

Evaluating Self-buffering Ionic Liquids for Biotechnological Applications

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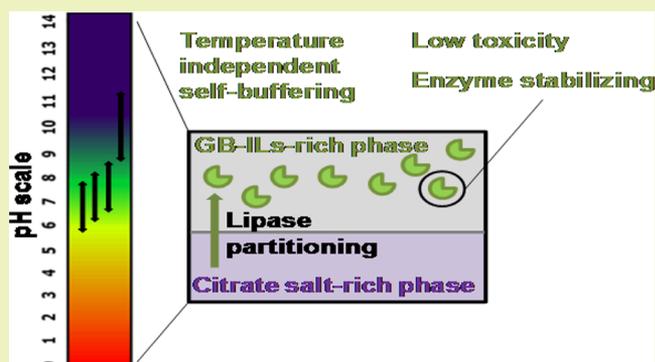
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S Supporting Information

ABSTRACT: A new range of Good's buffer ionic liquids (GB-ILs), displaying simultaneously the properties of ionic liquids and Good's buffers, were synthesized by a combination of Good's buffer anions (MOPSO, BES, TAPSO, and CAPSO) and tetrabutylammonium, tetrabutylphosphonium, and cholinium cations via an acid–base neutralization reaction. The activity and stability of a lipolytic enzyme from *Pseudomonas cepacia* in aqueous solutions of these buffers were evaluated, and the results show their advantage as media for enzymatic reactions when compared to conventional phosphate buffers. Moreover aqueous biphasic systems (ABS) composed by these GB-ILs and potassium citrate were investigated and shown to be highly effective and selective for the partitioning of the lipolytic enzyme into the GB-IL-rich phase. The results allow for the development of an efficient and biocompatible process combining the self-buffering and enzyme-stabilizing properties of the GB-ILs in the reaction step, with the advantages of GB-ILs as extraction solvents in ABS.

KEYWORDS: Good's buffer ionic liquids, Self-buffering GB-ILs, Aqueous biphasic systems, Enzyme-stabilizing systems, Biocatalysis and bioseparation



INTRODUCTION

Ionic liquids (ILs) have been proposed as environmentally benign solvents¹ to replace harmful volatile organic compounds due to their unique properties,² thus enhancing their recycling ability and product recovery, and improving the safety of (bio)chemical processes.³ The interest in biotechnological applications of ILs is growing not only for enzyme catalysis^{4–8} but also for bioseparation process.^{9,10} Their potential as media for biocatalytic reactions with remarkable yields¹¹ has been demonstrated, in addition to their enantioselectivity¹² and enzyme stability.¹³ In bioseparations, water miscible-ILs hold great potential as a feasible alternative for phase forming agents in aqueous biphasic systems (ABS) to replace conventional polymer and/or salt-rich phases,¹⁴ for the purification of various enzymes including alcohol dehydrogenases,¹⁵ lysozymes,¹⁶ cyclodextrin glycosyltransferases,¹⁷ peroxidases,¹⁸ and lipases.^{19–22}

The need to maintain a stable pH during an enzyme-catalyzed reaction is well established.^{23,24} Therefore, most applications of ILs in the biotechnological field involve the addition of buffers to stabilize hydrogen ion concentrations during enzymatic reaction. However, this is not adequate to maintain a precise pH control in IL systems as the intrinsic

acidity or basicity of hydrophilic ILs might destroy the buffer effect,²⁵ depending on the IL concentrations applied. Additionally, ionic liquid-based ABS (IL-based ABS) for enzyme extraction are always limited by the selection of a buffer in order to prevent the loss of activity during the extraction process. The phosphate buffer is most commonly used,²⁶ and the use of organic salts, such as citrates, is seldom considered when dealing with labile enzymes as these systems create a pH of 8 to 10 and lack buffering capacity to maintain the optimum pH during the extraction process,²⁷ thus limiting ABS formation²⁸ and the partitioning behavior of enzymes.¹⁶ It would, therefore, be important to find alternative approaches.

In this work, the design of ILs with self-buffering and enzyme-compatible behavior is addressed. The use of self-buffering ILs in IL-based ABS offers further advantages compared to those of traditional buffers, namely, the possibility of controlling salt precipitation/crystallization, since most ILs present low melting temperatures, exist in liquid state at room temperature, and are miscible with water in the entire range of

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concentration, allowing the tailoring of the polarity of both phases due to the wide array of IL structures available.²⁹ After the first report of the buffer-like ILs,³⁰ recently, some works reported the synthesis of ILs with self-buffering properties^{31–33} through the selection of anions derived from biological buffers, namely, Good's buffers (GBs). Besides the buffering ability, these GB anions³⁴ also conferred high biocompatibility to the synthesized ILs. The GB anions tricine, TES, CHES, HEPES, and MES have been paired with alkylimidazolium,³² tetraalkylammonium,³² and cholinium cations,^{31,33} and these new ILs demonstrated self-buffering behavior, low toxicity, and the ability to form ABS with inorganic and organic salts, thus revealing high protein stability. Yet, there is no literature available concerning these ILs in enzymatic systems.

