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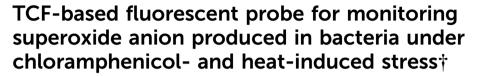


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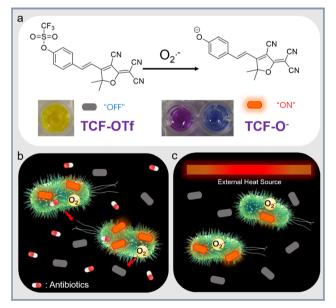


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We report on a superoxide anion (O2°-) responsive fluorescent probe called TCF-OTf. TCF-OTf is able to monitor  $O_2^{\bullet-}$  production when the bacterial species Pseudomonas aeruginosa, Staphylococcus aureus, Escherichia coli, and Enterococcus faecalis are exposed to chloramphenicol and heat shock at 50 and 58 °C.

Reactive oxygen species (ROS) are products of intracellular mitochondrial respiration and electron transport. 1,2 High levels of intracellular ROS can result in intracellular oxidative stress, generate cell aberrations, and thus result in cell dysfunction.<sup>3,4</sup> Prokaryotes have simple and fragile cytoplasmic structures, and as such ROS are toxic to bacteria, where even a low concentration can lead to the oxidation of biomolecules and the denaturation of proteins.5 As a type of highly-reactive ROS, the formation and fate of  $O_2^{\bullet-}$  in eukaryotic cells is associated with diverse physiological and pathological processes, including innate immunity and metabolic homeostasis.<sup>6</sup> However, its role in prokaryotic microorganisms has been much less investigated. Albesa et al. initially found that  $O_2^{\bullet-}$  is produced by bacteria under antibiotic treatment, especially chloramphenicol, vet this area of research has remained relatively unexplored. Furthermore, the production of unspecified ROS by Escherichia coli under heat stress has also been reported.8 However, the fluorescence-based detection of both ROS-related and heatrelated oxidative stress remain underexplored.9,10

Herein, we report on a fluorescent probe, TCF-OTf, for monitoring  $O_2^{\bullet-}$  generated during antibiotic treatment and from heat-stressed bacteria. TCF-OTf was readily prepared, and cleavage of the trifluoromethanesulfonate group by  $O_2^{\bullet-}$ resulted in the formation of the fluorescent anion TCF-O, which was confirmed via mass spectroscopic (MS) analysis (Fig. S1 and S2, ESI†). Reaction with O2. resulted in a fluorescence "turn-on" and colorimetric response from yellow to purple (aqueous phase) or blue (organic phase) (Scheme 1a). We then evaluated the probe using Gram-positive Staphylococcus aureus (NCTC 10788) and Enterococcus faecalis (NCTC 29212),



Scheme 1 (a) Superoxide anion  $(O_2^{\bullet-})$  responsive probe TCF-OTf and its proposed sensing response producing the anion  $\mathsf{TCF-O}^-$ , with fluorescence "turn-on" and a colorimetric change in aqueous (purple) and organic (DMSO, blue) phases. (b and c) Schematic illustration of **TCF-OTf** for detecting  $O_2^{\bullet-}$  upon oxidative stress stimulated by antibiotics (b) or heat (c).

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and Gram-negative Pseudomonas aeruginosa (PAO1) and Escherichia coli (BW 25113), which are of significant clinical concern and exhibit high infection rates.  $^{11,12}$  Significantly, we detected  ${\rm O_2}^{ullet-}$  stress under both antibiotic and heat treatment (Scheme 1b and c, respectively). Compared with available commercially available sensors, our probe contains a near-infrared (NIR) fluorophore to enable minimal biological damage and ensure biocompatibility, effective penetration through tissues, excellent stability of the measurement signals, and minimal interference from background fluorescence, resulting in improved clinical expectations. Moreover, our probe, unlike current commercial systems, enables the real-time and in situ mapping of O<sub>2</sub>•production within a range of complex biological environments.

