Production optimization of a heat-tolerant alkaline pectinase from *Bacillus subtilis* ZGL14 and its purification and characterization

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**ABSTRACT**
Alkaline pectinase has important applications in the pretreatment of waste water from food processing and in both the fabric and paper industries. In this study, a 2-level factorial design was used to screen significant factors that affect the activity of alkaline pectinase, and the response surface methodology (RSM) with a Box-Behnken design (BBD) was used to optimize their concentrations. Starch, peptone, KH$_2$PO$_4$ and K$_2$HPO$_4$·3H$_2$O were found to significantly affect the activity of alkaline pectinase. Their optimal concentrations were as follows: 4.68% starch, 1.6% peptone, 0.26% KH$_2$PO$_4$ and 0.68% K$_2$HPO$_4$·3H$_2$O. Under the above conditions, the activity of alkaline pectinase was significantly improved to 734.11 U/mL. Alkaline pectinase was purified to homogeneity with a recovery rate of 9.6% and a specific activity of 52372.52 U/mg. Its optimal temperature and pH were 50°C and 8.6, respectively. The purified enzyme showed strong thermostability and good alkali resistance. In addition, the activity of alkaline pectinase was improved in the presence of Mg$^{2+}$, Cu$^{2+}$, Mn$^{2+}$ and Co$^{2+}$ significantly inhibited its activity. This study provides an important basis for the future development and use of a heat-tolerant alkaline pectinase from *B. subtilis* ZGL14.

**KEYWORDS**
alkaline pectinase; *Bacillus subtilis* ZGL14; characterization; enzymatic properties; optimization; response surface methodology

**Introduction**

Pectic substances are a heterogeneous group of high molecular weight complex acidic structural polysaccharides that consist largely of D-galactopyranosyluronic acid, which is α-(1→4) glycosidically linked to polygalacturonic acid with small amounts of L-rhamnose (2–4%) and various side chains composed of L-arabinose, D-galactose and β-D-xylene.1–4 Recently, a significant interest in the degradation of pectic substances has been generated, which is evident from a vast range of industrial applications, such as the degumming of bast fibers,5–8 the treatment of alkaline pectic waste water,9 and the extraction, clarification and depectinization of fruit juices.10

The enzyme that hydrolyzes pectic substances is broadly known as pectinase, which includes polygalacturonase (exo-polygalacturonase and endo-polygalacturonase), pectin esterase, pectin lyase and pectate lyase on the basis of their modes of action. Pectinase is naturally produced by many organisms, including bacteria, fungi, yeasts, insects, nematodes, protozoan, and plants.11–15 Microbial pectinase is important in phyto-pathological processes, in plant-microorganism symbioses, and in the decomposition of dead plant materials, through which it contributes to the natural carbon cycle.8,16–21 Based on the pH requirement for optimal enzymatic activity, pectinase can be broadly classified into acidic and alkaline pectinase.22–25 Acidic pectinase has been extensively reported for the extraction and clarification of fruit juices and wines,10,26–28 the maceration of vegetables and fruits, the improvement of essential oil extraction and the enhancement of baby-food production,18,29 whereas alkaline pectinase is used for pretreating waste water from the food processing industry that contains pectinaceous waste9 and in the processing and degumming of bast fibers such as ramie (*Bochneria nivea*),8 sunn hemp (*Crotalaria juncea*),8 buel (*Grewia optiva*)12 and jute (*Chorchorus capsularis*).30 It is also widely used in the fabric industry for the retting of plant fibers such as flax, hemp and jute, the biopreparation of cotton fabrics and the enzymatic polishing of jute/cotton blended fabrics; in the paper industry to solve the retention problem in mechanical pulp bleaching and in the treatment of
pulp and paper mill effluents; and for improving the quality of black tea. However, the production of alkaline pectinase at a commercial scale remains quite low due to the paucity of cultures producing high enzyme yields.16,17,31-33

The conventional method for optimizing enzymatic production by the one factor at a time approach involves varying a single independent factor while maintaining the others at a constant level. This one-dimensional approach is laborious and time-consuming, especially for a large number of factors, and it does not consider interactions among factors. An alternative and more efficient approach is the use of statistical methods. Response surface methodology (RSM) involves a full factorial search by examining the simultaneous, systematic and efficient variation of all components, and it is useful for a large number of factors. It uses quantitative data from appropriate experiments to determine and simultaneously solve multivariate equations. It is a collection of statistical techniques that are used to design experiments, build models, evaluate the effects of factors, and analyze the optimum conditions of factors to obtain desirable responses.34-36 It has been successfully used to optimize compositions of fermentation medium, conditions of enzymatic hydrolysis, synthesis parameters for polymers, and parameters for food processes.37-39

In view of the industrial application of alkaline pectinase, the present study was performed to improve the production of a thermostable alkaline pectinase from *B. subtilis* ZGL14 isolated from the soil. 2-level factorial design and RSM were used to identify critical factors and optimize them for the maximum production of alkaline pectinase. The purification and characterization of the alkaline pectinase from fermentation broth were also investigated in detail. This study provides new insight into the future development and use of a heat-tolerant alkaline pectinase from *B. subtilis* ZGL14. To the best of our knowledge, this is the first report regarding the optimization of conditions for the production of a heat-tolerant alkaline pectinase from *B. subtilis* ZGL14 and its purification and characterization.