In the present study, the synthesis of ILs based on GB anions with pK_a values closer to the optimum pH of most enzymes was achieved. The anions chosen are MOPSO, BES, TAPSO, and CAPSO with pK_a values of 6.90, 7.09, 7.60, and 9.60, respectively, at 25 °C and 0.1 M. The selected anions were coupled with cations belonging to the tetrabutylammonium, tetrabutylphosphonium, and cholinium IL families in order to produce 12 Good's buffer ILs (GB-ILs). These cations were selected due to their characteristics reported in literature.^{35–37} To further assess the GB-ILs produced, the GB-IL buffering capacity was validated and compared to that of the respective GBs. The toxicity of each GB and GB-IL toward *Vibrio fischeri* marine bacteria was also characterized. The viability of GB-ILs as a benign medium to the enzymes was also investigated by measuring the activity of a lipase from *Pseudomonas cepacia* (recently reclassified as *Burkholderia cepacia*) in 0.05 and 1.0 M aqueous solution of GB-ILs at pH 7.0, after 4 h of contact, and comparing with the phosphate buffer. Finally, the ability of the synthesized GB-ILs to form ABS with a biodegradable citrate-based salt was investigated, and the partitioning behavior of the lipase in the selected GB-IL-based ABS was studied, aiming at evaluating the applicability of these GB-ILs in bioseparations.

EXPERIMENTAL SECTION

Materials. Tetrabutylammonium hydroxide solution (40 wt % in H₂O), tetrabutylphosphonium hydroxide solution (40 wt % in H₂O), choline hydroxide solution (46 wt % in H₂O), 2-hydroxy-3-morpholinopropanesulfonic acid (MOPSO, purity ≥99%), 2-[bis(2-hydroxyethyl)amino]ethanesulfonic acid (BES, purity ≥99%), N-[tris(hydroxymethyl)methyl]-3-amino-2-hydroxypropanesulfonic acid (TAPSO, purity ≥99%), 3-(cyclohexylamino)-2-hydroxy-1-propanesulfonic acid (CAPSO, purity ≥99%), potassium citrate tribasic monohydrate (C₆H₅K₃O₇·H₂O, analytical grade), potassium phosphate monobasic (KH₂PO₄, analytical grade), potassium phosphate dibasic (K₂HPO₄, analytical grade), 4-nitrophenyl laurate (*p*-NFL, purity ≥98.0%), 4-nitrophenol (*p*-NP, spectrophotometric grade), brilliant blue G-250, and the protein standard of bovine serum albumin (BSA) were supplied by Sigma-Aldrich. The enzyme used throughout this work was a lipase from *P. cepacia* (*Burkholderia cepacia*) powder purchased from Sigma-Aldrich. It should be highlighted that in this work *P. cepacia* is used, despite the fact that this microorganism was reclassified as *Burkholderia cepacia*, to maintain the commercial designation from Sigma-Aldrich, and thus the lipase name will be described throughout this work in accordance with the trade name, in order to allow the cross-reference of this commercial lipase. Methanol (HPLC grade, purity ≥99.9%), acetonitrile (HPLC grade, purity ≥99.9%), dimethyl sulfoxide (DMSO), hydrochloric acid (HCl), ethanol, acetic acid, and orthophosphoric acid were acquired from Fisher Scientific, and sodium hydroxide (NaOH) was purchased from Merck. Ultrapure water treated by a Milli-Q integral water purification system was used throughout the work. All of the stock solutions and

buffers for polyacrylamide gel electrophoresis and prestained protein standards (broad range) were obtained from Bio-Rad, and Proteosilver silver stain kits were purchased from Sigma-Aldrich.

Methods. Synthesis of GB-ILs. GB-ILs were synthesized by an acid–base neutralization reaction of hydroxide solution of cations and GBs as detailed in Section S1 from Supporting Information.

GB-ILs' Buffering Capacity. The pH titration profiles were established using a potentiometric titrator (Metrohm, model 904 Titrando) controlled using the Tiamo software, version 2.3, and equipped with a 801 stirrer with a stand and 805 dosimat with an exchange unit of 20 mL, a pH glass electrode 6.0262.100a, and a temperature sensor 6.1114.010. The pH electrode was previously calibrated with aqueous solutions of standard buffers of pH 4.0 and 7.0. The acid–base titration was performed with 10 mL of an aqueous solution of each GB-IL (0.05 M) in a double-walled glass vessel at two different temperatures, 25 and 37 °C, which were controlled using a thermostatic water bath. The titrants used were 0.05 M NaOH and 0.05 M HCl. The potentiometric titration measurements were performed in duplicate.

Ecotoxicity Analysis of GB-ILs. The GB-IL and GB ecotoxicity was investigated using the standard Microtox liquid-phase assay, which is based on the evaluation of the luminescence response of a marine bacterium, *Vibrio fischeri* strain NRRL B-11177, after exposure to each compound at 15 °C, as described elsewhere.³⁸ The decrease in the toxicant concentration and the light output of the bacteria produced a dose/response relationship, and the results were computed using the Microtox Omni Software version 4.1 to calculate the effective concentration associated with a 50% reduction in the luminescence from the marine bacteria *V. fischeri*, EC₅₀, at the exposure time of 5, 15, and 30 min for all of the GB-ILs and GBs.

Enzyme Activity of Lipase in GB-ILs. The lipolytic enzyme from *P. cepacia* was incubated in the aqueous solutions of phosphate buffers (K₂HPO₄/KH₂PO₄ mixture), GBs, and GB-ILs (pH 7.0, at both concentrations of 0.05 and 1.0 M) at 100 rpm at 25 ± 1 °C for 4 h, and the residual enzyme activity was assayed. The experiments were performed in triplicate for each compound. The results are presented in relative enzyme activity, as described in eq 1.

$$\text{Relative enzyme activity} = \frac{EA_{\text{GB-ILs/GBs}(0.05\text{M}/1.0\text{M})}}{EA_{\text{phosphate buffer}(0.05\text{M}/1.0\text{M})}} \quad (1)$$

