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Ultrasensitive detection of aromatic water pollutants through protein immobilization driven organic electrochemical transistors†

Subhankar Sahu, ^{‡a} Lokesh Kumar, ^{‡b} Sumita Das, ^a Dipti Gupta ^{*b} and Ruchi Anand ^{*a}

Xenobiotic aromatic water pollutants pose an extreme threat to environmental sustainability. Due to the lack of detectable functional groups in these compounds and scarcity of selective bio-recognition scaffolds, easy-to-use sensing strategies capable of on-site detection remain unavailable. Herein, to address this *lacune*, we entail a strategy that combines biosensor scaffolds with organic electronics to create a compact device for environmental aromatic pollution monitoring. As proof of principle, a sensor module capable of rapid, economic, reliable, and ultrasensitive detection of phenol down to 2 ppb (0.02 μ M) was designed wherein biosensing protein MopR was coupled with an organic electrochemical transistor (OEET). For effective interfacing of the sensing scaffold MopR, graphene oxide (GO) nanosheets were optimized as a host immobilization matrix. The MopR-GO immobilized sensor module was subsequently substituted as the gate electrode with PEDOT:PSS serving as an organic semiconductor material. The resulting OEET sensor provided a favourable microenvironment for protein activity, maintaining high specificity. Exclusive phenol detection with minimal loss of sensitivity (<5% error) could be achieved in both complex pollutant mixtures and real environmental samples. This fabrication strategy that amalgamates biological biosensors with organic electronics harnesses the potential to achieve detection of a host of emerging pollutants.

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Introduction

Access to clean water is indispensable for maintaining a healthy ecosystem as it is the key element for the survival and sustenance of life. According to an estimation by the World Health Organization (WHO) around 844 million people lack basic drinking water services and most importantly roughly around 2 billion people have used polluted drinking water since 2015 around the globe.¹ Specifically, the pollution caused by organic compounds such as benzene, chlorobenzene, phenol, catechol, and toluene produced from anthropogenic activities including catalytic reforming and cracking, petrochemical processing, petroleum refining, *etc.* leads to carcinogenicity and embryotoxicity.^{2,3} Traditional detection strategies such as GC-MS, LC-MS, or HPLC are typically employed for monitoring such monoaromatic xenobiotics. However, despite being sensitive, these techniques are costly to operate and demand complicated

sample-handling procedures.⁴ Additionally, it is hard to selectively monitor these compounds as they are essentially hydrophobic in nature and lack functional groups that can be exploited for detection. Thus, fast, sensitive, direct, and economically beneficial sensing alternatives remain scarce for these compounds.⁵ One of the potential choices to tackle this issue is to employ biosensing as an alternative. This is because due to the accumulation of these pollutants, interestingly a subset of bacteria in the environment has evolved to sense these xenobiotics and degrade them. Hence, these organisms possess natural sensors that can be used as a starting point to develop effective biosensing strategies.^{6,7}

Bioengineered sensing units amalgamated with electrochemical technologies serve as a lucrative prospect to create methods that possess several advantages such as ultrasensitivity, ease of operation, on-site detection, and cost-effectiveness.^{8–11} Against this background, in the last decade, organic electrochemical transistor (OEET) based platforms have become pivotal as next-generation devices due to their exemplary biocompatibility, ability to operate in an aqueous medium, high transconductance, cheap fabrication, and portable nature.^{12–14} In particular OEETs have become extensively popular for probing biological interaction/reactions through the modulation of drain current in aqueous environments making them an impressive platform for biosensing. In

^aDepartment of Chemistry, Indian Institute of Technology Bombay, Powai, Mumbai 400076, India. E-mail: ruchi@chem.iitb.ac.in

^bDepartment of Metallurgical Engineering and Materials Science, Indian Institute of Technology Bombay, Powai, Mumbai 400076, India. E-mail: diptig@iitb.ac.in

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‡ These authors contributed equally.



the past few years, OECTs have been employed for sensing a wide spectrum of analytes ranging from small molecules such as ascorbic acid¹⁵ and dopamine^{16,17} to large virus particles such as influenza¹⁸ and SARS-CoV-2.¹⁹ However, this sensing platform has remained less chartered and elusive in the fields of water pollutant detection and environmental monitoring. Several limiting factors including a lack of suitable selective and sensitive biosensing scaffolds, difficulty in protein immobilization, noise issues from real samples, *etc.*, have eluded its development.

Since there are several bacterial species that possess specific mechanisms for xenobiotic sensing, in recent times, whole cell biosensors (WCBs) have become a popular method for selective detection of mono-aromatics reaching highly sensitive LOD such as 1 ppb ($\sim 0.01 \mu\text{M}$) for phenol detection.²⁰ In several instances, using synthetic biology approaches genetically modified organisms have been generated. However, it poses limitations as it is a less practical approach and accidental release of modified organisms poses ethical issues.^{21,22} Alternatively colorimetric detection using *in vitro* protein-based solution has also been employed but it is marred by issues pertaining to a low limit of detection (LOD). In parallel, electrochemical biosensors for detection of compounds such as phenol have been constructed using generic enzymes such as tyrosinase²³ and laccase.²⁴ Unfortunately, these enzymes lack selectivity and respond to a multitude of compounds with phenolic hydroxyl embedded in them.^{11,25,26} Recently, for phenol sensing, MopR, which is a very specific and selective protein, has found success as an excellent biosensing candidate.⁶ The MopR sensor scaffold is malleable and it has been shown that the phenol binding pocket can be engineered to create a series of selective sensors for a host of mono-aromatic pollutants such as benzene, toluene, and xylenol classes of compounds.²⁷

The gap in MopR mediated biosensing remains twofold; first, the colorimetric detection system for *in vitro* sensing had a low LOD, and second, the setup was not in a translational-friendly compact version. Hence, here we have been successful in creating a compact and sensitive aromatic electrochemical biosensing platform using an OECT. As proof of principle, we show that phenol can be detected as low as 2 ppb ($0.02 \mu\text{M}$) in water, which is in the drinking water regime, thereby, surpassing the sensitivity achieved from the colorimetric MopR sensor by 5-fold. To create this setup a crucial step was to immobilize the protein on a suitable nano-support compatible with the OECT device. Graphene oxide (GO) was optimized as a host nano-interface to tackle protein immobilization. Since GO and its forms possess large surface area, abundant functional groups (such as epoxide, hydroxyl, and carboxylic groups), high dispersion ability in water, low toxicity under physiological conditions, *etc.*,²⁸⁻³⁰ they served as an excellent support matrix. The device employed PEDOT:PSS as both the channel and gate electrode material wherein the MopR-GO nano-bio module was immobilized onto the PEDOT:PSS gate electrode. The resulting electrochemical biosensor was specific for phenol and showed negligible response towards isostuctural compounds such as benzene, toluene, and 2,3-dimethylphenol (2,3-DMP). The setup

efficiently detected the target analyte even in a mixture of contaminants as well as in real environmental samples. Since the MopR scaffold is malleable, this approach can be extended to engineer a series of sensors for the xenobiotic class of aromatics such as benzene, toluene, *etc.*, and has the potential to be extended as a generic method for biosensing-based environmental pollution monitoring.