**Materials and methods**

**Chemicals**

Pectin, barbituric acid, 3,5-dinitrosalicylic acid, N-cyclohexyl-3-aminopropanesulfonic acid and coomassie brilliant blue G-250 were purchased from Sangon Biotech Co., Ltd., Shanghai, China. DEAE-cellulose 52 and Sephadex G-100 were purchased from Boyun Biotech Co., Ltd., Shanghai, China. All other chemicals used in the experiments were of analytical grade and were used as the routine method.

**Microorganism and culture conditions**

*B. subtilis* ZGL14, isolated from the soil, was used as the source of alkaline pectinase in this study. This strain was first grown for 24 h at 37°C and 200 rpm in a 250 mL shake flask containing 50 mL of seed medium. The seed medium contained 1% glucose, 0.3% beef extract, 1% peptone and 0.5% NaCl, pH 8.0. Nine percent of each culture was inoculated into 50 mL of fermentation medium prepared under pre-designed conditions by 2-level factorial design and RSM with a Box- Behnken design (BBD) and was cultured for 72 h at 40°C and 200 r/min. The samples were taken out to analyze the activity of alkaline pectinase. The basic fermentation medium contained 4% glucose, 2% peptone, 0.5% yeast extract, 0.6% K₂HPO₄·3H₂O and 0.3% KH₂PO₄, pH 9.0.

**2-level factorial design**

Nine factors, starch, peptone, K₂HPO₄·3H₂O, KH₂PO₄, temperature, pH, inoculation, liquid volume and fermentation time, were hypothesized to have an effect on the activity of alkaline pectinase. A 2-level factorial design to evaluate the main effect of the factors was performed as described by Bhunia and Dey.40

**Box-Behnken design**

Box-Behnken design was used to optimize the most significant factors to further improve the activity of alkaline pectinase. Each factor was studied at 3 different levels, and a set of 30 experiments was performed. For statistical calculations, the coding of factors was done according to the following equation:

\[
\chi_i = \frac{X_i - X_0}{\Delta X} \quad i = 1, 2, 3, \ldots k
\]  

where \(\chi_i\) is the coded value of an independent factor, \(X_i\) is the actual value of an independent factor, \(X_0\) is the actual value of an independent factor at the center point, and \(\Delta X\) is the step change. To predict the optimal point, a second-order polynomial equation was
fitted to correlate the relationship between factor and response. The model equation used for the analysis is given below:

\[
Y = \beta_0 + \sum \beta_i x_i + \sum \beta_{ij} x_i x_j + \sum \beta_{ii} x_i^2 \\
i = 1, 2, 3, \ldots k
\]  
(2)

where \(Y\) is the predicted response, \(\beta_0\) is the intercept, \(\beta_i\) is the linear coefficient, \(\beta_{ii}\) is the squared coefficient, and \(\beta_{ij}\) is the interaction coefficient. \(x_i, x_j\) represent independent factors in the form of coded values. The accuracy and general ability of the above polynomial model can be evaluated by the determination coefficient \(R^2\). Each design was performed in duplicate, and the mean value was given.

**Determination of the activity of alkaline pectinase**

The activity of alkaline pectinase was determined by the DNS method, using pectin as the substrate.\(^4\)\(^1\) Cultured cells were precipitated by centrifugation at 12000 rpm at 4°C for 20 min. The activity of alkaline pectinase in the supernatant was assayed by measuring the amount of D-galacturonic acid liberated from pectin. The reaction mixture containing 0.5 mL appropriately diluted alkaline pectinase and 0.5 mL of 0.5% pectin (pectin in N-cyclohexyl-3-aminopropanesulfonic acid buffer, pH 8.6) was incubated for 5 min at 50°C, and the end products were quantitated using the 3,5-dinitrosalicylic acid (DNS) reagent.\(^4\)\(^2\) One unit of the alkaline pectinase activity is defined as the enzyme required to liberate 1 \(\mu\)mol of D-galacturonic acid mL\(^{-1}\) min\(^{-1}\) under the assay conditions.

