

- Abstract¹
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 Antibodies are highly selective and sensitive, making them the gold standard for recognition affinity tools. However, their production cost is high and their downstream processing is time-consuming. Molecularly imprinted polymers (MIPs) are tailor-made by incorporating specific molecular recognition sites in their structure, thus translating into receptor-like activity mode of action. The interest in molecular imprinting technology, applied to biomacromolecules, has increased in the past decade. MIPs, produced using biomolecules as templates, commonly referred to as "plastic antibodies" or "artificial receptors", have been considered as suitable cheaper and easy to produce alternatives to antibodies. Research on MIPs, designed to recognize proteins or peptides is particularly important, with potential contributions towards biomedical applications, namely biosensors and targeted drug delivery systems. This mini review will cover recent advances on (bio)molecular imprinting technology, where proteins or peptides are targeted or mimicked for sensing and therapeutic applications. Polymerization methods are reviewed elsewhere, being out of the scope of this review. Template selection and immobilization approaches, monomers and applications will be discussed, highlighting possible drawbacks and gaps in research.

 Keywords: molecularly imprinted polymers, nanoparticles, artificial antibodies, biomimetics, biosensors, biomolecules, diagnostics, selective targeting, drug delivery

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1. Introduction

 The interest on molecular imprinting technology, studied since the 1970s, has grown exponentially between the 1970s and mid-2010s (Ansari and Masoum, 2019), with applications ranging from separation and purification, as selective adsorbers or 77 membranes, to sensors (Alexander et al., 2006; Whitcombe et al., 2014).

 Molecularly imprinted polymers (MIPs) are synthetic polymers that are tailor- made for specific recognition. As antibodies and enzymes, their three-dimensional structure and functional groups with a specific orientation are orchestrated to allow a selective molecular binding. Functional monomers interact with the template, or printed molecule, forming a template-monomer complex. Polymerization takes then place, in the presence the template, by reaction of a cross-linker and an initiator. After the polymer is formed, the template molecule is removed, leaving the MIP with empty cavities, that act as specific binding sites (Hui Lee and Doong, 2016; Scriba, 2016), as represented in a simplified scheme in Figure 1. This cavity will preferentially bind the template, as it matches the template's geometry, and it has affinity for its complementary functional groups. MIPs can be synthetized using virtually any molecule as a template, ranging from

 Abbreviations: AAm: acrylamide; AIBN: azo-bis isobutyronitrile; APMA: N-(3-aminopropyl) methacrylamide; APS: ammonium persulfate; BSA: bovine serum albumin; CEA: Carcinoembryonic antigen; DMAEM: 2-(dimethylamino)ethyl methacrylate; DMAPMA: N-(3- (dimethylamino)propyl)methacrylamide; EBA: N,N'-ethylenebis(acrylamide); EGDMA: ethylene glycol dimethylacrylate; EGFR: epidermal growth factor receptor; HPLC: high performance liquid chromatography; MAA – methacrylic acid; MBA: N,N' methylenebisacrylamide; MIPs: molecularly imprinted polymers; N/A: not available; NIPAm: Nisopropylacrylamide; PBS: phosphate buffered saline; PDA: 1,4-bis(acryloyl)piperazine; PEGDMA: poly(ethylene glycol)dimethacrylate; PSMA: prostate-specific membrane antigen; QCM – quartz crystal microbalance; SDS: sodium dodecyl sulphate; TBAm: N-tertbutylacrylamide; TEMED: N,N,N′,N′-Tetramethylethylenediamine.

 drugs, small molecules, amino acids, chiral enantiomers, DNA, peptides, proteins and even whole cells (Haginaka and Sakai, 2000; Hammam et al., 2018; Lin et al., 1997; Liustrovaite et al., 2023; Mohajeri and Ebrahimi, 2008; Ping Li et al., 2004; Suedee et al., 2002; Trinh et al., 2018), thus translating into a target molecule size range from a few Da (*e.g.* 126 Da melamine (Poma et al., 2013)) to several kDa (*e.g* 66.5 kDa bovine serum albumin (BSA) (Arabi et al., 2021b) and 180 kDa spike glycoprotein of SARS-CoV-2 (Ratautaite et al., 2023, 2022)).

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98 Figure 1. Schematic representation of protein MIP synthesis.

 Biomolecules such as antibodies, some receptors, and enzymes are the gold standard for affinity tools, since their target recognition capacity is highly selective and sensitive. However, the use of natural biomolecules presents disadvantages, including: limited working conditions, such as mild temperature, narrow pH range and low stability in organic solvents. Antibody production in mammalian cells has been optimized over last decades, still the associate production costs are high. Recombinant expression in bacterial or yeast systems still presents limitations like endotoxin production (Arbabi- Ghahroudi, 2022; Asaadi et al., 2021; Liu and Huang, 2018; Malaquias et al., 2021; Mark et al., 2022; Thompson et al., 2016). Therefore, given that MIPs are usually cheap, easy to synthetize in a reproducible way, and have shown robust performances in a variety of solvents (Hui Lee and Doong, 2016; Wackerlig and Schirhagl, 2016), MIPs, using biomolecules as templates, have been considered as suitable alternatives for medical diagnosis and theragnostics in the biomedicine field and as replacement of enzymes in catalytic processes or even used as bioelectrodes for energy harvesting based on microbial fuel cells in more advanced MIP applications (Ostovan et al., 2022).