Results and discussion

Graphene oxide as a biocompatible protein immobilization matrix for aromatic pollutant sensing

Biomolecule immobilization on nano-structures is one of the fundamental aspects of fabrication of ultrasensitive protein-based electrochemical sensors.^{31,32} As a first step, considering the views of rapid on-site detection process and potential for commercial translation, graphene oxide-based strips were designed for the effective immobilization of the protein (MopR) that is suitable for phenol detection without compromising protein activity. For this, GO was synthesized using modified Hummers' method³³ as detailed in the Experimental section and a stable aqueous dispersion of graphene oxide flakes was obtained. The phase purity of the synthesized GO was first characterized by XRD analysis (Fig. 1A). The strong and sharp peak at $2\theta = 9.8^\circ$ corresponds to the characteristic 001 plane of GO suggesting the formation of a long-range ordered structure.³⁴ The facile synthesis of GO yields a very thin and transparent sheet-like structure (Fig. 1B and S1A†) with lateral dimensions of several micrometres with an average height of around 1.4 nm (Fig. 1C) indicating the formation of mostly single-layer GO nanosheets. The functional groups identified for the synthesized GO nanosheets are O-H stretching (3412 cm^{-1}), C=C stretching (2135 cm^{-1}), C=C_(H₂O) stretching (1650 cm^{-1}), C-O-C stretching (1260 cm^{-1}), and C-O stretching (1040 cm^{-1}) vibrations as obtained from FT-IR study (Fig. S1B†).³⁵ Furthermore, Raman analysis of the synthesized GO nanosheets also showed the characteristic D (1355 cm^{-1}), G (1598 cm^{-1}), 2D (2711 cm^{-1}), and D + G (2916 cm^{-1}) bands, respectively (Fig. S1C†). The D band originates from the structural defects present in the lattice whereas the G band corresponds to the sp²-bonded graphitic carbon network.^{36,37} The contact angle values of $\sim 12^\circ$ (L) and $\sim 13^\circ$ (R) confirm the hydrophilic nature of the synthesized GO (Fig. S1C-inset†). In addition, XPS survey analysis of GO nanosheets reveals the relative amounts of carbon and oxygen (Fig. S2A†). Subsequently, the collected core spectra of C 1s contain three distinct types of carbon peaks at around 284.8 eV (C-C/C=C), 287 eV (C-O-C), and 287.4 eV (C=O), respectively (Fig. S2B†)³⁸ and Fig. S2C† represents the O 1s core spectra of GO sheets.

The design of MopR-coated GO strips and the steps adopted for the sensing strategy are entailed in Fig. 1D and S3.† Briefly, first, a uniform, air-dried, drop-casted layer of GO (Fig. S3A†) was created on a plastic surface. Subsequently, a pure MopR protein (Fig. S4A†) was immobilized on the GO nano-support by physical adsorption. Specifically, in our biosensor design, physical adsorption of the protein was preferred as MopR is reported to undergo dramatic



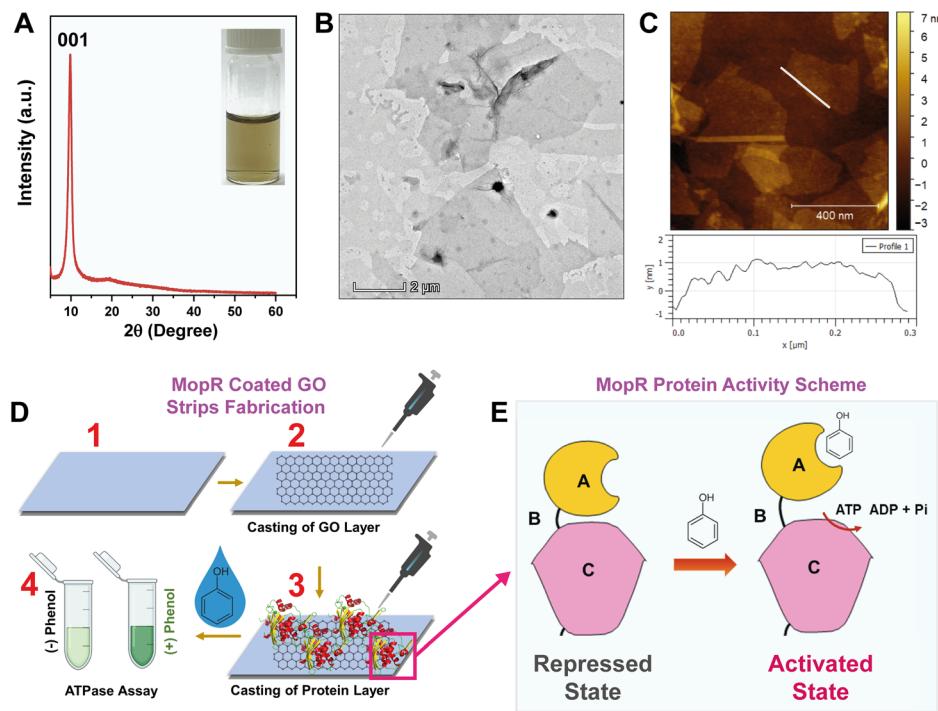


Fig. 1 Fabrication of MopR-GO strips for biosensing of phenol. (A) XRD pattern of GO. Inset: a stable aqueous dispersion of synthesized GO. (B) TEM images of GO nanosheets depicting the formation of thin layered sheets (scale bar 2 μ m). (C) AFM image and height profile of the GO nanosheets. (D) Schematic diagram depicting the steps for fabrication of protein coated GO strips. (E) Schematic representation of structural organization of the MopR protein showing the repressed state where ATP hydrolysis is shut, in the absence of phenol and its subsequent activation upon phenol binding.

reorganization and conformational changes during biosensing³⁹ which might be impeded by alternative approaches such as covalent tethering. The strips were then further treated with synthetically prepared water samples spiked with phenol and a colorimetric malachite green dye-based ATPase assay⁴⁰ was used to confirm the biosensing activity (Fig. S3B and S3C†). The amount of GO and the protein were both varied and an optimal concentration of both, where sensor performance was maximal, was finally chosen for further experimentation. By virtue of the biochemistry of the protein, phenol recognition by MopR is correlated with its ATP hydrolysis reaction. The MopR protein used here comprises two structural domains namely the A-phenol sensing domain and C-ATP hydrolysis domain connected by a B linker (Fig. 1E). The ATP hydrolysis reaction at the C domain is regulated and tightly controlled in a concentration dependent manner by phenol binding at the A domain.⁶ The hydrolysis of ATP results in production of ADP and inorganic phosphate. It is the phosphate generated by hydrolysis that complexes with malachite green producing the requisite signal (detailed in the Experimental section). It is important to note that the MopR biosensing scaffold employed for all phenol sensing assays is a truncated version of the full-length protein (which contains one additional DNA binding domain⁶ after the C domain), comprising only the A to C domain (first 500 amino acid residues). This engineered protein has enhanced thermostability and is hence an ideal candidate to pursue fabrication studies.⁴¹