**Purification of alkaline pectinase**

*B. subtilis* ZGL14 was first cultured in a 2000 mL shake flask with 1000 mL of fermentation medium prepared under the optimal conditions obtained by RSM. The purification of alkaline pectinase was performed as described by Bhunia et al.\(^4\)\(^3\) The purity of alkaline pectinase was determined by SDS-polyacrylamide gel electrophoresis (SDS-PAGE) with 12% separation gel and 5% concentrated gel.\(^4\)\(^4\)

**Effect of temperature and pH on the activity of the purified alkaline pectinase**

The optimal temperature for the activity of alkaline pectinase was investigated by incubating 0.5 mL of the reaction mixture for 30 min. The reaction mixture contained 0.2 mL of 20 mM Tris-HCl buffer (pH 8.6), 0.2 mL of 0.5% pectin and 0.1 mL of alkaline pectinase over the range of 25–80°C. The effect of pH on the activity of the purified alkaline pectinase was studied using 0.5 mL of the reaction mixture containing 0.1 mL of alkaline pectinase, 0.2 mL of buffer and 0.2 mL of 0.5% pectin as the substrate at 50°C for 30 min. The buffers used were a 20 mM phosphate buffer pH 6–7.5, a 20 mM Tris-HCl buffer pH 7.5–8.8, and a glycine-NaOH buffer pH 8.5–11. Water was used as a control, and the samples were analyzed in duplicate.

**Effect of metal ions on the activity of the purified alkaline pectinase**

The alkaline pectinase solutions were pre-incubated for 30 min at various temperatures ranging from 40 to 80°C and pH values ranging from 7 to 10 without the substrate and were immediately cooled to 4°C. The activity of alkaline pectinase was determined as described by Tang et al.\(^4\)\(^1\) The control sample was the purified alkaline pectinase solution at 4°C and pH 8.6, and its activity was defined as 100%. The relative activity was calculated as the ratio of the alkaline pectinase activity under a specific condition to the activity of the control sample.

**Statistical analysis**

Design expert, ver 8.0 (Statease Inc., Minneapolis, MN, USA) was used for experimental designs and the regression analysis of experimental data. Statistical analysis of the model was performed to evaluate the analysis of variance (ANOVA). The quality of the polynomial model equation was judged statistically by
the determination coefficient $R^2$, and its statistical significance was determined by $F$-test.

Results and discussion

2-level factorial design

The 2-level factorial design gives an effective screening method for finding the significant factors in one experiment. Table 1 shows 9 independent factors and their concentrations at different coded levels and gives the experimental results of 2-level factorial design for 20 runs using the activity of alkaline pectinase as the response. As shown in Table 1, the activity of alkaline pectinase showed considerable variation depending on the levels of the 9 factors. The regression results are shown in Table 2. Statistical analysis showed that starch, peptone, $\text{K}_2\text{HPO}_4 \cdot 3\text{H}_2\text{O}$, and $\text{KH}_2\text{PO}_4$ had a very significant impact on the activity of alkaline pectinase with the confidence level above 95% ($p < 0.05$).

Box-Behnken design and response surface methodology

Response surface methodology with a Box-Behnken design was used to determine the optimal concentrations of the 4 factors that were chosen by the 2-level factorial design. The coded and real values of the 4 factors and the results of the Box-Behnken design are shown in Table 3. The effects of the 4 factors on the activity of alkaline pectinase were predicted by the following second-order polynomial equation:

$$
Y = 705.29 + 62.24X_1 - 31.14X_2 + 36.24X_3
- 36.61X_4 - 10.84X_1X_2 + 35.8X_1X_3 + 24.96X_1X_4
+ 33X_2X_3 + 15.10X_2X_4 - 8.39X_3X_4
- 49.78X_1^2 - 45.23X_2^2 - 702.53X_3^2 - 34.19X_4^2 \quad (3)
$$

where $Y$ is the predicted activity of alkaline pectinase, $X_1$ is the concentration of starch, $X_2$ is the concentration of peptone, $X_3$ is the concentration of $\text{K}_2\text{HPO}_4 \cdot 3\text{H}_2\text{O}$, and $X_4$ is the concentration of $\text{KH}_2\text{PO}_4$.