 Due to their functional similarity to their natural counterparts, MIPs designed for biomolecules have been referred to as "plastic antibodies" or "artificial receptors". Research developed on MIPs selective for biomacromolecules, like proteins or peptides, is particularly important, since these have the potential to contribute towards biomedical applications aiming at biosensors, toxic analyte sequestration, or drug delivery systems, among others (Canfarotta et al., 2018; Chen et al., 2016; Hui Lee and Doong, 2016; Suedee et al., 2002).

 Molecularly imprinted technology has significantly contributed on the development of novel biosensors for disease detection. Namely, several cancer diagnostics sensors based on MIPs has been development, using prostate, breast, ovarian and hepatic cancer biomarkers as target molecules (Pilvenyte et al., 2023b). Polypyrrole-based electrochemical MIP sensors were developed targeting the CA-125 marker for

 epithelial ovarian cancer (Rebelo et al., 2019), the CA15-3 marker for breast cancer (Santos et al., 2018), and the PSA protein marker for prostate cancer (Yazdani et al., 2019). In the latter study, it was developed a sensor for PSA with a limit of detection (LOD) so low as 2.0 pg/mL, which is below the threshold "risk" values of 4.0-10.0 ng/mL of PSA concentration in blood, thus showing the potential competitiveness of this assay. Biomarkers for neurodegenerative diseases is another relevant focus area of MIP-based biosensors with recent advances in the development of MIP-based electrochemical sensors for Alzheimer's and Parkinson's diseases (Pilvenyte et al., 2023a). For instance, 135 a very competitive Alzheimer's disease MIP biosensor was designed for an amyloid- β peptide using a combination of polypyrrole and carbon nanotubes, reaching a LOD as low as 0.3 fg/mL (Özcan et al., 2020). A MIP-based biosensor was similarly developed for α- synuclein peptides with a LOD of 10 fg/mL, for Parkinson's disease, using aniline, the monomer of the conducting polymer polyaniline, and *m*-aminobenzenesulfonic acid (Lee et al., 2021). Overall, such studies suggest a growing tendency in developing MIP-based biosensors using such functional monomers to obtain an electroconductive polymeric structure in the MIP to enable high detection sensitivities (Canfarotta et al., 2016; Ramanavicius et al., 2022; Ramanavicius and Ramanavicius, 2022).

 This mini-review is focused on current advances on molecular imprinting technology where proteins, peptides and epitopes are used as templates, and/or MIPs that are designed to mimic biological (macro)molecules. Polymerization methods are reviewed elsewhere (Haupt et al., 2020), being out of the scope of this review. Template selection and immobilization approaches, functional monomers and applications will be discussed, highlighting possible drawbacks and gaps in the literature.

2. Protein imprinting

 Proteins are biomacromolecules of significant interest in research, with special focus on their detection and quantification, since they are often biomarkers of important human diseases, including viral infections (Cennamo et al., 2021; Liv et al., 2021; Raziq et al., 2021; Sukjee et al., 2022; Tai et al., 2005), hormonal and DNA regulation processes (Rachkov and Minoura, 2000; Zhang and Liu, 2018), and many cancer types (Canfarotta et al., 2018; Han et al., 2019; Tang et al., 2018; Zhang et al., 2015). However, detection and quantification of proteins requires labour and cost-intensive separation methods, often based on immunoassays.

 Table 1 shows a short overview of the common functional monomers, initiators, cross-linkers and solvents used on studies for protein imprinted MIPs. Apart from the polymerization method chosen, reagents for the polymerization reaction appear to be quite similar among studies with protein imprinted MIPs, without much novelty presented in terms of functional monomers, initiators and cross-linkers selected, in agreement with what has already been reviewed elsewhere (Teixeira et al., 2021; Yang et al., 2019).

168 **Table 1.** Short overview of polymerization reagents (monomers, cross-linkers, initiators) 169 and solvent for selected studies using protein imprinting technology.

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 AAm: acrylamide; AIBN: azo-bis isobutyronitrile; APMA: N-(3-aminopropyl) methacrylamide; APS: ammonium persulfate; DMAEM: 2-(dimethylamino)ethyl methacrylate; DMAPMA: N-(3-(dimethylamino)propyl)methacrylamide; EBA: N,N'- ethylenebis(acrylamide); EGDMA: ethylene glycol dimethylacrylate; MBA: N,N'- methylenebisacrylamide; N/A: not available; NIPAm: N-isopropylacrylamide; PBS: phosphate buffered saline; PDA: 1,4-bis(acryloyl)piperazine; PEGDMA: poly(ethylene glycol)dimethacrylate; SDS: sodium dodecyl sulphate; TBAm: N-tert-butylacrylamide; TEMED: N,N,N′,N′-Tetramethylethylenediamine.

