135.3, 128.0, 24.9, 5.4 (Figure 19).

c) Synthesis of compound 3 (4-(4-(4,4,5,5-tetramethyl-1,3,2-dioxaborolan-2-yl)benzyloxy)isophthalaldehyde.

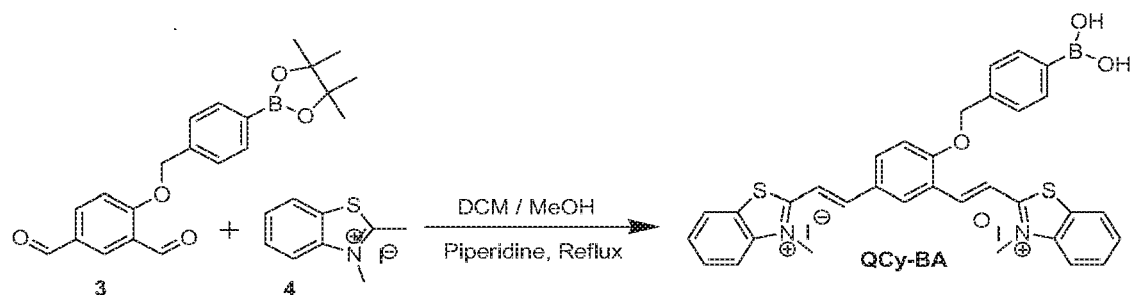


To a stirred solution of 4-hydroxyisipthalaldehyde (0.11 g, 0.73 mmol) in Dimethylformamide (DMF) (5 mL), K_2CO_3 (0.3 g, 2.17 mmol) is added and allowed to stir for about 20 minutes. After about 20 minutes, 2-(4-(iodomethyl)phenyl)-4,4,5,5-tetramethyl-1,3,2-dioxaborolane (**2**) (0.3 g, 0.87 mmol) is added and stirred overnight at room temperature (RT). The completion of reaction is monitored by TLC. After completion of the reaction, solvent is evaporated and product is extracted with diethylether (3×100 mL). The crude product is purified by column chromatography on silica gel using ethyl acetate/hexane (20:80) as an eluent to obtain compound **3** in good yield (60%).

^1H NMR (400 MHz, CDCl_3) δ_{ppm} 10.56 (s, 1H), 9.95 (s, 1H), 8.35 (d, $J = 2.4$ Hz, 1H), 8.08 (dd, $J = 2$ Hz, 8.8 Hz, 1H), 7.86 (d, $J = 8$ Hz, 2H), 7.44 (d, $J = 8$ Hz, 2H), 7.17 (d, $J = 8$ Hz, 1H), 5.32 (s, 2H), 1.35 (s, 12H) (Figure 20).

^{13}C NMR (100 MHz, CDCl_3) δ_{ppm} 190.1, 188.5, 164.9, 137.9, 135.5, 135.3, 131.9, 129.9, 126.5, 125.2, 113.7, 84.0, 71.0, 24.9 (Figure 21).

d) Synthesis of QCy-BA



To a stirred solution of N-methylated, 2-methyl benzothiazolinium (compound 4) (80 mg, 0.27 mmol) in methanol (10 mL) and dichloromethane (5 mL), piperidine (8 μ L) is added and allowed to stir for about 10 minutes. After about 10 minutes, 4-(4-(4,4,5,5-tetramethyl-1,3,2-dioxaborolan-2-yl)benzyloxy)isophthalaldehyde (compound 3) (45 mg, 0.12 mmol) in dichloromethane (DCM) (2 mL) is added and heated up to about 50°C for about 3 hours. After the completion of reaction, the solvent is evaporated. The crude brown color solid is washed with diethylether (50 mL) to remove the unreacted starting materials. The brown solid is dissolved in acetonitrile/water mixture and purified by reverse phase HPLC using 0.1% trifluoroacetic acid (TFA) in water/acetonitrile (50-100%) as a mobile phase to obtained boronic acid conjugate (QCy-BA) in moderate yield 30%.

$^1\text{H-NMR}$ (400 MHz, $\text{DMSO-}d_6$) δ_{ppm} 8.69 (d, $J = 2\text{ Hz}$, 1H), 8.44 (ddd, $J = 2.8\text{ Hz}$, $J = 4.0\text{ Hz}$, $J = 7.6\text{ Hz}$, 2H), 8.34-8.25 (m, 4H), 8.21 (d, $J = 4.2\text{ Hz}$, 1H), 8.16 (d, $J = 12.4\text{ Hz}$, 1H), 8.10 (d, $J = 8.2\text{ Hz}$, 1H), 7.92-7.88 (m, 4H), 7.81 (td, $J = 0.8\text{ Hz}$, $J = 7.6\text{ Hz}$, 2H), 7.56 (dd, $J = 4\text{ Hz}$, $J = 8.4\text{ Hz}$, 3H), 5.48 (s, 2H), 4.39 (s, 3H), 4.25 (s, 3H) (Figure 22).

