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Formation of amide bond catalyzed by lipase in aqueous phase for peptide synthesis

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Lipase

Graphical Abstract









N-acetyl-phenylalanine ethyl ester L-ty

L-tyrosinamide

N-Acetyl-L-phenylalanyl-L-tyrosinamide



1

2 Highlights

- 3 Lipase catalyzed amidation for peptide synthesis.
- 4 A high yield of dipeptide was obtained in aqueous solution.
- 5 RSM design for optimization of reaction parameters.
- 6 Using the kinetic model compared the catalytic efficiency.
- 7 Kinetic constants show the advantage of lipase on amidation.

8

9 ABSTRACT:

- 10 A dipeptide N-acetyl-L-phenylalanyl-L-tyrosinamide (N-Ac-Phe-Tyr-NH₂), with
- 11 angiotensin I converting enzyme (ACE) inhibitor activatity, was synthesized via
- 12 porcine pancreatic lipase catalyzed amidation of *N*-acetyl-phenylalanine ethyl ester
- 13 with L-tyrosinamide in an aqueous phase. Response surface methodology was

14	employed to evaluate the effects of synthesis parameters. The optimum synthesis
15	conditions obtained an 84.45% yield of N-Ac-Phe-Tyr-NH ₂ with a reaction time of
16	3.8 min, a temperature of 20.9 °C, an enzyme amount of 6.5 U, and a substrate molar
17	ratio of 2.5:1 (Tyr:Phe). The kinetics of lipase and α -chymotrypsin catalyzed
18	amidation was compared using the Ping-Pong mechanism. The lipase showed a lower
19	apparent kinetic constant than α -chymotrypsin indicating that the acyl lipase
20	intermediate had a higher affinity toward tyrosinamide in the amidation. In addition,
21	because the lipase can avoid the secondary hydrolysis of synthesized peptide, it is
22	expected to be an effective method for obtaining a good yield of dipeptide.
23	
24	
25	KEYWORDS: enzymatic peptide synthesis, amidation, porcine pancreatic lipase,
26	N-acetyl-L-phenylalanyl-L-tyrosinamide, kinetics
27	
28	1. Introduction
29	Protein synthesis involves a sequence of amino acid coupling reactions to form amide
30	bonds that link amino acids in the amino-to-carboxyl direction. In the cell, protein
31	synthesis is catalyzed by the peptidyl transferase which occurs between two
32	aminoacyl-tRNAs on the ribosome [1]. However, the whole process in the cell is very

33	complex as it is mediated by more than one hundred macromolecules, including
34	mRNAs, tRNAs, ribosome, activating enzymes, and protein factors. Hence, a number
35	of enzymes have been used as practical catalysts in peptide synthesis performed in test
36	tubes [2, 3]. Enzymatic catalyzed synthesis of peptides has several advantages, such
37	as mild conditions, no racemization, more rapidly, higher yield and good
38	regioselectivity [4]. α -Chymotrypsin is the most commonly used enzyme in peptide
39	synthesis [5-7], but some peptides using lipase as catalysts have been attempted [8].
40	Lipase and α -chymotrypsin belong to the serine hydrolase family (substrate
41	enzyme intermediate formation is related to the hydroxyl group of serines at the
42	enzyme active site) [9, 10]. Lipases hydrolytically cleave ester bonds while
43	α -chymotrypsin hydrolyzes amide bonds. The mechanisms of the lipase and
44	α -chymotrypsin-catalyzed reactions are fundamentally identical. Both reactions
45	proceed via a serine hydroxyl at the enzyme active site, which attacks the carbonyl
46	carbon atom of the substrate to form an acyl enzyme intermediate [10, 11].
47	Subsequently, the acyl enzyme intermediate (acyl donor) can be deacylated by the
48	nucleophilic amino group of an amino acid substrate (acyl acceptor) to provide the
49	desired peptide, whereas deacylation with water results in hydrolysis. Hence,
50	hydrolysis and amidation pathways occur competitively [12, 13].
51	α -Chymotrypsin-catalyzed synthesis of peptides has been performed in an aqueous

