

Co-immobilized Alcohol Dehydrogenase and Glucose Dehydrogenase with Resin Extraction for Continuous Production of Chiral Diaryl Alcohol

Jieyu Zhou

Jiangnan University

Yanfei Wu

Jiangnan University

Qingye Zhang

Jiangnan University

Guochao Xu

Jiangnan University

YE NI (✉ yni@jiangnan.edu.cn)

Jiangnan University <https://orcid.org/0000-0003-4887-7517>

Research Article

Keywords: flow reaction, alcohol dehydrogenase, co-immobilization, diaryl ketone, in situ recovery

Posted Date: February 17th, 2021

DOI: <https://doi.org/10.21203/rs.3.rs-238951/v1>

License:   This work is licensed under a Creative Commons Attribution 4.0 International License.

[Read Full License](#)

Version of Record: A version of this preprint was published at Applied Biochemistry and Biotechnology on April 7th, 2021. See the published version at <https://doi.org/10.1007/s12010-021-03561-5>.

1 **Co-immobilized Alcohol Dehydrogenase and Glucose Dehydrogenase with Resin**

2 **Extraction for Continuous Production of Chiral Diaryl Alcohol**

3

4 *Jieyu Zhou, Yanfei Wu, Qingye Zhang, Guochao Xu, Ye Ni**

5

6 *Key laboratory of industrial Biotechnology, Ministry of Education, School of*

7 *Biotechnology, Jiangnan University, Wuxi 214122, Jiangsu, China.*

8

9

10 **Corresponding author*

11 *Tel/Fax: +86-510-85329265*

12 *Email: yni@jiangnan.edu.cn*

13

14 **Abstract**

15 Ni²⁺-functionalized porous ceramic/agarose composite beads (Ni-NTA Cerase) can be
16 used as carrier materials to immobilize enzymes harboring a metal affinity tag. Here, a
17 6×His-tag fusion alcohol dehydrogenase Mu-S5 and glucose dehydrogenase from
18 *Bacillus megaterium* (*BmGDH*) were co-immobilized on Ni-NTA Cerase to construct
19 a packed bed reactor (PBR) for the continuous synthesis of the chiral intermediate (*S*)-
20 (4-chlorophenyl)-(pyridin-2-yl) methanol [(*S*)-CPMA]. NADPH recycling and *in situ*
21 product adsorption was achieved simultaneously by assembling a D101 macroporous
22 resin column after the PBR. Using an optimum enzyme activity ration of 2:1 (Mu-S5:
23 *BmGDH*) and hydroxypropyl- β -cyclodextrin as co-solvent, a space-time yield of 1,560
24 g/(L·d) could be achieved in the first three days at a flow rate of 5 mL/min and substrate
25 concentration of 10 mM. With simplified selective adsorption and extraction
26 procedures, (*S*)-CPMA was obtained in 84% isolated yield.

27 **Keywords:**

28 flow reaction; alcohol dehydrogenase; co-immobilization; diaryl ketone; *in situ*
29 recovery

30 **Abbreviations:**

31 ADH, alcohol dehydrogenase; GDH, glucose dehydrogenase; FDH, formate
32 dehydrogenase; CPMK, (4-chlorophenyl)(pyridine-2-yl)ketone; (*S*)-CPMA, (*S*)-(4-
33 Chlorophenyl)-(pyridin-2-yl) methanol; DNPH, 2,4-dinitrophenylhydrazine; PBR,
34 packed bed reactor; Cerase, ceramic/agarose composite beads

35

36 **Introduction**

37 In comparison to batch reactions, flow reactions are relatively new for biocatalytic
38 preparation of value-added chemicals and active pharmaceutical ingredients, such as
39 (*S*)-1-phenylethanol, (*R*)-flurbiprofen, fatty acid methyl esters etc. [1, 2]. In a flow
40 reaction, fluid containing substrates is pumped to the reactor to produce a product
41 stream [3-6]. When a flow process is applied to biocatalysis, it can be performed by
42 using immobilized cells or enzymes [7-11]. As a continuous process, flow reactions
43 have advantages of improved productivity, safety, ease of modifying scale, and are
44 especially useful for reactions with substrate/product inhibitory issues [12].
45 Additionally, downstream processing can become facile when using immobilized
46 biocatalysts [13].

47 For coupling or multienzymatic reaction systems, co-immobilization is an ideal strategy.
48 Co-immobilization could facilitate *in situ* removal of by-products, and is essential for
49 reactions that require cofactor regeneration [14, 15]. Co-immobilization of the main
50 and regeneration enzymes can also facilitate the mobility of the co-factor between two
51 enzymes. Therefore, it is critical to develop immobilization strategies favorable for
52 preserving the activity of all involved enzymes. Additionally, the distribution of the
53 enzyme on the support surface has a profound impact on the enzyme's overall
54 biocatalytic activity [16]. Guián's group has developed an elegant approach in which
55 they co-immobilize the main and recycling dehydrogenases on the same agarose-type
56 support (activated with glyoxyl groups and metal chelates) [17]. Recently, this approach
57 was adopted in a co-immobilized glycosyltransferase/sucrose synthase catalyzed

58 system for NDP-glucose recycling [18].

59 Immobilized metal affinity chromatography (IMAC) has also been regarded as a
60 potential strategy for enzyme co-immobilization. The IMAC strategy relies on
61 coordination bonds between transition metal ions (such as Ni²⁺, Cu²⁺, Zn²⁺, and Co²⁺)
62 and certain amino acids (such as His, Cys, and Trp) [19]. Co-expression of special
63 fusion peptides results in specific coordinations between metal ions and proteins, giving
64 advantages such as easy operation, target-specific purification, low activity loss, and
65 recyclable carrier materials [20-24]. His-tag is one of the most popular fusion peptide
66 for IMAC [25, 26]. In Böhmer and coworkers' study, a 6×His-tag fusion alcohol
67 dehydrogenase (ADH) and amine dehydrogenase (AmDH) were co-immobilized on
68 commercial metal-ion affinity glass beads EziG to produce chiral amine in batch
69 reactions [27]. Additionally, Heli-tag, a metal-binding site (His-Ile-His-Asn-Leu-Asp-
70 Cys-Pro-Asp-Cys) found in ATPase 439 from *Helicobacter pylori*, was identified to
71 have strong metal affinity [28, 29]. The Heli-tag was reported to exhibit higher affinity
72 than the 6×His-tag and has been successfully used for D-amino acid oxidase
73 immobilization to obtain improved operational stability [30].

74 Notably, the co-immobilization strategies discussed above have mainly been applied in
75 batch reactions. Additional aspects need to be considered when adopting co-
76 immobilized enzymes in continuous reactions, such as operational stability and
77 cofactor-recycling under flow mode. In a study by Dall'Oglio et al., chemical anchoring
78 was adopted for co-immobilization of ADH and GDH to construct a continuous flow
79 reactor. The stability of this system proved to be excellent during a 15-day operation

80 run at a flow rate of 50 $\mu\text{L}/\text{min}$ [31]. Peschke et al. constructed an all-enzyme hydrogel
81 consisting of ADH and GDH using a self-assembling strategy, in which a stable
82 bioconversion was operated for over 6 days at a flow rate of 10 $\mu\text{L}/\text{min}$ [32]. However,
83 activity loss caused by covalent anchors and insufficient structural rigidity of the all-
84 enzyme hydrogel restricted the flow rate of the continuous reaction. Moreover, a
85 continuous flow of cofactor was required in both studies. A regenerative
86 electrochemical method [33, 34] and cofactor immobilization [35] have been reported
87 for cofactor recycling in continuous flow biocatalysis; however, these studies were
88 limited by complex immobilization procedures or reusability of immobilized carriers
89 [36]. Given these limitations, we proposed a co-immobilization strategy for
90 $\text{NADP}^+/\text{NADPH}$ reutilization and regeneration with *in situ* product extraction in a
91 continuous reaction.

92 In this study, a non-ionic adsorptive resin was used for NADP^+ recycling due to its
93 specific adsorption of non-polar or low-polar organic compounds from aqueous
94 solutions. Herein, we designed an $\text{NADP}^+/\text{NADPH}$ regeneration strategy for a
95 continuous flow reaction catalyzed by co-immobilized ADH/GDH. The asymmetric
96 reduction of (4-chlorophenyl)(pyridine-2-yl)ketone (CPMK) catalyzed by Mu-S5 (a
97 mutant of *KpADH* from *Kluyveromyces polyspora*) [37, 38] was investigated as the
98 model reaction. The chiral product (*S*)-(4-Chlorophenyl)-(pyridin-2-yl) methanol [(*S*)-
99 CPMA] is an important intermediate for synthesizing the antiallergy drug bepotastine
100 [38, 39]. *BmGDH* from *Bacillus megaterium* was used for NADPH regeneration. The
101 6 \times His-tagged Mu-S5 and *BmGDH* were co-immobilized on nickel-nitrilotriacetic acid-

102 functionalized porous ceramic/agarose composite beads (Ni-NTA Cerase) to construct
103 a packed bed reactor (PBR), which was followed by a D101 macroporous resin column.
104 The feasibility of the proposed flow reaction process was validated by continuous
105 synthesis and *in situ* recovery of (*S*)-CPMA.

106

107 **Materials and Methods**

108 **Microorganisms and chemicals**

109 Recombinant *E. coli BL21* (DE3) strains harboring pET28-Mu-S5 and pET28-*BmGDH*
110 with *N*-terminal 6×His-tag were constructed in a previous study [38]. Ni-NTA Cerase
111 was purchased from Qianchun Bio (Yancheng, China). Macroporous resins of D101
112 and XDA-1 and chelating resin IRC-748 were purchased from Lebiochem, Co. Ltd.
113 (Xian, China). (4-Chlorophenyl)(pyridin-2-yl)methanone (CPMK) and all other
114 reagents and solvents of analytical grade and biochemical reagents were obtained from
115 Sinopharm Chemical Reagent Co. Ltd. (Shanghai, China).