Fig. 2A clearly depicts that after sequential dilution of GO, GO-D3 strips with a GO concentration of around 0.2 mg mL⁻¹ displayed significantly high sensor activity towards phenol detection compared to other dilutions of GO (GO-D1: 1.6 mg mL⁻¹, GO-D2: 0.4 mg mL⁻¹, and GO-D4: 0.1 mg mL⁻¹). To enable robust phenol sensing under optimal assay conditions, the concentration of the functional nano-support is, therefore, critical. The circular dichroism (CD) spectra confirm that the protein is folded in the presence of GO (Fig. S4B†) inferring high biocompatibility of the synthesized GO nanosheets. Since the main biorecognition module is the MopR protein, the concentration of the active MopR species immobilized onto the GO surface is another crucial factor for phenol detection. Experiments show that a concentration of 2 μ M protein exhibits the best phenol detection efficiency (Fig. S5†) and any protein concentration $> 2 \mu$ M results in a poor dynamic range because of the high absorbance value beyond the linear range (higher than 2). The colorimetric limit of detection of the 2 μ M MopR coated GO-D3 strips is 0.1 μ M phenol (LOD \sim 10 ppb) in water with an almost linear dynamic range of 0–1 μ M (Fig. 2B). After optimization of the sensor design the specificity of the MopR-GO sensor strip was ascertained by testing a wide range of other isotypic phenolics such as 3-chlorophenol (3-CP), catechol, 2,3-di-methylphenol (2,3-DMP), 2,4-dichlorophenol (2,3-DCP) and *m*-cresol. The MopR-GO biosensing strip exhibited the highest activity response towards phenol in comparison to other phenolics (activity was less than 20%) (Fig. 2C). Thus, it can be concluded that the sensor possesses high sensitivity and

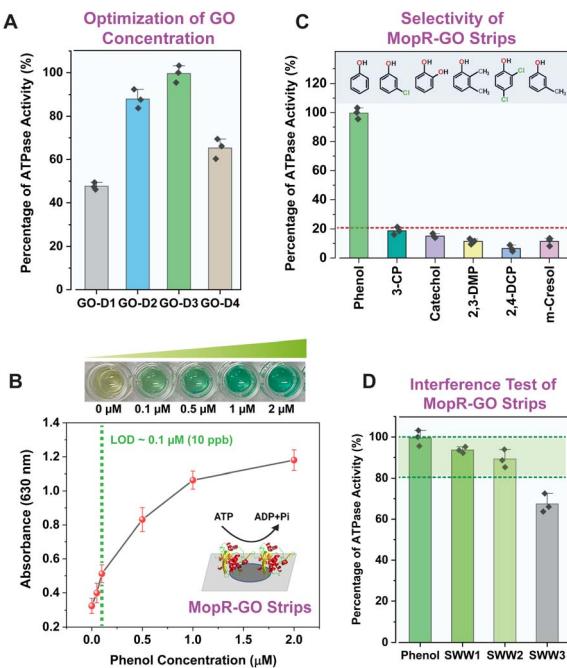


Fig. 2 (A) Optimization of GO concentration for biosensor design. Here, GO-D1: 1.6 mg mL^{-1} , GO-D2: 0.4 mg mL^{-1} , GO-D3: 0.2 mg mL^{-1} , and GO-D4: 0.1 mg mL^{-1} , respectively. (B) Sensitivity profile of MopR coated GO-D3 strips for phenol detection. The increase in colour intensity with higher phenol is visible through ATPase assay. (C) Selectivity profile of MopR coated GO-D3 strips. The response from the other non-targeted phenolics is less than 20%. (D) Test of SWW on MopR coated GO-D3 strips. The immobilized biosensing scaffold can detect phenol in a mixture of other pollutants. SWW1, SWW2, and SWW3 signify increasing concentrations of non-targeted pollutants. The regime with > 80% activity is highlighted.

selectivity in the developed strip format. The final optimized MopR-coated GO strips were similarly competent in detection of phenol (with 94% efficiency) when compared with the *in vitro* solution-based pure protein activity (Fig. S6 and S7†). Such an observation clearly emphasizes the success of GO as a host immobilization matrix for designing biosensing strips, as it can effectively retain the desired conformational plasticity of the protein. Furthermore, in order to screen the effect of surface functional groups on the sensor function, MopR immobilization on reduced graphene oxide (rGO) was also carried out. The results reveal that GO strips are a better nano-platform for sustaining the MopR protein activity (more details in the ESI†). Additionally, the MopR immobilized GO strip could also detect phenol with high confidence in simulated wastewater (SWW) samples containing other non-target pollutants (Fig. 2D). The response for SWW1 and SWW2 (comprising 1 μM and 5 μM of other aromatic pollutants respectively) was >80% thereby, showing negligible interference. SWW3 which contains a higher concentration, 10 μM , of other pollutants in the mixture also maintained a strong response of ~70% demonstrating the robustness of the sensor design. Mop-GO strip parameters such as shelf-life, thermostability, and operating pH range (Fig. S8†) were also measured to demonstrate its environmental durability for field application.

Fabrication of OECTs for phenol monitoring

Biomolecule immobilization on different electrodes of OECTs is useful for sensing purposes as the device architecture is capable of monitoring biosensing reactions through current/voltage change as an output signal. The schematic of the designed OECT with the nano-bio module is presented in Fig. 3A, wherein the MopR-coated GO is immobilized on the gate electrode as a phenol sensing scaffold. The steps involved in device fabrication entail first the deposition process creating a screen printed PEDOT:PSS layer followed by optimization of nano-bio module immobilization on the gate electrode. PEDOT:PSS is used as a semiconducting channel between the source (S) and drain (D) and also as a gate (G) electrode. A similar kind of device architecture was exploited in the past by Gualandi *et al.* where the redox reaction at the PEDOT:PSS layer was monitored to detect ascorbic acid.⁴² However, here we aim to probe phenol *via* GO nanosheets coated with the protein. Briefly, the PEDOT:PSS gate electrode was primed for protein immobilization by first drop-casting and allowing physical adsorption of the GO layer onto the gate electrode before completing the functionalization. The integration of the nano-bio module on the gate electrode was next undertaken under well-controlled conditions of temperature and humidity to achieve the best quality of the electrode.

