

Supporting Information

Phosphate-binding protein-loaded iron oxide particles: Adsorption performance for phosphorus removal and recovery from water

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The supporting information includes four sections:

- S1 provides characteristics of the tertiary wastewater effluent sample.
- S2 provides a transmission electron microscopy (TEM) image showing the NHS-activated magnetic IOPs.
- S3 describes the adsorption kinetic models for P_i using PBP-IOPs.
- S4 details the theoretical calculations used to explore the influence of protein and substrate parameters on phosphate adsorption capacity.

S1. Wastewater Effluent Characterization

The wastewater effluent parameters are shown in Table S1.

Table S1. Tertiary wastewater effluent parameters from the South Shore Water Reclamation Facility in Oak Creek, WI. Each measurement was performed in triplicate.

Parameter*	Average \pm 1 Standard Deviation	Unit
Total Suspended Solid (TSS)	16 \pm 3	mg L ⁻¹
Dissolved Organic Carbon (DOC)	9.2 \pm 0.1	mg-C L ⁻¹
Total Hardness	327 \pm 10	mg L ⁻¹ as CaCO ₃
Alkalinity	180 \pm 10	mg L ⁻¹ as CaCO ₃
Chloride	460 \pm 10	mg L ⁻¹
Phosphate	1.2 \pm 0.1	mg PO ₄ ³⁻ L ⁻¹
Nitrate	8.0 \pm 0.05	mg NO ₃ ⁻ L ⁻¹
Sulfate	224 \pm 0.13	mg SO ₄ ²⁻ L ⁻¹
Hydrogen Sulfide	0	mg S ²⁻ L ⁻¹
pH	7.13 \pm 0.1	---

* TSS was measured in accordance with Method 2540D from Standard Methods for the Examination of Water and Wastewater (1). DOC was measured in accordance with U.S. EPA Method 415.3 using a Shimadzu TOC-V_{CSN}. Other parameters were measured using Single Parameter Test Kits from the Hach Company (Loveland, CO).

S2. Transmission Electron Microscopy (TEM) of NHS-IOPs

A TEM image showing the NHS-activated magnetic IOPs is shown in Figure S1.



Figure S1. Transmission electron microscopy image of BcMagTM NHS-activated magnetic IOPs (image shared by the material supplier, Bioclone Inc., USA). The TEM image depicts the silica shell surrounding the iron oxide particles.

S3. Adsorption Kinetics

Figure S2 shows the linearized pseudo second-order phosphate adsorption kinetics using PBP-IOPs. Figure S3 shows the linearized pseudo first-order phosphate adsorption kinetics using PBP-IOPs.

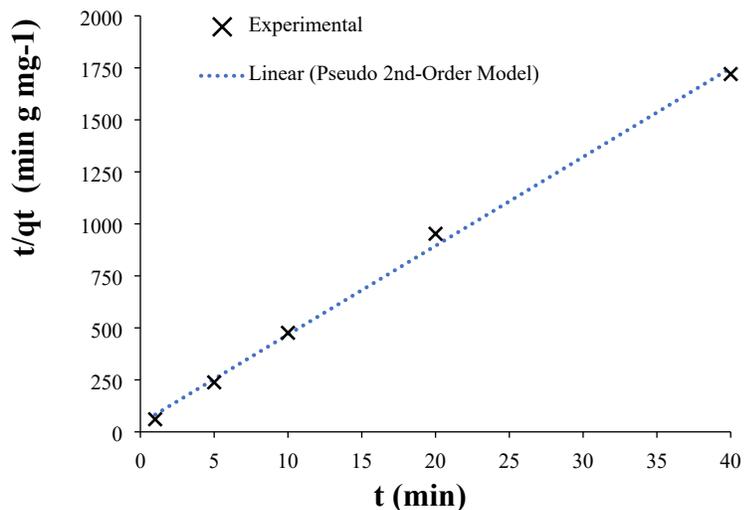


Figure S2. Linearized pseudo second-order P_i adsorption kinetics using the PBP-IOPs at neutral pH and 20 °C. Best fit model parameters: $k_2 = 45.4 \text{ g mg}^{-1}\text{min}^{-1}$, $q_e = 0.023 \text{ mg g}^{-1}$.