52	solution [12]. Cbz-Asp-Phe-OMe has been synthesized in a monophasic
53	organic-aqueous (50% DMSO) solvent [14]. In contrast, the use of lipases to generate
54	amide bonds in organic solvents has been reported [15, 16]. Lipase catalyzed
55	aminolysis of dialkyl carbonates were run in <i>tert</i> -butyl alcohol [17]. Huang et al have
56	synthesized tetrapeptide Bz-Arg-Gly-Asp-Ser-NH2 using lipase in
57	dimethylformamide (DMF) and dimethyl sulfoxide (DMSO) [18]. Recently, lipase
58	catalyzed synthesis of peptidic hydrogels in an aqueous solution has been
59	demonstrated [19]. The use of lipases to catalyze amide bond formation is an
60	interesting alternative to conventional methods using proteases, as lipases generally
61	do not cleave amide bonds that are able to avoid secondary hydrolysis of peptides. So
62	far, lipase catalyzed synthesis of peptides in an aqueous system has rarely been
63	reported.
64	Hypertension is one of the major risk factors for development of cardiovascular
65	diseases, stroke and the end stage of renal disease [20]. The Angiotensin I converting
66	enzyme (ACE) plays an important physiological role in the regulation of hypertension
67	[21]. Inhibition of ACE activity leads to a decrease in the concentration of angiotensin
68	II and consequently reduces blood pressure [22]. Several dipeptides separated from
69	garlic with tyrosine or phenylalanine residue at the C terminus have been shown to
70	inhibit ACE activity. Among these dipeptides, Phe-Tyr is the most potent ACE

71	inhibitor [23]. The antihypertensive effect of Phe-Tyr has been demonstrated in
72	spontaneously hypertensive rats where blood pressure significantly decreased after
73	oral administration [24].
74	The present work focused on lipase-catalyzed synthesis of dipeptide
75	<i>N</i> -Ac-Phe-Tyr-NH ₂ in an aqueous solution. Our purpose was to better understand the
76	solvent effect and the relationships between reaction variables (reaction time,
77	temperature, enzyme amount, and substrate molar ratio) and the response (yield of
78	<i>N</i> -Ac-Phe-Tyr-NH ₂), as well as to obtain the conditions for dipeptide
79	<i>N</i> -Ac-Phe-Tyr-NH ₂ synthesis using 5-level-4-factor composite rotatable design
80	(CCRD) and response surface methodology (RSM). In addition, a kinetic study of
81	amidation was also performed to determine the apparent kinetic parameters in order to
82	compare the specific activity and specificity of the lipase and α -chymotrypsin as
83	catalysts.
84	2. Experiment
85	2.1. Materials
86	N-acetyl-phenylalanine ethyl ester (N-Ac-Phe-OEt), L-tyrosinamide (Tyr-NH ₂),
87	acetonitrile (ACN), trifluoroacetic acid (TFA), Trizma base buffer (Tris buffer) and
88	porcine pancreatic lipase type II (lipase from porcine pancreas, EC 3.1.1.3, 30-90
89	Units/mg) purchased from Sigma Chemical Co. (St. Louis, MO). All other reagents

90 and chemicals, unless otherwise noted, were of analytic grade.

- 91 2.2. Experimental design
- 92 A four-factor, five-level CCRD consisting of 27 treatments was employed in this
- 93 study. The manipulated (independent) variables and their respective levels selected for
- 94 *N*-Ac-Phe-Tyr-NH₂ synthesis included reaction time (2–10 min), temperature (20–

95 40 °C), enzyme amount (2-10 U) and substrate molar ratio (Tyr :Phe = 1:1-3:1).