From our previous research we determined that the 2dicyanomethylene-3-cyano-4,5,5-trimethyl-2,5-dihydrofuran (TCF-OH) fluorophore was an attractive platform for the study of enzymes and ROS in cancer cell-based applications. 13-16 Here, we incorporated a trifluoromethanesulfonate ester, as a known selectively cleaved  $O_2^{\bullet-}$  responsive group, to study the specific ROS (O<sub>2</sub>•-) production from a range of bacterial species that has not previously been reported. The TCF core was prepared using 3-hydroxy-3-methyl-2-butanone, malononitrile and NaOEt in EtOH. Subsequently, TCF, and 4-hydroxybenzaldehyde in EtOH with piperidine (cat.) afforded TCF-OH in 75% yield. Trifluoromethanesulfonate anhydride was then linked to TCF-OH (see ESI†). TCF-OTf was then evaluated during the addition of 0-50 equiv. of  $O_2^{\bullet-}$  in pH = 7.40 PBS buffer (10% DMSO). The fluorescence of TCF-OTf gradually increased at 606 nm, suggesting the formation of TCF-OH, while no response was observed in a pure organic phase (Fig. S3, ESI†). The UV-vis absorption changes for TCF-OTf upon the addition of  $O_2^{\bullet-}$  resulted in a colour change (Fig. S4, ESI†), which was different in the two phases. Based on the fluorescence observation, purple is believed to be the colour of the TCF-O- aggregates, while the dispersed form was blue (Fig. S3 and S4, ESI†). We then determined the sensitivity and response time, resulting in an acceptable sensing performance and full response within 30 min (Fig. S5, ESI†).

We also evaluated the effect of pH on the probes. We observed that TCF-OTf is only fluorescent at pH values above 7 (neutral to alkaline conditions) (Fig. S6, ESI†), which may indicate that the fluorescence originates from TCF-O rather than TCF-OH. Fortunately, the conditions for bacterial growth are neutral or weakly alkaline, and under those conditions the probe functions well. As mentioned above, chemo-selectivity is important for the study of ROS in biological systems. Therefore, we challenged TCF-OTf with other ROS and other common species found in bacteria, such as glutathione, <sup>17</sup> and Fe(III). <sup>18</sup> **TCF-OTf** was found to be selective for  $O_2^{\bullet -}$  (Fig. 1a and b). However, TCF-OTf also responds somewhat to hypochlorite (ClO<sup>-</sup>), but the ClO<sup>-</sup> concentration used was 10-fold greater than  $O_2^{\bullet -}$ , and  $ClO^-$  is unlikely to be found in bacteria under normal conditions.  $^{19}$  Using a common  $O_2^{\bullet-}$  decomposition catalyst FeTMPyP, 20 we confirmed that the O2. signal can be specifically quenched, which makes it an important tool for us to differentiate O<sub>2</sub>• signals from ClO (Fig. 1c and d).

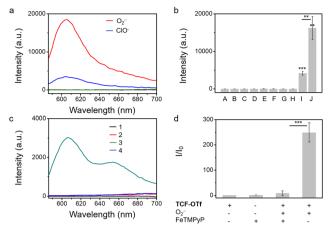


Fig. 1 (a and b) Selectivity of TCF-OTf (10  $\mu$ M) towards a range of ROS and other substances existing in bacteria, after reacting for 2 h. A, probe only; B, ONOO- (100 equiv.); C, HO• (100 equiv.); D, <sup>1</sup>O<sub>2</sub> (100 equiv.); E, H<sub>2</sub>O<sub>2</sub> (100 equiv.); F, GSH (100 equiv.); G, Fe<sup>3+</sup> (100 equiv.); H, ROO<sup>4</sup> (100 equiv.); I,  $ClO^-$  (100 equiv.); and J,  $O_2^{\bullet-}$  (10 equiv.). (c and d) Fluorescence emission spectrum of TCF-OTf (10  $\mu$ M) (c) and the corresponding change in fluorescence (d) in the absence or presence of O2 • and FeTMPyP (100  $\mu$ M) in PBS buffer solution (10% DMSO, pH = 7.40). 1, TCF-OTf; 2, FeTMPyP; 3, TCF-OTf +  $O_2^{\bullet-}$  (pos.); and 4, TCF-OTf +  $O_2^{\bullet-}$  and FeTMPyP (neg.). \*\*P < 0.01, \*\*\*P < 0.001.  $\lambda_{ex} = 560$  nm,  $\lambda_{em} = 606$  nm.