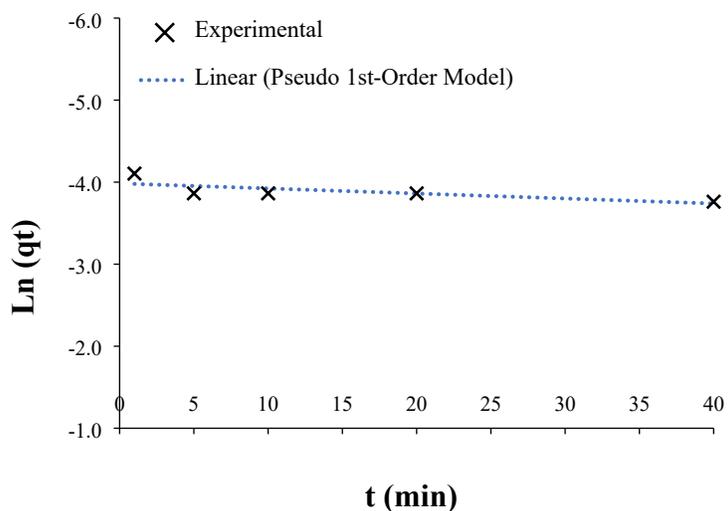


Figure S3. Linearized pseudo first-order P_i adsorption kinetics using the PBP-IOPs at neutral pH and 20 °C. Best fit model parameters: $k_1 = 0.0061 \text{ min}^{-1}$, $q_e = 0.019 \text{ mg g}^{-1}$.

S4. Calculation of theoretical capacity

Theoretical estimates of P_i adsorption were performed to assess the relative influence of the protein (or peptide) and the particle on which the protein was immobilized. All estimates were made assuming the particle and the PBP or peptide could be modeled as spherical particles. We recognize that these modeling scenarios do not fully represent reality (e.g., spherical particle modeling, neglecting particle curvature, etc.); however, these back-of-the-envelope theoretical estimations enable scenario comparisons and inform directions for fruitful future developments.

S4.1. Calculation of increases in surface area per mass

This set of calculations modeled the impact of changes in the surface area per mass ratio of the PBP-based adsorbent. Two different scenarios were explored: 1) P_i capacity limited by theoretical NHS ligand density (ligand-based) on the particle surface and 2) P_i capacity limited by the physical space taken up by the PBP (footprint-based).

S4.1.1. Ligand-based calculations

Ligand-based calculations were made based on the maximum NHS ligand density reported by the particle manufacturer (either for IOPs or Sepharose resin). This approach assumed that the number of NHS ligands (and therefore coupled PBP) was directly proportional to changes in the surface area to mass ratio (and that all other properties remained the same as the baseline IOP or NHS scenarios). The surface area to mass ratio of the particles was calculated as the surface area divided by particle volume and density, simplified as Eq. S1.

$$\left(\frac{SA}{m}\right) = \frac{3 \cdot 10^6}{r_{particle} \cdot \rho_{particle}} \quad \text{Eq. S1}$$

Where (SA/m) = surface area to mass ratio (m² g⁻¹), r_{particle} = radius of the particle (μm), ρ_{particle} = density of the particle (5.24x10⁶ g m⁻³ for IOP, 0.7x10⁶ g m⁻³ for Sepharose resin) (g m⁻³), and 10⁶ accounts for unit conversions between μmol and mol.

The theoretical maximum number of PBPs on the particle was calculated using Eq. S2, assuming proportional NHS density to particle surface area and 1:1 molar PBP binding to NHS.

$$q_{PBP} = \frac{MW_{PBP} \cdot \rho_{NHS}}{10^6} \left(\frac{\left(\frac{SA}{m}\right)}{\left(\frac{SA}{m}\right)_{baseline}} \right) \quad \text{Eq. S2}$$

Where q_{PBP} = capacity of PBP (g PBP (g particle)⁻¹), MW_{PBP} = molecular weight of the PBP = 35,000 g PBP (mol PBP)⁻¹, ρ_{NHS} = maximum NHS ligand density reported by the particle manufacturer = 250 μmol NHS (g IOP)⁻¹ or 32.8 μmol NHS (g Sepharose bead)⁻¹, (SA/m) is the ratio of surface area to mass compared to the baseline ratio of surface area to mass for the particle (m² g⁻¹), and 10⁶ accounts for unit conversions between μmol and mol.