116

117 **Construction of N-terminal Heli-tagged Mu-S5**

118 Plasmid pET28-Helitag-Mu-S5 was constructed by replacing the *N*-terminal 6×His-tag
119 sequence in pET28-Mu-S5. Plasmid pET28-Mu-S5 was used as the template, 6×His-
120 tag at *N*-terminal was replaced with Heli-tag by whole-plasmid PCR using KOD-Plus-
121 Neo DNA polymerase purchased from Toyobo (Osaka, Japan). The following upstream
122 and downstream primers were used:

123 Heli-tag-F:

124 CATATTCATAATCTTGATTGTCCTGATTGTAGCAGCGGCCTGGTG

125 Heli-tag-R:

126 ACAATCAGGACAATCAAGATTATGAATATGGCTGCTGCCCATGGT

127 The PCR protocol was as follows: 95°C for 2 min, followed by 15 cycles of 94°C for
128 30 s, 55°C for 30 s, and 68°C for 2.5 min. After *Dpn* I digestion at 37°C for 30 min, the
129 PCR products were transformed into competent *E. coli* BL21 (DE3) cells and plated
130 onto Luria-Bertani agar plates supplemented with 50 µg/mL kanamycin.

131

132 **Screening of supporting materials for enzyme immobilization**

133 *Preparation of the NTA Cerase column*

134 Each column ($\Phi=72$ mm, $h=24$ mm) was packed with 1 mL of NTA Cerase and washed
135 with 10 mL of deionized water followed by a solution of NiSO₄ or MnCl₂ (10 mL, 100
136 mM). The columns were then washed with 10 mL of deionized water to remove the
137 unbound metal ions, and equilibrated with 100 mM potassium phosphate buffer (pH
138 7.5). A ten-fold scale-up was used for the 10 mL Ni-NTA column.

139 *Preparation of reaction solution*

140 Solution A: For 1 mL PBR, a mixture solution containing 20% ethanol (v: v), 2 mM
141 CPMK, 2 mM glucose and 2 mM NADP⁺ was prepared with 100 mM sodium
142 phosphate buffer at pH 7.0. The mixture was warmed to 30°C before use.

143 Solution B: For 10 mL PBR (after optimization); a mixture solution containing 20 mM
144 hydroxypropyl-beta-cyclodextrin (HP- β -CD), 10 mM glucose and 10 mM NADP⁺ was
145 prepared with 100 mM Tris-HCl buffer at pH 7.5. The mixture was incubated at 30°C

146 for 1 h, and then CPMK (1M in ethanol) was added to a final concentration of 10 mM.

147 *Determination of the amount of immobilized enzyme and dilution rate*

148 Mn-NTA and Ni-NTA Cereose columns (1-mL) were prepared as described above. First,
149 Mu-S5 was purified by Ni affinity chromatography and desalinated by ultrafiltration
150 centrifugation. The resultant purified Mu-S5 was obtained with specific activity of 6.64
151 U/mg (**Table S1**), and then diluted to around 10 U/mL (the actual value was determined
152 to be 11.7 U/mL).

153 The amount of immobilized enzyme was measured by calculating the difference in
154 enzyme activity between inflows and outflows. It is presumed that saturation adsorption
155 was reached when total activity in outflows was 10% of the initial activity. The enzyme
156 activities of inflows and outflows were measured following assay method described in
157 analysis methods. The total activity of the immobilized enzyme was equal to the
158 difference in total activity between inflows and outflows, and the amount of
159 immobilized enzyme could be calculated by dividing by the specific activity of Mu-S5
160 (6.64 U/mg). The detailed data can be found in supplemental data (**Table S2**).

161 Dilution rate: According to the optimized results, the enzyme loading was 20 mg
162 protein per mL of Ni-NTA Cereose (eg. 16.7 mg protein/g Ni-NTA Cereose). Thus, 14.11
163 mg of purified Mu-S5 (94 U) and 5.89 mg of purified *BmGDH* (47 U) were needed to
164 construct a 1-mL PBR based on Ni-NTA/Mn-NTA Cereose. For operational
165 convenience, 10 mg/mL solutions of both Mu-S5 and *BmGDH* were prepared,
166 corresponding to 1.41 mL and 0.596 mL loading volume, respectively. Then, a
167 peristaltic pump was used to pump the reaction solution through the reactor at flow

168 rates of 0.2, 0.4, 0.6, 0.8, and 1.0 mL/min. For a 10-mL Ni-NTA PBR, 141.1 mg (940
169 U) of Mu-S5 and 58.9 mg (470 U) of *BmGDH* were mixed and loaded, and the flow
170 rates were set at 2.0, 4.0, 6.0, 8.0, and 10 mL/min. All of the post-column effluents were
171 collected separately and detected by HPLC.

$$172 \text{ Dilution rate} = \frac{\text{Volumetric flow rate (mL/min)}}{\text{Reactor volume (mL)}}$$

173

174 **Analysis methods**

175 *Enzymatic activity assay*

176 The activity of Mu-S5 and *BmGDH* was spectrophotometrically determined according
177 to the changes in absorption of NADPH or NADP⁺ at 340 nm and 30°C with a UV-Vis
178 spectrophotometer using a molar extinction coefficient of 6,220 M⁻¹cm⁻¹. For Mu-S5,
179 the assay mixture contained 1.0 mM CPMK, 1.0 mM NADPH, and a proper amount
180 enzyme in PBS buffer (100 mM, pH 7.0). For *BmGDH*, the assay mixture contained 1
181 mM D-glucose, 1 mM NADP⁺, and 10 μL enzyme in PBS buffer (100 mM, pH 7.0).
182 One unit of activity was defined as the amount of enzyme required for the depletion or
183 production of 1.0 μmol of NADPH under the aforementioned conditions.

184 *HPLC analysis*

185 The substrate conversion and *ee* value was determined using an Agilent 1100 HPLC
186 system (USA) equipped with a Chiralcel OB-H column (0.46 mm × 250 mm, 5 μm,
187 Diacel, Japan). The HPLC was performed at 254 nm using hexane:ethanol (95:5, v/v)
188 as eluent at a flow rate of 0.8 mL/min.

189 *Activity recovery of immobilized Mu-S5*

190 Since spectrophotometric determination was not applicable for immobilized Mu-S5,
191 the initial velocities of all entries were calculated using the substrate conversion from
192 the first 10 min (substrate conversion < 20%) using an HPLC assay.

193 A purified Mu-S5 solution of 10 U/mL was prepared as described above. The Mu-S5
194 solutions (40 mL) were loaded on 1 mL Ni-NTA and Mn-NTA PBR columns,
195 respectively. Enzymatic activity of inflows was performed in a 2 mL reaction mixture
196 consisting of 10 mM CPMK, 10 mM NADPH, and 250 μ L inflows in 100 mM PBS
197 (pH 7.0) at 30°C and 200 rpm for 10 min (**Fig. S1 & Table S3**).

198 The outflows were collected and the enzymatic activity was determined in a 2 mL
199 reaction mixture consisting of 10 mM CPMK, 10 mM NADPH, and 1 mL outflows in
200 100 mM PBS (pH 7.0) at 30°C and 200 rpm for 10 min (**Fig. S2 & Table S4**).

201 The activity of immobilized Mu-S5 was determined as follows. Weight (in gram) rather
202 than volume (in mL) was used to quantify immobilized Mu-S5. The total weights of 1
203 mL of immobilized enzymes (Mn-NTA and Ni-NTA Creose) were measured to be 1.23
204 and 1.20 g, respectively. Immobilized Mu-S5 was placed in a beaker containing 9 mL
205 100 mM PBS (pH 7.0) buffer. After mixing, 100 μ L homogenized solution was
206 transferred into 2.0 mL tubes followed by centrifugation. The liquid was then carefully
207 removed, and the immobilized Mu-S5 were weighed. The activity of immobilized Mu-
208 S5 was determined in a 2 mL reaction mixture consisting of 10 mM CPMK, 10 mM
209 NADPH, and about 0.01g of immobilized Mu-S5 in 100 mM PBS (pH 7.0) at 30°C and
210 200 rpm for 10 min (**Fig. S3 & Table S5**). The data are summarized in **Table S6**.

211 Activity recovery = $\frac{\text{Activity of immobilized enzyme}}{\text{Inflows activity-Outflows activity}} \times 100\%$

212 ***2,4-Dinitrophenylhydrazine (DNPH) assay***

213 DNPH (2,4-dinitrophenylhydrazine) is a reagent used to detect the carbonyl of ketone
214 or aldehyde functional groups. In this study, the formation of red
215 dinitrophenylhydrazone with CPMK and DNPH substrates can be regarded as a
216 positive result. First, DNPH was dissolved at 20 mM in ethanol containing 3% sulfuric
217 acid and stored in darkness. Then, the chromogenic reaction was performed by addition
218 of 100 μ L of the DNPH solution into 100 μ L of effluent (after packed bed reactor).
219 After standing for 15 min at 30°C, 1 mL of KOH (0.5 M) was added, and the absorbance
220 of the solution was determined at 500 nm by a microplate reader (Biotek, USA).
221 Samples without CPMK were used as the control.

222

223 **Bio-reduction catalyzed by immobilized enzyme**

224 ***Optimal pH and temperature***

225 Substrate conversion was performed in a 5-mL mixture containing Mu-S5 and *BmGDH*
226 (2 U each), 50 mM CPMK, and 50 mM glucose. The optimal pH of the reaction was
227 determined by measuring the substrate conversion in the following buffers: sodium
228 citrate-citric acid buffer (SCC buffer, pH 5.0–6.0, 100 mM), sodium phosphate buffer
229 (pH 6.0–8.0, 100 mM), and Tris-HCl (pH 7.5–9.0, 100 mM). The optimal temperature
230 was determined by measuring substrate conversion at 25–40°C.

231 ***Optimal ratio of Mu-S5 and BmGDH***

232 A total dry mass of 2 mg purified Mu-S5 and *BmGDH* were mixed at activity ratios of
233 10:1 to 1:10. The reaction mixture containing 100 mM CPMK, 100 mM glucose,

234 premixed solutions of Mu-S5 and *BmGDH*, and 100 mM Tris-HCl (pH 7.5) in a final
235 volume of 5 mL, was conducted at 30°C for 30 min. The substrate conversion was
236 analyzed by HPLC.