Given that treatment of S. aureus with chloramphenicol was known to increase  $O_2^{\bullet-}$  production, we evaluated chloramphenicol as a model antibiotic with S. aureus, P. aeruginosa, E. coli, and E. faecalis. To determine the efficiency of  $O_2^{\bullet}$ production, we used standard minimum inhibitory concentration (MIC) assays, which determine the minimum concentration of a compound required to completely inhibit bacterial growth (Fig. S7 and S8, ESI†), then the sub-MIC wells (with antibiotic concentrations just below the full-inhibitory concentration) were selected and incubated with TCF-OTf (Fig. 2 and Fig. S9, ESI†). It was found that S. aureus produced the most O2. (further confirmed using an FeTMPyP inhibition assay (Fig. S10, ESI†) that removes the possibility of interference from ClO<sup>-</sup>). Clear fluorescence signal changes from the baseline were also detected when using P. aeruginosa. However, for E. coli and E. faecalis, no O<sub>2</sub>• was detected, even though clear therapeutic effects were seen via the MIC.

After assessing O2. in relation to chloramphenical treatment, we screened an additional four antibiotics, including streptomycin, amikacin, tetracycline, and ciprofloxacin. Initially, we mixed TCF-OTf with the antibiotics in solution to evaluate the stability of the probes (Fig. S11, ESI†). MIC assays were performed for the four species (Fig. S12-S15, ESI†), where the sub-MIC wells with each antibiotic towards each bacterium with TCF-OTf were tested (Fig. S16, ESI†). Whilst modest levels of O2. were reported for ciprofloxacin-treated E. coli and E. faecalis, no signal was observed for ciprofloxacin or tetracycline towards all four species P. aeruginosa, S. aureus, E. coli, and E. faecalis. The absence of signals with ciprofloxacin may be because ciprofloxacin is a strong antibiotic and the MICs of ciprofloxacin (drug concentrations) towards the four species are

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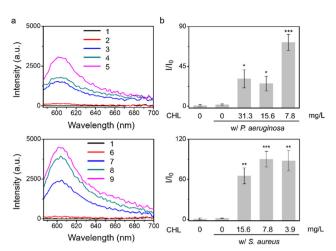


Fig. 2 TCF-OTf based fluorescence detection (a) and corresponding change in fluorescence (b) using P. aeruginosa and S. aureus without antibiotic treatment (control), and P. aeruginosa, and S. aureus picked from the MIC plate wells under treatment at sub-MIC concentrations with chloramphenicol (CHL). 1, TCF-OTf; 2, P. aeruginosa with (w/) TCF-OTf; 3, P. aeruginosa + 31.3 mg  $L^{-1}$  CHL w/ **TCF-OTf**; 4, P. aeruginosa + 15.6 mg  $L^{-1}$  CHL w/ **TCF-OTf**; 5, *P. aeruginosa* + 7.8 mg  $L^{-1}$  CHL w/ **TCF-**OTf; 6, S. aureus w/ TCF-OTf; 7, S. aureus + 15.6 mg  $L^{-1}$  CHL w/ TCF-OTf; 8, S. aureus + 7.8 mg L<sup>-1</sup> CHL w/ **TCF-OTf**; and 9, S. aureus + 3.9 mg L<sup>-1</sup> CHL w/ **TCF-OTf**. \*P < 0.02, \*\*P < 0.002, \*\*\*P < 0.001.  $\lambda_{ex} = 560$  nm,  $\lambda_{em} = 606 \text{ nm}.$ 

very low (generally below 1 mg L<sup>-1</sup>). However, amikacin and streptomycin with P. aeruginosa and S. aureus exhibit some fluorescence signal, although it was still insufficient to observe the characteristic peak of TCF-OH. By contrast, E. coli and E. faecalis exhibited no fluorescence. Similarly, the screening was also confirmed using an FeTMPyP assay (Fig. S17, ESI†) which ensures that  $O_2^{\bullet-}$  was quenched. From these results, chloramphenicol is the most significant O2. -inducing antibiotic for P. aeruginosa and S. aureus.

After determining that the probe was stable at higher temperatures (Fig. S18, ESI†) we explored the effect of heating on bacterial stress and the production of O2. Marcen et al. reported that E. coli (BW 25113) was a bacterial strain that generates ROS under heat-stress.8 Hence, this strain served as a model system among three other additional strains, including P. aeruginosa, S. aureus, and E. faecalis, where ROS generation under heating was unknown. Gratifyingly, signals were observed amongst all the strains using TCF-OTf (Fig. 3), which was in accordance with bacterial mortality induced by heat from 37 °C to 50 °C and 58 °C (Fig. S19, ESI†), and the production of O2. • was further confirmed using an FeTMPyP assay (Fig. S20, ESI†) -. Furthermore, time-dependent detection confirmed that the signals gradually increase with time (Fig. S21, ESI†) and correlated with bacterial cell death (Fig. S22, ESI†).