- 96 Table 1 shows the independent factors (xi), levels and experimental design, both
- 97 coded and uncoded. All reactions were carried out in duplicate.
- 98 2.3. Lipase catalyzed synthesis of N-Ac-Phe-Tyr-NH₂
- 99 N-Ac-Phe-OEt (50 mM) with various amounts of L-tyrosinamide and porcine
- 100 pancreatic lipase were mixed with 80mM pH 9.0 Tris-HCl buffer in screw-capped
- 101 tubes. The reaction mixture was then placed in an orbital sharking bath (150 rpm) at
- designed reaction temperature for the times shown in Table 1. The reaction of porcine
- 103 pancreatic lipase catalyzed synthesis of *N*-Ac-Phe-Tyr-NH₂ is represented in scheme
- 104 1. After reaction, a 4-fold volume of termination reagent containing ACN and acetic
- acid was added into the reaction mixture to deactivate the enzyme. Twenty microliters
- 106 of the reaction mixture was injected into a high-performance liquid chromatography
- 107 (HPLC) for composition analysis.
- 108 *2.4. Analysis*

109	The samples we	ere analyzed by a Gil	son HPLC System (C	Gilson 322 pump, UV-Vis
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- 110 152 detector; L.M.I. Company) equipped with an ultraviolet detector and a Hypersil
- 111 ODS-2 C18 column (Thermo Instrument Systems, Runcorn, UK, 25 cm 4.6 mm). The
- elution solvents used were 0.1% TFA of water and ACN. The flow rate was set at 1.0
- 113 mL/min and gradient elution was performed as follows: ACN was set at 30% for the
- first 5 min, gradually increased to 70% between 5 and 10 min, and then returned to
- 115 30% for the last 5 minutes. An ultraviolet detector was set at a wave-length of 254 nm.
- 116 The yields were calculated from the peak areas of the substrate and dipeptide.
- 117 2.5. Statistical analysis
- Using SAS software, the experimental data (Table 1) were then analyzed by the
 response surface regression (RSREG) procedure to fit the following second-order
- 120 polynomial Eq. 1:

$$Y = \beta_0 + \sum_{i=1}^4 \beta_i x_i + \sum_{i=1}^4 \beta_{ii} x_i^2 + \sum_{i=1}^3 \sum_{j=i+1}^4 \beta_{ij} x_i x_j$$
(1)

121

where Y is the yield of *N*-Ac-Phe-Tyr-NH₂; β_0 , β_i , β_{ij} , are constant coefficients; *xi* and *xj* are uncoded independent variables. The option of ridge max option in the SAS software was employed to compute the estimated ridge of maximum response for increasing the radius from the center of the original design.

126 2.6. Kinetic study

127 To determine the kinetics of the amidation, reaction mixtures were prepared by a fixed

128 <i>IV</i> -AC-PRE-OEL CONCENTRATION (10 mWL) COMDINED WITH VARIOUS L-tyrosinal
--

- 129 concentrations (5-25 mM). The amount of enzyme was 55 U in Tris buffer at 20 °C
- 130 for 1min. Initial reaction rates were expressed as mM produced *N*-Ac-Phe-Tyr-NH₂

131 per min.

- 132 **3. Results and discussion**
- 133 *3.1. Solvent selection for synthesis of N-Ac-Phe-Tyr-NH*₂

134 The substrates, *N*-Ac-Phe-OEt and Tyr-NH₂, have a reasonable solubility in the

135 aqueous Tris-HCL buffer (80mM, pH=9) and also in the organic solvents: ACN and