$^{13}\text{C-NMR}$ (100 MHz, $\text{DMSO-}d_6$) δ_{ppm} 171.9, 171.8, 160.6, 158.3, 158.0, 147.0, 142.1, 142.0, 141.6, 137.5, 135.0, 134.5, 132.3, 129.5, 129.4, 128.6, 128.4, 127.9, 127.8, 127.3, 127.0, 124.3, 124.2, 123.1, 118.1, 117.0, 116.9, 115.6, 115.1, 114.4, 113.0, 71.0, 36.4, 36.2 (Figure 23).

HRMS (ESI-MS): found 288.0875, calcd $m/z = 288.0851$ for $\text{C}_{33}\text{H}_{29}\text{BN}_2\text{O}_3\text{S}_2$ $[\text{M}-2\text{I}]^{2+}$ (Figure 24).

Example 2: NMR-analysis of H₂O₂ triggered release of DNA minor groove binder

In a preliminary study, time-dependent NMR spectroscopy analysis of QCy-BA is carried out in the presence of H₂O₂ to assess the stimuli-responsive slicing of phenyl boronic acid functionality (Figure 1(a)). The ¹H NMR spectrum of QCy-BA (2mM) alone in D₂O (0.5 mL) showed a single peak at 5.05 ppm corresponding to the O-CH₂(C-H_a)-bearing phenyl boronic acid group and peaks at 8.2-7.2 ppm corresponding to aromatic protons of the parent QCy-DT.

The chemical shifts of O-CH₂, aromatic region of QCy-BA and appearance of possible new peaks for *p*-quinonemethide (sliced byproduct corresponding to phenyl boronic acid functionality) upon sequential addition of H₂O₂ is monitored. After about 1 hour of H₂O₂ (10mM, 5 μL from the stock H₂O₂ of 1M) addition, the peak intensity at 5.05 ppm, i.e., C-H_a(O-CH₂) gradually decreased and new peaks appeared at 5.20 ppm and 6.5-7.0 ppm regions, suggesting the coexistence of both phenyl boronic acid protected and deprotected forms of QCy-BA. The peaks at 5.20 ppm and aromatic region 6.5-7.0 ppm correspond to the newly-formed exocyclic C-H_b protons of *p*-quinone-methide and QCy-DT moieties, respectively. After about 2 hours, a single peak at 5.20 ppm and prominent new peaks at 6.5-7.0 ppm are observed, indicating the complete conversion of QCy-BA to QCy-DT and *p*-quinonemethide (Figure 1(b)). This study confirmed the H₂O₂ stimulus-triggered slicing of phenyl boronic acid functionality of QCy-BA to release QCy-DT, a DNA minor groove binding probe. It is observed that the color of the solution changes from yellow to brown after the addition of H₂O₂ to QCy-BA, a naked eye detection of the formation of *p*-quinone-methide and QCy-DT (Figure 1(c)).

Example 3: Photo-physical properties of QCy-BA in presence of H₂O₂

Next, the photophysical properties of QCy-BA in the absence and presence of H₂O₂ are studied using UV-vis absorption and emission studies in PBS-buffer solution (10mM, pH = 7.4) under ambient conditions.

UV-vis absorption spectrum of QCy-BA(5 μM) showed broad absorbance in the 300-500 nm region with absorption maximum (λ_{max}) at 400 nm. Upon excitation at 400 nm, emission spectrum of QCy-BA(5 μM) showed weak fluorescence with emission maximum (E_{max}) at 565 nm (Figure 6(b)). As expected, QCy-BA did not emit in the NIR region due to phenyl boronic acid protection of backbone-phenolic hydroxyl moiety.