 To obtain MIPs for biomacromolecules, two main protein imprinting strategies have been developed: protein imprinting, including i) non-oriented surface imprinting and ii) oriented surface imprinting, where the entire protein works as template; and epitope imprinting, where the template will be a part of the structure of the target protein. Representative examples found in literature of protein and epitope imprinting strategies discussed in this review are represented in Figure 2.

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Figure 2. Representative synthesis methods: non-oriented surface imprinting – electropolymerization (reprinted from (Shumyantseva et al., 2016), Copyright 2023, with permission from Elsevier) and precipitation (reprinted with permission from (Hoshino et al., 2008), Copyright 2023 American Chemical Society); oriented surface imprinting – sol-gel (reprinted from (Guoning et al., 2020), Copyright 2023, with permission from

 Elsevier); epitope imprinting: solid-phase synthesis (reprinted with permission from (Canfarotta et al., 2018), Copyright 2023 American Chemical Society).

2.1. Protein imprinting

 In the protein imprinting approach, the whole protein is the template. Such strategy, following the traditional MIP concept, could be argued as the most appropriate biomimetic approach in terms of binding affinity, since it would retain the tertiary structure of the target protein, as well as, affinity groups for weak protein interactions, such as hydrogen bonds, electrostatic and van der Waals interactions (Boysen, 2019). However, epitope imprinting, which will be reviewed further ahead, may provide a recognition mechanism more similar to natural receptors. Depending on the application, it may be useful to have a binding site for only a specific peptide motif of the biomolecule. This is particularly relevant for cell membrane proteins, when considering specific biological variants detection, or to promote for cost-effective MIPs development and manufacture strategies. The literature reports MIPs targeting common proteins, such as BSA, albumin, ribonuclease A, horseradish peroxidase (HRP), or cytochrome c, using molecular imprinting methods based on different strategies for the immobilization of the target, and varied functional monomers (Kryscio and Peppas, 2012; Li et al., 2006; Wang et al., 2019). Additional examples are resumed in Table 2, including methods for removal of the template after MIP synthesis. Template removal techniques include protein denaturation steps, so that the template changes its conformation and will be released from the imprinted cavity, followed by washing steps to promote the elution of the denatured template. A drawback of these methods is that they do not allow recovery of the template molecule for reuse. Another methodology is based solely in washing with mild solvents, like aqueous solutions, thus relying on disruption of weak interactions, such as electrostatic and hydrogen bonds, and the slow diffusion of the protein through the polymer network. Additionally, the use of thermo-responsive monomers (e.g. N- isopropylacrylamide) in the structure of the MIP allows the release of the protein simply by increasing the temperature of the washing solution above the lower critical solution temperature (LCST), which leads to increase space between polymer chains, allowing the protein to be released from the MIP's cavity.

227 **Table 2.** Summary of protein templates and template removal procedures for selected 228 studies using protein imprinting technology.

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 Some studies report the development of MIPS for the whole tertiary protein structure in liquid solution, by addition of the target biomolecule to the reaction mixture (Wang et al., 2014; Yang et al., 2023). Some studies have even reported the use of whole cells, like bacteria, to develop MIP-based sensors reaching relevant LOD values, a MIP- based electrochemical sensor developed for Listeria monocytogenes has a LOD of 70 CFU/mL (Liustrovaite et al., 2023). However, the most common approach reported resorts to solid phase synthesis, in which the template protein is first immobilized in a solid support such as silica beads (Canfarotta et al., 2016; Zhang et al., 2009), glass surfaces (El Kirat et al., 2009; Kryscio and Peppas, 2012) and other silica moulds (Dabrowski et al., 2019; Li et al., 2006) and an affinity chromatography step is used for the synthesis and purification of the molecularly imprinted nanoparticles (MIP-NPs) (Figure 2 and Figure 3).

 Immobilization of the protein to the solid support could be achieved using an affinity ligand of the protein (Ambrosini et al., 2013), or by chemical functionalization of the surface of the solid support (Canfarotta et al., 2016). The use of an affinity ligand enables the orientation of the immobilized protein, meaning that all binding sites are constructed with similar orientation, thus improving binding site homogeneity.

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Figure 3. Schematic representation of protein imprinting with MIP synthesis with the target protein immobilized on the surface of a silica bead.