After successful modulation of the gate electrode, the stability of the device was tested by immersing it in the buffer optimal for ATP hydrolysis of MopR. The device was then tested and it exhibited a drain current (I_D) vs. gate voltage (V_G) graph as typically observed for conventional OECT devices in depletion mode (Fig. 3B) at $V_D = -0.7$ V. The maximum transconductance (g_m) was achieved at $V_G = +0.4$ V for the resulting OECT device (Fig. 3B). Increasing the V_G influences the cations from the electrolyte to accumulate into the bulk of PEDOT:PSS and compensates for the negative sulfonate anions; this is equivalent to de-doping of the semiconducting material reducing the hole density and subsequently the I_D . The output characteristics (I_D vs. V_D) of the designed OECT at varying V_G of -1 V to $+1$ V are depicted in Fig. 3C. A typical decrease in the response of I_D is observed for increasing values of V_G which is a signature of depletion mode behaviour of PEDOT:PSS based OECTs.⁴³ From the drain and transfer characteristics of the device, gate and channel bias were optimized and the ideal specifications for the operation of the OECT were set at $V_G = +0.4$ V and $V_D = -0.7$ V.

The working principle of the MopR-OECT biosensor is represented in Fig. 4A wherein the phenol concentration regulates the extent of ATP hydrolysis which in turn controls the negatively charged inorganic phosphate ions generated near the vicinity of the gate electrode (Fig. 4A). This influences the effective gate voltage (V_G^{eff}) and the proportional increase in the V_G with the addition of phenol results in a reduction of I_D . Immobilized MopR on the gate is inherently negatively charged in the electrolyte medium of pH ~ 7.4 (MopR pI = 5.6). The negative charge arising from phosphate ions due to the ATPase activity alters the surface potential of the gate. In order to maintain a similar effective gate voltage, the V_G increases resulting in a specific decrease in the I_D (Fig. 5A). Moreover, the

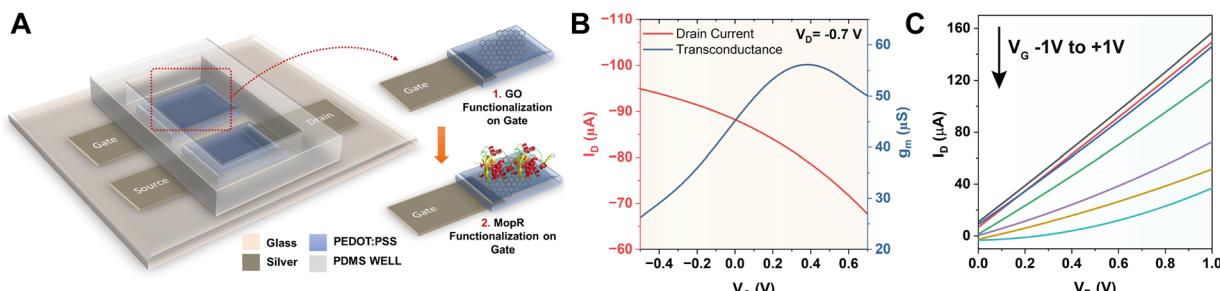


Fig. 3 (A) Schematic diagram of the fabricated OECT. The PEDOT:PSS gate electrode is first coated with GO-D3; subsequently, the MopR protein is immobilized on GO rendering the device ready for phenol sensing. (B) Transfer characteristic of the OECT (I_D vs. V_G) measured in 1× ATPase buffer of MopR exhibiting depletion mode of operation ($V_D = -0.7$ V). (C) Output characteristics of the device (I_D vs. V_D) for different V_G ranging from -1 V to $+1$ V.

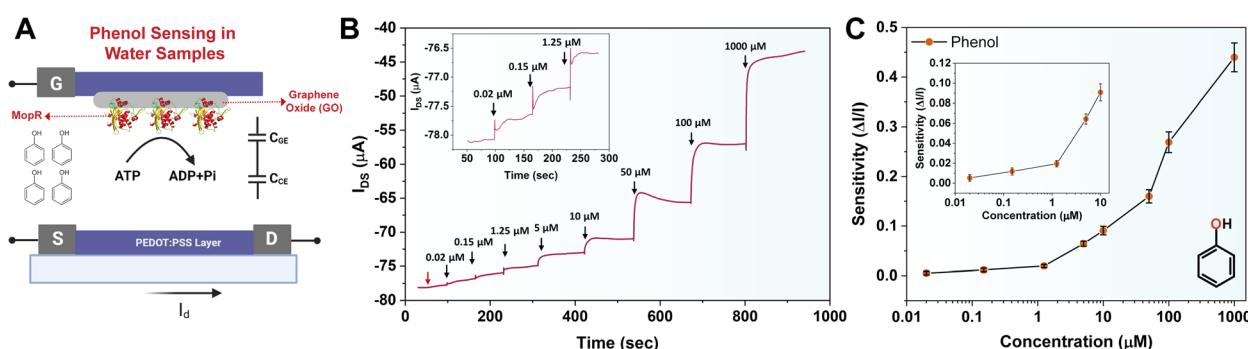


Fig. 4 (A) Schematic of the operation of the phenol sensing OECT. The immobilized MopR at the gate electrode selectively binds phenol triggering an ATP hydrolysis reaction. (B) Time- I_D response of the OECT with an increasing concentration of phenol. The device starts producing a signal from 0.02 μM of phenol; the first arrow represents 0.008 μM phenol showing no significant change in I_D . (C) Normalized sensitivity plot of the channel current at different concentrations of phenol; inset: sensitivity, 0.01 – 10 μM range.

interplay of the capacitance at the gate/electrolyte (C_{GE}) and channel/electrolyte (C_{CE}) interface impacts the charge distribution at the two interfaces and thus influences the total current flow in the active channel.⁴⁴ The amount of phenol in a particular sample is quantitated by employing this OECT device by serial addition of increasing concentrations of phenol and continuous monitoring of the I_D . Real-time evolution of the I_D as obtained showcases a staircase-like signal (Fig. 4B) which in general is perceived as a signature for the successful detection of analytes at the gate electrode.⁴⁴ The decrease in the I_D proportionally increases with increasing phenol concentration and the normalized current response is then converted to sensitivity. A titration of ATP concentrations yielded 5 mM as the optimal working concentration for the biosensor module (Fig. S9A†) and in the absence of ATP the device shows negligible sensitivity (Fig. 5B). Furthermore, it was established that no appreciable response was observed in the absence of the biosensing scaffold (Fig. S9B†). Moreover, Fig. 5B-inset, demonstrates that for 1 μM phenol the sensitivity of the device is around 4 times lower than that in the absence of ATP (sensitivity^{+ATP} – 0.020 and sensitivity^{-ATP} – 0.005).