Interestingly, absorption spectrum of QCy-BA (5 μ M) showed a gradual decrease in absorption maxima at 400 nm in presence of H_2O_2 (1 mM); this is accompanied by the appearance of a new absorption band at 465 nm with a shoulder at 530 nm and an isosbestic point at 442 nm (Figure 6(a)). The new absorption bands at 465 nm and 530 nm revealed the transformation of QCy-BA to the phenolate form of QCy-DT. In agreement with the NMR study (Figure 1(b)), UV-vis absorption data confirmed the generation of QCy-DT through H_2O_2 -assisted oxidation of boronic acid in QCy-BA followed by the hydrolysis and 1,6-elimination of *p*-quinone-methide group. Evidently, UV-vis absorption spectral characteristics clearly support the observed change in solution color from yellow to brown, as a result of the newly formed QCy-DT (λ_{max} at 465 and 530 nm) from QCy-BA (λ_{max} = 400 nm) as seen in Figure 1 (c).

The emission spectra of QCy-BA (5 μ M) in the presence of H_2O_2 (1 mM) displayed gradual decrease in fluorescence intensity at 565 nm and a weak basal level fluorescence band centered around 680 nm with a large Stokes shift ($\Delta\lambda_{max}$ = ~280 nm) upon excitation at 400 nm (Figure 6(b)). Therefore, H_2O_2 -triggered slicing of phenyl boronic acid functionality of QCy-BA is a highly useful transformation for the generation of stimuli-responsive switch-on DNA binding fluorescence probe QCy-DT owing to its large Stokes shift and non-fluorescence in the unbound state.

Example 4: Concentration-dependent fluorescence study of QCy-BA

Further, concentration-dependent fluorescence study on slicing of the phenyl boronic acid functionality of QCy-BA (5 μ M) in response to sequential addition of H_2O_2 (5 to 100 μ M) is performed. The fluorescence intensity of QCy-BA at 565 nm is decreased in response to added H_2O_2 in the concentration range of 5 to 50 μ M and subsequently reached saturation at 100 μ M. A linear relationship ($R^2 = 0.9877$) is observed with increasing concentration of H_2O_2 in the concentration range of 5-50 μ M. Based on $3\sigma/\text{slope}$ the limit of detection (LOD) of H_2O_2 , using the decrease in fluorescence of QCy-BA at 565 nm, is found to be 5.3 μ M (Figure 7).

H_2O_2 is one of the many ROS present in the biological system and it is necessary to test the probe QCy-BA against all of them to assess the selectivity and specificity. Therefore, the response of QCy-BA towards H_2O_2 (100 μ M) in the presence of other ROS (100 μ M) is examined, including tertbutyl hydroperoxide (TBHP), superoxide (O_2^-), hydroxyl radical ($HO\cdot$),

tert-butoxy radical ($^t\text{BuO}^\bullet$) and hypochlorite (OCl^-). Remarkably, H_2O_2 efficiently decreased the fluorescence emission at 565 nm owing to selective slicing of phenyl boronic acid functionality of the QCy-BA (5 μM). On the other hand, very minimal or no effect on the probe response in the presence of superoxide (O_2^-), hydroxyl radical ($\text{HO}^\bullet$), tert-butoxy radical ($^t\text{BuO}^\bullet$) and hypochlorite (OCl^-) (Figure 1(d)) is observed. These results are in agreement with the selective 1,6-elimination of phenyl boronic acid functionality of QCy-BA in the presence of H_2O_2 to liberate *p*-quinone-methide moiety and QCy-DT.

Example 5: Photo-physical properties of combination probe QCy-BA/Drew-AT in presence of H_2O_2

To further validate the H_2O_2 - stimulated conversion of QCy-BA to QCy-DT, a DNA minor groove binder, the transformation is monitored using UV-vis absorption and emission studies in the presence of an AT-rich DNA strand (Drew-AT: 5'-GCGCAAATTTGCGC-3'). QCy-DT binds AT-rich DNA minor groove with high sequence-specificity (5'-AAATTT-3'), which reflects in the strong NIR-fluorescence. Thus, Drew-AT is chosen, a self-complementary 14-base pair (bp) sequence containing central 5'-AAATTT-3' sequence for fluorescence reporting of QCy-DT released in response to H_2O_2 stimulus, by means of strong emission in the NIR region. The absorption spectrum of QCy-BA (2 μM) in the presence of Drew-AT (2 μM) duplex showed an increase in absorption maxima at 416 nm with bathochromic shift ($\Delta\lambda_{\text{max}} = 16 \text{ nm}$) (Figure 8 (a)). On the other hand, the fluorescence spectrum of QCy-BA (2 μM) in the presence of Drew-AT showed emission maxima at 500 nm with hypsochromic shift ($\Delta\lambda_{\text{max}} = \sim 50 \text{ nm}$) (Figure 8(b)). These changes in absorption and emission spectra are attributed to weak interactions between QCy-BA and Drew-AT duplex through electrostatic and hydrophobic interactions.