 Ambrosini and co-workers reported the use of solid-phase synthesis of MIP-NPs for protein recognition using the model protein trypsin (23 kDa) (Ambrosini et al., 2013), adapting the solid-phase synthesis method previously developed for synthesis of a MIP for melamine, a small molecule (<1 kDa) (Poma et al., 2013). Trypsin was immobilized on the surface of glass beads, according with the approach reported for melamine, and the beads were then packed into a column, where the functional monomers (NIPAM and EBA) were added, and the reaction took place. Finally, several washing steps were performed for purification of the MIPs. Despite the difference in size of the target molecules, the immobilization on the solid support appears to depend mostly on the functional groups present in the target molecule, thus rendering solid-phase synthesis a versatile method for MIP synthesis. Still, one has to consider that large biomolecules are more complex and they often have similar reactive groups (e.g. amines or carboxylic acids) on different locations, which makes more challenging to obtain specificity on immobilization of the template biomolecule with uniform orientations. One advantage of the solid-phase synthesis strategy is the decrease of the cost of the process due to the solid phase being reused several times for MIP synthesis, thereby saving template molecules. 269 One study reported the maintenance of the size and K_D of the MIPs for over 30 batches of template reuse (Poma et al., 2013), that value being inferior to the standard protein A chromatography performance for antibody production, which can be reused for 100 cycles. However, stability of proteins under reaction conditions for several cycles should be assessed, as literature is scarce concerning this information and there is a severe lack of studies presenting a process design and model along with realistic economic analyses.

 To selectively separate lysozyme from a mixture of proteins in aqueous solution, acrylamide and acryloyl-β-cyclodextrin were used as functional monomers (Zhang et al., 2009). Here, the target protein was immobilized on the surface of silica beads and the polymerization reaction took place around the immobilized lysozyme. After removal of the template and consequent detachment from the beads, the MIPs were packed in a column. A successful high performance liquid chromatography (HPLC) separation was achieved, with lysozyme selectively separated from cytochrome c, BSA and avidin, with a maximum adsorption capacity for lysozyme of 44.6 mg/g, being 4 times higher than for the remaining proteins (Zhang et al., 2009). In a different study, instead of using solid phase synthesis, the same acrylamide and methacrylic acid were also used as functional monomers to obtain a lysozyme imprinted MIP. Impressively, the obtained MIP was

 successfully used to purify this protein from egg white using a chromatographic column, 287 with maximum adsorption capacity reaching 94.8 mg/g , purity close to 100% and mass recovery of 98.2% (Wang et al., 2014). Overall, these studies highlight the high values of maximum adsorption capacities achieved through solid-phase synthesis of MIPs for proteins, thus suggesting that immobilization and consequent orientation of the template molecule might contribute to higher selectivity of the MIPs.

 A different approach that also does not require the immobilization of the template is MIP electrosynthesis, where cyclic voltammetry is used for electropolymerisation of the MIP. Here, a pre-polymerisation mixture of template protein and monomer is prepared, which is then deposited onto an electrode surface by applying cyclic potential sweeps. A study on the electrochemical quantification of troponin T (37 kDa), a biomarker of myocardial injury, used o-phenylenediamine as a monomer and a gold electrode for deposition of the MIP to build a biosensor based on a redox probe (Karimian et al., 2014). A similar study for recognition and electrochemical detection of myoglobin (18 kDa) used screen-printed electrodes where the pre-polymerisation mixture of myoglobin and o-phenylenediamine was electrodeposited (Shumyantseva et al., 2016). Even so, it is necessary to consider the difficulty to scale-up electrosynthesis processes due to potential difficulties to increase the electrode active area and electronic transport in the bulk of the reaction mixture. Furthermore, recyclability of the template is not considered, thus limiting the applicability of this method to less expensive targets, or resulting in process with prohibitive costs.

 To the best of our knowledge, only one molecular imprinting methodology was reported with the objective to synthetize a replica of the protein, which is based on a two- step imprinting process. The strategy follows the approach to: i) firstly, to obtain a molecular cast of the target antibody, synthetized as a MIP particle, and then ii) to perform the second imprinting stage, analogous to a stamping method, in which polymerization occurs by compression of the pre-synthetized MIPs particles onto a pre‐polymerized layer placed on the surface of a quartz crystal microbalance (QCM) electrode. Therefore, the polymer layer will be covered with molecularly imprinted antibody replicas after removing the stamp (Hussain et al., 2013; Jenik et al., 2009; Latif et al., 2014; Schirhagl et al., 2010). This methodology limits the range of applications to those that are usually based on immobilized antibodies on surfaces, such as biosensors or immunoassays that are often performed on chips. However, it is difficult to gather if the production of such immunoassay platforms could be improved with this strategy instead of using actual antibodies. Again, literature is found lacking on an actual economic analysis of the cost of production.

 Although direct imprinting for detection and separation of proteins seems to be fairly well explored in the literature, there is still a call for designing MIPs for other challenging proteins with biomedical interest, particularly disease biomarkers, like surface membrane proteins expressed in cancer. One point of concern is the fact that cross-selectivity between similar proteins may impair a MIP performance, resulting on false positives, as several studies look at the selectivity of the MIPs against proteins that are not the target, but share similar structural characteristics. Indeed, in a competitive assay, using the previously mentioned lysozyme MIPs, no statistical difference was found in adsorption capacity for lysozyme, trypsin and cytochrome c, three proteins of high isoelectric point. While such MIP bound preferentially to lysozyme, the maximum adsorption capacity was close to 800 mg/g for the three proteins (Culver et al., 2016). This result raises a concern over the fact that imprinting alone may not account for the selectivity of certain classes of proteins.