136 DMSO. To understand the relation of the solvent effects on the lipase catalyzed

- 137 synthesis of *N*-Ac-Phe-Tyr-NH₂, these solvents were employed to investigate the
- 138 effects of solvent on the dipeptide yield. The solvent compositions and the resulting
- 139 *N*-Ac-Phe-Tyr-NH₂ yields from the tests are listed in Table 2. In the single solvent
- 140 environments, the 60.7% yield of *N*-Ac-Phe-Tyr-NH₂ was obtained in a Tris-HCL
- 141 buffer. However, the dipeptide was not synthesized in pure ACN or DMSO solvent.
- 142 In addition, the yield of *N*-Ac-Phe-Tyr-NH₂ in aqueous-ACN and aqueous-DMSO
- 143 was decreased to 12.6 and 40.6%, respectively. As the pK_2 of tyrosine is at 9.1, this
- 144 means that $\frac{1}{2}$ of the amino acid molecules have lost the H⁺ from the NH₃ group at pH
- 145 9. Therefore, the α -amino group of Tyr-NH₂ contains the nitrogen lone pair that
- 146 attacks the carbonyl carbon atom of *N*-Ac-Phe-OEt-enzyme intermediate to form an

147	amide bond. These results indicate that the aqueous Tri-HCL buffer at pH 9 was
148	required for deacylation of acyl-enzyme intermediate with a nucleophile reactant
149	(Tyr-NH ₂).
150	3.2. Model fitting
151	The RSM combined with a four-factor/five-level CCRD employed in this study was
152	more efficient in reducing the experimental runs and time needed to investigate the
153	relationship between the synthesis conditions and dipeptide yield. The RSREG
154	procedure of SAS was employed to fit the second-order polynomial equation (1) to
155	the experimental data-percent mole conversions (Table 1). Among the various
156	treatments, the greatest mole conversion (83.6%) was treatment 11 (4 min, 25°C,
157	enzyme amount 4 U, substrate mole ratio 2.5:1), and the smallest conversion (only
158	60.5%) was treatment 18 (8 min, 35 $^{\circ}$ C , enzyme amount 8 U, substrate mole ratio
159	1.5:1). From the SAS output of RSREG, the second-order polynomial equation (2) is
160	given below:
161	$Y = 75.23333 - 0.21875X_1 - 0.99583X_2 - 0.47708X_3$
162	$+19.59167X_4 - 0.01X_1X_2 - 0.1125X_1X_3 - 0.025X_1X_4$
163	$-0.03125X_2X_3 + 0.235X_2X_4 + 0.6125X_3X_4 + 0.053646X_1^2$
164	$+2.58333 \times 10^{-3} X_2^2 + 0.013021 X_3^2 - 4.99167 X_4^2 $ ⁽²⁾
165	The analysis of variance (ANOVA) results indicate that this second-order polynomial

166	model was highly significant and adequately represents the actual relationship
167	between the response (percentage molar conversion) and the significant variables with
168	very a small p value (0.0001) and a satisfactory coefficient of determination ($R^2 =$
169	0.9744). Furthermore, the overall effect of the four synthesis variables on the
170	percentage molar conversion was analyzed further by a joint test. The results reported
171	in Table 3 reveal that temperature (X_2) and substrate molar ratio (X_4) are the most
172	important factors, exerting a statistically significant overall effect ($p < 0.0001$) on the
173	response molar conversion.
174	3.3.Mutual effect of parameters
175	The substrate molar ratio and temperature were investigated in the ranges 1:1–3:1 and
176	20-40 $^{\circ}$ C, respectively. Figure 1 shows the effect of varying the substrate molar ratio,
177	temperature and their mutual interaction on N-Ac-Phe-Tyr-NH ₂ synthesis in 4 min
178	using a constant enzyme amount of 4 U. The lowest molar conversion was 60% at the
179	lowest substrate molar ratio of 1:1 and at the highest temperature of 40 $^{\circ}$ C. A reaction
180	condition at substrate molar ratio of 2.5:1 and a temperature of 20 $^{\rm o}{\rm C}$ obtained ~ 85 %
181	mole conversion. These results indicate that both substrate molar ratio and
182	temperature were important parameters. In the N-Ac-Phe-Tyr-NH ₂ synthesis reaction,
183	L-tyrosinamide was competed with water to react with acyl enzyme intermediate.
184	Therefore, overloading of L-tyrosinamide benefitted the reaction process in forming