Potential of GB-ILs in ABS Formation. The ability of GB-ILs to form ABS with potassium citrate (K₃C₆H₅O₇) was investigated and the experimental binodal curves determined using the cloud point titration method at 25 ± 1 °C and atmospheric pressure. The experimental procedure of determining ABS with ILs was adopted from other works performed and validated by us,^{26,39,40} using an analytical balance (Mettler Toledo, model AB204-S/FACT). All of the calculations considering the mass fractions or molarity of the citrate-based salt were carried out discounting the complexed water presented in the commercial citrate-based salt. The experimental coexisting curves were fitted to eq 2:

$$[\text{IL}] = A \exp[(B \times [\text{K}_3\text{C}_6\text{H}_5\text{O}_7]^{0.5}) - (C \times [\text{K}_3\text{C}_6\text{H}_5\text{O}_7]^3)] \quad (2)$$

where [IL] and [K₃C₆H₅O₇] are the weight fraction percentages of IL and potassium citrate, and A, B, and C are constants obtained by the regression of the experimental binodal curves. The tie-lines (TLs) and respective tie-line lengths (TLL) of several selected ternary phase diagrams were determined as described elsewhere.^{26,27,40}

Partitioning Behavior of Lipolytic Enzyme in GB-IL-Based ABS. The total mass of GB-IL-based ABS was 5.0 g, and 0.1 wt % of commercial lipase from *P. cepacia* was added. The mixture pH was adjusted to 7.0 and measured at 25 ± 1 °C using a pH meter (Fisher scientific, model accumet AB15 basic). The mixture was stirred in a vortex until all of the components were dissolved. Then, the enzymatic solutions were centrifuged at 2500 rpm for 15 min by a centrifuge (Eppendorf, model 5810 R), and placed in an incubator at 25 °C for at least 12 h to reach equilibrium. After this period, both phases were carefully separated, and their volume and weight were measured and

prepared for enzyme activity assay and the protein assay. Triplicate independent runs were performed for each partitioning test, and the results were expressed as the average of the three assays.

The partitioning behavior of the lipolytic enzyme was studied by determining the partition coefficient of the enzyme, K_E , and total proteins, K_P , as defined by eqs 3 and 4, and additionally, the enzyme recovery, R_T^E , and protein recovery, R_T^P , in the top GB-IL-rich phase, as well as the selectivity parameter, $S_{E/P}$, were determined, as described by eqs 5 to 7).

$$K_E = \frac{EA_T}{EA_B} \quad (3)$$

$$K_P = \frac{C_T}{C_B} \quad (4)$$

$$R_T^E = \frac{100}{1 + \frac{1}{R_V K_E}} \quad (5)$$

$$R_T^P = \frac{100}{1 + \frac{1}{R_V K_P}} \quad (6)$$

$$S_{E/P} = \frac{K_E}{K_P} \quad (7)$$

where EA_T and EA_B are the enzyme activity at the top and bottom phases, respectively, C_T and C_B are the total protein concentration at the top and bottom phases, respectively, and R_V is the volume ratio of the top and bottom phases.

Enzyme Activity Assay. The lipase activity was spectrophotometrically assayed using a UV-vis spectrophotometer (Shimadzu, model UV-1800), as described in our earlier work.⁴¹ The substrate solution was prepared by dissolving 0.018 g of *p*-NFL in 1 mL of DMSO and diluted 100 times in a phosphate buffer solution $\text{KH}_2\text{PO}_4/\text{K}_2\text{HPO}_4$ (50 mM and pH 7.0). The substrate solution was incubated at 37 °C for 30 min. Then, 200 μL of each sample was added to 2.8 mL of substrate solution, and the linear range of absorbance variation at 410 nm during 150 s ($\Delta\text{Abs}\cdot\text{min}^{-1}$) was analyzed. The enzyme activity is expressed in $\text{U}\cdot\text{L}^{-1}$, one unit (U) of enzyme activity defined as the amount of enzyme that produces 1 μmol of *p*-NP per minute under standard assay conditions.

Protein Assay. The protein concentration was determined by the Bradford's method,⁴² using a UV-vis spectrophotometer (Shimadzu, UV-1800) at 595 nm and a standard calibration curve established using the standard protein BSA. To eliminate the influence of the IL presence on the protein concentration analysis, a blank control system for each partition system without enzyme was prepared under the same conditions used as the reference for measurement.

Sodium Dodecyl Sulfate–Polyacrylamide Gel Electrophoresis (SDS–PAGE) Analysis. The SDS–PAGE analysis was performed using hand cast polyacrylamide gel made of a 12% revolving gel and a 4% stacking gel, thus added to a vertical gel electrophoresis unit (Bio-Rad, model mini-PROTEAN tetra cell), as described by Laemmli,⁴³ at 160 V for 1 h. After the electrophoresis process, the gel was stained following the direct silver staining procedure.

RESULTS AND DISCUSSION

Synthesis and Characterization of GB-ILs. The chemical structures of the synthesized GB-ILs depicted in Figure S1 in Supporting Information, were confirmed by NMR analysis as reported in Table S1. Unlike the previous work,³² all of the GB-ILs in this study are liquids at room temperature, except for $[\text{N}_{4,4,4,4}][\text{TAPSO}]$, $[\text{P}_{4,4,4,4}][\text{TAPSO}]$, and $[\text{Ch}][\text{CAPSO}]$.