Next, absorption and emission spectra of QCy-BA are recorded in the presence of Drew-AT duplex and H_2O_2 (100 μM). The absorption spectrum showed a gradual decrease in absorption at 416 nm with corresponding increase in the absorption at 564 nm with an isosbestic point at 456 nm, which is in agreement with the absorption characteristics observed for QCy-DT/Drew-AT complex (Figure 2(a)). Similarly, the emission spectrum of QCy-BA (excitation at $\lambda_{\text{max}} = 400 \text{ nm}$) in the presence of Drew-AT duplex and H_2O_2 showed fluorescence decrease at 500 nm and corresponding increase at 650 nm (Figure 2(b)). This remarkable ratiometric emission at 500 nm

and 650 nm ($\Delta\lambda_{\text{max}} = \sim 250$ nm) is a desirable property of a fluorescence probe to increasing signal-to-noise ratio; measurement at low wavelengths minimizes the error arising from various environmental factors. Further, upon excitation at 564 nm (λ_{max} of QCy-DT bound to Drew-ATduplex), strong fluorescence enhancement at 650 nm is observed (Figure 2(c)). These results reiterated that the H_2O_2 -triggered conversion of QCy-BA to a DNA minor groove binder QCy-DT is a promising ratiometric fluorescence platform for H_2O_2 detection in the presence of exogenous DNA (Drew-AT).

Time-dependent fluorescence study to analyse release kinetics of QCy-BA to QCy-DT in response to H_2O_2

The time-dependent fluorescence study is carried out to evaluate the release kinetics of QCy-BA (2 μM) to QCy-DT in response to H_2O_2 (100 μM) stimulus, in presence of Drew-AT duplex. The change in fluorescence intensities at 500 nm and 650 nm corresponding to emission maxima (E_{max}) of QCy-BA and QCy-DT in presence of Drew-AT is monitored. Upon excitation at 400 nm, fluorescence intensity of QCy-BA gradually decreased at 500 nm while that of QCy-DT increased at 650 nm (Figure 2(d)).

Similarly, the fluorescence spectra recorded upon excitation at 564 nm showed an exponential increase in emission intensity at 650 nm as a function of time and reached saturation ≥ 4 hours (Figure 9). The calculation of kinetics parameter using pseudo-first-order conditions for conversion of QCy-BA (2 μM) to QCy-DT in the presence of H_2O_2 (1 mM) and Drew-AT (2 μM) gave the rate constant of $k_{\text{obs}} = 1.0 \times 10^{-3} \text{ s}^{-1}$ (Figure 10).

Overall, photo-physical (absorption and emission) studies demonstrated that H_2O_2 triggers the slicing of phenyl boronic acid functionality of QCy-BA to generate QCy-DT, a DNA minor binding probe that shows switch-on NIR-fluorescence in the presence of Drew-AT duplex (Fig. 2e).

Example 6: Probing of *in situ* generated H₂O₂ using combination probe QCy-BA $\subset$ Drew-AT

In biological systems, enzymes such as, oxidases generate H₂O₂ by the oxidation of numerous biochemicals. Glucose oxidase (GOx) is one of the most important enzymes known to selectively catalyze the oxidation of glucose to gluconic acid in the presence of oxygen to generate H₂O₂. In this context, *in situ* generation of H₂O₂ by the oxidation of glucose in the presence of GOx using the combination probe QCy-BA $\subset$ Drew-AT (Figure 3 (a)) is investigated. To monitor the *in situ* generation of H₂O₂, glucose is added to PBS buffer (10mM, pH = 7.4) containing GOx (4 U/mL) and QCy-BA $\subset$ Drew-AT (2 μ M). The reaction mixture showed a gradual decrease in fluorescence at 500 nm (λ_{max} = 400 nm) and corresponding increase in fluorescence intensity at 650 nm (Figure 11(a)).

Similarly, upon excitation at 564 nm, the fluorescence spectra showed strong enhancement in fluorescence emission at 650 nm, which may be attributed to the release and binding of QCy-DT to Drew-AT (Figure 3(b)). Next, the reaction kinetics of *in situ* generation of H₂O₂ through the oxidation of glucose by GOx is investigated using the combination probe, upon excitation at 564 nm. The plot of fluorescence intensity at 650 nm as a function of time, after addition of glucose, is shown in Figure 3(c). Upon addition of glucose (1 mM) in the presence of GOx, QCy-BA $\subset$ Drew-AT showed gradual increase in fluorescence intensity at 650 nm and reached saturation at 1 hour. However, in the absence of glucose, GOx and QCy-BA $\subset$ Drew-AT did not show such increase in fluorescence intensity. Further, the fluorescence is monitored by adding increasing concentration of glucose (0 to 1mM) to the mixture of GOx and QCy-BA $\subset$ Drew-AT. The fluorescence emission at 650 nm increases and showed a linear relationship in the concentration range of 0 to 0.2 mM (Figure 3(d), and Figure 11 (b)). Based on $3\sigma/\text{slope}$, the LOD of H₂O₂ is found to be 6.11 μ M (from the concentration of glucose) and is in good agreement with LOD of H₂O₂ (5.33 μ M) using the combination probe (Figure 12).