Confocal laser-scanning microscopy (CLSM) was subsequently used to examine the bioimaging feasibility of TCF-OTf among living bacterial species with antibiotic/heat TCF-OTf-stained chloramphenicol S. aureus exhibited clear and bright fluorescent images, but

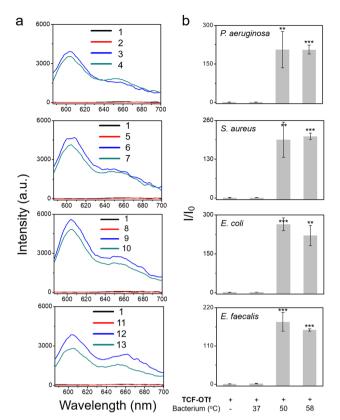


Fig. 3 TCF-OTf based fluorescence detection (a) and corresponding change in fluorescence (b) with P. aeruginosa, S. aureus, E. coli, and E. faecalis grown under normal conditions (37 °C), and P. aeruginosa, S. aureus, E. coli, and E. faecalis after 1 h heating at 50 °C and 58 °C, respectively. 1, TCF-OTf; 2, TCF-OTf w/ P. aeruginosa 37 °C; 3, TCF-OTf w/ P. aeruginosa 50 °C; 4, TCF-OTf w/ P. aeruginosa 58 °C; 5, TCF-OTf w/ S. aureus 37 °C; 6, TCF-OTf w/ S. aureus 50 °C; 7, TCF-OTf w/ S. aureus 58 °C; 8, TCF-OTf w/ E. coli 37 °C; 9, TCF-OTf w/ E. coli 50 °C; 10, TCF-OTf w/ E. coli 58 °C; 11, TCF-OTf w/ E. faecalis 37 °C; 12, TCF-OTf w/ E. faecalis 50 °C; and 13, TCF-OTf w/ E. faecalis 58 °C. \*\*P < 0.01, \*\*\*P < 0.001.  $\lambda_{ex} = 560$  nm,  $\lambda_{em} = 606$  nm.

the three other species also exhibit some "turn-on" fluorescence, which could be quenched by FeTMPyP (Fig. 4, Fig. S23, and S24, ESI†). Significantly, we found that compared with P. aeruginosa and S. aureus, the E. coli and E. faecalis systems, which fail to display any spectroscopic output (Fig. S9, ESI†), exhibited some signal when imaged (Fig. S24, ESI†). TCF-OTfstained and heated bacterial species, on the other hand, were modestly activated under imaging conditions and then quenched by FeTMPyP (Fig. S25–S27, ESI†).

In summary, a new trifluoromethanesulfonate-functionalized TCF-based long-wavelength fluorescent probe TCF-OTf for O<sub>2</sub>• was developed. The TCF-OTf probe was used to monitor O2. production by clinically relevant bacteria under chloramphenicol treatment. In addition, we monitored the production of  $O_2^{\bullet}$ during the heat-treatment of bacteria. Pointedly, long-wavelength TCF-OTf enabled fluorescence cell staining/imaging. We are currently working to develop probes with improved selectivity in order to avoid interference by other highly-reactive ROS such as ClO-. However, in the meantime we anticipate that TCF-OTf could be Communication ChemComm

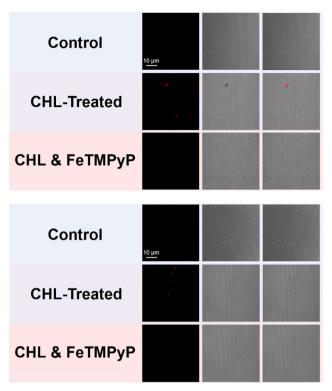


Fig. 4 Confocal laser-scanning microscopy (CLSM) imaging of the intracellular signal of TCF-OTf-stained S. aureus (above) and E. coli (below) (control), and that with treatment of sub-MIC concentrations of chloramphenicol (CHL) (15.6 mg  $L^{-1}$  for S. aureus, 1.95 mg  $L^{-1}$  for E. coli), without or with subsequent treatment with FeTMPyP.  $\lambda_{ex}$  = 561 nm (laser source),  $\lambda_{em}$  = 606 nm.

used to assist the understanding of O2 • -- related pathological/ therapeutic processes and the development of new treatments (antibiotics) for life-threatening bacterial pathogens.

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## Conflicts of interest

There are no conflicts to declare.

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