2.2. Epitope imprinting

 When entire proteins are used as templates for MIPs preparation, their efficient removal after polymerization is impaired, due to difficult diffusion through the MIP network. Additionally, proteins' tertiary conformations, which depend on conditions, such as pH, solvent and temperature (Kryscio et al., 2012), are unstable, contributing to lack of MIPs selectivity. To overcome such drawbacks, using only a part of the protein as template has been purposed as an imprinting strategy. Indeed, in the epitope imprinting approach, short linear peptides are used as the target molecules for the MIP. Therefore, in this case, the selective recognition neglects the protein 3D conformational specificity and relies on amino acid recognition, since imprinting is based only on the amino acid sequence, the primary structure of the peptide, instead of secondary and tertiary structures of proteins. Epitope design and selection strategies have been extensively discussed elsewhere (Tse Sum Bui et al., 2023), covering computational tools for selection of appropriate amino acids sequences and peptide length to maximize affinity of the MIP developed.

 A possible strategy for proteins that have their C- or N-terminus exposed, such extremity is used as the site around which the MIP will be formed and the protein selectively captured. Nonapeptides, peptides with nine amino acid sequence length, as target molecules have been selected the length large enough to allow the unique identification of a particular protein (Nishino et al., 2006). Such strategy was used for the cases of MIPs development for cytochrome c, BSA and alcohol dehydrogenase (ADH), in which as nine amino acid sequence peptides of the C-terminus of these proteins were successfully used as templates in the synthesis of molecular imprinted films (Nishino et al., 2006). In these examples, the authors also used solid phase synthesis to facilitate the polymerization reaction around the template peptides.

 MIP-NPs were also synthetized for an exposed C-terminus peptide of green fluorescent protein (GFP) by inverse microemulsion polymerization, using AAm and MBA as monomers and APS and TEMED as initiators (Zeng et al., 2011). Surfactants were also added so that polymerization occurred in inverse microemulsion, where the peptide was correctly oriented at the interface of water and oil domains, without the need for previous template surface immobilization. In another study an exposed antigenic domain of Lpp20, an outer membrane lipoprotein antigen specifically expressed by all the *H. pylori* strains, was used as a template in inverse mini-emulsion polymerization, using AAm as functional monomer. The obtained MIP was successfully assessed to capture the bacteria *H. pylori* (Han et al., 2015). While examples of epitope imprinting by inverse emulsion polymerization are still scarcely reported, possibly due to poor stability of epitopes on the water/oil interface, this is an innovative and apparent simple method for MIPs preparation, as it skips the steps of template immobilization required on solid phase synthesis, *i.e.* avoids the steps of activation and functionalization of the solid support, immobilization of template and several washing steps.

 It is worth noting that most studies overviewed here use acrylate-based functional monomers, without discussion further progresses in the literature in the selection and use of alternatives types of monomers. The broader use of different functional monomers is reviewed elsewhere (Teixeira et al., 2021). Again, for the most studies reviewed, MBA and APS are the recurrent choices for cross-linker and initiator, respectively. Such selection has several advantages in terms of polymerization techniques efficiency and obtained MIP performance. However, considering today's concerns on promoting the development of sustainable and greener processes in MIP design (Arabi et al., 2021a),

 alternative reagents could be procured, such as, for example, itaconic acid obtained from fermentation of Aspergillus species.

 Overall, epitope imprinting has the advantage of relying only on the use of short peptides as templates instead of the whole protein, which are potentially easy to synthetize, and thus more available than their protein counterparts. However, not all proteins meet the criteria set for this approach.

2.2.1. Conformational epitope imprinting

 Epitope imprinting based on the primary structure of an exposed peptide sequence, although efficient, is limited, as not all proteins meet the required criteria, such as the target protein to include an exposed C-terminus or an amino acid sequence both specific and short enough. A possible solution for these limitations is the use of conformational epitope imprinting, this approach uses as template the primary, secondary and tertiary structure of the selected epitope, thus, incorporating the recognition of the specific 3D conformation of the epitope into the MIP.

 This strategy was applied in the design of a MIP for the p32 protein, by inverse microemulsion polymerization (Zhang et al., 2015). p32, a membrane protein that is overexpressed on the surface of varied tumour cell types, has the potential to target and mediate drug delivery and thus the developed MIP has the potential to actively targeting 405 tumours. This protein has an N-terminal α -helix in the extracellular domain, which was set as the target site for the MIP. The peptide apamin mimics the extracellular domain of p32, as it has a sequence of seven aa residues identical to the one present on this domain, and it was successfully used as the imprinting template (Zhang et al., 2015).