Several other parameters such as device geometry and protein concentration were also tuned to optimize the biosensor device. Since PEDOT:PSS can take up the ions from the electrolyte into its bulk volume which results in volumetric

doping, the optimization of the device geometry and thickness of the electrodes plays a crucial role in the sensitivity of the OECT geometry. Schematic of the optimized OECT geometry is shown in Fig. S10A† where the channel ($L_{CH} \times W_{CH} = 8$ mm × 2 mm) and the gate ($L_G \times W_G = 8$ mm × 2 mm) dimensions are the same and channel to gated lateral separation is kept at 2 mm. The surface topology of the screen printed PEDOT:PSS layer can be observed from a top view of FEG-SEM (Fig. S10B†) and the cross-section of the PEDOT:PSS layer indicated that it has an average thickness of ~ 0.7 μm (Fig. S9C†). The GO layer with thickness ~ 2.5 μm on PEDOT acts as an effective nano-platform for protein immobilization (Fig. S10C†). Furthermore, the EDS spectrum of the GO-coated gate indicates the desired functionalization of the PEDOT:PSS layer in combination with GO nanosheets (Fig. S10D†).

To achieve maximal sensing activity the protein concentration immobilized at the gate electrode was also optimized as clustering of the biomolecule at the nano-junction is a fundamental aspect of activity (schematic shown in Fig. S11A†). It was observed that a slightly higher protein concentration of 10 μM is optimum for achieving better sensitivity (Fig. S11B†). Here, it was noticed that for effective phenol probing through the OECT device, a higher protein concentration was required in comparison to the strip-based colorimetric biosensor. This could be because, in the case of the OECT, at least 10 μM of

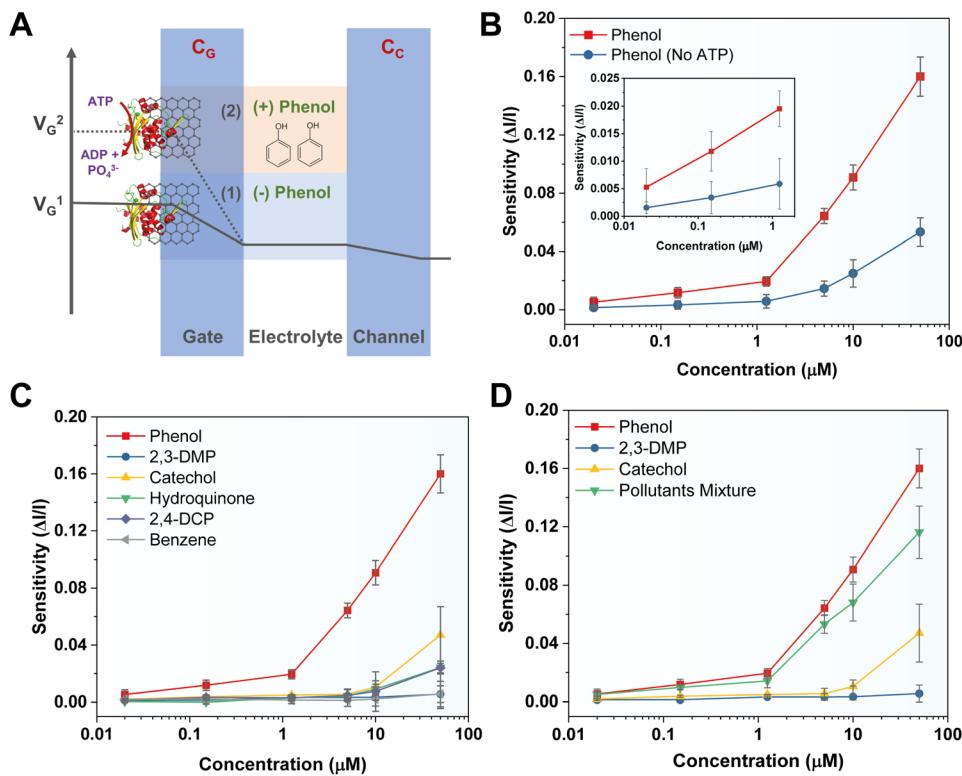


Fig. 5 (A) Schematic illustration showing the potential drop of the device at two electric double layers of the gate and channel. The change in the V_G before (solid) and after (dashed line) the biosensing reaction is represented. (B) Sensitivity plot for phenol sensing in the presence and absence of ATP. In the absence of ATP the biosensor shows a significantly low response. (C) Selectivity profile of the OECT based phenol biosensor; other iso-typical aromatic pollutants have negligible sensitivity. (D) Interference studies of the biosensor. The MopR-OECT sensor can detect phenol in a mixture of other contaminants. 'Pollutants mixture' contains phenol, 2,3-DMP, and catechol at an equivalent concentration.

protein is required to produce sufficient phosphate anions to influence the V_G^{eff} . However, increasing the concentration further does not enhance device performance as the protein crowding in the interface⁴⁵ might impact the conformational flexibility required by MopR affecting target molecule recognition and activity.

The chronoamperometric response shows that the device can detect phenol as low as 0.02 μM (Fig. 4B, inset). The LOD of the device was confirmed to be 0.02 μM of phenol. A lower concentration of 0.008 μM (first arrow-red, Fig. 4B) doesn't manifest any significant I_D response. The sensitivity ($\Delta I/I$) response plot for different concentrations of the phenol (Fig. 4C) depicts that the biosensor exhibits a wide dynamic range. Previously Mayorga-Martinez *et al.*⁴⁶ reported an electrochemical biosensor for phenol detection using tyrosinase enzyme immobilized on a p-nitrophenol-based substrate and the device reports a LOD level of 0.5 μM (500 nM). A higher LOD in this case could be because despite the well-optimized design of the accompanying immobilization and sensing scaffold, the tyrosinase and laccase enzymes are less selective. In contrast, the MopR-based biosensing unit is highly selective for phenol. The MopR scaffold has additional advantages as the ligand binding pocket can be structurally engineered to bind a battery of compounds and resulting sensors exhibit selective sensing for an array of monoaromatic (both phenolic and non-phenolic)

ligands.^{27,47} Thus not only can this sensor design be easily multiplexed in the future where each pollutant can be reported with high precision but also the optimized GO nano-platform achieves a LOD of 0.02 μM which is 25-fold better than that of the tyrosinase system under similar electrochemical-based readout setting making this a robust and versatile setup. The device stability is comparable to the stability exhibited by the MopR-GO strip-based colorimetric sensor (Fig. S12†).