185	an amidation product and in quicker completion. Figure 2 represents the effect of
186	varying the substrate molar ratio and reaction time on synthesis, with a constant
187	enzyme activity of 2 U at 20 °C. At any substrate molar ratio, the molar conversion
188	was not affected significantly by reaction time. As shown in the results contained in
189	Table 3, that reaction time (X ₁) has a less significant effect ($p > 0.05$) on the synthesis
190	of <i>N</i> -Ac-Phe-Tyr-NH ₂ . The relationships between the reaction factors and the
191	response can be better understood by examining the planned series of contour plots
192	(Figure 3). These were generated from the predicted model (Eq. 2) by holding both
193	the substrate molar ratio (1:1, 2:1, or 3:1) and the enzyme amount (2, 6, or 10 U)
194	constant. Figure 3 A, B and C represent the same substrate molar ratio (1:1); and A, D
195	and G represent the same enzyme activity (2 U). Such an application could be adopted
196	to study the synthesis variables simultaneously in a five-dimensional space. In general,
197	all nine contour plots shown in Fig. 3 exhibit similar behaviors; the predicted molar
198	conversion increased with an increasing substrate molar ratio, whereas the molar
199	conversion decreased with corresponding increases in temperature. Therefore,
200	temperature (X_2) and substrate molar ratio (X_4) were the most important variables for
201	<i>N</i> -Ac-Phe-Tyr-NH ₂ synthesis, with small p values (Table 3), and can be considered as
202	indicators of effectiveness and economic performance.

203 *3.4. Optimum synthesis conditions*

204	The optimum conditions were determined by ridge max analysis. This method
205	computes the estimated ridge of maximum response for increasing radii from the
206	center of original design. The ridge max analysis (Table 4) was verified that the
207	maximum molar conversion of <i>N</i> -Ac-Phe-Tyr-NH ₂ dipeptide was $85.0\pm1.6\%$ at a
208	reaction time of 3.8 min, a temperature of 20.9 °C, a substrate molar ratio of 2.53:1
209	and an enzyme amount of 6.5 U. The validity of the predicted model was tested by
210	repeating the experiments three times. At the suggested optimal synthesis condition,
211	the actual molar conversion of N-Ac-Phe-OEt was 84.40±0.98%. The results indicated
212	that the observed value was almost identical to the predicted value.
213	3.5. Kinetic study
214	α -Chymotrypsin is most used enzyme in peptide synthesis. In order to compare the
215	catalytic efficiency of lipase and α -chymotrypsin in N-Ac-Phe-Tyr-NH ₂ synthesis, the
216	kinetic constants were used as an indicator in this study. Most of lipases catalyzed
217	transesterification has been proposed to follow the Ping-Pong Bi-Bi mechanism [25,
218	26], supposing the initial amount of products to be zero with no inhibition of either
219	substrates or products. The general Ping-Pong Bi-Bi mechanism rate equation derived
220	by Albert is given as [27]:
221	$v_{0} = \frac{V_{m}[A_{0}][B_{0}]}{K_{m}^{B}[A_{0}] + K_{m}^{A}[B_{0}] + [A_{0}][B_{0}]} $ (3)

222 where v_0 is the initial reaction rate, Vm is the maximum rate of reaction, [A₀] is the