237 *Effect of various co-solvents*

238 In the pre-experiments, 10 mM CPMK was dissolved in different co-solvents. A mother
239 solution of CPMK (1.0 M) was prepared in ethanol by heating. Different amounts of
240 co-solvents were added into 10 µL of the CPMK mother solution (1.0 M), then different
241 volumes of water were added to a final volume of 1 mL (**Table S7 & S8**). According
242 to preliminary results, 5% Tween 80, 2.5% Triton X-100, and 20–40 mM HP-β-CD
243 were tested as co-solvents. Substrate conversion was performed in a 5-mL mixture
244 containing 0.2 U of Mu-S5 and 0.1 U of *BmGDH*, 10 mM glucose, and 10 mM CPMK.
245 The substrate conversion was analyzed by HPLC.

246

247 **Extraction with macroporous resins**

248 *Static adsorption/desorption rate of macroporous resins toward CPMA*

249 Adsorption rate: Substrate CPMK in Solution B was replaced with the product (*S*)-
250 CPMA to prepare Solution C. A mass of 0.5 g resin D101 or XDA-1 was added into 10
251 mL of Solution C; the adsorption process was then carried out at 30°C and 150 rpm for
252 4 h, and the non-adsorbed (*S*)-CPMA was analyzed by HPLC.

253 Adsorption rate =
$$\frac{\text{Initial concentration}-\text{Residual concentration}}{\text{Initial concentration}} \times 100\%$$

254 Desorption rate: The 0.5 g masses of resin D101 or XDA-1 that had adsorbed (*S*)-
255 CPMA were washed with water and filtered on a Buchner funnel. A volume of 10 mL

256 ethanol or ethyl acetate was added, and the desorption process was carried out at 30°C
257 and 150 rpm for 4 h. The concentration of (S)-CPMA in the desorbing agent was
258 analyzed by HPLC.

$$259 \text{ Desorption rate} = \frac{\text{Concentration in the desorbing agent}}{\text{Initial concentration}} \times 100\%$$

260 *Adsorption capacity of macroporous resins toward NADP⁺*

261 A mass of 0.5 g resin D101 or XDA-1 was added to 10 mL of solution containing 10
262 mM NADP⁺. The adsorption process was carried out at 30°C and 150 rpm for 4 h and
263 the adsorption capacity was determined using an EnzyChrom NADP⁺/NADPH assay
264 kit from BioAssay Systems (Hayward, CA).

265

266 **Continuous flow reaction in a 10-mL PBR**

267 A 12-mL polypropylene column was packed with 10 mL of Ni-NTA Cerase to prepare
268 a PBR. A total of 200 mg of purified enzyme was loaded (141.1 mg Mu-S5 (940 U) and
269 58.9 mg *BmGDH* (470 U)), and the enzyme loading was 20 mg protein/mL Ni-NTA
270 Cerase (eg. 16.7 mg protein/g Ni-NTA Cerase). For operational convenience, 50
271 mg/mL solutions of both Mu-S5 and *BmGDH* were prepared, corresponding to 5.08 mL
272 and 1.99 mL loading volumes, respectively. A peristaltic pump was used to pump
273 Solution B through the PBR at 5 mL/min. A glass column filled with 28 g (40 mL) of
274 resin D101 was connected to the PBR to extract CPMA from the effluent of the PBR.
275 A total of 4 L of reaction solution was prepared as described above. To ensure the
276 continuity of the process, the 4 L initial reaction solution was divided into two portions
277 (2 L each) for rotation. According to the dynamic adsorption capacity of macroporous

278 resin D101, 5 columns (40 mL) filled with D101 were required to adsorb (*S*)-CPMA
279 from 2 L of effluent. In this study, a fresh D101 column was changed every 80 min.
280 Notably, D101 resin is reproducible and recyclable. After 400 min, 2 L of effluent
281 solution was collected, and 2 mL CPMK solution (dissolved in ethanol at 1 M) was
282 added as reaction solution for recycling. Meanwhile, the resin 101 that adsorbed (*S*)-
283 CPMA in each column was collected and filtered with a Buchner funnel. A double
284 volume (80 mL) of ethyl acetate was added, and the desorption process was carried out
285 at 30°C and 150 rpm for 4 h. The product (*S*)-CPMA was concentrated by rotary
286 evaporation.

287 The space-time-yield (STY) was defined as the mass of (*S*)-CPMA produced per
288 milliliter of PBR enzyme per day.

$$289 \text{ STY} = \frac{\text{Amount of } (S)\text{-CPMA (g)}}{\text{Volume of PBR (L)} \times \text{Reaction time (d)}}$$

290

291 **Treatment of column packing for recovery and reuse**

292 After each batch process, the immobilized Ni-NTA Cerase was washed with 10 column
293 volumes of 50 mM EDTA and water sequentially. After 30 min incubation with a
294 solution of 100 mM NiSO₄, the column was washed with 10 column volumes of water,
295 and then reused for the next batch.

296 For the product adsorbent column, (*S*)-CPMA was collected by soaking with ethyl
297 acetate, then the D101 macroporous resin was sequentially washed using a Buchner
298 funnel with 2 column volumes of the following: a solution containing 95% ethanol,
299 water, and 3% HCl; water (rinsed to neutral); a 3% NaOH solution; water (rinsed to

300 neutral).

301

302 **Results and discussion**

303 **Device for enzyme immobilization and product recovery**

304 For continuous synthesis of (*S*)-CPMA, a prototype combined device consisting of four
305 main parts was designed (**Fig. 1**), including a bottle of flow reaction solution
306 (containing dissolved CPMK, NADP⁺, and glucose), a PBR containing Ni-NTA Cerase
307 with immobilized Mu-S5 and *BmGDH*, a macroporous resin column for adsorbing
308 hydrophobic products, and a beaker for effluent recovery. When the reaction solution
309 flowed through the PBR, the ketone substrate CPMK was reduced into the
310 corresponding chiral alcohol (*S*)-CPMA. After that, the hydrophobic (*S*)-CPMA would
311 be adsorbed by the resin column, whereas the effluent solution containing residual
312 water-soluble NAD(P)H/NAD(P)⁺ and by-product gluconate would be collected for
313 recovery. After replenishing CPMK and glucose, the effluent solution could be reused
314 as reaction solution. The post-column samples of the PBR and the resin column were
315 determined by using a previously established DNPH assay [37] to ensure complete bio-
316 reduction and adsorption. When the resin column reached saturation, it could easily be
317 replaced with a new one, and the adsorbed (*S*)-CPMA could be separated from the resin
318 by static or dynamic elution. This operation mode not only avoids the emulsification
319 issue caused by extraction with organic solvents in biocatalytic reaction system, but
320 also allows for the reuse of expensive cofactors. Moreover, Ni-NTA Cerase could be
321 recycled and reused by treatment with EDTA and NiSO₄ solutions after each batch. For

322 the product adsorbent column, D101 macroporous resin is also recyclable and can be
323 reused by treatment with dilute acid and alkali solutions. The whole process is simple
324 and easy to operate with low material and downstream costs. Optimization of reaction
325 conditions was performed to verify the feasibility of the continuous reaction process.

326

327 **Screening of supports for the immobilization of Mu-S5 and *BmGDH***

328 Immobilized metal affinity chromatography (IMAC) is commonly used for His-tagged
329 protein purification and can also be applied to the immobilization of enzymes [25].

330 Herein, a commercially available NTA Cerase was chosen as the immobilization carrier
331 material. Compared with the NTA agarose (0.05 Mpa/cm) commonly used in
332 purification, NTA Cerase (1 Mpa/cm) possesses a stronger stability under system
333 pressure and can withstand relatively higher flow rates. Ni²⁺ and Mn²⁺ were selected
334 due to their positive effects on the catalytic activity of Mu-S5 (104% for Ni²⁺ and 116%
335 for Mn²⁺) (**Fig. 2A**). After immobilization, protein loading per gram carrier was
336 calculated to be 61.9 mg for Ni-NTA Cerase and 42.2 mg for Mn-NTA Cerase (**Table**
337 **S2**). Additionally, Ni-NTA Cerase and Mn-NTA Cerase maintained 58.0% and 66.4%
338 relative catalytic activity compare with that of free enzyme in shake flasks, respectively
339 (**Table S6**).

340 The dilution ratio was defined as the reactor volume divided by the volumetric flow
341 rate of substrate, and it was used to evaluate the reactor performance. For Ni-NTA
342 Cerase immobilized Mu-S5, the substrate conversion remained over 99% with
343 increased substrate flow rate, whereas a consistent decline of conversion was observed

344 for Mn-NTA Cerase. The conversion decreased to less than 50% at a flow rate of 1
345 mL/min for Mn-NTA (**Fig. 2B**). Due to the larger ionic radius of the Mn^{2+} ion (91 nm)
346 compared to the Ni^{2+} ion (73 nm), the metal coordination bond strength of Ni^{2+} is
347 stronger than that of Mn^{2+} . This difference in coordination bond strength is likely
348 responsible for the poor performance of Mn-NTA. Notably, no protein leaching was
349 detected at dilution rates of 0.2–1.0 min^{-1} as evaluated by a Bradford assay. Therefore,
350 insufficient enzyme immobilized on Mn-NTA is likely responsible for the low
351 conversion. To further reduce the cost, a cheaper chelating resin IRC-748 was
352 functionalized with Ni^{2+} and employed for enzyme immobilization. However, activity
353 above 85% was lost after immobilization due to the poor biocompatibility of the resin.
354 Therefore, Ni-NTA Cerase was selected for protein immobilization.

355

356 **Comparison of 6×His-tag and Heli-tag in protein immobilization**

357 In addition to the 6×His- tag, the Heli-tag fusion peptide with higher affinity was
358 attempted for immobilization [28, 29]. Since the C-terminal portion of Mu-S5 is buried
359 and solvent inaccessible, the Heli-tag was fused to the N-terminal of Mu-S5.

360 The affinity of 6×His- and Heli-tagged Mu-S5 with Ni-NTA was compared under the
361 same elution conditions (**Fig. 3**). The two fused Mu-S5s showed similar solubility as
362 confirmed by SDS-PAGE. Both fusion proteins were barely detected in the flow-
363 through (lane 3). The majority of the 6×His-tag fused Mu-S5 eluted at 100 mM
364 imidazole, and some was eluted at 50 mM imidazole. (**Fig. 3B**). In contrast, almost all
365 of the Heli-tag fused Mu-S5 was eluted at 20-50 mM imidazole (**Fig. 3 A**). After elution

366 with 50 mM imidazole, over 95% of enzymatic activity was lost with the of Heli-tag
367 fused Mu-S5, whereas only 3.9% activity loss was observed with the 6×His-tag fused
368 Mu-S5 (**Table S9**). It is speculated that the Heli-tag may not be fully exposed on the
369 surface of Mu-S5, which may affect the structure and conformation of the protein [40].
370 Collectively, the Heli-tag that has been proven to be effective in D-amino acid oxidase
371 immobilization [30] is not applicable for Mu-S5, and the 6×His-tag was used for
372 enzyme immobilization for further study.