Sensitivity and selectivity of the biosensor

The MopR-GO biosensing strips can detect phenol with high precision. This is apparent from its selectivity profile. A similar observation was also found in the case of the OECT device; iso-structural compounds such as benzene, catechol, hydroquinone, 2,3-DMP, and 2,4-DCP show negligible sensitivity (Fig. 5C). Though at high concentrations (50 μM) of analytes such as catechol, hydroquinone, and 2,4-DCP, the device shows increased sensitivity ~ 0.04 , but it is significantly less than that for phenol (sensitivity ~ 0.16 at 50 μM). The time-channel current response profile also validates these observations (Fig. S9C†). Furthermore, the OECT was tested for its ability to detect phenol in a simulated pollutant mixture. Samples comprising phenol, 2,3-DMP, and catechol (referred to as 'Pollutants Mixture') were tested and the MopR-based OECT device could detect phenol selectively without loss of sensitivity (less than 5% error) in

Table 1 Comparison of available phenol biosensing technologies ['+++ highly selective', '++ moderate selective', and '+ less selective']

	Biosensing protein	Details	Selectivity	LOD	Reference
1	Tyrosinase	Electrochemical biosensor. A graphene-decorated gold nanoparticle/chitosan (Gr-Au-Chit/Tyr) nanocomposite-modified screen-printed carbon electrode (SPCE) is used for protein immobilization and detection of phenolic compounds	+	Phenol, 0.05 μ M	54
2	Tyrosinase	Electrochemical biosensor. Screen-printed carbon electrodes were modified with electrodeposition of Au nanoparticles (AuNPs) and tyrosinase was covalently immobilized	++	Phenol, 0.5 μ M	55
3	Tyrosinase	Electrochemical biosensor. A glassy carbon electrode (GCE) was modified with enzyme-immobilized carbon nanotubes (CNTs)	++	Phenol, 0.035 μ M	56
4	Laccase & tyrosinase	Electrochemical biosensor. Tyrosinase and laccase on graphite screen printed electrodes modified with ferrocene	+	Phenol, 2 μ M	57
5	Tyrosinase	Electrochemical biosensor. Two-dimensional pniotogen nano-sheets are used for protein immobilization on glassy carbon electrodes	+++	Phenol, 0.5 μ M	58
6	Tyrosinase	Electrochemical biosensor. The enzyme was immobilized using a glutaraldehyde linker on gold nanoparticle modified screen-printed electrodes (Tyr-AuNPs-SPCEs)	+	Phenol, 1.2 μ M	59
7	Tyrosinase	Optical biosensor. Tyrosinase was immobilized on a chitosan film coated on microscopy glass slides	+	Phenol, 1 μ M	60
8	Laccase	Optical biosensor. Auto-indicating optical properties of laccase including intrinsic molecular absorption and fluorescence are used for phenol sensing	++	Phenol, 9 μ M	61
9	Tyrosinase	Optical biosensor. A long-period fiber grating coated with an enzyme-entrapped polyacrylamide gel is used	+	Phenol, 81 μ M	62
10	MopR	Electrochemical biosensor. An organic electrochemical transistor (OECT) based device employing protein immobilized graphene oxide nano-sheets is used	+++	Phenol, 0.02 μ M	This work

a lower concentration range of 0.02 to 1.25 μ M (Fig. 5D). At higher concentrations of pollutants (example 10 μ M and 50 μ M) the sensitivity is affected with respect to the water spiked with pure phenol samples with an error of 20%; however, the sensitivity of phenol remains substantially higher than that of the non-targeted pollutants. Next, in order to validate the operation of the electrochemical biosensor for environmental samples, a locally collected water sample (EVS) from Vihar Lake, Mumbai, was spiked with a similar concentration range of phenol and experimented on the device. The sensitivity of the OECT was very similar in comparison to distilled water spiked with lab-grade phenol (0.02–0.15 μ M) with an error of less than 3% (Fig. S13[†]) inferring that the OECT device was responsive for real samples and didn't undergo any degradation. A comparison of the MopR-OECT prototype biosensor with other phenol-detecting biosensing platforms such as those constructed from tyrosinase and

laccases (Table 1) clearly demonstrates the high sensitivity and specificity of the MopR-OECT-based detection module. As mentioned earlier, both laccase and tyrosinase display non-specific activity, often generating signals for various phenolic compounds such as catechol or chlorophenols,⁴⁸ diminishing their selectivity. Conversely, the high selectivity for phenol, exceptional thermal stability, and biocompatibility of MopR make it conducive to integration within the transistor architecture. Thus, it positions the MopR-GO-OECT module as a more promising scaffold for probing mono-aromatics in water.

Conclusion

In this work, an OECT-based device was fabricated for phenol monitoring by utilizing an MopR biosensing scaffold immobilized on the GO-modified gate electrode. Both the daunting



aspects of protein immobilization and selective electrochemical sensing of hard-to-probe mono-aromatic xenobiotics were addressed. The resulting sensor yielded a LOD of less than 2 ppb towards phenol detection. In contrast to earlier reports, where enzymes such as tyrosinase and laccase were employed, MopR-based phenol biosensors were highly selective and could distinguish between isostructural ligands. The high thermal durability and biocompatibility allowed for integration with the OECT's modern architecture which is appealing due to the transistor-based amplification of the signal,⁴⁹ making it more useful for probing biochemical interactions. In comparison to cyclic voltammetry and amperometry,⁵⁰ OECTs are more suitable for bioelectronics and allow effective miniaturization of the sensing platform,⁵¹ which is useful for creation of low-cost and portable devices. Although the LOD of the MopR-OECT platform is comparable to that of synthetic biology inspired sensors,²⁰ it circumvents most of the limitations associated with bio-contamination, membrane transport and other biohazard risks, thereby, making the MopR-OECT design superior. The graphene oxide-based sensor strips developed here are also easy-to-fabricate, cheap, portable, and accurate for phenol detection in complex water samples without the need for any further modification to the sensor surface. Though the application of GO-based composite materials for bioremediation and sensing is widespread,^{52,53} the reported easy-to-fabricate MopR-GO design is versatile and is effective both in a colorimetric strip format as well as in an electrochemical sensing platform. Overall, the MopR-OECT platform established here holds the potential to be translated to other compounds and a panel of sensors that can individually sense a particular mono-aromatic can be designed. Thus, this design provides a future possibility of developing a water monitoring system that can accurately read multiple pollutants simultaneously. To conclude, here, a rational merging of MopR's biochemistry with an OECT device yielded a functional platform for ultrasensitive and selective detection of phenol. The work presented here opens doors for building systems that require amalgamation of bio-scaffolds with materials and device electronics. Our research sits at the crossroads of environmental sustainability and human health proving a promising approach for selective detection of aromatics.

Experimental section

MopR-GO strip fabrication and colorimetric ATPase assay for phenol detection

The MopR protein was purified *via* a previously established protocol⁴⁷ and detailed in the ESI.† Subsequently, different concentrations of the GO substrate were tested for MopR protein immobilization and activity testing. Plastic coverslips (Hampton Research, Cat. No. HR8-074) of 6 × 6 mm were used as a base and 10 µL of series of different concentrations of GO (GO-D1: 1.6 mg mL⁻¹, GO-D2: 0.4 mg mL⁻¹, GO-D3: 0.2 mg mL⁻¹, and GO-D4: 0.1 mg mL⁻¹) were drop-cast and dried for 30 min at 40 °C. The strips were then kept at room temperature for 5 min and 1 µL of 20 µM protein was drop-cast; the strips were incubated at RT for 20 min and finally, water samples containing the phenol samples were tested by colorimetric ATPase

assay. After optimization of the GO concentration in the strip version, all further experiments were carried out using GO-D3 strips. Next, the MopR protein of different concentrations (1 µM, 2 µM, and 4 µM) was tested and the 2 µM protein was chosen due to its maximal performance. The ATPase assay employed herein is typically a malachite green dye based assay for the detection of phosphate in an *in vitro* solution.⁴⁰ In an *in vitro* reaction, MopR protein, ATP, and water containing phenol are supplied for sensing (more details in the ESI†).