223 initial concentration of N-Ac-Phe-OEt, $[B_0]$ is the initial concentration of tyrosinamide, K^{A}_{m} is the Michaelis constant for N-Ac-Phe-OEt, and K^{B}_{m} is the 224 225 Michaelis constant for tyrosinamide. At constant but not saturating $[A_0]$, the equation can be rearranged as Eq. 4, which is the form of the Michaelis–Menten equation. 226 $\boldsymbol{v}_0 = \frac{\boldsymbol{v}_m \boldsymbol{\kappa}_1[\boldsymbol{\mathsf{B}}_0]}{[\boldsymbol{\mathsf{B}}_0] + \boldsymbol{\kappa}_2} = \frac{\boldsymbol{v}_m'[\boldsymbol{\mathsf{B}}_0]}{[\boldsymbol{\mathsf{B}}_0] + \boldsymbol{\kappa}_2}$ 227 (4) where $K_1 = \frac{[A_0]}{K_m^A + [A_0]}$; $K_2 = \frac{K_m^B[A_0]}{K_m^A + [A_0]}$ and $V'_m = V_m K_1$ 228 In previous study, the α -chymotrypsin catalyzed synthesized dipeptide using the 229 230 substrate concentrate of 10 mM N-Ac-Phe-OEt and enzyme amount of 71.5 U [7]. In order to compare the apparent kinetic constants of lipase and α -chymotrypsin, the 231 232 *N*-Ac-Phe-OEt concentration $[A_0]$ was fixed at 10 mM with various tyrosinamide 233 concentrations (5-25 mM) using an enzyme amount of 55 U. At the sufficient enzyme 234 amount situation, increase in enzyme amount increases the reaction rate, but the final 235 molar conversion doesn't increase significantly [28-29]. The Lineweaver–Burk plot of the reciprocal initial rate versus the reciprocal concentration of tyrosinamide was 236 linear (Figure 4). The curve shows good linearity even at a high tyrosinamide (25 mM) 237 238 concentration; this indicates that no tyrosinamide inhibition occurred. In addition, the initial rates were determined by the tyrosinamide concentration [B₀], which was fixed 239 240 at 10 mM with various N-Ac-Phe-OEt concentrations (5-15 mM). That the initial rate increased with increasing N-Ac-Phe-OEt concentrations indicates no N-Ac-Phe-OEt 241

242	inhibition occurred, so Eqs. (3) and (4) can be applied in this study. According to the
243	Ping-Pong Bi-Bi mechanism, the ester substrate N-Ac-Phe-OEt (acyl donor) first
244	combines with enzyme to form a N-Ac-Phe-OEt–enzyme complex, which then
245	intermediately transfers to acyl (N-Ac-Phe) enzyme intermediate and releases ethanol.
246	The acyl enzyme intermediate can then be deacylated by the nucleophilic
247	tyrosinamide (nucleophile) to yield <i>N</i> -Ac-Phe-Tyr-NH ₂ and free enzyme is released.
248	The values of apparent kinetic constant (K_2), apparent V'm, and V'm/ K_2 were 7.74 mM,
249	13.14 mM/min, and 1.69 min for lipase and 11.56 mM, 20.24 mM/min, and 1.75 min
250	for α -chymotrypsin. The V'm is directly proportional to K_{I} , and the K^{A}_{m} is inversely
251	proportional to K_I . As compared with lipase, α -chymotrypsin exhibits a higher $V'm$
252	value, representing its K^{A}_{m} value is less than that of lipase. A lower K^{A}_{m} value
253	indicates a higher affinity of α -chymotrypsin for <i>N</i> -Ac-Phe-OEt. This is probably due
254	to phenylalanine is one of specific amino acid for α -chymotrypsin. On the other hand,
255	the value of the apparent kinetic constant K_2 is also a measure of the strength of the
256	<i>N</i> -Ac-Phe-OEt–enzyme complex; the lower value of K_2 in Eq. 4, indicates the higher
257	affinity of acyl enzyme intermediate for tyrosinamide. As compared with
258	α -chymotrypsin, lipase exhibits a lower K_2 value, thus indicating a higher affinity of
259	acyl lipase intermediate for tyrosinamide. The $V'm/K_2$ is often considered to be a
260	suitable parameter to represent enzyme performance [30]. The apparent kinetic