To convert the number of PBP on the particle to the P_i-binding capacity, Eq. S3 was used.

$$q_{PO_4} = q_{PBP} * \frac{MW_{PO_4}}{MW_{PBP}} * 10^3 \quad \text{Eq. S3}$$

Where q_{PO_4} = capacity of PO₄ (mg PO₄ (g particle)⁻¹), q_{PBP} = capacity of PBP (g PBP (g particle)⁻¹), MW_{PO_4} = 94.97 g PO₄ mol⁻¹, MW_{PBP} = molecular weight of the PBP = 35,000 g PBP (mol PBP)⁻¹, and 10³ accounts for unit conversions between mg and g.

S4.1.2. Footprint-based calculations

Eq. S4 was used to estimate the number of proteins or peptides that could physically fit on the surface of the particle assuming that the spherical particle's surface was completely filled with peptides (neglecting the impact of curvature).

$$N_{PBP} = \frac{SA_{particle}}{A_{X,PBP}} = \frac{4\pi r_{particle}^2}{\pi r_{PBP}^2} \quad \text{Eq. S4}$$

Where N_{PBP} = number of PBP on the particle surface, $SA_{particle}$ = surface area of the particle (μm²), $A_{X,PBP}$ = cross-sectional area of PBP protein (μm²), $r_{particle}$ = radius of the particle (μm), r_{PBP} = radius of PBP protein (3.45x10⁻³ μm).

To convert the number of peptides on the particle surface to the P_i-binding capacity (assuming 1:1 molar peptide binding to NHS), Eq. S5 was used.

$$q_{PO_4} = \frac{N_{PBP} * MW_{PO_4} * 10^3}{6.02 * 10^{23} \frac{PBP}{mol} * \left(\frac{4}{3} \pi r_{particle}^3\right) * \rho_{particle}} \quad \text{Eq. S5}$$

Where q_{PO_4} = capacity of PO₄ (mg PO₄ (g particle)⁻¹), N_{PBP} = number of PBP on the particle surface, MW_{PO_4} = 94.97 g PO₄ mol⁻¹, $r_{particle}$ = radius of the particle (m), $\rho_{particle}$ = density of the particle (5.24x10⁶ g m⁻³ for IOP, 0.7x10⁶ g m⁻³ for Sepharose resin) (g m⁻³), and 10³ accounts for unit conversions between mg and g.

S4.2. Calculation of the effects of PBP size reduction

The PBP size reduction scenario modeled the impact of substituting a PO₄-selective peptide sequence for the full PBP. This approach assumed that peptides (ranging in size from PBP's PO₄-binding site diameter, 4 Å, up to the size of the PBP at 69 Å) 1) retained 1:1 PO₄-binding and 2) the mass of the peptide binding sequence decreased proportionally to the change in size (radius).

Two different scenarios were explored: 1) P_i capacity limited by theoretical NHS ligand density (ligand-based) on the particle surface and 2) P_i capacity limited by the physical space taken up by the protein/peptide binding sequence (footprint-based).

S4.2.1. Ligand-based calculations

Ligand-based calculations were made based on the maximum NHS ligand density reported by the particle manufacturer (either for IOPs or Sepharose resin). The theoretical maximum number of PBP on the particle was calculated using Eq. S6, assuming 1:1 molar PBP binding to NHS.

$$q_{PBP} = \frac{MW_{PBP} * \rho_{NHS}}{10^6} \quad \text{Eq. S6}$$

Where q_{PBP} = capacity of PBP (g PBP (g particle)⁻¹), MW_{PBP} = molecular weight of the PBP = 35,000 g PBP (mol PBP)⁻¹, ρ_{NHS} = maximum NHS ligand density reported by the particle manufacturer = 250 $\mu\text{mol NHS (g IOP)}^{-1}$ or 32.8 $\mu\text{mol NHS (g Sepharose bead)}^{-1}$, and 10^6 accounts for unit conversions between μmol and mol .

To convert the number of PBP on the particle to the P_i-binding capacity, Eq. S7 was used.

$$q_{PO4} = MW_{PBP} * \frac{r_{pep}}{r_{PBP}} * MW_{PO4} * 10^3 \quad \text{Eq. S7}$$

Where q_{PO4} = capacity of PO₄ (mg PO₄ (g particle)⁻¹), MW_{PBP} = molecular weight of the PBP = 35,000 g PBP (mol PBP)⁻¹, r_{pep} = radius of the peptide (nm), r_{PBP} = radius of PBP = 3.45 nm, MW_{PO4} = 94.97 g PO₄ mol⁻¹, and 10^3 accounts for unit conversions between mg and g.