Fabrication of OECTs

For making the OECT devices, the source, drain, and gate silver metal contacts (LADD, Cat. No. 60805) were screen printed on a cleaned glass slide. The substrate was printed with commercially available PEDOT:PSS (Sigma-Aldrich, Cat. No. 655201). Conductive screen printable ink was used as the active material (channel and gate) for the fabrication of the OECTs. The silver metal contact points are of dimensions 2 mm × 2 mm. The channel ($L_{CH} \times W_{CH} = 8 \text{ mm} \times 2 \text{ mm}$) and the gate ($L_G \times W_G = 8 \text{ mm} \times 2 \text{ mm}$) with a channel to gate lateral separation of 2 mm (D) were obtained by screen-printing PEDOT:PSS over a polyester film. For both steps, screen-printing was performed with a polyester 51T/130-71 PW mesh screen and 3M™ Screen Printing UV Ink for masking/patterning the mesh. Controlling the thickness of the PEDOT:PSS layer is crucial for the protein immobilization of the device. The average thickness of the PEDOT:PSS layer is optimized at ~0.7 µM. For the thickness measurement, the PEDOT:PSS layer is coated on a silicon substrate (Sigma-Aldrich, Cat. No. 647675) and dried at 60 °C for 30 min. Then, at first the top image is captured showing the uniform formation of the PEDOT:PSS layer. Furthermore, the 0.2 mg per mL GO layer is coated on top of the PEDOT:PSS layer and the cross-section image is collected on a JEOL JSM-7600F FEG-SEM instrument. A Keithley 4200A parameter analyzer is used for measurement purposes. The sample testing and device operation are detailed in the ESI.†

Further experimental details on protein purification, GO, rGO synthesis, and electrical characterization can be found in the ESI.†

Data availability

All the experimental data related to this study are available from the corresponding author upon reasonable request.

Author contributions

Subhankar Sahu: conceptualization, data curation, formal analysis, methodology, investigation and writing – original draft. Lokesh Kumar: conceptualization, data curation, methodology, investigation and writing – original draft. Sumita Das: data curation, methodology, investigation and writing – original draft. Dipti Gupta: funding acquisition, supervision, resources and writing – review & editing. Ruchi Anand: project administration, funding acquisition, resources and writing – review & editing.



Conflicts of interest

Authors declare no conflict of interest.

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References

- W. H. Organization and U. N. C. F. (UNICEF), *Progress on drinking water, sanitation and hygiene: 2017 update and SDG baselines*, World Health Organization, 2017.
- O. Al-Madanat, Y. Alsalka, W. Ramadan and D. W. Bahnemann, *Catalysts*, 2021, **11**, 1–45.
- L. G. C. Villegas, N. Mashhadi, M. Chen, D. Mukherjee, K. E. Taylor and N. Biswas, *Curr. Pollut. Rep.*, 2016, **2**, 157–167.
- E. Matisová and S. Hrouzková, *Int. J. Environ. Res. Public Health*, 2012, **9**(9), 3166–3196.
- S. Sahu and R. Anand, in *The World Scientific Reference of Water Science: Volume 1 Molecular Engineering of Water Sensors*, World Scientific, 2023, pp. 101–136.
- S. Ray, M. J. Gunzburg, M. Wilce, S. Panjikar and R. Anand, *ACS Chem. Biol.*, 2016, **11**, 2357–2365.
- R. Fernandez-López, R. Ruiz, F. de la Cruz and G. Moncalián, *Front. Microbiol.*, 2015, **6**, 1–21.
- F. Arduini, S. Cinti, V. Scognamiglio, D. Moscone and G. Palleschi, *Anal. Chim. Acta*, 2017, **959**, 15–42.
- G. Hernandez-Vargas, J. Sosa-Hernández, S. Saldarriaga-Hernandez, A. Villalba-Rodríguez, R. Parra-Saldivar and H. Iqbal, *Biosensors*, 2018, **8**, 29.
- G. K. Mishra, V. Sharma and R. K. Mishra, *Biosensors*, 2018, **8**, 1–13.
- E. Moczko, G. Istamboulie, C. Calas-Blanchard, R. Rouillon and T. Noguer, *J. Polym. Sci., Part A: Polym. Chem.*, 2012, **50**, 2286–2292.
- R. A. Picca, K. Manoli, E. Macchia, L. Sarcina, C. Di Franco, N. Cioffi, D. Blasi, R. Österbacka, F. Torricelli, G. Scamarcio and L. Torsi, *Adv. Funct. Mater.*, 2020, **30**, 1904513.
- B. Burtscher, P. A. M. Urbina, C. Diacci, S. Borghi, M. Pinti, A. Cossarizza, C. Salvarani, M. Berggren, F. Biscarini, D. T. Simon and C. A. Bortolotti, *Adv. Healthcare Mater.*, 2021, **10**, 2100955.
- J. Rivnay, S. Inal, A. Salleo, R. M. Owens, M. Berggren and G. G. Malliaras, *Nat. Rev. Mater.*, 2018, **3**, 17086.
- I. Gualandi, M. Marzocchi, E. Scavetta, M. Calianni, A. Bonfiglio and B. Fraboni, *J. Mater. Chem. B*, 2015, **3**, 6753–6762.
- I. Gualandi, D. Tonelli, F. Mariani, E. Scavetta, M. Marzocchi and B. Fraboni, *Sci. Rep.*, 2016, **6**, 1–10.
- C. Liao, M. Zhang, L. Niu, Z. Zheng and F. Yan, *J. Mater. Chem. B*, 2014, **2**, 191–200.
- W. Hai, T. Goda, H. Takeuchi, S. Yamaoka, Y. Horiguchi, A. Matsumoto and Y. Miyahara, *Sens. Actuators, B*, 2018, **260**, 635–641.
- K. Guo, S. Wustoni, A. Koklu, E. Díaz-Galicia, M. Moser, A. Hama, A. A. Alqahtani, A. N. Ahmad, F. S. Alhamlan, M. Shuaib, A. Pain, I. McCulloch, S. T. Arold, R. Grünberg and S. Inal, *Nat. Biomed. Eng.*, 2021, **5**, 666–677.
- R. Roy, S. Ray, A. Chowdhury and R. Anand, *ACS Sens.*, 2021, **6**, 1933–1939.
- J. C. Gutiérrez, F. Amaro and A. Martínez-González, *Front. Microbiol.*, 2015, **6**, 1–8.
- S. Sahu, R. Roy and R. Anand, *ACS Sens.*, 2022, **7**, 704–715.
- A. V. Bounegru and C. Apetrei, *Nanomaterials*, 2023, **13**, 760.
- M. C. Castrovilli, P. Bolognesi, J. Chiarinelli, L. Avaldi, P. Calandra, A. Antonacci and V. Scognamiglio, *TrAC, Trends Anal. Chem.*, 2019, **119**, 115615.
- J. Adamski, P. Nowak and J. Kochana, *Electrochim. Acta*, 2010, **55**, 2363–2367.
- A. Tarasov, N. Stozhko, M. Bukharinova and E. Khamzina, *Life*, 2023, **13**, 291.
- S. Ray, S. Panjikar and R. Anand, *ACS Sens.*, 2018, **3**, 1632–1638.
- K. Chaudhary, K. Kumar, P. Venkatesu and D. T. Masram, *Adv. Colloid Interface Sci.*, 2021, **289**, 102367.
- G. Hernández-Canel, D. Suazo-Dávila, A. J. Ojeda-Cruzado, D. García-Torres, C. R. Cabrera and K. Griebenow, *J. Nanobiotechnol.*, 2015, **13**, 70.
- M. Mathesh, B. Luan, T. O. Akanbi, J. K. Weber, J. Liu, C. J. Barrow, R. Zhou and W. Yang, *ACS Catal.*, 2016, **6**, 4760–4768.
- V. Ganesh and A. Muthurasu, *J. Phys.: Conf. Ser.*, 2012, **358**, 012003.
- A. A. Homaei, R. Sariri, F. Vianello and R. Stevanato, *J. Chem. Biol.*, 2013, **6**, 185–205.
- K. Krishnamoorthy, M. Veerapandian, K. Yun and S. Kim, *Carbon*, 2012, **53**, 38–49.
- D. C. Marcano, D. V. Kosynkin, J. M. Berlin, A. Sinitskii, Z. Sun, A. Slesarev, L. B. Alemany, W. Lu and J. M. Tour, *ACS Nano*, 2010, **4**(8), 4806–4814.
- H. Yuan, J. Ye, C. Ye, S. Yin, J. Li, K. Su, G. Fang, Y. Wu, Y. Zheng, M. Ge, R. Tang, G. Feng, Y. Qu and Y. Zhu, *Chem. Mater.*, 2021, **33**, 1731–1739.
- Å. Björkman, *Schweiz. Z. Hydrol.*, 1969, **31**, 632–645.