The statistical significance of the above equation (3) was checked by $F$-test. The analysis of variance (ANOVA) for the second-order polynomial model is shown in Table 4. The analysis of factors showed that it was a highly significant model, as suggested by the model’s $F$-value and low probability value ($p < 0.0001$). The analysis of factors also showed that the second-order polynomial equation was well adjusted to the experimental data. The low value, 2.41, of the coefficient of variation (CV) indicated a high degree of precision and a good deal of reliability of the experimental values because the coefficient of variation could indicate the degree of precision to which treatments are compared. Usually, a higher CV value indicates that the reliability of the experiment is lower. The precision of a model can be checked by the

| Run | $X_1$ (%) | $X_2$ (%) | $X_3$ (%) | $X_4$ (%) | $X_5$ ($^\circ$C) | $X_6$ (mL) | $X_7$ (mL) | $X_8$ (mL) | alkaline pectinase activity (U/mL) |
|-----|-----------|-----------|-----------|-----------|-----------------|-----------|-----------|-----------|----------------------------------|
| 1   | 5         | 1         | 0.9       | 0.2       | 37              | 9.5       | 8         | 75        | 84                              | 579.14 |
| 2   | 5         | 3         | 0.3       | 0.2       | 37              | 9.5       | 10        | 25        | 84                              | 674.4  |
| 3   | 3         | 3         | 0.9       | 0.2       | 37              | 8.5       | 10        | 75        | 84                              | 447.76 |
| 4   | 5         | 3         | 0.3       | 0.2       | 43              | 8.5       | 10        | 75        | 60                              | 579.36 |
| 5   | 3         | 3         | 0.3       | 0.4       | 37              | 9.5       | 10        | 75        | 60                              | 633.86 |
| 6   | 5         | 1         | 0.9       | 0.2       | 43              | 8.5       | 8         | 25        | 84                              | 669.43 |
| 7   | 3         | 1         | 0.3       | 0.2       | 37              | 8.5       | 8         | 75        | 60                              | 652.34 |
| 8   | 5         | 3         | 0.3       | 0.4       | 37              | 8.5       | 8         | 75        | 60                              | 612.95 |
| 9   | 5         | 1         | 0.9       | 0.4       | 37              | 9.5       | 10        | 25        | 60                              | 259.45 |
| 10  | 3         | 1         | 0.9       | 0.2       | 43              | 9.5       | 10        | 25        | 60                              | 666.44 |
| 11  | 4         | 2         | 0.6       | 0.3       | 40              | 9         | 9         | 50        | 72                              | 684.43 |
| 12  | 5         | 3         | 0.9       | 0.4       | 43              | 9.5       | 10        | 75        | 84                              | 576.34 |
| 13  | 3         | 3         | 0.3       | 0.2       | 43              | 9.5       | 8         | 75        | 60                              | 520.79 |
| 14  | 4         | 2         | 0.6       | 0.3       | 40              | 9         | 9         | 50        | 72                              | 693.69 |
| 15  | 5         | 1         | 0.3       | 0.4       | 43              | 9.5       | 8         | 25        | 84                              | 269.47 |
| 16  | 4         | 2         | 0.6       | 0.4       | 40              | 9         | 9         | 50        | 72                              | 676.32 |
| 17  | 3         | 3         | 0.9       | 0.4       | 37              | 9.5       | 8         | 25        | 60                              | 622.97 |
| 18  | 3         | 1         | 0.9       | 0.4       | 43              | 8.5       | 8         | 75        | 84                              | 610.38 |
| 19  | 3         | 3         | 0.3       | 0.4       | 43              | 8.5       | 10        | 25        | 84                              | 671.99 |
| 20  | 4         | 2         | 0.6       | 0.3       | 40              | 9         | 9         | 50        | 60                              | 699.06 |

$X_1$: starch (%), $X_2$: peptone (%), $X_3$: $\text{K}_2\text{HPO}_4 \cdot 3\text{H}_2\text{O}$ (%), $X_4$: $\text{KH}_2\text{PO}_4$ (%), $X_5$: temperature ($^\circ$C), $X_6$: pH, $X_7$: inoculation (mL), $X_8$: liquid volume (mL) and $X_9$: fermentation time (h)
determination coefficient ($R^2$) and correlation coefficient ($R$). Here, $R^2$ was calculated to be 0.9921, indicating that 99.21% of the variability in the response could be explained by this model. Normally, a regression model with an $R$ value higher than 0.9 is considered to exhibit very high correlation. It was considered that the closer the value of the correlation coefficient is to 1, the better the correlation is between the experimental values and predicted ones. $R$ (0.9821) for the above equation (3) indicated close agreement between the experimental results and the predicted theoretical ones. $p$-values are used to check the significance of each coefficient, which may, in turn, indicate the pattern of interaction among the factors. The smaller the value of $p$ is, the more significant the corresponding coefficient is. It can be seen that $X_1$, $X_2$, $X_3$, $X_4$, $X_1X_2$, $X_1X_3$, $X_1X_4$, $X_2X_3$, $X_2X_4$, $X_3X_4$, $X_1^2$, $X_2^2$, $X_3^2$, and $X_4^2$ were significant, with a small $p$-value ($p < 0.05$).