All GB-ILs studied are completely miscible with water. The potential of aqueous solutions of GB-ILs to act as buffer media was investigated by determining their pH profiles, which correspond to the titration of acids and bases, at 25 and 37 °C. All of the buffering property data of GB-ILs and GBs are

presented in Table S2. As depicted in Figure 1a, GB-ILs with the same anionic species have identical buffering capacity at

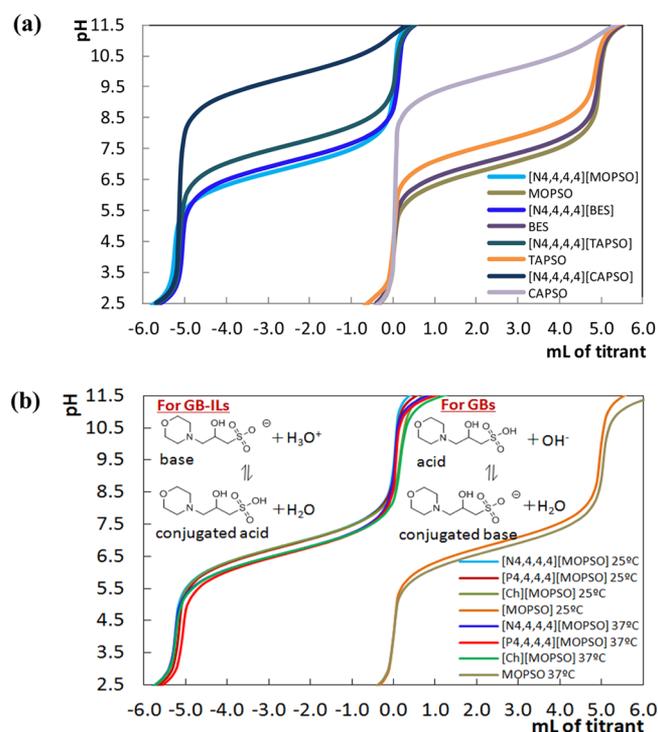


Figure 1. The pH acid-based titration profiles of (a) tetrabutylammonium-based GB-ILs and the GBs at 25 °C; (b) GB-ILs based on the MOPSO anion and the respective GB, MOPSO, at 25 and 37 °C. The titrant volumes from 0 to 6 mL are for the titration of the 0.05 M GB-ILs/GBs aqueous solutions with 0.05 M NaOH, whereas the -1 mL entries correspond to the volumes of 0.05 M HCl added to the aqueous solution of GB-ILs/GBs (reverse titration).

every single acid/base titration points as the respective GB, including their buffer region, midpoint of buffer region, and buffer capacity.

One of the important characteristics of the GBs is the temperature stability of their buffering capacity. This temperature independent character of the buffering ability is also displayed by GB-ILs, for which there is only a slight change in their buffering behavior between 25 and 37 °C, as depicted in Figure 1b for the GB-ILs based on the MOPSO anion. This GB-IL behavior shows their potential as buffer media to be used at the optimum temperature of a specific enzyme.

The (eco)toxicity of each GB-IL for 5, 15, and 30 min of exposure was assessed using the Microtox bioassay and further compared with the respective GB toxicity (EC_{50} data in units of $\text{mg}\cdot\text{L}^{-1}$ and $\text{mmol}\cdot\text{L}^{-1}$, respectively, in Tables S3 and S4). In general, GBs demonstrate a toxicity decrease with the increase of exposure time from 5 to 30 min, which is explained by the marine bacteria's adaptation to the solution. An inverse trend of toxicity–exposure time relationship was observed for tetrabutylammonium- and tetrabutylphosphonium-based GB-ILs, exhibiting at the same time, an increase of toxicity with exposure time, which is justified by the fact that longer time periods of contact are essential for the toxicity mechanism to take place.^{44,45} However, cholinium-based GB-ILs show a nonlinear relationship of toxicity–exposure time, and this complexity might be explained by the different trends found for

this family⁴⁴ and also probably due to the toxicity mechanism in which both cationic and anionic parts participate.

To carry the analysis at the maximum effect of the GB-ILs toward bacteria, the discussion will be based on 30 min of exposure time (Figure 2). Taking into account the 30 min-EC₅₀

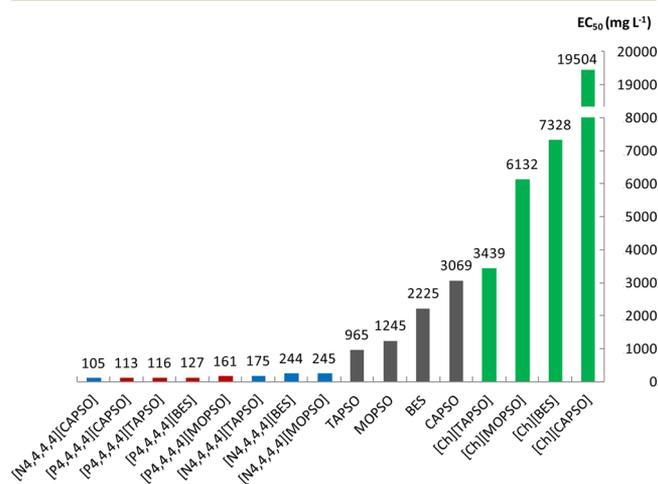


Figure 2. EC₅₀ experimental values toward the bioluminescent marine bacteria, *Vibrio fischeri*, at 30 min of exposure time to cholinium-based GB-ILs (green bars), tetrabutylammonium-based GB-ILs (blue bars), tetrabutylphosphonium-based GB-ILs (red bars), and GBs (gray bars).