261	constant $V'm/K_2$ ratio of lipase and α -chymotrypsin were almost the same indicating
262	that the catalytic efficiency is quite close, but the use of lipase in peptide synthesis can
263	avoid secondary hydrolysis of peptides.
264	4. Conclusions
265	In this study, the optimal conditions for lipase catalyzed synthesis of
266	<i>N</i> -Ac-Phe-Tyr-NH ₂ was successfully assessed and demonstrated. The pH plays an
267	important role in peptide synthesis in the aqueous phase. The dipeptide molar
268	conversion can be further enhanced by the central composite design and response
269	surface methodology. A RSM model was obtained to describe the relationship
270	between the molar conversion of N-Ac-Phe-Tyr-NH ₂ and the four parameters of
271	reaction time, temperature, enzyme amount and substrate molar ratio. The results
272	indicated the temperature and substrate molar ratio significantly affected the
273	enzymatic synthesis. An optimal conversion of <i>N</i> -Ac-Phe-Tyr-NH ₂ (around 85%) was
274	observed at a reaction time of 3.8 min, a reaction temperature of 20.9 °C, a substrate
275	molar ratio (Tyr:Phe) of 2.53:1 and an enzyme amount of 6.5 U. In the kinetic study,
276	lipase shows a similar catalytic efficiency as α -chymotrypsin in N-Ac-Phe-Tyr-NH ₂
277	synthesis. The present lipase catalyzed amidation procedure offers some important
278	advantages over the chemical methods or proteases for application in industrial
279	processes.

280

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- 286

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337	Figure captions:	
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- **Scheme 1.** Enzymatic synthesis of *N*-Ac-Phe-Tyr-NH₂ by porcine pancreatic lipase.
- **Figure 1.** Response surface plots showing the effect of substrate molar ratio,
- 340 temperature and their mutual interaction on *N*-Ac-Phe-Tyr-NH₂ synthesis.
- 341 Figure 2. Response surface plots showing the effect of substrate molar ratio,
- 342 reaction time and their mutual interaction on *N*-Ac-Phe-Tyr-NH₂ synthesis.
- **Figure 3.** Contour plots of yields for *N*-Ac-Phe-Tyr-NH2. The number associated
- 344 with the plots indicates the predicted yield at the given reaction condition. Numbers
- 345 within contour plots indicate molar conversions at given reaction conditions.
- 346 SR, substrate molar ratio (Tyr:Phe).
- **Figure 4.** Lineweaver–Burk plot of reciprocal initial reaction rate versus reciprocal
- 348 tyrosinamide (S_0) concentrations at fixed *N*-Ac-Phe-OEt concentration.
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- 350