### 3.3. Response surface plot and contour plot

3D response surface plots and 2D contour plots were drawn to illustrate the interactive effect of each independent factor on the activity of alkaline pectinase, and the optimal concentration of each component required to maximize the activity of alkaline pectinase (Fig. 1). Each figure shows the effect of 2 factors, and the third factor is set at 0. As shown in Fig. 1a and b, when $K_2HPO_4$·$3H_2O$ ($X_3$) and KH$_2$PO$_4$ ($X_4$) were fixed at 0, starch ($X_1$) and peptone ($X_2$) had a significant impact on the activity of alkaline pectinase. The activity of

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**Table 2.** ANOVA for the 2-level factorial design.

| Source            | df | Sum of squares | Mean of square | $F$-value | $p$-value |
|-------------------|----|----------------|----------------|-----------|-----------|
| $X_1$, starch     | 1  | 22936.35       | 22936.35       | 133.44    | 0.0003** |
| $X_2$, peptone    | 1  | 18658.88       | 18658.88       | 108.59    | 0.0005** |
| $X_3$, $K_2HPO_4$·$3H_2O$ | 1  | 2103.37        | 2103.37        | 12.24     | 0.0249*  |
| $X_4$, $KH_2PO_4$ | 1  | 17718.94       | 17718.94       | 103.09    | 0.0005** |
| $X_5$, temperature| 1  | 411.38         | 411.38         | 2.39      | 0.1968   |
| $X_6$, pH         | 1  | 99.75          | 99.75          | 0.58      | 0.4886   |
| $X_7$, inoculation| 1  | 47.85          | 47.85          | 0.28      | 0.5346   |
| $X_8$, liquid volume| 1  | 341.23         | 341.23         | 1.99      | 0.6257   |
| $X_9$, fermentation time | 1  | 432.95         | 432.95         | 2.32      | 0.2316   |
| Lack of fit       |    | 394.12         | 394.12         | 4.03      | 0.1383   |
| Pure error        | 3  | 293.42         | 97.81          |           |           |

*Statistically significant at 95% confidence level ($p < 0.05$).

**Table 3.** Experimental results of the Box-Behnken design.

| Run | $X_1$ | $X_2$ | $X_3$ | $X_4$ | $X_1$ | $X_2$ | $X_3$ | $X_4$ | alkaline pectinase activity (U/mL) |
|-----|-------|-------|-------|-------|-------|-------|-------|-------|---------------------------------|
| 1   | 0     | 0     | 0     | 0     | 2     | 1     | 0.3   | 0.15  | 694.69                         |
| 2   | 0     | 0     | 0     | 0     | 2     | 1     | 0.3   | 0.15  | 712.25                         |
| 3   | −1    | −1    | 1     | −1    | 1.5   | 0.5   | 0.45  | 0.1   | 476.65                         |
| 4   | 1     | −1    | −1    | 1     | 2.5   | 0.5   | 0.15  | 0.1   | 521.01                         |
| 5   | 0     | 0     | 0     | 0     | 2     | 1     | 0.3   | 0.15  | 707.65                         |
| 6   | 0     | −2    | 0     | 0     | 2     | 0     | 0.3   | 0.15  | 587.88                         |
| 7   | 1     | −1    | −1    | −1    | 2.5   | 0.5   | 0.15  | 0.1   | 552.12                         |
| 8   | 0     | 0     | −2    | 0     | 2     | 1     | 0     | 0.15  | 228.56                         |
| 9   | 0     | 0     | 0     | 2     | 2     | 1     | 0.3   | 0.25  | 521.6                          |
| 10  | 0     | 0     | 0     | 0     | 2     | 1     | 0.3   | 0.15  | 689.58                         |
| 11  | −1    | −1    | 1     | 1     | 1.5   | 0.5   | 0.45  | 0.2   | 317.34                         |
| 12  | 1     | 1     | 1     | 1     | 2.5   | 1.5   | 0.45  | 0.2   | 590.61                         |
| 13  | 0     | 0     | 0     | −2    | 2     | 1     | 0.3   | 0.05  | 628.28                         |
| 14  | −2    | 0     | 0     | 0     | 1     | 1     | 0.3   | 0.15  | 387.33                         |
| 15  | 1     | 1     | −1    | −1    | 2.5   | 1.5   | 0.15  | 0.1   | 363.5                          |
| 16  | 1     | −1    | 1     | −1    | 2.5   | 0.5   | 0.45  | 0.1   | 653.44                         |
| 17  | 2     | 0     | 0     | 0     | 3     | 1     | 0.3   | 0.15  | 637.81                         |
| 18  | −1    | 1     | 1     | 1     | 1.5   | 1.5   | 0.45  | 0.2   | 371.83                         |
| 19  | −1    | −1    | −1    | −1    | 1.5   | 1.5   | 0.15  | 0.1   | 398.46                         |
| 20  | 1     | −1    | −1    | 1     | 2.5   | 1.5   | 0.15  | 0.2   | 405.35                         |
| 21  | 0     | 2     | 0     | 0     | 2     | 2     | 0.3   | 0.15  | 473.69                         |
| 22  | 0     | 0     | 0     | 0     | 2     | 1     | 0.3   | 0.15  | 729.48                         |
| 23  | −1    | 1     | 1     | −1    | 1.5   | 1.5   | 0.45  | 0.1   | 468.59                         |
| 24  | −1    | −1    | −1    | −1    | 1.5   | 1.5   | 0.15  | 0.2   | 307.9                          |
| 25  | 0     | 0     | 2     | 0     | 2     | 1     | 0.6   | 0.15  | 374.62                         |
| 26  | −1    | −1    | −1    | −1    | 1.5   | 0.5   | 0.15  | 0.1   | 525.63                         |
| 27  | 1     | −1    | 1     | 1     | 2.5   | 0.5   | 0.15  | 0.4   | 576.12                         |
| 28  | 0     | 0     | 0     | 0     | 2     | 1     | 0.3   | 0.15  | 697.58                         |
| 29  | 1     | 1     | 1     | −1    | 2.5   | 1.5   | 0.45  | 0.1   | 597.01                         |
| 30  | −1    | −1    | −1    | 1     | 1.5   | 0.5   | 0.15  | 0.2   | 399.98                         |
alkaline pectinase was dramatically improved with an increase in starch content from 1.5% to 2.34%, and with a decrease in peptone content from 1.5% to 0.8%. Subsequently, it dropped slightly as the content of starch increased further and the content of peptone decreased further. The 2D contour plot showed that the interaction between starch and peptone was significant. It can thus be concluded

