Research Article
Culture Conditions Affect the Category and Production of Ubiquinones in a Recombinant Escherichia Coli with an Exogenous Decaprenyl Diphosphate Synthase

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Abstract: Ubiquinones (UQ) are important electron transporters and play lot of important roles in most organisms. In different species, UQ was classified to be UQ-6, 7, 8, 9, 10 according to their polyprenyl side chain length. The side chain's length is determined by the enzyme named Poly-Prenyl diphosphate Synthases (PPPS). Bacteria are usually reconstructed to producing UQ-10 used in human's food additive, medicine or cosmetics, such as using decaprenyl Diphosphate Synthase (DPS) gene from *R. radiobacter* to substitute *E. coli*'s octaprenyl diphosphate synthase gene, just like *E. coli* BL21 (Δ*ispB::ddsA*) used in this study. It is interesting that not only in these reconstructed bacteria, but in human-being, DPS can synthesize UQ-9 besides UQ-10. The mechanism of this phenomenon is still unknown. In this study, the effects of culture conditions, including the temperature, dissolved oxygen, pH and culture medium, on the DPS characteristics in *E. coli* BL21 (Δ*ispB::ddsA*) were examined. Results show that temperature greatly affects the ratio of UQ-9/UQ-10, but not the total ubiquinone's production. Increasing dissolved oxygen and protein concentration in culture medium can promote total ubiquinone's production, but not the ratio of UQ-9/UQ-10. These results may give reference for UQ-10's industrial produce and the mechanism of these conditions' effect on DPS will be discussed.

Keywords: Decaprenyl diphosphate synthase, *Escherichia coli*, *Rhizobium radiobacter*, ubiquinone

INTRODUCTION

Ubiquinones (Coenzyme Q, UQ or CoQ) are essential components for organisms. Their basic function is to transfer electrons in respiratory chain from complex I/II to the cytochrome bc_1 complex in the inner membrane of eukaryote mitochondrion or the plasma membrane of prokaryote (Battino *et al*., 1990; Brandt and Trumpower, 1994; Lenaz and Genova, 2009). Additionally, ubiquinones have lots of other important functions as the following: UQ can function as the lipid-soluble antioxidants preventing the oxidation of DNA, proteins and lipids (Maroz *et al*., 2009); UQ regulates the cytosolic ratio of NAD+/NADH ratio involved in cell growth and differention (Gomezdiaz *et al*., 1997); UQ prevents the collapse of mitochondrial function by maintaining the integrity of mitochondrial membrane transition pore (Papucci *et al*., 2003); The oxidized UQ is an obligatory cofactor to assist uncoupling protein in the inner mitochondrial membrane to translocate protons (Echtay *et al*., 2001); UQ influences the expression of NFκB1-dependent genes involved in anti-inflammation (Schmelzer *et al*., 2007); CoQ can protect LDL from

oxidation, which has anti-atherosclerotic properties (Thomas *et al*., 1996; Turunen *et al*., 2002); UQ can stimulate endothelial release of nitric oxide to counteract endothelial dysfunction (Hamilton *et al*., 2007); UQ mediates the disulfide bonds formation in bacterial and fungal proteins (Bader *et al*., 2000). In the past years, the research on UQ synthesis was intensively carried out and it is used as food additive, medicine or cosmetics universally (Clarke, 2000; Littarru and Tiano, 2007).

Ubiquinone consists of a redox active benzoquinone ring (4-hydroxybenzoic acid, 4-HB) attached to a polyprenyl side chain. After covalent bond formation between 4-HB and polypreny tail, the resulting 3-polyprenyl-4-Hydroxybenzoate (HHB) undergoes several modifications such as hydroxylation, methylation and decarboxylation (Kawamukai, 2002). According to the length of poly-prenyl side chain, ubiquinones could be classified into UQ-6, UQ-7, UQ-8, UQ-9 and UQ-10 and different UQ exists in different species. For example, UQ in mice is UQ-9, while in humans UQ-10 is the dominant UQ. In microorganisms, *Saccharomyces cerevisiae* UQ is UQ-6, *E. coli* UQ is UQ-8. The different UQ species

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execute similar functions and could be artificially substituted with each other (Huang *et al*., 2011; Li *et al*., 2012).