2.2.2. Peptide imprinting

 Peptide imprinting has not only been used for protein recognition, but also for recognition of the peptides themselves. In this strategy, a short peptide sequence is again used as template. Synthesis of peptide-selective MIPs has been reported for varied peptides. A MIP for the recognition of the hormone oxytocin was synthetized by bulk polymerization, using as template a tetrapeptide with the same three amino-acid C- terminal section of the structure of oxytocin, and using methacrylic acid (MAA) as functional monomer and ethylene glycol dimethacrylate (EGDMA) as crosslinker (Rachkov and Minoura, 2000).

 A MIP for the recognition of the hormone angiotensin II, with a detection limit of 8 pM, was achieved by free radical polymerization using as functional monomer sodium acrylate, and as cross-linker poly(ethylene glycol) diacrylate. In this case, the whole peptide was used as template since the it has a short eight aa sequence (Rachkov et al., 2004).

 The synthesis of MIP-NPs for the specific binding of melittin, which is a bee venom biotoxin, was achieved by precipitation polymerization (Hoshino et al., 2008). Melittin has 26 amino acids, of which 6 are positively charged, 6 residues at the C- terminus are hydrophilic, and the remainder is mostly composed of non-polar amino acid residues. In this study, the authors established a rationale for monomer selection considering the overall polarity of the peptide and the individual charge of each amino acid residue of the peptide sequence. Hence, the optimum monomer combination should comprise a mix of hydrophobic and negatively charged monomers to bind to the opposing charges of the residues of the target peptide. The two most successful monomer combinations contain 40% of hydrophobic monomers (TBAm) and 5% of negatively

 charged functional monomers (acrylic acid, AAc), or 5% hydrogen bonding monomer (AAm), 5% negatively charged monomers (AAc) and 40% of hydrophobic monomers 437 (TBAm). Dissociation constants (K_D) obtained by nonlinear fitting of Langmuir isotherms for these MIPs were in the range of 7.3-25 pM, which are comparable to the 439 dissociation constant of a natural antibody (17 pM). The values of K_D of the MIPs are of the same order of magnitude as the ones found for antibodies, which range from 10 pM to 5 nM depending on antibody (Friguet et al., 1985; Landry et al., 2015; Pan et al., 2016), thus suggesting that the same affinities can be achieved for the MIP as the natural counterpart, showing the importance of functional monomer selection for successful biomacromolecule MIP synthesis. These strategies illustrate a more holistic approach where the global charge and hydrophobic/hydrophilic balance of molecules is considered on MIP design, to potentiating more probable success on tailoring host site formation to template specific structural properties.

3. Applications

 MIPs for biomacromolecules have the potential for numerous applications, from bioseparation and purification processes in biotechnology and pharmaceutical industries to biomedical applications. Many of the studies described in this section target selective recognition of proteins aiming at the potential application of MIPs in separation and purification of proteins within pharma or food industry processes. In this context, the use of MIPs has been explored in affinity chromatography as it has been demonstrated for lysozyme, BSA, haemoglobin, and cytochrome c (Li et al., 2006; Ouyang et al., 2010; Wang et al., 2019, 2014; Zhang et al., 2009).

 The biomedical field covers a broad spectrum of potential applications for biomacromolecule MIPs, spanning from biosensors and targeted drug delivery. MIP- based biosensors have been extensively explored in the literature, with protein quantification method being the prime target for MIP application, as the current technology relies on expensive immunoassays. An especially motivating application of MIP biosensors is their use as diagnostic tools. Figure 4 and Table 3 resume some examples of recent studies using MIPs for proteins, coupled to varied sensing units, as biosensor diagnostic tests with clinical relevance. As previously mentioned, MIP-based biosensors for detection of biomarkers of cancer and neurodegenerative diseases, often using electrochemical detection, have been a focus of recent research. Nevertheless, MIP- based electrochemical biosensors have also been developed for other clinically relevant biomarkers such as troponin T (Karimian et al., 2014) or myoglobin (Shumyantseva et al., 2016) in cardiac disease. Several studies have also explored MIPs for detection of viral detection systems, for example for poliovirus (Wang et al., 2010), bovine leukaemia virus (Ramanaviciene and Ramanavicius, 2004) and dengue virus (Tai et al., 2005).

 Of particular relevance in recent years are MIP-based sensors for detection of SARS-CoV-2, for instance using a disposable electrochemical chip with high sensitivity (Raziq et al., 2021), or a surface plasmon resonance optical sensor (Cennamo et al., 2021), capable of detecting SARS-CoV-2 in nasopharyngeal swab samples of COVID-19 positive patients (Figure 4). Further works using MIP-based electrochemical sensors for SARS-CoV-2 have been developed using electroconductive polymers, namely polypyrrole, with significant sensitivity of detection with calibration curves with protein concentrations ranging 0-25 µg/mL (Ratautaite et al., 2023, 2022). A study worth noting is the design of an electrochemical sensor using two alkane thiols (11- mercaptoundecanoic acid and 6-mercapto-1-hexanol) to form a self-assembled monolayer MIP for the spike protein of SARS-CoV-2 with a reported LOD as low as 0.34

 nM and a limit of quantification around 1 nM (Zukauskas et al., 2023). Overall, electrochemical detection coupled to MIP for proteins appears to be a promising method to reach low LODs for relevant clinical targets.