Table 4. ANOVA of the Box-Behnken design.

| Source      | df | Sum of squares | Mean of square | $F$-value | $p$-value |
|-------------|----|----------------|----------------|-----------|-----------|
| Model       | 14 | 5.730E+005     | 40929.98       | 261.17    | <0.0001*  |
| $X_1$       | 1  | 92967.89       | 92967.89       | 593.23    | <0.0001*  |
| $X_2$       | 1  | 23277.15       | 23277.15       | 148.53    | <0.0001*  |
| $X_3$       | 1  | 31519.38       | 31519.38       | 201.12    | <0.0001*  |
| $X_4$       | 1  | 23978.71       | 23978.71       | 153.01    | <0.0001*  |
| $X_1X_2$    | 1  | 1879.44        | 1879.44        | 11.99     | 0.0035*   |
| $X_1X_3$    | 1  | 20502.66       | 20502.66       | 130.83    | <0.0001*  |
| $X_1X_4$    | 1  | 9965.53        | 9965.53        | 63.59     | <0.0001*  |
| $X_2X_3$    | 1  | 17424.66       | 17424.66       | 111.19    | <0.0001*  |
| $X_2X_4$    | 1  | 3646.05        | 3646.05        | 23.27     | <0.0001*  |
| $X_3X_4$    | 1  | 1127.45        | 1127.45        | 7.19      | 0.0171*   |
| $X_1^2$     | 1  | 67979.85       | 67979.85       | 433.78    | <0.0001*  |
| $X_2^2$     | 1  | 56112.34       | 56112.34       | 358.05    | <0.0001*  |
| $X_3^2$     | 1  | 2.883E+005     | 2.883E+005     | 1839.85   | <0.0001*  |
| $X_4^2$     | 1  | 32065.34       | 32065.34       | 204.61    | <0.0001*  |
| Residual    | 15 | 2350.74        | 156.72         |           |           |
| Lack of Fit | 10 | 1285.75        | 128.57         | 0.60      | 0.7678    |
| Pure Error  | 5  | 1064.99        | 213.00         |           |           |
| Cor Total   | 29 | 5.75E+005      |                |           |           |

$^*$Statistically significant at 95% confidence level ($p < 0.05$).

Figure 1. 3D response surface plots and 2D contour plots showing the effect of starch ($X_1$), peptone ($X_2$), K$_2$HPO$_4$·3H$_2$O ($X_3$) and KH$_2$PO$_4$ ($X_4$) on the activity of alkaline pectinase. (A) $Y = f(X_1, X_2)$, (B) $Y = f(X_1, X_3)$, (E) $Y = f(X_1, X_4)$, (G) $Y = f(X_2, X_3)$, (I) $Y = f(X_2, X_4)$ and (K) $Y = f(X_3, X_4)$. 