S4.2.2. Footprint-based calculations

Eq. S8 was used to estimate the number of proteins or peptides that could physically fit on the surface of the particle assuming that the spherical particle's surface was completely filled with peptides (neglecting the impact of curvature).

$$N_{pep} = \frac{SA_{particle}}{A_{X,pep}} = \frac{4\pi r_{particle}^2}{\pi r_{pep}^2} \quad \text{Eq. S8}$$

Where N_{pep} = number of peptides (or PBP proteins) on the particle surface, $SA_{particle}$ = surface area of the particle (μm^2), $A_{X,pep}$ = cross-sectional area of the peptide (or PBP protein) (μm^2), $r_{particle}$ = radius of the particle (μm), r_{pep} = radius of the peptide (or PBP protein) (μm).

To convert the number of peptides on the particle surface to the P_i-binding capacity (assuming 1:1 molar peptide binding to NHS), Eq. S9 was used.

$$q_{PO4} = \frac{N_{pep} * MW_{PO4} * 10^3}{6.02 * 10^{23} \frac{pep}{mol} * \left(\frac{4}{3} \pi r_{particle}^3\right) * \rho_{particle}} \quad \text{Eq. S9}$$

Where q_{PO4} = capacity of PO₄ (mg PO₄ (g particle)⁻¹), N_{pep} = number of peptides (or PBP proteins) on the particle surface, MW_{PO4} = 94.97 g PO₄ mol⁻¹, $r_{particle}$ = radius of the particle (m), $\rho_{particle}$ = density of the particle ($5.24 \times 10^6 \text{ g m}^{-3}$ for IOP, $0.7 \times 10^6 \text{ g m}^{-3}$ for Sepharose resin) (g m^{-3}), and 10^3 accounts for unit conversions between mg and g.

S4.2.3. Capacity comparisons

To contextualize our theoretical calculations of PBP adsorbent capacity relative to P_i adsorption performance using other adsorbents reported in the literature, we compared values to those shown in Table S2. Figure 6 in the main text illustrates the comparisons.

Table S2. Literature reports of P_i adsorption capacity (reported based on Langmuir isotherm modeling).

Quartile	Adsorbent	Capacity, mg PO_4 (g bead) ⁻¹	Reference
1	RC-BOFS (0.8 – 2.3 mm)	0.4	(2)
	NFS	0.5	(3)
	HA-MNP	3	(4)
	FMS-0.1 La	6.1	(5)
	NLZ	7.7	(6)
	La-Z	8	(6)
	Fe-Al	8.2	(7)
	Hydrogel beads	11.7	(8)
	ZrO ₂ @SiO ₂ @Fe ₃ O ₄	12	(9)
	Mg-Biochar	13.2	(10)
	GO-Zr	13.2	(11)
2	MG@La	16.6	(12)
	Aluminum hydroxide	19.8	(13)
	RHB	19.9	(14)
	Bauxite	20.6	(15)
	Fe-GAC	21.8	(16)
	ACF-Zr-Fe	29.1	(17)
	ZnFeZR-adsorbent	32.2	(18)
	Diethylamine Zr-OH	38	(19)
Magnetic Fe-Zr	41.9	(20)	
3	Zr hydroxide	46.9	(21)
	am-ZrO ₂	48.5	(22)
	N-Ethylmethylamine Zr-OH	51.8	(19)
	Akaganeite	51.8	(23)
	Magnetic Fe-Zr	54.8	(20)
	Fe-Al (Mesoporous spheres)	61.5	(7)
	Akaganeite (β -FeOOH)	71.5	(23)
	Fe-Zr	76.4	(24)
	Pure-Zr-OH	79.1	(19)
Fe-La	90.5	(25)	

	CuFe ₂ O ₄ -2N-La	91.4	(26)
	Fe-Zr	102.4	(24)
	Fe-Cu	107.9	(27)
	Dimethylamine Zr-OH	110	(19)
4	Fe-Mn	110.4	(28)
	Fe-Cu	122.0	(27)
	Fe ₃ O ₄ @ASC	133	(29)
	MOD	149	(30)
	Al-Mn	183.4	(31)
	l-Y(OH) ₃	244	(32)

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