From the pseudo-first-order calculations, combination probe QCy-BA $\subset$ Drew-AT (2 μ M) showed the rate constant of $k_{\text{obs}} = 6.87 \times 10^{-4} \text{ s}^{-1}$ in the presence of Gox (4 U/mL) and glucose (1 mM) (Figure 13). Overall, Gox assay demonstrated the *in situ* monitoring of H₂O₂ generated from the oxidation of glucose.

Example 7: Study of effect of Catalase on the combination probe QCy-BA $\subset$ Drew-AT

Effect of enzyme that spontaneously decomposes H_2O_2 , resulting in the conversion of QCy-BA to QCy-DT by the action H_2O_2 is studied. Catalase is one of the most efficient enzymes that convert H_2O_2 to water and oxygen to protect cells from oxidative damage and ROS. Catalase exhibits highest turnover number for H_2O_2 and capable of decomposing almost 10^6 molecules per second to water and oxygen. Interestingly, the fluorescence emission is not observed at 650 nm upon addition of H_2O_2 (1 mM) to a solution of QCy-BA $\subset$ Drew-AT (2 μM) containing catalase (4 U/mL). The seized fluorescence emission at 650 nm can be attributed to the prevention of QCy-BA to QCy-DT conversion, as the added H_2O_2 was used as a substrate by the catalase (Figure 14). These results validate that the combination probe QCy-BA $\subset$ Drew-AT is a promising molecular tool for monitoring the *in situ* turnover of H_2O_2 involving oxidase and catalases.

Example 8: Fluorescence imaging and cytotoxicity studies of QCy-BA in presence of H_2O_2

Remarkable selectivity of QCy-BA towards H_2O_2 and its detection through DNA-assisted switch-on NIR fluorescence led to evaluation of the uptake and application of the probe to detect H_2O_2 in cells. For this purpose, confocal fluorescence imaging of HeLa cells treated with H_2O_2 (exogenous) is carried out. First, the HeLa cells are incubated with QCy-BA (5 μM) for about 30 minutes and imaged under a confocal microscope. Fluorescence images of HeLa cells showed weak fluorescence under the blue channel and no emission in the red channel (Figure 4(a) and 4(b)). HeLa cells containing QCy-BA are then treated with H_2O_2 (100 μM) for about 15 minutes, after which the cells are again scanned under a confocal microscope. Confocal images of these cells showed strong fluorescence in the red channel with maximum localization in the cell nucleus (Figure 4(c) and 4(d)). Interestingly, cells also showed the pattern of black nucleoli, a characteristic feature of specific DNA minor groove binders over single-strand DNA and RNAs.

In order to check the cytotoxicity of probe QCy-BA cell viability assay is performed in HeLa cells. Upon incubation with QCy-BA, more than 80% of the cells are viable even at 25 μM concentration after about 24 hours (Figure 15).

In general, above results confirm the permeability and non-toxicity (at standard working concentration and time of about 5 μM and 24 hours, respectively) of QCy-BA, and detection of

exogenously added H_2O_2 in HeLa cells through selective fluorescence staining of the cell nucleus. Therefore, the probe of the present disclosure is permeable, non-toxin and is capable of detecting H_2O_2 in cells.

Example 9: Monitoring of *in situ* generated H_2O_2 levels by EGF/Nox pathways and post-genotoxic stress in live cells

NMR, photophysical study, GOx-assay and confocal fluorescence imaging of HeLa cells showed the detection of exogenously added H_2O_2 using QCy-BA. Next, QCy-BA for probing cellular (physiologically generated) H_2O_2 levels in live cells is employed. HeLa cells are incubated with QCy-BA (about 5 μM) for about 30 minutes in the absence and presence of N-acetyl-L-cysteine (NAC), a well-known H_2O_2 scavenger. In the absence of NAC, flow cytometry analysis of cells treated with the probe (about 5 μM) showed an increase in mean fluorescence intensity of PerCP as compared to control cells (Figure 16). Upon addition of NAC (8 mM), fluorescence intensity of PerCP decreased significantly (Figure 4(e)). In a control experiment, flow cytometry analysis of live HeLa cells treated with QCy-BA and H_2O_2 (100 μM) for about 30 minutes at 37°C showed an increase in the mean fluorescence intensity of PerCP (Figure 4(e)). Thus, probe QCy-BA is also capable of detecting the cellular H_2O_2 levels in live cells.