The side chain is synthesized by Poly-Prenyl diphosphate Synthases (PPPS), which belong to trans-prenyltransferases family and catalyze chain elongation of Farnesyl Pyrophosphate (FPP) substrates via consecutive condensation reactions with Isopentenyl Pyrophosphate (IPP) to generate linear polymers with defined chain length (Sanchez *et al*., 2006). The FPP is synthesized by other poly-prenyl diphosphate synthase, namely farnesyl diphosphate synthase, from IPP and its isomer dimethaylallyl pyrophosphate (Yoon *et al*., 2006; Gu *et al*., 2011). The crystal structures of some PPPS had been reported (Guo *et al*., 2004; Chang *et al*., 2006; Sasaki *et al*., 2011). Results shows that two Asp-rich motifs, DD $(XX)_n$ D (X represents any amino acid and n = 2 or 4) exist in these enzymes amino sequences, the first one responsible for FPP binding and the second for IPP (Kellogg and Poulter, 1997; Ogura and Koyama, 1998). After condensation, the hydrophobic tail of poly-prenyl chain extends into the cavity of the PPPS. The depth of cavity determines the length of chain. Tarshis *et al*. (1996) showed the side chains of residues corresponding to F112 and F113 of avian FPP synthase are very important for determining the ultimate length of poly-prenyl chain. By site directed mutagenesis, the avian FPP synthase was transformed into synthases capable of producing geranylgeranyl diphosphate (F112A), gernylfarnesyl diphosphate (F113S) and longer prenyl chains (F112A/F113S). By X-ray analysis, the cavity of the F112A/F113S mutant increases in depth by 5.8Å compared with that for the wild-type enzyme. Guo *et al*. (2004) also proved the F132A mutant of octaprenyl diphosphate synthase from *Thermotoga maritima* can produce decaprenyl diphosphate.

The prenyl chains length are believed to be determined by PPPS, but it is interesting that the PPPS in some species can produce more than one prenyl chain with different length. For example, in human, the decaprenyl diphosphate synthase can synthesize UQ-9 besides UQ-10 (Aberg *et al*., 1992). In *E. coli* the octaprenyl diphosphate synthase can synthesize UQ-7 besides UQ-8. In our previous study when *E. coli* BL21 (Δ *ispB*::*ddsA*) was constructed by substituting *E. coli* octaprenyl diphosphate synthase with decaprenyl Diphosphate Synthase (DPS) from *R. radiobacter* to produce UQ-10 (Li *et al*., 2012), we also found that *E. coli* BL21 (△*ispB*::*ddsA*) can synthesize UQ-9 besides UQ-10. What factors affect the synthesis of different length UQ is still unknown.

In this study, to see if the culture conditions affect the production of different length UQ, the ratio of UQ-9 and UQ-10 from *E. coli* BL21 (Δ*ispB*::*ddsA*) cultured under different temperatures, dissolved oxygen, pH and culture medium were tested. This study may give

reference for promoting UQ-10 production in industry. The possible reason why these conditions affect the characteristics of DPS will also be discussed.

MATERIALS AND METHODS

Microbial used in this study was *E. coli* BL21 (Δ*ispB*::*ddsA*) constructed by our lab (Datsenko and Wanner, 2000). The DPS gene was cloned from *R. radiobacter*. Luria-Bertani (LB), tryptone and yeast extract were purchased from Oxoid LTD. Ubiquinone-10 and -9 were purchased from Sigma Chemical Co. (Shanghai) used as standard samples. Ethanol, methanol and *n*-hexane were chromatographic grade purchased from Yongda chemical reagent Co. (Tianjin). All the other chemicals were analytic grade.

Methods: *Escherichia coli* BL21 (Δ*ispB*::*ddsA*) was pre-cultivated with LB medium in shake flask for overnight at 220 rpm and 37°C to be used as seed culture. For exploring the temperature effect on characteristics of DPS *in vivo*, the temperature 24, 27, 30, 33, 36 and 39°C were chosen for test, respectively. Under each temperature, triple 50 mL LB broth (pH 7.2) in 250 mL flasks were inoculated with 2 mL seed culture and cultivated at 220 rpm for 12 h.

For exploring the DO effect on characteristics of DPS *in vivo*, different shaking speed of shaker was chosen. Under 0, 40, 80, 120, 160, 200 and 240 rpm, triple 50 mL LB broth (pH7.2) in 250 mL flasks were inoculated with 2 mL seed culture and incubated for 12 h at 37°C, respectively.

With the initial pH of 6.0, 6.5, 7.0 and 7.5, 2 mL seed culture was transferred into the 50 mL LB medium in 250 mL flask, respectively. The flasks were shaked in a rotary shaker at 220 rpm and 37°C for 12 h.

Peptone was chosen as limited protein nutrition in LB medium. Changing the peptone concentration, nutrition condition effect on characteristics of DPS was studied. As preceding methods, experiments were carried out under different peptone concentration as 0.25, 0.50, 0.75, 1.00 and 1.25%, respectively. Each was repeated three times.

Analysis methods: Cell density of fermentation broth was measured with optical density at 600 nm and converted to dry cell mass (g/L) according to a pre-prepared standard curve. Cells were collected from 50 mL fermentation broth by centrifuging at 3214 g for 8 min and stored at -20°C for UQ extraction. Cells pellet was suspended in 5 mL *n*-hexane and sonicated for 15 min. Cells debris was collected by centrifuge and extracted with *n*-hexane one more time. These tow supernatants were mixed and vacuum evaporation was used to remove the solvent, n-hexane. The dry ubiquinone sample was resolved in 1 mL chromatographic ethanol for further test.

CoQ was detected according to the methods described before (Yen and Shih, 2009). High Performance Liquid Chromatograph (HPLC) was carried out with Prominence LC-20A (SHIMADZU, Janpan) equipped with shodex C18M 4E (250×4.6 mm) column. Mixed liquid of ethanol and methanol (7:3) was used as mobile phase at a flow rate of 1 mL/min. UV detector at 275 nm was used.