	Time	Temperature	Enzyme	Substrate	Yield ^b
Treatment	(min)	(°C)	activity	Molar ratio	(%)
NO. ^a			(U)	(Tyr/Phe)	
	\mathbf{X}_1	X_2	X_3	X_4	Y
1	0 (6) ^c	0 (30)	0 (6)	0 (2)	75.3±0.32
2	0 (6)	-2 (20)	0 (6)	0 (2)	81.5±1.25
3	0 (6)	0 (30)	-2 (2)	0 (2)	77.4 ± 0.86
4	0 (6)	0 (30)	2 (10)	0 (2)	72.4±2.36
5	-1 (4)	1 (35)	1 (8)	-1 (1.5)	66.4±1.89
6	1 (8)	1 (35)	-1 (4)	-1 (1.5)	67.6±2.01
7	-1 (4)	-1 (25)	1 (8)	-1 (1.5)	72.2±3.06
8	-1 (4)	1 (35)	-1 (4)	-1 (1.5)	68.3±0.99
9	0 (6)	0 (30)	0 (6)	-2 (1)	57.7±1.54
10	0 (6)	2 (40)	0 (6)	0 (2)	68.4 ± 0.54
11	-1 (4)	-1 (25)	-1 (4)	1 (2.5)	83.6±0.69
12	0 (6)	0 (30)	0 (6)	0 (2)	75.9±0.21
13	1 (8)	-1 (25)	-1 (4)	-1 (1.5)	74.1±0.13
14	1 (8)	-1 (25)	1 (8)	-1 (1.5)	69.6±0.88
15	1 (8)	1 (35)	-1 (4)	1 (2.5)	77.2±2.16
16	-1 (4)	1 (35)	1 (8)	1 (2.5)	$77.0{\pm}2.81$
17	-1 (4)	1 (35)	-1 (4)	1 (2.5)	77.9±1.67
18	1 (8)	1 (35)	1 (8)	-1 (1.5)	60.5±1.53
19	-2 (2)	0 (30)	0 (6)	0 (2)	77.3±1.49
20	-1 (4)	-1 (25)	-1 (4)	-1 (1.5)	75.6±0.12
21	1 (8)	-1 (25)	-1 (4)	1 (2.5)	79.2±0.22
22	0 (6)	0 (30)	0 (6)	2 (3)	81.7 ± 0.88
23	0 (6)	0 (30)	0 (6)	0 (2)	73.2±0.54
24	-1 (4)	-1 (25)	1 (8)	1 (2.5)	81.4±1.64
25	1 (8)	-1 (25)	1 (8)	1 (2.5)	79.8±0.32
26	2 (10)	0 (30)	0 (6)	0 (2)	73.8±0.75
27	1 (8)	1 (35)	1 (8)	1 (2.5)	72.6±1.78

Table 1. Four-factor, five-level CCRD and experimental results of dipeptidederivative yield in response surface analysis.

353 ^a The treatments were run in random order.

354 ^b Yield was the average (\pm SD) of duplicated experiments.

- **Table 2.** Solvent effect on the peptide (*N*-Ac-Phe-Tyr-NH₂) yield from
- 357 *N*-Ac-Phe-OEt and Tyr-NH₂.

Solvent Component(a)	Yield		
Solvent Component(s)	Peptide (%)		
Tris-HCL buffer	60.7		
ACN			
DMSO			
Tris-HCL : ACN(1:1)	12.6		
Tris-HCL: DMSO (1:1)	40.6		
ACN:DMSO (1:1)			

358 — No detected for HPLC analysis.

•						
	Factor	Degrees of freedom	Sum of squares	Prob.>F ^a		
	Reaction $Time(X_1)$	5	38.952315	0.0334		
	Temperature(X ₂)	5	236.735648	< 0.0001		
	Enzyme amount (X ₃)	5	59.029537	0.0079		
	Substrate molar ratio(X ₄)	5	668.997315	< 0.0001		

Table 3. Analysis of variance for joint test for dipeptide derivative.

361 ^aProb. > F, level of significance.

т	production.						
		Estimated	Actual				
	Coded	Response	Response	\mathbf{X}_1	X_2	X_3	X_4
	Radius	(% conversion)	(% conversion)	(min)	(°C)	(U)	(Tyr/Phe)
	0.1	76.0 ± 0.84	79.6±1.01	5.9	29.5	5.9	2.07
	0.4	79.1±0.77	80.4±1.43	5.6	27.6	5.7	2.29
	0.8	82.2±0.83	81.4±0.81	4.8	24.2	5.8	2.46
	1.2	85.0±1.60	84.4±0.98	3.8	20.9	6.5	2.53

363 Table 4. Estimated ridge of maximum response for variable percentages of molar364 production.

365



371 lipase.

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Figure 1. Response surface plots showing the effect of substrate molar ratio,

temperature and their mutual interaction on *N*-Ac-Phe-Tyr-NH₂ synthesis.

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Figure 2. Response surface plots showing the effect of substrate molar ratio,

reaction time and their mutual interaction on *N*-Ac-Phe-Tyr-NH₂ synthesis.







400



402 tyrosinamide (S_0) concentrations at fixed *N*-Ac-Phe-OEt concentration.

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