373

374 **Optimization of reaction conditions of PBR**

375 The reaction conditions were optimized to obtain improved catalytic performance. The
376 pH, temperature, ratio of Mu-S5/*BmGDH*, and co-solvent were optimized in shake
377 flasks, and the dilution rate was determined in a PBR. As shown in **Fig. 4A**, Mu-S5
378 showed optimal activity at pH 7.5 in Tris-HCl buffer. The optimal temperature of the
379 reaction was 30°C (**Fig. 4B**), and the reaction can be conveniently performed at room
380 temperature. Under the optimized pH and temperature, the highest conversion (75.1%)
381 was achieved using immobilized Mu-S5/*BmGDH* at an activity ratio of 2:1 (**Fig. 4C**),
382 which may be related to the addition of an equal amount of glucose as co-substrate. As
383 there were not significant differences in the large-scale fermentation costs between Mu-
384 S5 and *BmGDH* (**Table S1**), the optimized activity ratio of 2:1 is acceptable.

385 Additionally, an aqueous solution of the insoluble substrate CPMK should be prepared
386 for the continuous process. Several commonly used dispersants were screened, and a
387 10 mM aqueous solution of CPMK could be obtained when supplemented with 5%

388 Tween 80, 2.5% Triton X-100, and 20 mM hydroxypropyl-beta-cyclodextrin (HP- β -CD)
389 in preliminary experiments (**Table S7 & S8**). **Fig. 4D** shows the effect of co-solvents
390 on catalytic activity of immobilized Mu-S5, and a reaction system with the addition of
391 20% ethanol served as a control. A decrease in substrate conversions was observed
392 when using Tween 80 and Triton X-100 as co-solvents, and this may be attributed to
393 the disrupted protein conformation that results from the interactions between the
394 nonionic surfactant and the hydrophobic region of the enzyme. Surprisingly, HP- β -CD
395 not only improved the substrate solubility but also exhibited excellent biocompatibility.
396 Benefiting from the hydrophilic exterior and hydrophobic interior structure of HP- β -
397 CD, mass transfer was improved, and almost complete conversions were achieved with
398 the addition of 20–40 mM HP- β -CD, whereas only 76.4% conversion was observed in
399 the control reaction (20% ethanol). Finally, the dilution rate of 10 mL PBR ($\Phi=1.6$ cm,
400 $h=5.5$ cm) was determined under the optimized reaction conditions. As shown in **Fig.**
401 **4E**, the substrate conversion began to decrease when the volumetric flow rate of CPMK
402 exceeded 6 mL/min, and only 93.1% conversion was detected at an increased flow rate
403 of 10 mL/min without protein leaching. Considering possible damage to the
404 immobilized materials and enzymes caused by the long-term continuous reaction, a
405 volumetric flow rate of 5 mL/min was chosen.

406

407 **Comparison of properties of D101 and XDA-1**

408 Macroporous resins possess stable physical and chemical properties, and can be
409 classified into three types: polar, non-polar, and weakly polar. According to the design,

410 the hydrophobic diaryl alcohol product is expected to be adsorbed whereas the water-
411 soluble cofactor should be recycled and reused for consecutive reaction cycles.
412 Therefore, two commonly used non-polar macroporous resins D101 and XDA-1 were
413 selected for (*S*)-CPMA enrichment (**Table S10**). The adsorption rate is an important
414 index to evaluate the adsorption performance of macroporous resins. As shown in **Table**
415 **1 & S11**, the two resins maintained almost the same adsorption capacity toward the
416 product (*S*)-CPMA. Resin XDA-1 with a larger surface area exhibited more than a 90%
417 static adsorption rate. The desorption rate of the two resins toward (*S*)-CPMA was also
418 compared (**Table S12**). Firstly, different proportions of recommended aqueous-ethanol
419 were used as washing solutions, however, only about 20% of the product could be
420 eluted even when pure ethanol (100%) was used. Then, ethyl acetate with lower polarity
421 was tested, and the desorption rate increased rapidly. In the case of D101, more than
422 95% of CPMA could be eluted with the addition of ethyl acetate, whereas only 82.6%
423 of CPMA could be eluted when XDA-1 was selected for extraction. Afterward, the
424 residual NADP⁺ concentration was measured using an NADP⁺/NADPH assay kit to
425 evaluate whether the selected resin could be used for coenzyme recycling. The results
426 showed that neither D101 nor XDA-1 could adsorb NADP⁺ (**Table S13, Fig S4**). Thus,
427 D101 was selected as the supporting material in the extraction column due to the higher
428 desorption rate. Using a 40 mL column ($\Phi=1.6$ cm, $h=20$ cm) filled with 28g of D101
429 resin, fractions were collected sequentially every 10 mL, and a trace amount of (*S*)-
430 CPMA was detected in the 47th sample. The dynamic adsorption capacity of D101 was
431 calculated to be 36.3 mg/g, which is roughly equal to its static adsorption capacity (38.6

432 mg/g).

433

434 **Asymmetric synthesis and in situ extraction of (S)-CPMA in a continuous device**

435 Asymmetric synthesis of (S)-CPMA was carried out under optimized immobilization
436 and reaction conditions. **Fig. 5** shows the conversion ratio of CPMK catalyzed by
437 immobilized Mu-S5/*BmGDH* in a 10-mL PBR. A complete conversion of CPMK was
438 observed in the first 3 days followed by a gradual decline. When the reaction proceeded
439 to day 4, an obvious red-brown color formed in DNPH assay, indicating an incomplete
440 conversion of the ketone substrate. On day 7, only 54.3% conversion was observed by
441 HPLC, and conversion continued to decreased sharply to 22.5% on day 8. In view of
442 the negligible effect of the His-tag on protein structure, the insufficient stability of the
443 Mu-S5 (half-life of 152 h at 30°C, **Fig. S5**) might be responsible for the decreased
444 substrate conversion after day 3. Finally, the STY was calculated to be 1,560 g/(L·d)
445 based on data from day 1 to 3. A total volume of 4 L initial reaction solution was
446 recycled up to 6 times. After 400 min of reaction, the D101 resin in the first 5 adsorption
447 columns was collected, and a total of 3.63 g (S)-CPMA with 84% isolated yield was
448 obtained by ethyl acetate extraction.

449 Peschke et al. reported an emerging co-immobilization approach using *Spy*-tag and *Spy*-
450 catcher fusion enzymes [32]. However, hydrogels self-assembled from free enzymes
451 alone are mechanically fragile, which limit the operation strength and flow rate of
452 continuous reactions. In comparison, immobilized *KpADH* Mu-S5 and *BmGDH*
453 maintained excellent catalytic activity and operational stability under a 5 min⁻¹ dilution

454 rate during the first 3 days of the reaction. In a study by Dall'Oglio and coworkers,
455 KRED1-Pglu and GDH were covalently bound to aldehyde activated agarose [31].
456 Although the stability and solvent tolerance were improved, alkaline operating
457 conditions ($\text{pH} \geq 10$) pose a challenge to the catalytic activity of enzymes. For example,
458 Tt-ADH2 and formate dehydrogenase (FDH) had to be immobilized by covalent
459 binding and ionic adsorption, respectively, due to the inactivation of FDH after
460 aldehyde immobilization [35]. In our study, ADH and GDH were both coordination
461 bound to Ni-NTA Cerase, and over 60% of catalytic activity could be maintained under
462 near-neutral conditions ($\text{pH} 7.5$). Using commercially available enzyme membrane
463 reactor, (*R*)-2-octanol could be continuously synthesized by co-immobilized *Lb*ADH
464 and GDH, and produced a fairly high STY of 454 g/(L·d). Additionally, product
465 separation and cofactor recycling were achieved by supercritical carbon dioxide (scCO_2)
466 extraction [41] (**Table 2**).

467 In our study, the macroporous resin adsorption strategy with low equipment
468 requirements was facily used for product recovery, and in the subsequent resin
469 extraction step, much less organic solvent was used than in traditional liquid-liquid
470 extractions. Additionally, emulsification caused by proteins in free-enzyme systems
471 was completely avoided, and a satisfactory yield was obtained by the simple operations
472 of soaking and rotary evaporation.

473 **Conclusions**

474 Continuous flow biocatalysis can improve the overall reaction process by protecting

475 fragile enzymes from mechanical stirring or agitation, as well as simplifying the
476 downstream product isolation and treatment. Herein, a 6×His-tag fused Mu-S5 and
477 *BmGDH* were co-immobilized on Ni-NTA Cerase to construct a PBR for continuous
478 asymmetric synthesis of (*S*)-CPMA. As a support material for enzyme immobilization,
479 the commercially available Ni-NTA Cerase is recyclable in process operation,
480 Furthermore, the D101 macroporous resin column was used after the PBR to adsorb the
481 (*S*)-CPMA product specifically, and recycling and reuse of the reaction solution
482 containing hydrophilic NADP⁺/NADPH was easily achieved. During 400 min of flow
483 reaction, a 4 L initial reaction solution could be recycled up to 6 times, demonstrating
484 the feasibility of reducing the cost of cofactors. Additionally, the (*S*)-CPMA product
485 was enriched on the D101 resin and could be separated conveniently and effectively by
486 soaking in ethyl acetate. In summary, the designed continuous biocatalytic process
487 using coupled PBR and *in situ* product recovery could be potentially applied in
488 preparation of chiral alcohols. This continuous flow biocatalysis system is especially
489 suitable for enzymatic reactions involving cofactor recycling and substrate/product
490 inhibitory issues.

491

492 **Acknowledgements**

493 This work was supported by National Key R&D Program [2018YFA0901700],
494 National Natural Science Foundation of China [21907040, 21776112, 22077054],
495 China Postdoctoral Science Foundation [2019M651703], National First-Class
496 Discipline Program of Light Industry Technology and Engineering [LITE2018-07],

497 and the Program of Introducing Talents of Discipline to Universities [111-2-06].