> 100 mg·L⁻¹ obtained, all GB-ILs studied can be classified as “practically harmless”.⁴⁴ The results here reported suggest that the cation has a dominant impact on GB-IL ecotoxicity, following an increasing tendency of [Ch]⁺ < [N_{4,4,4,4}]⁺ < [P_{4,4,4,4}]⁺. It is noteworthy to mention that the incorporation of the cholinium into the structure of GBs contributes, in this case, to drastically reduce the toxicity of the salts while simultaneously magnifying the effect of the anions, following the identical trend of increasing ecotoxicity exhibited by their respective GBs in the order: [CAPSO]⁻ < [BES]⁻ < [MOPSO]⁻ < [TAPSO]⁻. These results are consistent with the model prediction conducted by Couling et al.⁴⁶ where IL toxicity decreases as the number of negatively charged atoms in the cation increases and further strengthened by the experimental work stating that the cholinium cation (with negatively charged oxygen atoms) is relatively nontoxic. In addition, several works also reported that the toxicity increases with the elongation of the cation’s alkyl side chain,^{47,48} and this is supported by the findings that hydrophobic compounds will attach or penetrate into membranes of aquatic organisms, resulting in the disruption of their metabolic activity.⁴⁹ Besides, the results suggest that the phosphonium salts were shown to have a more deleterious effect on the bacterial population compared to that of the ammonium counterpart, which is in good agreement with our earlier work.⁴⁸

In contrast to the least toxic cholinium-based GB-ILs, the variation of the anionic species has a minimal effect on the toxicity of the GB-ILs comprising [N_{4,4,4,4}]⁺ and [P_{4,4,4,4}]⁺ cations, which suggests that, for these particular GB-ILs, the toxicity is largely driven by the cation instead of the anion. In addition, those GB-ILs have different toxicity mechanisms since they do not follow the same ecotoxicity sequence of the respective GBs, described by the following trend: [MOPSO]⁻ < [BES]⁻ < [TAPSO]⁻ < [CAPSO]⁻. Likewise, when compared to those reported earlier,³² for the GB-ILs with the same cation,

the presence of the β-hydroxyl group in the anionic species of these GB-ILs in particular [MOPSO]⁻, [TAPSO]⁻, and [CAPSO]⁻ contributes to a higher toxicity, although this functional group exerts a decreasing effect on the GBs’ toxicity.

Lipase Enzyme Activity in GB-ILs. In the applications dealing with labile enzymes, it is important that the enzyme–solvent system is capable of establishing balanced interactions allowing the enzyme to retain its function and conformational structure.⁵⁰ Phosphate buffer is commonly used in enzyme applications to keep the system’s pH constant, preventing the alteration/disruption of the enzyme’s surface properties. Moreover, the selection of the salting-out agent in IL-based ABS is limited by its effect on pH, and thus, often a phosphate buffer aiming at controlling the pH value of the coexisting aqueous phases is used.^{19,51} Therefore, in order to examine the viability of self-buffering GB-ILs as media for enzyme applications, the enzyme activity and stability parameters were investigated regarding the contact effect of 0.05 and 1.0 M of water–GB-ILs mixtures (pH 7.0) after 4 h of exposure, and the results were expressed in relative enzyme activity. Phosphate buffer in the same concentrations is used as control. The effects in water–GB mixtures to enzyme were also studied. In this work, a lipase from *P. cepacia* was used as a model enzyme. The class of lipases is widely recognized as a key enzyme as biocatalyst in biotechnological applications due to its multiplicity of catalytic applications.^{41,52,53}

As depicted in Figure 3a, the lipase exhibits a higher activity after 4 h of exposure to some of the 0.05 M GB-IL solutions, in particular GB-ILs based on the BES anion and [Ch][MOPSO], while a slightly lower relative activity was observed for the remaining GB-ILs, when compared with the results attained with the phosphate buffer. Besides, the lipase displays a lower relative activity after being in contact for 4 h in both 0.05 and 1.0 M of GBs’ aqueous solution. This behavior can be attributed to the kosmotropic/chaotropic effects,⁵⁴ in which GBs comprising the sulfonate anion are more chaotropic than the phosphate anion and thus destabilize the enzyme.⁵⁵ However, the incorporation of a bulky and hydrophobic cation into the GBs’ structure has a significant effect in improving the enzyme-stabilizing properties of the GB-ILs. This effect is even more pronounced at high concentration of GB-ILs. At 1.0 M, tetrabutylammonium and tetrabutylphosphonium families demonstrate a higher capacity to enhance the enzyme’s catalytic activity, with approximately 1.7 to 3.0-fold higher activities when compared to those of the phosphate buffer at the same concentration, as presented in Figure 3b. These results clearly reveal that the enzyme activity is correlated with the hydrophilic/hydrophobic nature of the enzyme–solvent system on the water hydration level of microaqueous phase at the enzyme surface.⁵⁶ The presence of tetrabutyl-ammonium and -phosphonium cations, comprising hydrophobic alkyl side chains, reduces the tendency to strip off the hydration water from the enzyme surface. However, cholinium-based GB-ILs with polar cations, especially at high concentrations, dehydrate the microenvironment at the enzyme surface causing enzyme deactivation. This behavior explains the lower relative activity observed for the cholinium-based GB-ILs compared to that of their counterparts. The results here reported suggest that more hydrophobic GB-ILs stabilize the enzyme, whereas hydrophilic GB-ILs act as an enzyme destabilizer as reported previously for other ILs.⁵⁷ Additionally, the experimental results ascertain that tetrabutylammonium- and tetrabutylphosphonium-based GB-ILs contribute to the preservation of the enzyme active catalytic