Imprinting on capillary tubes surface coated with Au nanostars for SERS detection

Electropolymerization on electrode surface for electrochemical detection

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490 Figure 4. Representative surface mediated MIPs detection methods, including surface plasmon resonance (SPR) using a plastic optical fiber (POF) (reprinted with permission from (Cennamo et al., 2021) under Creative Commons License – Attribution 4.0 International – CC BY 4.0 – https://creativecommons.org/licenses/by/4.0/legalcode), 494 OCM (reprinted from (Lim et al., 2023), Copyright 2023, with permission from Elsevier), surface-enhanced Raman scattering (SERS) (reprinted from (Arabi et al., 2021b), Copyright 2023, with permission from Elsevier), and an electrochemical method (reprinted from (Raziq et al., 2021), Copyright 2023, with permission from Elsevier).

499 **Table 3.** MIPs for proteins with clinical relevance and detection units as biosensor 500 diagnostic tools.

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MIPs may also be explored as targeting agents, for example, of a tumour, by 504 binding to the membrane surface protein of the tumour cells, namely of a protein that is 505 overexpressed on the surface of cancer cells. In this line of work, MIP-NPs have been

 developed to target the extracellular α-helix domain of the p32 protein in *in vitro* and *in vivo* models (Zhang et al., 2015). Such type of MIPs, loaded with fluorescent probe (IR- 783 dye), were assessed for tumour imaging in mice, and when encapsulating a photosensitizer compound, such as methylene blue, that was used for photodynamic treatment of tumours. The encapsulation process took place during polymerization by simply adding the desired compounds (IR-783 and methylene blue) to the aqueous phase before initiating the inverse microemulsion polymerization, where AAm and MBA were used, respectively, as functional monomer and cross-linker (Zhang et al., 2015). This study demonstrated that MIP-NPs can simultaneously function as targeting tools and nanocarriers for drugs.

 In another study, the C-terminal linear peptide of EGFR (amino acids 418−435: SLNITSLGLRSLKEISDG), an overexpressed receptor on the cell surface of many tumours, was used as the template for production of MIP-NPs by solid phase synthesis. This MIP was assessed for targeted drug delivery to MDA-MB-468 breast cancer cells (Canfarotta et al., 2018). In this case, a dual imprinting strategy was followed, in which the chosen drug (doxorubicin) was used as a secondary template present in solution with the monomers (NIPAm, TBAm and APMA) and the cross-linker (MBA) (Canfarotta et al., 2018), as represented in [Figure](#page-15-0) *5*5. After binding of the MIP to the EGFR receptor, the anticancer drug would be released by diffusion and accumulate around cancer cells, promoting cancer cell death. This study illustrates a strategy where the MIP presents binding sites for the drug and for the membrane receptor, demonstrating that MIP-NPs can selectively deliver a drug, by response to the tumour microenvironment, to specific cell targets.

 The potential of using MIPs for therapies in humans is an extremely important feature that need to be addressed. Some studies have demonstrated efficacy in cell culture and some have used *in vivo* models, such as mice. For instance, a MIP targeting the folate receptor in cancer cells have been tested in mice bearing tumours, showing both the safety of using the MIPs in an organism, while showing the effectiveness of the targeting of the tumour and drug delivery (Liu et al., 2017).

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537 Figure 5. Schematic representation of dual imprinting strategy using solid-phase synthesis.

 An important feature in clinical applications is the successful detection of a therapeutic targeting agent by imaging techniques. A synergistic chemo- and photo- dynamic cancer therapy with a dual imaging agent relying on dual-template MIP-NPs was recently demonstrated in *in vitro* and *in vivo* models. In this study, the MIPs were synthetized for the CD59 epitope, as this protein is overexpressed in solid tumours, and the secondary template was doxorubicin as the chemotherapy agent. For the dual fluorescent/magnetic resonance imaging, gadolinium-doped silicon quantum dots were first prepared and used as the core of the MIP-NPs, and photosensitizer chlorin e6, the photo-dynamic cancer therapeutic, was embedded in the silica core. The polymerization took place on the surface using NIPAm, Aam and TBAm as functional monomers and MBA as cross-linker (Peng et al., 2020). The synthesis process complexity increases the number of variables to be considered over the synthesis, *i.e.* functional monomers that will match functional groups in both peptide and drug templates, cross-linkers, initiators and solvents compatible for both molecules. This work successfully showed that the already evidenced targeting ability of MIPs for biomarkers can be further explored to comprise several therapeutic options. Furthermore, it also evidences the potential of protein MIP-NPs to be coupled to imaging tools, allowing a more precise guided cancer treatment.