498

499 **Authors' contributions**

500 Jieyu Zhou: Conceptualization, Investigation, Methodology, Writing-original draft.

501 Yanfei Wu: Data curation. Qingye Zhang: Data curation, Formal analysis. Guochao

502 Xu: Writing-review & editing. Ye Ni: Supervision, Writing-review & editing, Project

503 administration, Funding acquisition.

504

505 **Compliance with Ethical Standards**

506 **Competing Interests** The authors declare that they have no known competing

507 financial interests or personal relationships that could have appeared to influence the

508 work reported in this paper

509

510 **Supplementary**

511 Supplementary material related to this article can be found in the online version.

512

513 **References**

- 514 1. Britton, J., Majumdar, S. and Weiss, G.A. (2018) *Chem. Soc. Rev.* 47, 5891-
515 5918.
- 516 2. Adamo, A., Beingsner, R.L., Behnam, M., Chen, J., Jamison, T.F., Jensen,
517 K.F., Monbaliu, J.M., Myerson, A.S., Revalor, E.M., Snead, D.R., Stelzer, T.,
518 Weeranoppanant, N., Wong, S.Y. and Zhang, P. (2016) *Science* 352, 54-61.
- 519 3. Webb, D. and Jamison, T.F. (2010) *Chem. Sci.* 1, 675-680.
- 520 4. Cuong, N.P., Lee, W., Oh, I., Thuy, N.M., Kim, D., Park, J. and Park, K. (2016)
521 *Process Biochem.* 51, 282-287.
- 522 5. Cimporescu, A., Todea, A., Badea, V., Paul, C. and Peter, F. (2016) *Process*
523 *Biochem.* 51, 2076-2083.
- 524 6. Jia, C., Wang, H., Zhang, W., Zhang, X. and Feng, B. (2018) *Process Biochem.*
525 66, 28-32.
- 526 7. de Oliveira Lopes, R., Ribeiro, J.B., Silva De Miranda, A., Vieira Da Silva, G.V.,
527 Miranda, L.S.M., Ramos Leal, I.C. and Mendonça Alves De Souza, R.O. (2014).
528 *Tetrahedron* 70, 3239-3242.
- 529 8. Tamborini, L., Romano, D., Pinto, A., Contente, M., Iannuzzi, M.C., Conti, P.
530 and Molinari, F. (2013). *Tetrahedron Lett.* 54, 6090-6093.
- 531 9. J. Döbber, T.G.H.O. (2018). *Green Chem.* 20, 544-552.
- 532 10. Xiao, M., Qi, C. and Obbard, J.P. (2011) *Bioenergy* 3, 293-298.
- 533 11. Tan, A.W., Fischbach, M., Huebner, H., Buchholz, R., Hummel, W., Dausmann,
534 T., Wandrey, C. and Liese, A. (2006). *Appl. Microbiol. Biotechnol.* 71, 289-93.

- 535 12. Thompson, M.P., Peñafiel, I., Cosgrove, S.C. and Turner, N.J. (2018). *Org.*
536 *Process Res. Dev.* 23, 9-18.
- 537 13. Li, F., Zheng, Y., Li, H., Chen, F., Yu, H. and Xu, J. (2019). *Tetrahedron* 75,
538 1706-1710.
- 539 14. Orrego, A.H., López-Gallego, F., Espaillet, A., Cava, F., M., J. and A, G.A.A.J.
540 (2018). *ChemCatChem* 10, 3002-3011.
- 541 15. García-García, P., Rocha-Martin, J., Fernandez-Lorente, G. and Guisan, J.M.
542 (2018) *Enzyme Microb. Tech.* 115, 73-80.
- 543 16. Arana-Pena, S., Carballares, D., Morellon-Sterlling, R., Berenguer-Murcia, A.,
544 Alcantara, A.R., Rodrigues, R.C. and Fernandez-Lafuente, R. (2020). *Biotechnol.*
545 *Adv.* 107584.
- 546 17. Rocha-Martín, J., Rivas, B.D.L., Muñoz, R., Guisán, J.M. and López-Gallego, F.
547 (2012). *ChemCatChem* 4, 1279-1288.
- 548 18. Trobo-Maseda, L., Orrego, A.H., Guisan, J.M. and Rocha-Martin, J. (2020). *Int.*
549 *J. Biol. Macromol.* 157, 510–521.
- 550 19. Hearon, J.Z., Sundberg, L., and Malmström, B.G. (1975). *Nature* 258, 598-599.
- 551 20. Planchestainer, M., Contente, M.L., Cassidy, J., Molinari, F., Tamborini, L. and
552 Paradisi, F. (2017) *Green Chem.* 19, 372-375.
- 553 21. Liu, J., Pang, B.Q.W., Adams, J.P., Snajdrova, R. and Li, Z. (2017).
554 *ChemCatChem* 9, 425-431.
- 555 22. Vahidi, A.K., Yang, Y., Ngo, T.P.N. and Li, Z. (2015). *ACS Catal.* 5, 3157-3161.
- 556 23. Yang, J., Ni, K., Wei, D. and Ren, Y. (2015). *Biotechnol. Bioproc. E.* 20, 901-

- 557 907.
- 558 24. Engelmark Cassimjee, K., Kadow, M., Wikmark, Y., Svedendahl Humble, M.,
559 Rothstein, M.L., Rothstein, D.M. and Bäckvall, J.E. (2014). *Chem. Commun.* 50,
560 9134.
- 561 25. Ueda, E.K.M., Gout, P.W. and Morganti, L. (2003). *J. Chromatogr. A.* 988, 1-23.
- 562 26. Chou, Y., Ko, C., Chen, L.O. and Shaw, C.Y. (2015). *Molecules* 20, 3744-3757.
- 563 27. Böhmer, W., Knaus, T. and Mutti, F.G. (2018). *ChemCatChem* 10, 731-735.
- 564 28. Melchers, K., Herrmann, L., Mauch, F., Bayle, D., Heuermann, D.,
565 Weitzenegger, T., Schuhmacher, A., Sachs, G., Haas, R., Bode, G., Bensch, K.
566 and Schäfer, K.P. (1998). *Acta. Physiol. Scand. Suppl.* 643, 123-135.
- 567 29. Melchers, K., Weitzenegger, T., Buhmann, A., Steinhilber, W., Sachs, G. and
568 Schafer, K.P. (1996). *J. Biol. Chem.* 271, 446-57.
- 569 30. Hou, J., Jin, Q., Du, J., Li, Q., Yuan, Q. and Yang, J. (2014). *Bioproc. Biosyst.*
570 *Eng.* 37, 857-864.
- 571 31. Dall'Oglio, F., Contente, M.L., Conti, P., Molinari, F., Monfredi, D., Pinto, A.,
572 Romano, D., Ubiali, D., Tamborini, L. and Serra, I. (2017). *Catal. Commun.* 93,
573 29-32.
- 574 32. Peschke, T., Bitterwolf, P., Gallus, S., Hu, Y., Oelschlaeger, C., Willenbacher,
575 N., Rabe, K.S. and Niemeyer, C.M. (2018). *Angew. Chem., int. Edit.* 57, 17028-
576 17032.
- 577 33. Fassouane, A., Laval, J.M., Moiroux, J. and Bourdillon, C. (1990). *Biotechnol.*
578 *Bioeng.* 35, 935-939.

- 579 34. Ruinatscha, R., Buehler, K. and Schmid, A. (2014). *J. Mol. Catal. B-Enzym.* 103,
580 100-105.
- 581 35. Velasco-Lozano, S., Benítez-Mateos, A.I. and López-Gallego, F. (2017). *Angew.*
582 *Chem., int. Edit.* 56, 771-775.
- 583 36. Benítez-Mateos, A.I., San Sebastian, E., Ríos-Lombardía, N., Morís, F.,
584 González-Sabín, J. and López-Gallego, F. (2017). *Chem-Eur. J.* 23, 16843-16852.
- 585 37. Zhou, J., Xu, G., Han, R., Dong, J., Zhang, W., Zhang, R. and Ni, Y. (2016).
586 *Catal. Sci. Technol.* 6, 6320-6327.
- 587 38. Zhou, J., Wang, Y., Xu, G., Wu, L., Han, R., Schwaneberg, U., Rao, Y., Zhao,
588 Y., Zhou, J. and Ni, Y. (2018). *J. Am. Chem. Soc.* 140, 12645-12654.
- 589 39. Ni, Y., Zhou, J. and Sun, Z. (2012). *Process Biochem.* 47, 1042-1048.
- 590 40. Gaberc-Porekar, V. and Menart, V. (2001). *J. Biochem. Biophys. Methods* 49,
591 335–360.
- 592 41. Kohlmann, C., Leuchs, S., Greiner, L. and Leitner, W. (2011). *Green Chem.* 13,
593 1430-1437.
594

595 **Table and Figure Legends**

596 **Table 1.** Comparison of properties of D101 and XDA-1 resins.

597 **Table 2.** Comparison of asymmetric reductions catalyzed by various alcohol
598 dehydrogenases (ADHs) and glucose dehydrogenases (GDHs)/formate
599 dehydrogenases (FDHs) in flow reactions.

600 **Fig. 1.** Continuous biosynthesis of (*S*)-(4-chlorophenyl)-(pyridin-2-yl) methanol (*S*-
601 CPMA) with immobilized Mu-S5 and *Bm*GDH and *in situ* product recovery.

602 **Fig. 2.** Effect of metal ions on catalytic activity of Mu-S5 (A). Effect of dilution ratio
603 on conversion ratio of Ni-and Mn-functionalized porous ceramic/agarose composite
604 beads (Cerase) immobilized enzymes (B).

605 **Fig. 3.** SDS-PAGE of recombinant Mu-S5 fused with Heli-Tag (A) or 6×His Tag (B)
606 eluted at different Imidazole Concentrations. (A) M: protein ladder; lane 1:
607 supernatant of crude Heli-tag fused Mu-S5; lane 2: sediment of crude Heli-tag fused
608 Mu-S5; lane 3: Flow-through of crude Heli-tag fused Mu-S5; lane 4-7: eluent from
609 washing Heli-tag fused Mu-S5 with 20, 50, 100 and 300 mM imidazole, respectively;
610 (B) M: protein ladder; lane 1: supernatant of crude 6×His-tag fused Mu-S5; lane 2:
611 sediment of crude 6×His-tag fused Mu-S5; lane 3: Flow-through of crude 6×His-tag
612 fused Mu-S5; lane 4-7: eluent from washing 6×His-tag fused Mu-S5 with 20, 50, 100
613 and 300 mM imidazole, respectively.