37 M. A. Pimenta, G. Dresselhaus, M. S. Dresselhaus, L. G. Cançado, A. Jorio and R. Saito, *Phys. Chem. Chem. Phys.*, 2007, **9**, 1276–1290.

38 W. Xu, Z. Jin, X. Pang, Y. Zeng, X. Jiang, Y. Lu and L. Shen, *ACS Appl. Bio Mater.*, 2020, **3**, 4263–4272.

39 J. Singh, R. Anand and A. Horovitz, *J. Bacteriol.*, 2022, **204**(8), e00179.

40 A. A. Baykov, O. A. Evtushenko and S. M. Avaeva, *Anal. Biochem.*, 1988, **171**, 266–270.

41 S. Ray, T. Senapati, S. Sahu, R. Bandyopadhyaya and R. Anand, *Anal. Chem.*, 2018, **90**, 8960–8968.

42 I. Gualandi, M. Marzocchi, E. Scavetta, M. Calienni, A. Bonfiglio and B. Fraboni, *J. Mater. Chem. B*, 2015, **3**, 6753–6762.

43 N. Saraf, E. R. Woods, M. Peppler and S. Seal, *Biosens. Bioelectron.*, 2018, **117**, 40–46.

44 K. Tang, C. Turner, L. Case, A. Mehrehjedy, X. He, W. Miao and S. Guo, *ACS Appl. Polym. Mater.*, 2022, **4**, 2337–2345.

45 R. García-Álvarez and M. Vallet-Regí, *Nanomaterials*, 2021, **11**, 888.

46 C. C. Mayorga-Martínez, R. Gusmão, Z. Sofer and M. Pumera, *Angew. Chem., Int. Ed.*, 2019, **58**, 134–138.

47 S. Ray, S. Panjikar and R. Anand, *ACS Sens.*, 2017, **2**, 411–418.

48 P. A. Raymundo-Pereira, T. A. Silva, F. R. Caetano, L. Ribovski, E. Zapp, D. Brondani, M. F. Bergamini, L. H. Marcolino, C. E. Banks, O. N. Oliveira, B. C. Janegitz and O. Fatibello-Filho, *Anal. Chim. Acta*, 2020, **1139**, 198–221.

49 D. Khodagholy, J. Rivnay, M. Sessolo, M. Gurfinkel, P. Leleux, L. H. Jimison, E. Stavrinidou, T. Herve, S. Sanaur, R. M. Owens and G. G. Malliaras, *Nat. Commun.*, 2013, **4**, 2133.

50 G. Alarcon-Angeles, G. A. Álvarez-Romero and A. Merkoçi, in *Encyclopedia of Interfacial Chemistry*, Elsevier, 2018, pp. 140–155.

51 H. Liu, A. Yang, J. Song, N. Wang, P. Lam, Y. Li, H. K. Law and F. Yan, *Sci. Adv.*, 2021, **7**, 1–10.

52 D. Tyagi, H. Wang, W. Huang, L. Hu, Y. Tang, Z. Guo, Z. Ouyang and H. Zhang, *Nanoscale*, 2020, **12**, 3535–3559.

53 K. S. Obayomi, S. Y. Lau, M. Danquah, T. Chiong and M. Takeo, *Environ. Nanotechnol., Monit. Manage.*, 2022, **17**, 100647.

54 F. Fartas, J. Abdullah, N. Yusof, Y. Sulaiman and M. Saiman, *Sensors*, 2017, **17**, 1132.

55 M. N. Karim and H. J. Lee, *Talanta*, 2013, **116**, 991–996.

56 Y. Wee, S. Park, Y. H. Kwon, Y. Ju, K.-M. Yeon and J. Kim, *Biosens. Bioelectron.*, 2019, **132**, 279–285.

57 M. R. Montereali, L. Della Seta, W. Vastarella and R. Pilloton, *J. Mol. Catal. B: Enzym.*, 2010, **64**, 189–194.

58 C. C. Mayorga-Martínez, R. Gusmão, Z. Sofer and M. Pumera, *Angew. Chem., Int. Ed.*, 2019, **58**, 134–138.

59 M. Cerrato-Alvarez, E. Bernalte, M. J. Bernalte-García and E. Pinilla-Gil, *Talanta*, 2019, **193**, 93–99.

60 J. Abdullah, M. Ahmad, N. Karuppiah, L. Y. Heng and H. Sidek, *Sens. Actuators, B*, 2006, **114**, 604–609.

61 J. Sanz, S. de Marcos and J. Galbán, *Anal. Bioanal. Chem.*, 2012, **404**, 351–359.

62 S. K. Mishra and K. S. Chiang, *Opt. Laser Technol.*, 2020, **131**, 106464.