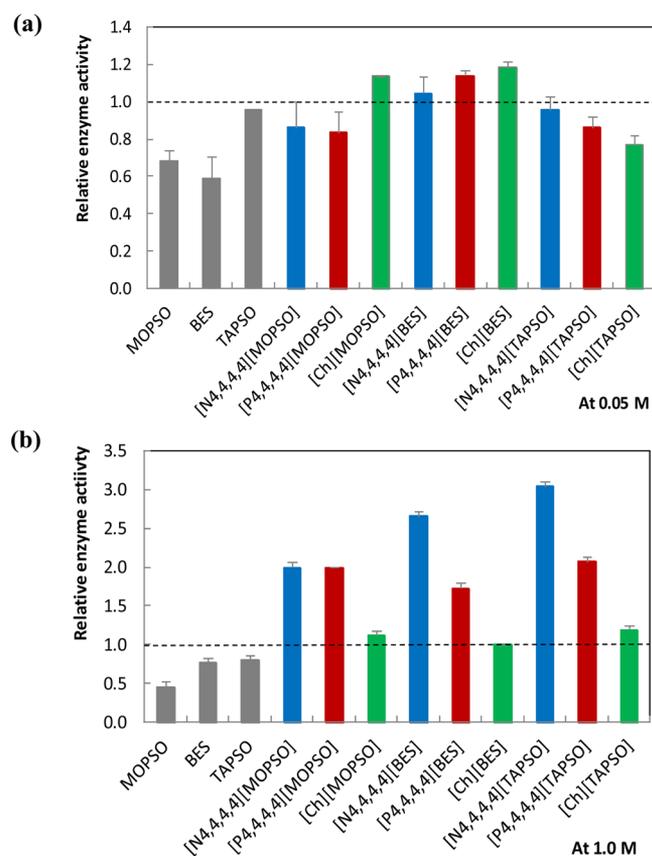


Figure 3. Relative enzyme activity of a lipase from *P. cepacia* after being in contact for 4 h with (a) 0.05 M and (b) 1.0 M of aqueous solutions of cholinium-based GB-ILs (green bars), tetrabutylammonium-based GB-ILs (blue bars), tetrabutylphosphonium-based GB-ILs (red bars), and GBs (gray bars). A phosphate buffer solution was used as the control, regarding the same conditions described for the GB's and GB-ILs samples, namely, 4 h of contact and (a) 0.05 M and (b) 1.0 M.

conformation in aqueous solution through balanced IL–enzyme interactions. Furthermore, the outstanding solvation ability normally found in ILs can enhance enzyme solubility and establish a better enzyme–medium–substrate relationship¹⁵ and thus improve the enzyme's catalytic efficiency.

GB-IL-Based ABS. The ability of GB-ILs to create ABS with biodegradable potassium citrate was investigated in order to assess the potential of the GB-ILs to create more efficient and sustainable IL-based ABS. The experimental weight fraction data for the studied systems are provided in Table S5. The experimental binodal data of these systems was adequately fitted to the mathematical model described by eq 2. The corresponding parameters and the respective standard deviations and correlations are presented in Table S6. The general results of the ABS formation depicted in Figure 4 show that, apart from the cholinium-based GB-ILs, all of the tetrabutylammonium- and tetrabutylphosphonium-based GB-ILs are able to undergo liquid–liquid demixing in the presence of potassium citrate. The GB-ILs' ability to form ABS with potassium citrate follows the rank: $[N_{4,4,4,4}][TAPSO] < [P_{4,4,4,4}][TAPSO] < [N_{4,4,4,4}][BES] < [N_{4,4,4,4}][MOPSO] < [P_{4,4,4,4}][BES] < [P_{4,4,4,4}][MOPSO] < [N_{4,4,4,4}][CAPSO] < [P_{4,4,4,4}][CAPSO]$. The ability of quaternary ammonium and phosphonium salts for ABS formation can be attributed to their higher hydrophobicity resulting from the four butyl side chains

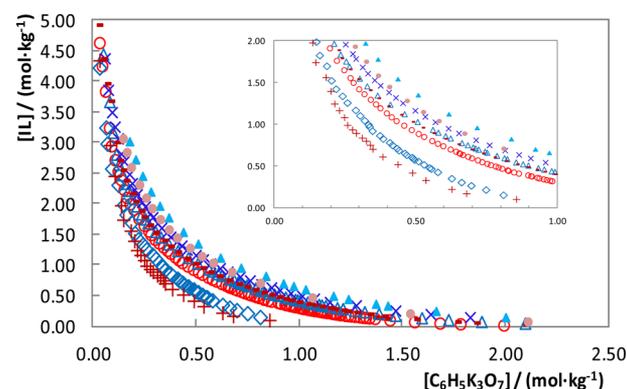


Figure 4. Experimental binodal curves of the systems composed of GB-IL + $K_3C_6H_5O_7$ + water at 25 ± 1 ($^{\circ}C$): (blue closed triangle) $[N_{4,4,4,4}][TAPSO]$, (red closed circle) $[P_{4,4,4,4}][TAPSO]$, (blue cross sign) $[N_{4,4,4,4}][BES]$, (blue open triangle) $[N_{4,4,4,4}][MOPSO]$, (red rectangle) $[P_{4,4,4,4}][BES]$, (red open circle) $[P_{4,4,4,4}][MOPSO]$, (blue open diamond) $[N_{4,4,4,4}][CAPSO]$, and (red plus sign) $[P_{4,4,4,4}][CAPSO]$.