614 **Fig. 4.** Optimization of reaction catalyzed by immobilized Mu-S5 and *Bm*GDH. (A)
615 pH; (B) temperature; (C) ratio of Mu-S5 and *Bm*GDH; (D) co-solvent; (E) flow rate
616 optimum of (4-chlorophenyl)(pyridine-2-yl)ketone (CPMK) in a 10-mL Ni-

617 functionalized porous ceramic/agarose composite beads (Ni-NTA Cerase) packed bed

618 reactor.

619 **Fig. 5.** Continuous production of (*S*)-(4-chlorophenyl)-(pyridin-2-yl) methanol (*S*-

620 CPMA) in a 10-mL packed bed reactor.

621

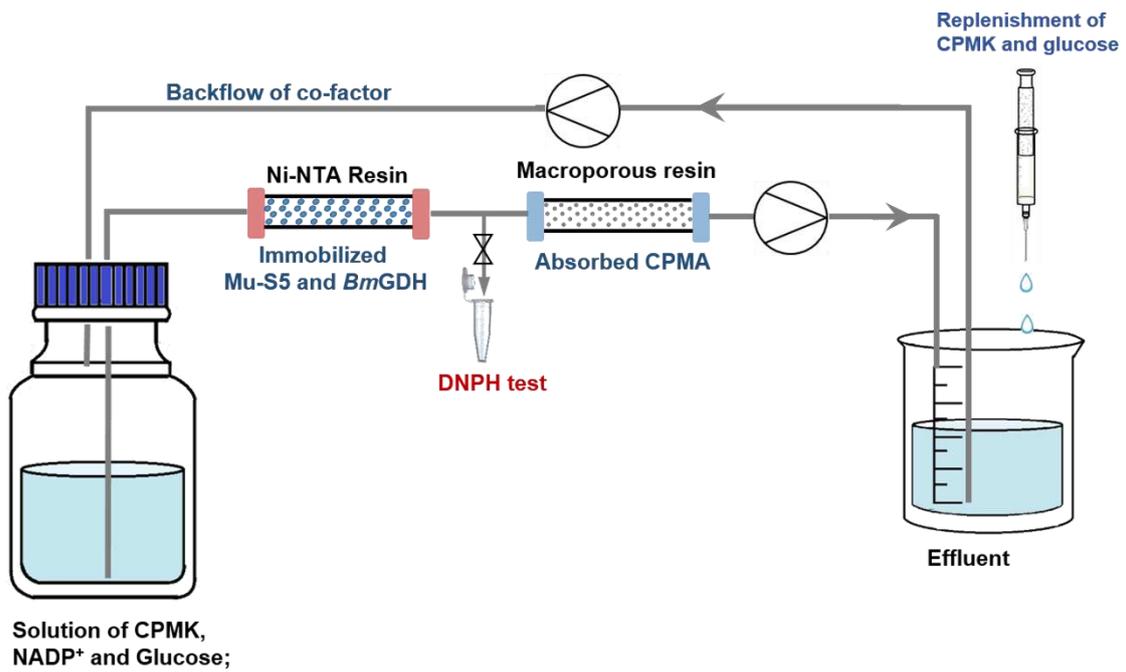
622 **Table 1.**

Resin	Constituent materials	Average aperture size (nm)	Specific surface area (m ² /g)	Static adsorption rate (%)	Desorption rate (%)
D101	Polystyrene	25–28	480–520	88.1±0.13	96.1±0.22
XDA-1	Polyacrylic acid	85–89	1000–1100	91.3±0.08	82.7±0.31

623

Table 2.

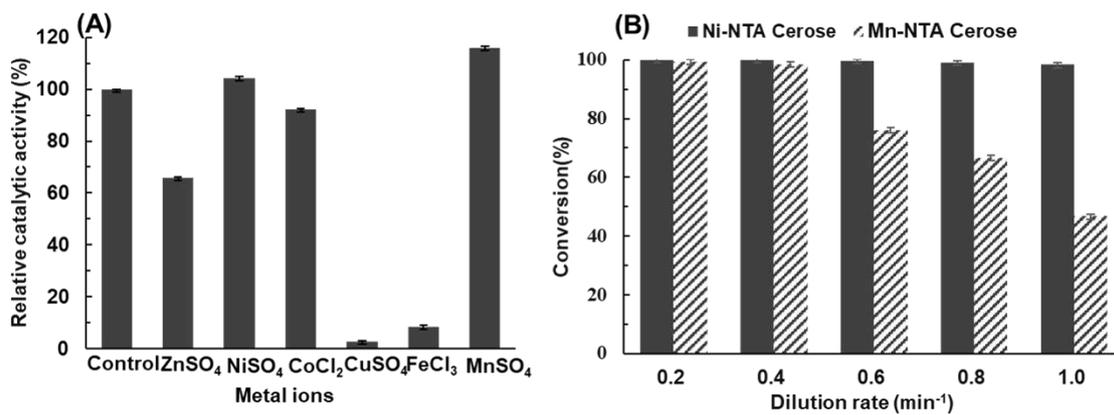
	This study	Peschke et al. [32]	Dall'Oglio et al.[31]	Kohlmann et al.[41]	Velasco-Lozano et al [35]
Ketone reductase	Mu-S5 from <i>Kluyveromyces polyspora</i>	<i>Lb</i> ADH from <i>Lactobacillus brevis</i>	KRED1-Pglu from <i>Pichia glucozyma</i>	<i>Lb</i> ADH from <i>Lactobacillus brevis</i>	Tt-ADH2 from <i>Thermus thermophilus</i>
Cofactor recycling	GDH	GDH	GDH	GDH	FDH
Immobilized strategy	IMAC	Self-assembling;	Covalent binding	Interception by semi-permeable membrane	Covalent binding & ionic adsorption
Support material	Cerose	Hydrogel	Aldehyde activated agarose	Enzyme membrane reactor	Aldehyde activated agarose & polyethyleneimine agarose
Substrate	CPMK	5-nitrononane-2,8-dione	1-phenylpropane-1,2-dione	octan-2-one	2,2,2-trifluoro-1-phenylethanone
Conc.(mM)	10	5.0	3.0	6.0	5.0
Reactor volume (mL)	10	0.15	0.90	16	1.2
Flow rate (mL/min)	5.0	0.010	0.050	0.10	0.050
SYT (g/(L·d))	1560	≈100	4.4–81.5	454.5	62.7
<i>In situ</i> product recovery	Macroporous resins adsorption	No	No	scCO ₂ extraction.	No



626

627

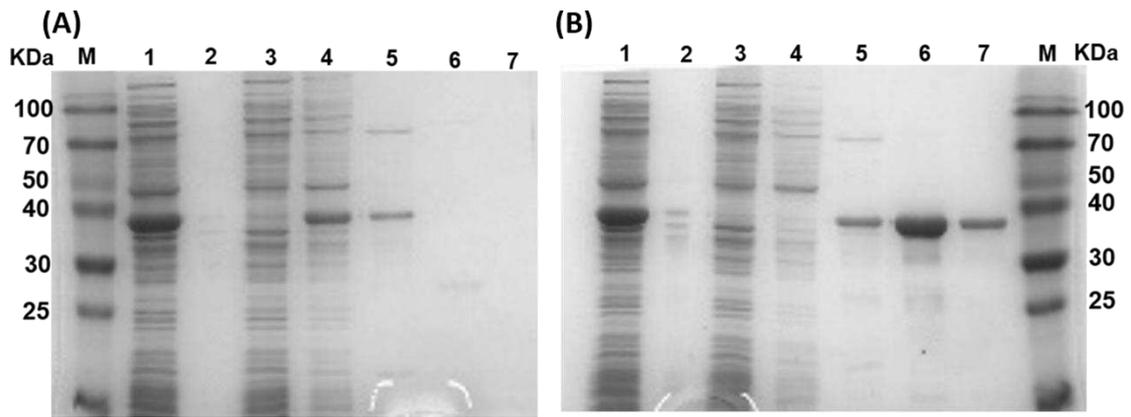
Fig. 1.



628

629

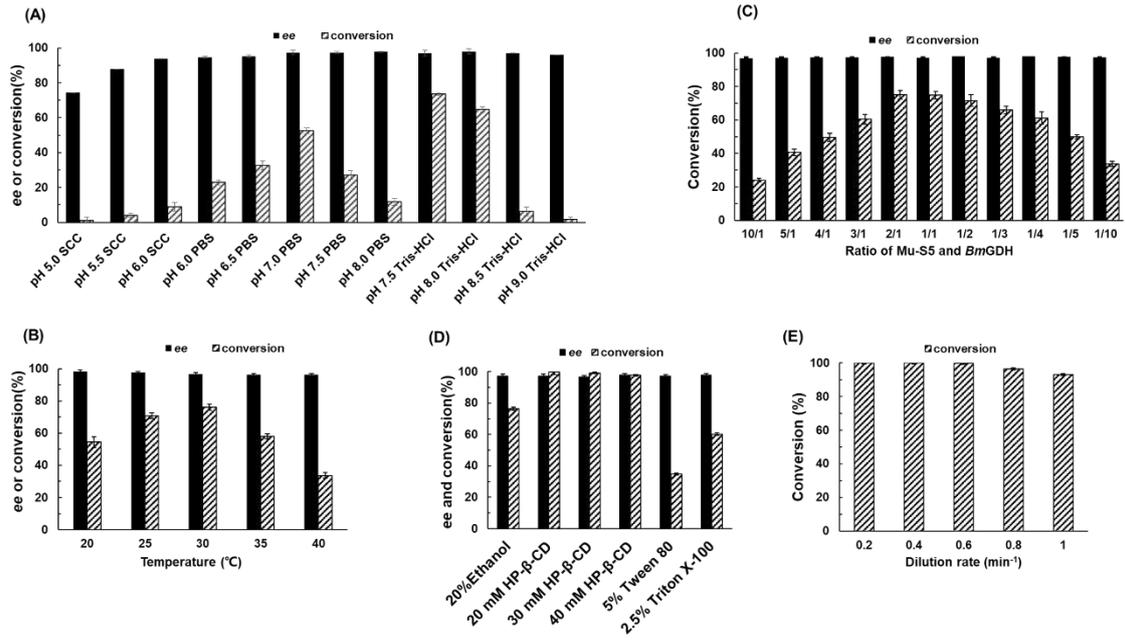
Fig. 2.