Further, the study is extended to visualize the *in situ* H_2O_2 generation by a known signaling pathway in live cells. Well-known epidermal growth factor (EGF) binding to epidermal growth factor receptor (EGFR) signaling pathway is selected, which stimulates the production of H_2O_2 in cells by activating the NOX/PI3K pathways. In this experiment, live HeLa cells are incubated with the epidermal growth factor (EGF) (500 ng/mL) for about 40 minutes under physiological conditions (37°C, pH = 7.4). EGF-treated live HeLa cells are incubated with QCy-BA (5 μM) for 30 minutes and flow cytometry analysis of these cells showed strong fluorescence intensity in the PerCP region (Figure 4(f)). On the other hand, the control experiment performed on live HeLa cells without EGF stimulation showed modest fluorescence due to the presence of cellular H_2O_2 level. In contrast, NAC-treated cells showed a decrease in the fluorescence even in the presence of EGF (Figure 4(f)). These results provided concrete evidence that QCy-BA is a versatile and practically viable molecular probe for monitoring concentration levels of H_2O_2 in live cells.

In order to detect the *in situ* generated H_2O_2 in other physiological conditions, fluorescent plate reader-based studies for cellular senescence in primary and cancer cells are performed using probe QCy-BA. First, the confocal fluorescence imaging of primary cells using probe QCy-BA in the presence of H_2O_2 is performed. Live cell imaging of MRC5 cells showed NIR fluorescence in the nucleus compared with control cells incubated with probe QCy-BA (5 μ M) for about 30 minutes after treating with H_2O_2 (100 μ M) (Figure 5 (a-d)). It is well-established that genotoxic stress causes accumulation of DNA damage in cells that can trigger the generation of H_2O_2 inside the cells. It has also been shown that DNA damage induced cell cycle arrest, termed as cellular senescence where ROS played an integral role.

To measure ROS generated concomitant to the dose of the DNA damage, HeLa cells are treated with increasing doses (0 to 200 μ M) of 5-bromo-2'-deoxyuridine (BrdU). BrdU is a thymidine analog, which gets directly incorporated into DNA and triggers DNA damage response. From previous studies in the art it is known that 48 hours of treatment with BrdU (100 μ M) or another DNA damaging agent, doxorubicin at a concentration of 0.1 μ M can lead to the induction of cellular senescence. After about 48 hours of treatment with BrdU (100 μ M), HeLa cells showed a 3-fold increase in fluorescence of 2',7'-dichlorofluorescein diacetate (DCFDA) compared to control cells; DCFDA is a known ROS probe for live cells (Figure 17). Interestingly, probe QCy-BA showed almost 10-fold increase in fluorescence compared to control cells unlike DCFDA, which showed only 3–4 fold change, suggesting that QCy-BA dye has a much better dynamic range than DCFDA (Figure 5(e)).

Further, similar experiments are performed in primary MRC5 cells, which are human lung primary fibroblasts. To induce DNA damage, MRC5 cells are similarly treated with various doses of BrdU and doxorubicin (0.1 μ M) for 72 h. After 72 h, probe QCy-BA showed increase in fluorescence compared to control cells in a dose-dependent manner, indicating that the probe can be used to monitor the *in situ* generated H_2O_2 on primary cells as well (Figure 5(f)). Therefore, above results reveal that QCy-BA is a versatile probe to monitor the elevated levels of H_2O_2 in both primary and cancer cells in senescence state.

In addition to QCy-BA, other Formula I compounds also show similar physical & chemical characteristics, and biological activity results when studies related to photophysical properties,

switch-on NIR-fluorescence in the presence of DNA, base pair-specific recognition and switch-on fluorescence in the presence of DNA, AT-rich DNA Recognition, Sequence-Specific Recognition of DNA, Fluorescence Imaging, Cytotoxicity Studies, Exogenous and endogenous detection of H_2O_2 in HeLa cells, Epidermal growth factor (EGF) produced H_2O_2 detection in HeLa cells, Immunofluorescence studies with Formula I compounds for detection of H_2O_2 in HeLa cells, Detection limit of H_2O_2 in presence of Formula I compounds, detection of ROS using fluorescence plate reader, live cell imaging of MRC5 cells, Fluorescence imaging and cytotoxicity studies of Formula I compounds in presence of H_2O_2 , monitoring of *in situ* generated H_2O_2 levels by EGF/Nox pathways and post-genotoxic stress in live cells are performed.