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from Fig. 1c and d that there was an obvious mutual interaction between starch (X1) and K2HPO4·3H2O (X3) when peptone (X2) and KH2PO4 (X4) were fixed at 0. The activity of alkaline pectinase increased with an increase in starch content and in the concentration of K2HPO4·3H2O. It reached a maximum when the concentrations of the 2 factors were approximately 2.34% and 0.34%, respectively. The 2D contour plot showed that the interaction between starch and K2HPO4·3H2O was significant. Fig. 1e and f show that starch (X1) and KH2PO4 (X4) had a significant mutual interaction. When the content of starch increased from 1.5% to 2.34% and the concentration of KH2PO4 decreased from 0.2% to 0.13%, the activity of alkaline pectinase increased. Fig. 1g and h show an obvious interaction between peptone (X2) and K2HPO4·3H2O (X3). When starch (X1) and KH2PO4 (X4) were fixed at 0, the activity of alkaline pectinase increased gradually with an increase in peptone content from 0.5% to 0.8%, and with an increase in K2HPO4·3H2O concentration from 0.15% to 0.34%. However, higher concentrations led to a decrease in the alkaline pectinase activity. The 2D contour plot showed that the interaction between peptone and K2HPO4·3H2O was significant. When starch (X1) and K2HPO4·3H2O (X3) were fixed at 0, as shown in Fig. 1i and j, the interaction of peptone (X2) and KH2PO4 (X4) was significant. When the concentrations of peptone and KH2PO4 were reduced from 1.5% to 0.8% and from 0.2% to 0.13%, respectively, the activity of alkaline pectinase increased gradually. Thus, lower concentrations were not conducive to improving the activity of alkaline pectinase. From Fig. 1k and l, it can be seen that K2HPO4·3H2O (X3) and KH2PO4 (X4) were significantly positive to improve the activity of alkaline pectinase.

Figure 2. Elution curves of alkaline pectinase and the SDS-PAGE analysis. (A) Elution curve of the DE-52 ion-exchange chromatography purification step of the sample. (B) Elution curve of the Sephadex G-100 size-exclusion chromatography purification step of the sample. (C) SDS-PAGE analysis of alkaline pectinase. Lane M: protein molecular weight marker; lane 1: the sample after PEG20000 concentration; lane 2: the sample after DE-52 ion-exchange chromatography; lane 3: the sample after Sephadex G-100 size-exclusion chromatography.
pectinase. When starch ($X_1$) and peptone ($X_2$) were fixed at 0, an increase in the content of $K_2HPO_4\cdot3H_2O$ and a decrease in the content of $KH_2PO_4$ led to a rise in the alkaline pectinase activity. The activity of alkaline pectinase reached a maximum when the concentrations of $K_2HPO_4\cdot3H_2O$ and $KH_2PO_4$ were 0.34% and 0.13%, respectively.

Table 5. Purification results for alkaline pectinase.

| Steps                      | Total volume (mL) | Total protein (mg) | Enzyme activity (U) | Specific activity (U/mg) | Purification fold | Recovery (%) |
|----------------------------|-------------------|--------------------|---------------------|-------------------------|-------------------|--------------|
| Crude enzyme               | 940               | 364.41             | 665632.8            | 1826.60                 | 1                 | 100          |
| (NH₄)₂SO₄ precipitation    | 100               | 159.76             | 425722.64           | 2664.76                 | 1.46              | 64           |
| PEG20000 concentration     | 22                | 52.31              | 293967.46           | 5619.72                 | 3.08              | 44.2         |
| Cellulose DE-52 chromatography | 4               | 4.05               | 97873.53            | 24166.30                | 13.23             | 14.7         |
| Sephadex G-100 chromatography | 4               | 1.22               | 63894.47            | 52372.52                | 28.67             | 9.6          |

![Figure 3](image_url)

**Figure 3.** Enzymatic properties of the purified alkaline pectinase. (A) Effect of temperature on the activity of the purified alkaline pectinase. (B) Effect of pH on the activity of the purified alkaline pectinase. (C) Thermal stability of the purified alkaline pectinase. (D) pH stability of the purified alkaline pectinase. (E) Effect of different metal ions on the activity of the purified alkaline pectinase.
By using Design Expert 8.0, the optimal concentrations of the 4 key factors predicted by the model were as follows: 4.68% starch, 1.6% peptone, 0.26% KH$_2$PO$_4$, 0.68% K$_2$HPO$_4$·3H$_2$O. Under the above conditions, the predicted maximal activity of alkaline pectinase was 742 U/mL. To verify the predicted value, 3 tests in shake flasks were performed under the optimal medium composition. The mean value of the activity of alkaline pectinase in these experiments was 734.11 U/mL, which is in good agreement with the predicted value.