630

631

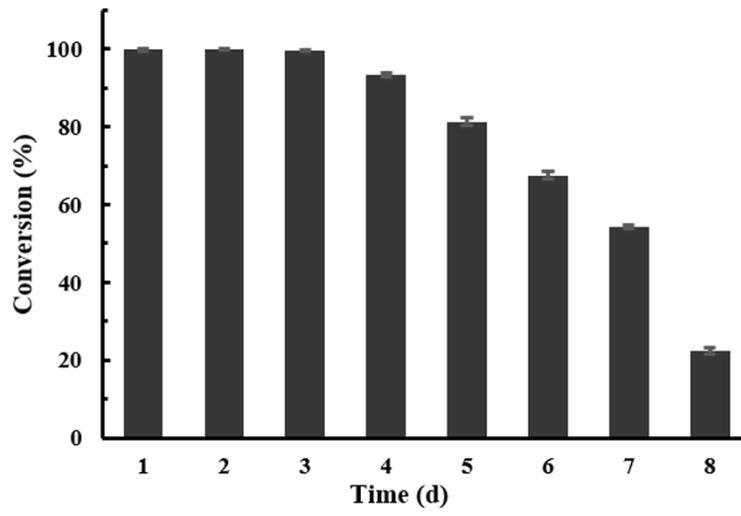
Fig. 3.



632

633

Fig. 4.



634

635

Fig. 5

Figures

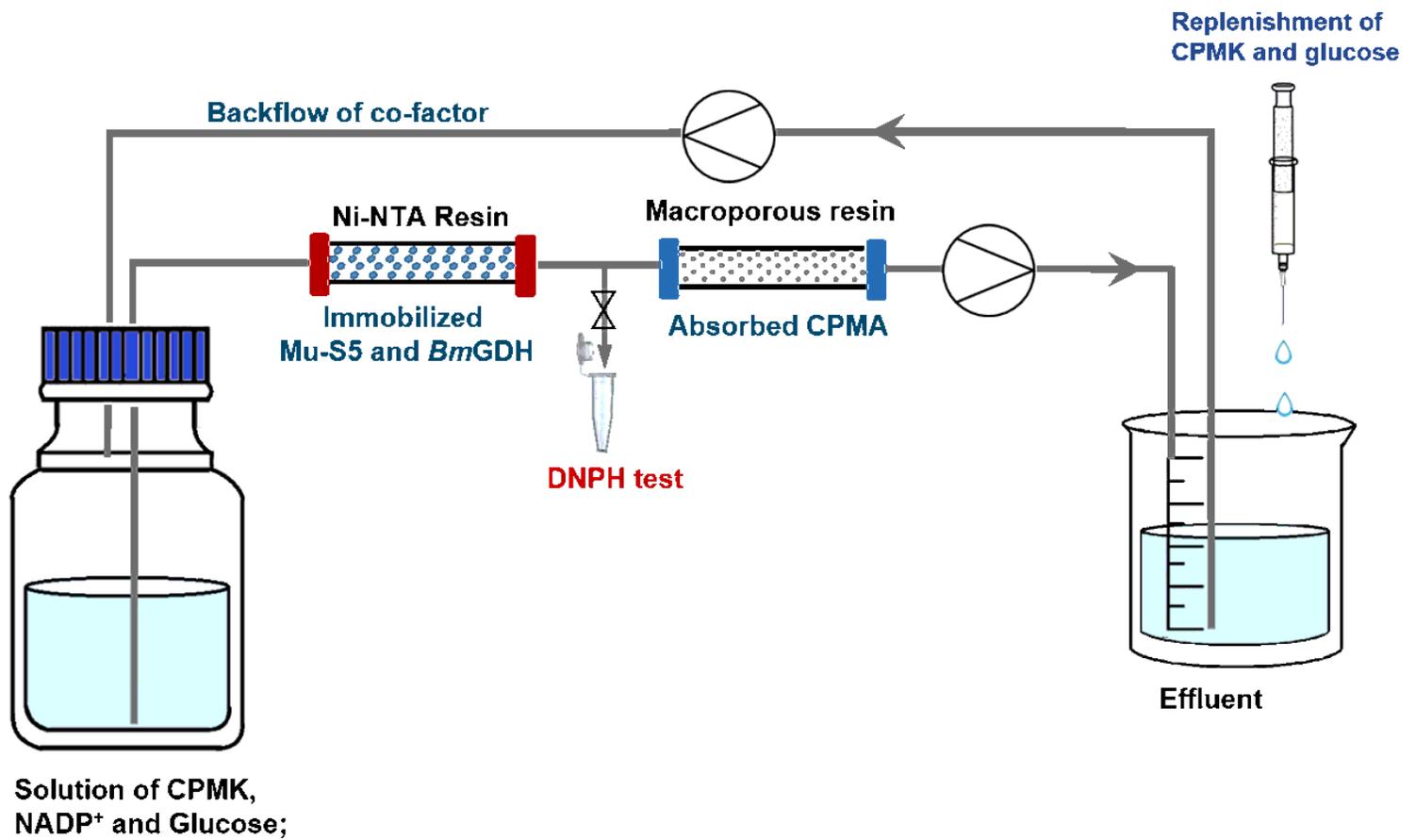


Figure 1

Continuous biosynthesis of (S)-(4-chlorophenyl)-(pyridin-2-yl) methanol (S-CPMA) with immobilized Mu-S5 and BmGDH and in situ product recovery.

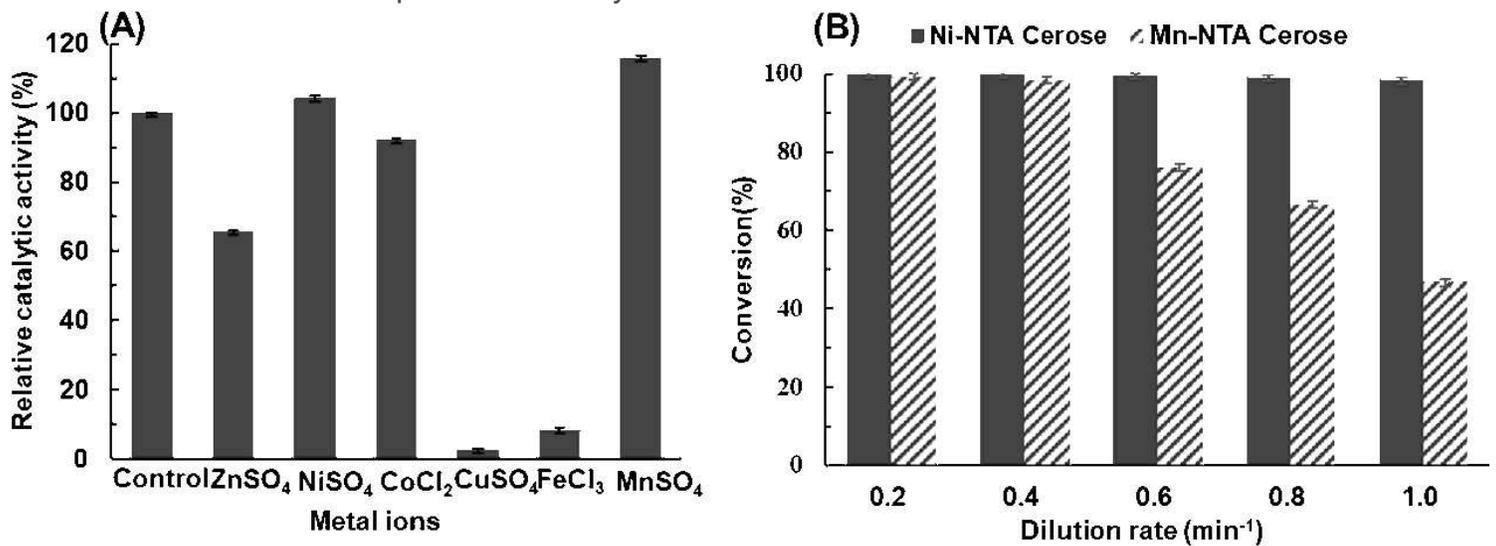


Figure 2

Effect of metal ions on catalytic activity of Mu-S5 (A). Effect of dilution ratio on conversion ratio of Ni- and Mn-functionalized porous ceramic/agarose composite beads (Cerose) immobilized enzymes (B).