In conclusion, a stimuli-responsive, colorimetric and switch-on NIR fluorescence combination probe (compounds of formula I, e.g. QCy-BA in combination with exogenous AT-rich DNA or endogenous nuclear DNA) for H_2O_2 is provided by the present disclosure. It is also evident that in QCy-BA, the phenyl boronic acid functionality effectively suppressed the NIR fluorescence of QCy-DT, a DNA minor groove binder and restored selectively in the presence of H_2O_2 .

NMR and UV-vis absorption study showed selective conversion of QCy-BA to QCy-DT and quinine methide in response to H_2O_2 while the solution color changed from yellow to brown for naked eye detection of H_2O_2 over other ROS. The fluorescence study demonstrated selective conversion of QCy-BA to QCy-DT in response to H_2O_2 stimulus that showed NIR-fluorescence in the presence of AT-rich DNA duplex (Drew-AT). Further, glucose oxidase assay confirmed the use of combination probe QCy-BA/Drew-AT for probing *in situ* generated H_2O_2 by the oxidation of glucose to gluconic acid. Cell viability and confocal fluorescence imaging of HeLa cells showed the cell permeability, non-toxicity and preferential nuclear staining selectively of the probe in the presence of H_2O_2 . Furthermore, QCy-BA is sensitive probe to detect normal and *in situ* generated levels of H_2O_2 by EGF/Nox pathways in live cells. Probe QCy-BA is also found to be effective in detection of H_2O_2 in the primary cells as well as senescent cancer cells.

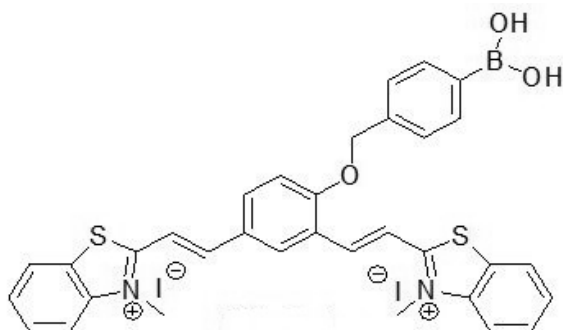
The approach of conjugating DNA fluorescence probes with stimuli-responsive appendages opens up a new approach in the development of DNA targeting theranostic prodrugs for targeting

disease-related cells. This approach is further expanded to create new stimuli-responsive probes for various biochemical processes including enzymatic activities.

Some of the non-limiting applications of the present compound of Formula I are as follows:

- Diagnostic and therapeutic tool for Reactive Oxidative Species, particularly Hydrogen peroxide.
- Used as a Disease marker and for live cell imaging.
- The probe can be used as (NIR) fluorescence binding markers for biomolecules like DNA and protein.
- The probe is useful for sequence specific recognition of dsDNA in treating gene-related human diseases especially cancer, parasitic and viral infections.
- The intrinsic fluorescence property of the probe makes it a versatile fluorescence marker for molecular biology and immunohistochemistry, fluorescence spectroscopy and microscopy, flow cytometry and DNA quantification applications.
- By choosing suitable donors, formation of an efficient FRET-pair can be arrived at which is useful for monitoring the conformational changes in nucleic acids and proteins.
- Detection and inhibition of parasites.
- This approach of conjugating DNA fluorescence probes with stimuli responsive appendages open up a new approach in the development of DNA targeting theranostic prodrugs for spatiotemporal targeting of disease related cells. This approach can be further expanded to create new stimuli responsive probes for various biochemical processes including enzymatic activities.
- Monitor enzyme activity (Glucose oxidase, Cholinesterase and Sarcosine oxidase *etc*), multiple applications, chemical biology and research related ROS in primary and cancer cells.