**Purification of alkaline pectinase**

The purification of alkaline pectinase was performed sequentially using ammonium sulfate precipitation, PEG20000 concentration, DE-52 anion-exchange column chromatography and Sephadex G-100 size-exclusion chromatography. The purification results are presented in Fig. 2 and Table 5. After (NH$_4$)$_2$SO$_4$ precipitation, 56% of the total protein was removed. The residual protein was concentrated by PEG20000, giving 14.4% of the total protein. The concentrated protein was subject to DE-52 anion-exchange column chromatography (Fig. 2a), and the protein from the main peak was collected and pooled, giving 1.1% of the total protein. The protein was subjected to further purification using Sephadex G-100 column chromatography (Fig. 2b). A big peak (fractions 8–16) with higher activity of alkaline pectinase was presented (Fig. 2b). The protein from this peak was collected and pooled, giving 0.3% of the total protein. The fold purification of alkaline pectinase from the above 4 steps was 1.46, 3.08, 13.23 and 28.67, respectively. SDS-PAGE analysis of the purified protein is shown in Fig. 2c. Only a single protein band with an estimated molecular weight of approximately 65 kDa was presented after Sephadex G-100 column chromatography. This indicates that the alkaline pectinase obtained is electrophoretically pure and may be used to analyze its enzymatic properties.

**Enzymatic properties of the purified alkaline pectinase**

The enzymatic properties of the purified alkaline pectinase are shown in Fig. 3. The activity of alkaline pectinase increased when the temperature was increased from 25 to 50°C, and reached a maximum at 50°C. Then, it decreased rapidly as the temperature increased beyond 50°C. This shows that the optimal temperature of the purified alkaline pectinase is 50°C (Fig. 3a). The activity of alkaline pectinase varied as a function of pH. The highest activity of alkaline pectinase was observed at pH 8.6 (Fig. 3b). For alkaline pectinase from *B. subtilis* ZGL14, strong thermal stability was observed at 40 and 50°C. The activity of alkaline pectinase showed a negligible change when incubated at either of these temperatures for 40 min. The residual activity of alkaline pectinase was 51, 40 and 22% when it was incubated at 80°C for 60, 80 and 100 min, respectively (Fig. 3c), which suggested that alkaline pectinase had good thermal stability. High pH stability of alkaline pectinase was observed when it was incubated at pH 8.6 and 10.0. The enzymatic activity declined rapidly to 12.2% of the maximal activity at pH 7.0 for 120 min. When pH was 8.0, the enzymatic activity was 93.9% of the maximum after 60 min. The activity of alkaline pectinase was retained at approximately 49.4% after 120 min at pH 10.0, which indicated that the purified pectinase had good alkali resistance (Fig. 3d).

The effect of different metal ions on the activity of the purified alkaline pectinase is shown in Fig. 3e. It was obvious that metal ions had a certain effect on the activity of alkaline pectinase. The addition of Mg$^{2+}$ could obviously promote the activity of alkaline pectinase, which was increased by 18.9% in the presence of 1 mM Mg$^{2+}$. Ca$^{2+}$ had no significant effect on the activity of alkaline pectinase, while Cu$^{2+}$, Mn$^{2+}$, and Co$^{2+}$ significantly inhibited the activity of alkaline pectinase. Zn$^{2+}$, Fe$^{2+}$ and Ba$^{2+}$ had a moderate inhibitory effect on the activity of alkaline pectinase.

**Conclusions**

The conditions for the production of alkaline pectinase from *B. subtilis* ZGL14 were optimized by RSM. The optimal concentrations of the 4 key factors screened were as follows: 4.68% starch, 1.6% peptone, 0.26% KH$_2$PO$_4$, and 0.68% K$_2$HPO$_4$·3H$_2$O. Under the above conditions, the activity of alkaline pectinase reached 734.11 U/mL. The molecular weight of the purified alkaline pectinase was approximately 65 kDa. The final specific activity of the purified alkaline pectinase was 52372.52 U/mg. The optimal temperature and pH was 50°C and 8.6, respectively. The purified alkaline pectinase had strong thermo-stability and good alkali resistance. This enzyme had an absolute requirement for Mg$^{2+}$ to stimulate its enzymatic activity, but it did not need other metal ions such as Cu$^{2+}$, Mn$^{2+}$, Co$^{2+}$, Fe$^{2+}$, Zn$^{2+}$ and Ba$^{2+}$. The study provides
an important basis for the future industrial production of heat-tolerant alkaline pectinase.

**Disclosure of potential conflicts of interest**

No potential conflicts of interest were disclosed.

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