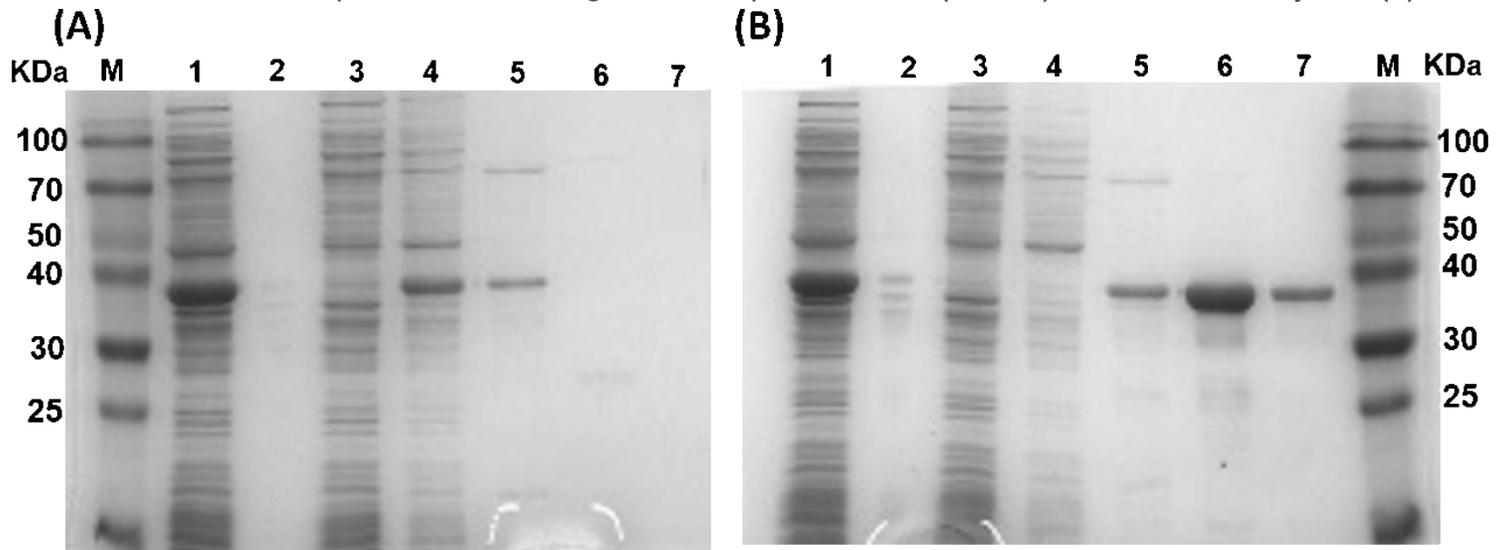


Figure 3

SDS-PAGE of recombinant Mu-S5 fused with Heli-Tag (A) or 6×His Tag (B) eluted at different Imidazole Concentrations. (A) M: protein ladder; lane 1: supernatant of crude Heli-tag fused Mu-S5; lane 2: sediment of crude Heli-tag fused Mu-S5; lane 3: Flow-through of crude Heli-tag fused Mu-S5; lane 4-7: eluent from washing Heli-tag fused Mu-S5 with 20, 50, 100 and 300 mM imidazole, respectively; (B) M: protein ladder; lane 1: supernatant of crude 6×His-tag fused Mu-S5; lane 2: sediment of crude 6×His-tag fused Mu-S5; lane 3: Flow-through of crude 6×His-tag fused Mu-S5; lane 4-7: eluent from washing 6×His-tag fused Mu-S5 with 20, 50, 100 and 300 mM imidazole, respectively.

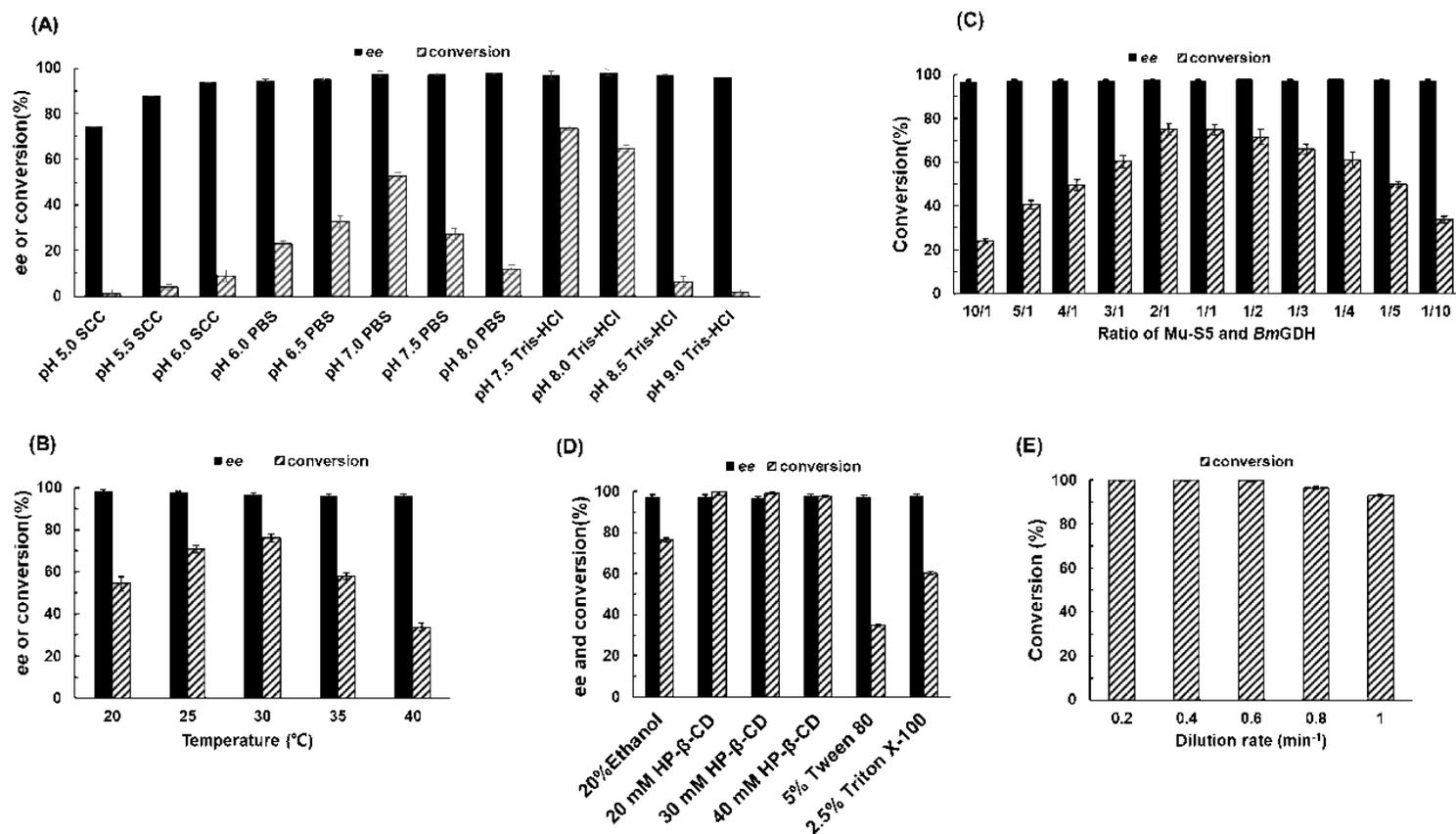


Figure 4

Optimization of reaction catalyzed by immobilized Mu-S5 and BmGDH. (A) pH; (B) temperature; (C) ratio of Mu-S5 and BmGDH; (D) co-solvent; (E) flow rate optimum of (4-chlorophenyl)(pyridine-2-yl)ketone (CPMK) in a 10-mL Ni-functionalized porous ceramic/agarose composite beads (Ni-NTA Cereose) packed bed reactor.

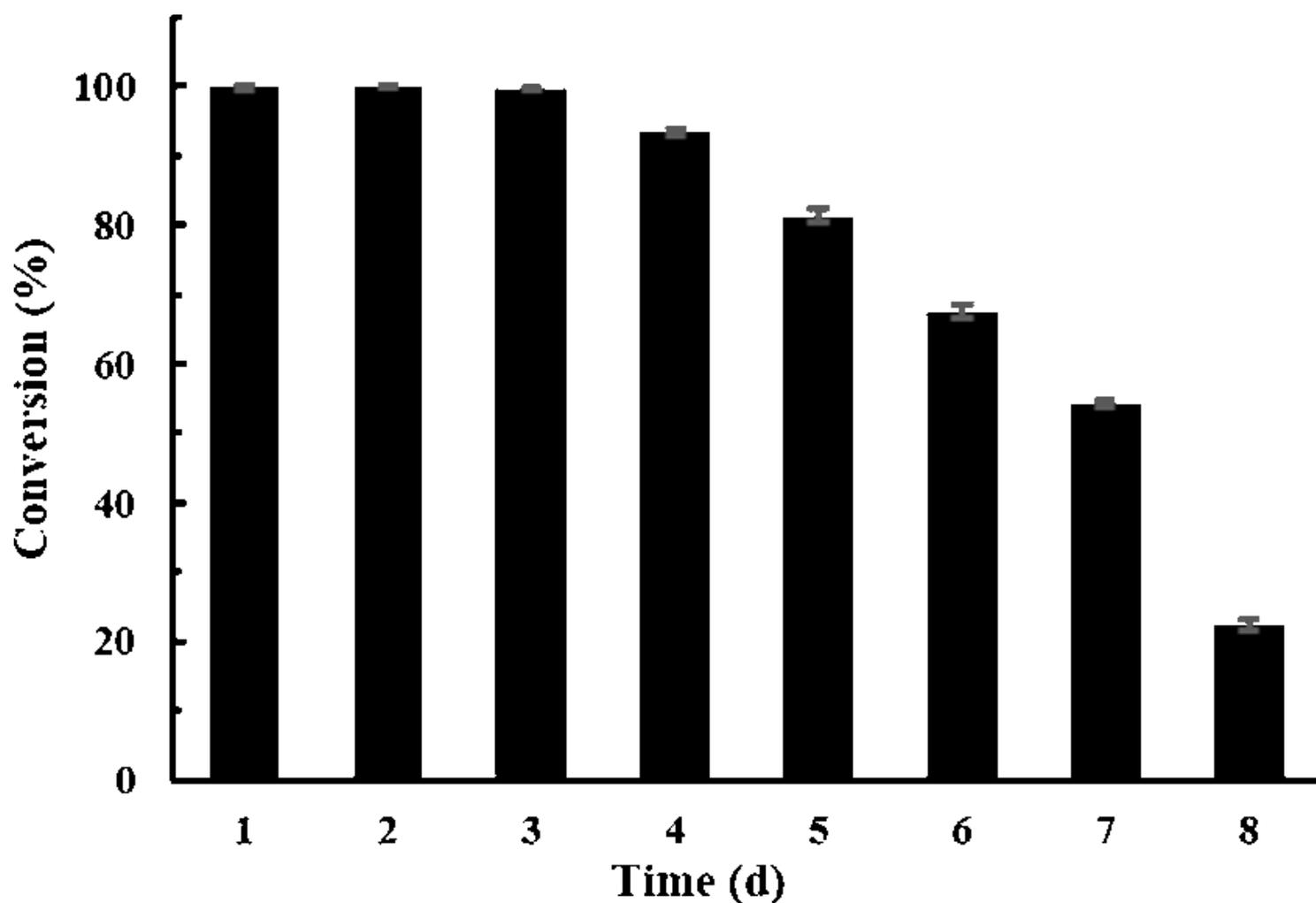


Figure 5

Continuous production of (S)-(4-chlorophenyl)-(pyridin-2-yl) methanol (S-CPMA) in a 10-mL packed bed reactor.

Supplementary Files

This is a list of supplementary files associated with this preprint. Click to download.

- [2021.2.5supportinginformation.docx](#)