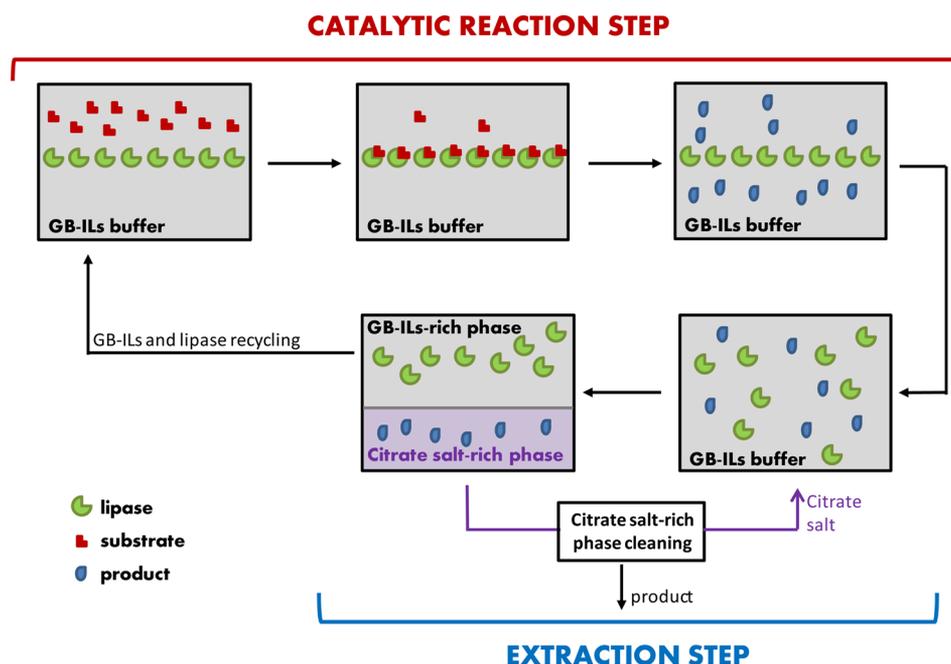
attached to the central heteroatom, leading to a low affinity for water and, consequently, displaying a higher tendency to be salted-out from the aqueous solution. Quaternary salts with phosphorus as the cation's central atom have demonstrated a higher hydrophobic nature when compared to ammonium salts⁴⁸ and thus are more effective in promoting the ABS formation.²⁶ Cholinium cations, which are smaller in size and present a polar hydroxyl group at the side chain, have a higher affinity for water which reduces their ability to induce ABS formation with citrates. Nevertheless, cholinium-based GB-ILs are capable of forming ABS with more hydrophobic phase components.^{31,33}

The influence of the IL anion's nature on ABS formation is also dominated by the system's hydrophobicity/hydrophilicity.³⁹ Therefore, the GB-IL's aptitude to induce ABS, for the same cation, follows the anions hydrophobic sequence: $[TAPSO]^- < [BES]^- < [MOPSO]^- < [CAPSO]^-$, reflected by the octanol–water partition coefficient: TAPSO ($\log P = -2.81$) < BES ($\log P = -2.52$) < MOPSO ($\log P = -2.05$) < CAPSO ($\log P = -0.36$).⁵⁸ Additionally, the effect of the anion's nature on ABS phase behavior is correlated with the relative hydrogen bond accepting strength of the IL's anion.⁵⁹ The anionic species with greater hydrogen bond accepting strength, in this case $[TAPSO]^-$, present larger ability to form stronger interactions with water and to create hydration complexes, and therefore, they are not easily salted-out (for more details, see Table S7).

Partitioning Studies of Lipase from *P. cepacia* in GB-IL-Based ABS. The applicability of GB-IL-based ABS in a purification system dealing with labile enzymes was evaluated through the determination of the partitioning of lipase from *P. cepacia* in GB-IL-based ABS. Several GB-IL-based ABS were selected considering their buffering capacity at pH 7.0. For all of the six selected GB-ILs/potassium citrate ABS, the biphasic mixture chosen for the lipase partitioning corresponds to the TLs with similar TLL, namely, between 40.23 and 42.65, in order to guarantee identical compositions of each component and similar surrounding environment conditions for the enzyme partition. The shorter tie-line was chosen as the operation biphasic mixture for the partitioning studies to minimize the difference in physical properties between the two phases and maximize the water content, providing a gentle

Table 1. Experimental Results of the Partition, Recovery, and Selectivity Parameters of Lipase from *P. cepacia* between Both Aqueous Phases of GB-IL-Based ABS and Respective Standard Deviations, σ

GB-ILs	IL (wt %)	K ₃ C ₆ H ₅ O ₇ (wt %)	TLL	R _V ± std	K _E ± std	R _T ^E ± std (%)	K _P ± std	R _T ^P ± std (%)	S _{E/P}
[N _{4,4,4,4}][MOPSO]	24.40	20.00	42.08	0.65 ± 0.02	34.0 ± 3.9	95.7 ± 0.3	25.1 ± 2.3	94.2 ± 0.7	1.35
[P _{4,4,4,4}][MOPSO]	29.68	15.00	40.23	1.58 ± 0.00	29.3 ± 1.9	97.9 ± 0.3	18.3 ± 0.6	96.6 ± 0.3	1.60
[N _{4,4,4,4}][BES]	27.00	20.00	41.44	1.05 ± 0.00	115.7 ± 6.0	99.2 ± 0.0	32.4 ± 0.9	97.1 ± 0.1	3.57
[P _{4,4,4,4}][BES]	24.00	20.00	42.65	0.68 ± 0.00	119.5 ± 7.0	98.8 ± 0.1	36.9 ± 1.6	96.2 ± 0.2	3.24
[N _{4,4,4,4}][TAPSO]	31.69	20.00	41.08	0.93 ± 0.03	82.1 ± 3.5	98.7 ± 0.1	35.1 ± 2.7	97.0 ± 0.2	2.34
[P _{4,4,4,4}][TAPSO]	29.20	20.00	41.97	0.60 ± 0.00	41.7 ± 3.0	96.1 ± 0.3	23.6 ± 2.8	93.4 ± 0.7	1.77