1. The compound of Formula



- 5 QCy-BA, or its tautomer, polymorph, or solvate thereof.
2. A process for preparing 4-((N,N-dimethyl,2,4-bis((E)-2-(benzo[d]thiazol-2-
ylinium)vinyl)phenoxy)methyl)phenylboronic acid [QCy-BA] comprising the steps:
- 10 a. reacting 2-methyl benzothiazole with methyl iodide to obtain N-methyl-2-
methylbenzothiazole; and
- b. reacting the N-methyl-2-methylbenzothiazole with 4-(4-(4,4,5,5-tetramethyl-1,3,2-
dioxaborolan-2-yl)benzyloxy)isophthalaldehyde in presence of piperidine and ethanol.
3. The process as claimed in claim 2, wherein said process is carried out at a temperature
15 ranging from about 30 °C to 80°C, and for a time period ranging from about 2 hours to 24
hours.
4. The process as claimed in claim 2 or claim 3, wherein the steps a) and b) further comprise
isolation, purification or a combination thereof of the corresponding product; wherein said
20 isolation and purification is carried out by acts selected from a group comprising addition of
solvent, washing with solvent, quenching, filtration, extraction, chromatography and
combinations thereof.

5. A pharmaceutical composition comprising the compound of claim 1 or its tautomeric form, polymorph and solvate thereof, optionally along with at least one pharmaceutically acceptable excipient.

5 6. The pharmaceutical composition as claimed in claim 5, wherein the pharmaceutically acceptable excipient is selected from a group comprising adjuvant, diluent, carrier, granulating agents, binding agents, lubricating agents, disintegrating agent, sweetening agents, glidant, anti-adherent, anti-static agent, surfactant, anti-oxidant, gum, coating agent, coloring agent, flavouring agent, coating agent, plasticizer, preservative, suspending agent, 10 emulsifying agent, plant cellulosic material, spheronization agent, or any combination thereof.

7. A method of detecting or quantifying the presence of reactive oxygen species (ROS) in a biological sample, said method comprising the act of contacting the compound of claim 1 or 15 its tautomer, polymorph or solvate thereof, or the composition of claim 5 or 6; with the sample and detecting the fluorescence indicative of the presence of ROS in the biological sample.

8. The method as claimed in claim 7, wherein said ROS compound is selected from a group 20 comprising hydrogen peroxide, tertbutyl hydroperoxide, superoxide, hydroxyl radical, tert-butoxy radical, hypochlorite and ONOO^- .

9. The method as claimed in claim 7, wherein the biological sample is cells, tissue, biological fluids, or combinations thereof.

25 10. The method as claimed in claim 7, wherein the detection or quantifying is by technique selected from fluorescence microscopy, fluorescence spectroscopy, confocal laser scanning microscopy, total internal reflection fluorescence microscopy, Near infra-red fluorescence and combinations thereof.

11. The method as claimed in any one of claims 7 to 10, wherein the compound or the composition is provided as a combination probe with DNA sequences for detecting and quantifying the ROS, and wherein the DNA sequence is exogenous DNA or endogenous nuclear DNA, or a combination thereof.
- 5 12. The method as claimed in any one of claims 7 to 11, wherein the compound in the presence of ROS is cleaved to release a fluorescent probe quinone cyanine–dithiazole (QCy-DT), which is capable of binding to the AT-rich DNA for fluourometric detection and quantification of the ROS.
- 10 13. A method of diagnosing a disease condition characterized by abnormal reactive oxygen species (ROS) in a subject, said method comprising contacting the compound of claim 1 or its tautomer, polymorph or solvate thereof, or the composition of claim 5 or 6 with a sample obtained from the subject.
- 15 14. The method as claimed in claim 13, wherein the subject is a mammal or a plant, and wherein the sample from mammal is selected from group comprising blood, serum, in-vitro sample, synthetic sample, any bodily fluid and combinations thereof.
- 20 15. The method as claimed in claim 13, wherein the disease is selected from a group comprising cancer, cardiovascular dysfunction, neurodegenerative disease, gastroduodenal pathogenesis, inflammatory disorder, metabolic dysfunction of organ, premature aging, any disorder related to ROS, and combinations thereof.
- 25 16. The method as claimed in claim 13, wherein the disease is diagnosed by detecting and optionally quantifying reactive oxygen species (ROS) in the sample.
- 30 17. Compound as claimed in claim 1 or its tautomer, polymorph or solvate thereof, or the composition as claimed in claim 5 or 6 for use in treating a disease characterized by abnormal levels of ROS in a subject.

18. Use of the compound as claimed in claim 1 or its tautomer, polymorph or solvate thereof, or the composition as claimed in claim 5 in the preparation of a medicament for treating a disease characterized by abnormal levels of ROS in a subject.
- 5 19. A kit for detecting reactive oxygen species (ROS) in a sample, wherein the kit comprising the compound of claim 1 or its tautomer, polymorph or solvate thereof; or a composition of claim 5, wherein the said compound is present in an amount effective to detect the presence of ROS.