- **4. Conclusions**
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 The field of molecular imprinting of biomacromolecules has seen significant growth over the past decade despite the inherent challenges, such as the need of their synthesis to be compatible with the use of aqueous media and allow for the immobilization of the target molecule, to ensure that proteins and peptides maintain their native conformations during polymerization.

 In light of the advantages of MIPs, such as being cheap, easy to synthetize in a reproducible way, and showing robust performances in a variety of solvents, it is not surprising that MIPs for biomolecules have been considered as suitable for medical diagnosis and therapeutics, but also as replacement of enzymes in catalytic processes or even used as bioelectrodes for energy harvesting.

 Concerning the template molecule, most proof-of-concept studies used as templates the readily available and "low-cost" proteins, such as BSA or cytochrome c. This could be somewhat expected as more interesting protein targets are expensive, as is the case of human disease biomarkers, such as receptors that are overexpressed on the surface of tumour cells. However, the impact of using MIPs, instead of antibodies, is based on the cost effectiveness of the process to obtain MIPs, therefore it is important to maintain low costs when expensive templates are used. A possible strategy could be the reuse of the template in several polymerization reactions, when it is immobilized on a solid surface, similarly to what is seen for catalytic enzymes in bioprocesses, that are reused throughout batches until they lose enzymatic activity. Unfortunately, the literature is severely lacking on studies reporting for how many batches, can the templates be reused in MIPs synthesis, with only one available study reporting the maintenance of the MIPs 583 properties (size and K_D) for over 30 batches (Poma et al., 2013). Therefore, there is the need of additional studies on templates recyclability, scaling up of synthesis and purification of MIPs for proteins. It is fundamental that process design and model accompanied by economic analyses to be performed for such systems in order to assess the applicability of MIPs for proteins in industry, concerning MIPs economic competitiveness with the use of antibodies.

 Epitope imprinting appears to be a promising approach for the near future as it not only minimizes the complications of dealing with very large and condition-sensitive templates during polymerization, but also decreases the difficulty in template removal after syntheses, due to diffusion constraints in the polymer network. This strategy could contribute to decrease the cost associated with the template as well. However, this also

 comes with limitations as these epitopes need to comply with certain characteristics, namely an appropriate length to ensure a specific recognition mechanism, and loss of the tertiary structure contributions to recognition specificity.

 In terms of applications, the potential of using MIPs as biosensors seems to be a consensus in the literature, with most studies focusing on diagnostics, ranging from cardiovascular and neurodegenerative diseases to cancer. There is a continuous call for new MIP development as new pathogens are identified, and more reliable disease biomarkers are being assessed in biology and medicine. The targeting potential for directed drug delivery has also been reported showing that MIPs for biomarkers can target tumours and function as drug carriers, for example. However, it was not until recently that MIPs with dual affinity sites (biomarker and drug) were synthetized and tested in *in vitro* and *in vivo* models. Indeed, this adds complexity to the synthesis process and possible cross interference in binding site formation for the protein should be further assessed. So far, studies have relied on MIP-NP endocytosis by the cells for drug delivery or laser incidence for photodynamic therapy. Thus, it would be interesting to assess the incorporation of stimuli responsive characteristics on MIPs, such as pH-responsive or electro-responsive, for enhanced controlled targeted drug delivery.

 Furthermore, it is extremely important to ensure the safety of use of MIPs in human patients. However, very few studies have been performed using animal models in pre-clinical settings, to show the safety and efficacy of MIPs for therapeutic applications.

 Overall, moving forward to clinical approval and commercialization of MIPs for proteins and peptides for biomedical applications, it would be interesting to see further studies reporting biocompatibility, safety and immunogenic responses using in vivo models to ease bench to bedside transition. Additionally, for studies claiming reusability of the template and downsizing of production costs, it will be important to have experimental data and economic analysis supporting claims that would ease the path to commercial success of MIPs.

Declaration of competing interest

 The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

- **Acknowledgments**
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 This work is financed by national funds from FCT - Fundação para a Ciência e a Tecnologia, I.P., with dedicated funds from the project eOnco (2022.07252.PTDC) and the PhD scholarship (SFRH/BD/145057/2019), iBB (UIDB/04565/2020 and UIDP/04565/2020), i4HB (LA/P/0140/2020). This publication is part of the I+D+i project PID2021-125767OB-I00 funded by MCIN/AEI/ 10.13039/501100011033 and, as appropriate, by "ERDF A way of making Europe", by the European Union. Authors are thankful to the Agència de Gestió d'Ajuts Universitaris i de Recerca (2021 SGR 00387) for financial support. Support for the research of C.A. was also received through the prize "ICREA Academia" for excellence in research funded by the Generalitat de Catalunya.

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