**Figure 5.** Conceptual process diagram of the integrated use of GB-ILs as buffers in a general catalytic reaction and as ABS formers promoting the recovery of the lipase.

environment for the enzyme. All of the ternary phase diagrams for the selected ABS composed of GB-ILs with the TLs determined and the respective operation two-phase points are presented in Figure S2. It should be highlighted that for all ABS studied, the top layer is the GB-IL-rich phase, whereas the bottom represents the salt-rich phase. As aforementioned, a commercial lipase from *P. cepacia* was applied as a model enzyme, and the respective SDS–PAGE analysis is reported in Figure S3. We can clearly see the presence of the target lipase from *P. cepacia* at ca. 33 kDa, which is consistent with the literature,⁶⁰ and several less pronounced bands representing the contaminant proteins. Since the lipase used in this study is of commercial purity, the expectable purification in the GB-IL-based ABS is limited.

Table 1 presents the biphasic mixture compositions of GB-IL/potassium citrate ABS, the respective TLL, partition coefficients, and recovery yields of both enzyme and total proteins, and the selectivity parameters. In all of the investigated systems, the lipolytic enzyme partitions preferentially to the GB-IL-rich (top) phase. The enzyme displays partition coefficients, between 29.3 ± 1.9 and 119.5 ± 7.0 , while the enzyme recovery yields at the GB-IL-rich phase ranges from 95.7 ± 0.3 (%) to 99.2 ± 0.0 (%). The total protein recovery yields achieved at the GB-IL-rich phase range between 93.4 ± 0.7 (%) and 97.1 ± 0.1 (%), based on the partition coefficients ranging from 18.3 ± 0.6 to 36.9 ± 1.6 .

The results for enzyme and total protein recovery yields at the GB-IL-rich (top) phase are similar since almost all of the lipolytic enzyme (protein exhibiting lipolytic activity) migrates to the top phase. As described before, the main goal of using ABS based on the GB-ILs was the evaluation of their potential as purification systems, meaning their capacity to isolate the target enzyme from the remaining proteins (here considered as the contaminants present in the system). As the protein partitioning in ABS can be described as a surface-dependent phenomenon,⁶¹ it can be proposed that the preferential partition of the lipolytic enzyme into the GB-IL-rich phase is mainly due to the hydrophobic interaction with the hydrophobic groups of GB-ILs. Furthermore, bigger proteins with negative charges on their surface will tend to have higher affinity to the IL-rich phase, mainly because of the driving force established by the interactions between negatively charged amino-acid residues and the IL's cation.¹⁵ Thus, the extent of the enzyme isolation from the contaminant proteins is here discussed by considering the S_{E/P} presented in Table 1.

The results obtained demonstrate that all of these systems, especially those based on the BES anion, in particular [N_{4,4,4,4}][BES] and [P_{4,4,4,4}][BES], are more selective to the target enzyme than to total proteins, thus showing a selective capacity to isolate the target lipase from the main contaminants. It should be highlighted that, despite the limited selectivity parameters observed in this study, these results suggest that,

with the adequate optimization of different conditions, it will be possible to increase the purification capacity of these ABS. In this context, this work proposes the development of an integrated process scheme considering the industrialization of this process, combining the self-buffering and enzyme-stabilizing properties of the GB-ILs as media of the catalytic reaction step. In this process, the advantages of GB-ILs as extraction solvents in ABS, thus allowing enzyme recycling and the product purification, are envisaged, as depicted in Figure 5. This process will allow the recovery of most of the GB-ILs since the recyclability of the top layer enriched in GB-ILs can be totally achieved, which will be decreasing the economic impact and environmental footprint of the process.⁶²

CONCLUSIONS

GB-ILs were synthesized with anions derived from Good's buffers (MOPSO, BES, TAPSO, and CAPSO) and coupled with tetrabutylammonium, tetrabutylphosphonium, and cholinium cations. It was shown that these "practically harmless" GB-ILs possess temperature independent self-buffering characteristics and exhibit an enzyme-stabilizing effect for retaining the catalytic activity of lipase from *P. cepacia*. Besides, these GB-ILs are capable of forming ABS with a biodegradable citrate salt and present a significant selectivity on the isolation of the target lipolytic enzyme from *P. cepacia* into GB-ILs ascertaining the potential of these GB-IL-based ABS as purification technologies. The ABS technique based on GB-ILs can be a feasible alternative to replace the conventional ABS using phosphate buffer for the extraction of the biomolecules in particular enzymes. The experimental data reported suggest that GB-ILs should be further explored for their application in enzyme catalysis and/or bioseparation technology, due to their great advantageous properties.

ASSOCIATED CONTENT

Supporting Information

The Supporting Information is available free of charge on the ACS Publications website at DOI: 10.1021/acssuschemeng.5b01155.

Experimental details for GB-IL synthesis and characterization, NMR analysis of GB-ILs, buffering property data of GB-ILs/GBs, EC₅₀-5,15, and 30 min data of GB-ILs/GBs, experimental weight fraction data for GB-IL-based ABS, the correlation constants by the regression of the experimental binodal data through the application of eq 2 of the studied systems, H-bond formation tendency of GB-IL anions, chemical structures, full names and acronyms of the synthesized GB-ILs, ternary phase diagrams for the selected systems for enzyme partition study, and SDS-PAGE patterns of commercial lipase from *P. cepacia* (PDF)

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Notes

The authors declare no competing financial interest.

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