RESULTS AND DISCUSSION

The *E. coli* BL21 (Δ*ispB*::*ddsA*) was cultured under different temperatures and the UQ was tested by HPLC. The results were shown in Fig. 1.

In this figure, we can see when the temperature was less than 30°C, only UQ-10 was detected. While when temperature was above 30°C, the UQ-9 began to be synthesized and the ratio of UQ-9 to UQ-10 increased with the rise of temperature. At 39°C, the concentration of UQ-9 is 0.21 mmol/g DCM (mmol per gram dry cell mass), higher than the concentration of UQ-10 (0.12 mmol/g DCM). While the total UQ at different temperature showed no great difference and sustained about 0.33 mmol/g DCM. These results show the temperature may not affect the catalytic efficiency of DPS, while affect the dissociation of product from the enzyme. When the temperature above 30°C, as the synthesis proceeded to be UQ-9, some of them dissociated from DPS enzyme before the last IPP polymerized.

Naturally, there are three kinds of polyprenyl diphosphate synthases, namely cis-, trans-, cis- and trans-mixed (Kera *et al.*, 2012). Cis-forms appear intensively in microorganisms and plants and produce long chain cis-polyprenyl diphosphate with eleven isopentenyl units, such as undecaprenyl diphosphate synthase for peptidoglycan biosynthesis (Teng and Liang, 2012), to thousands units, such as natural rubber (Takahashi *et al.*, 2012). Trans-forms are responsible for biosynthesis of relatively short polyprenyl chains, such as UQ and lycopene (Bhataya *et al.*, 2009). The physiological functions of cis- and trans- mixed are not elucidated until now. About the length determination mechanism of polyprenyl chain, all researches were based on the structures of the synthases. About the trans-form, it was been reported that the length of polyprenyl chain of UQ is determined by the depth of the cavity formed by two DPS homo-dimmer subunit. The bulge side chains of two Phe in this enzyme cavity stop the elongation of the polyprenyl tail and then the products dissociated from the enzyme. But this cannot be used to explain why the same synthase can produce different length UQ *in vivo*, such as that in human tissues; UQ-9 occupies 2-5% of total ubiquinone. Then the environment conditions may be the most suspect factors.

In our study, we found the *E. coli* BL21 (Δ *ispB*::*ddsA*), which octaprenyl diphosphate synthase gene was replaced by *ddsA* from *R. radiobacter*, could produce UQ-9 besides UQ-10 under temperature over 30°C. Schmelzer *et al.* (2007) had reported that CoQ_{10} affects NFκB-related genes' expression (such as IL5, thrombin, vitronectin receptor and cAMP receptor protein) and CoQ_{10}-inducible genes play an important role in inflammatory response. There may be an implication that human's temperature rise in inflammation situation affects DPS' UQ synthesis and that the ratio change of UQ-9/UQ-10 affects inflammatory response. The ratio of UQ-9/UQ-10 may be looked as a sign of inflammation of body. All proteins' functions were connected with them structures tightly. We can conclude that the high temperature would affected the structure of enzyme active center and promote the UQ-9'dissociation. But no such obvious phenomenon was reported about DPS from other species, such as *Agrobacterium tumefaciens* (Ha *et al.*, 2007), *Rhodobacter sphaeroides* (Zahiri *et al.*, 2006), *Gluconobacter suboxydans* (Okada *et al.*, 1998) and so on. These mean DPS from different species have different stabilities. DPS from *R. radiobacter* is more unstable than others and more sensitive to temperature. Our result also indicated that temperature is important factor in bacteria producing UQ-10 in industry. From the stability of molar sum of ubiquinones content, it can be concluded that the feedback regulation of ubiquinone synthesis is irrelevant with the kind of ubiquinones.

According to the curve in Fig. 1, the following equations can be got to show the relationship between the concentration of UQ9 (y, mmol/g DCM), UQ10 (x, mmol/g DCM) and the temperature (T, °C):

$$\begin{cases} y = 0.0243T - 0.7273 \\ x = -0.0236T + 1.0386 \\ y + x = 0.335113 \end{cases} \tag{1}$$

Ubiquinone is important in electron transport of cell respiration. DO concentration's effect on *E. coli* BL21 (Δ*ispB*::*ddsA*) producing ubiquinones at 37°C was shown in Fig. 2.

We can see with the increase of DO concentration, both the production of UQ-9 and UQ-10 in *E. coli* BL21 (Δ*ispB*::*ddsA*) increase. These results indicated increasing DO concentration in these bacteria promotes the bacteria's respiration and the ubiquinone production increases. The production of total ubiquinone was promoted in *E. coli* BL21 (Δ*ispB*::*ddsA*) with OD concentration increasing, while the ratio of UQ-9/UQ-10 remain steadily, which indicates OD concentration does not affect the DPS catalysis properties.

Yoshida *et al.* (1998) reported *R. sphaeroides* could increase yield of ubiquinone 10 under limited supply of air, but *R. radiobacter* is quite the opposite. It has been reported when DO concentration increases in a certain range, the $CoQ10$ production decreases in *R. radiobacter* just like our results (Wu *et al.*, 2003).
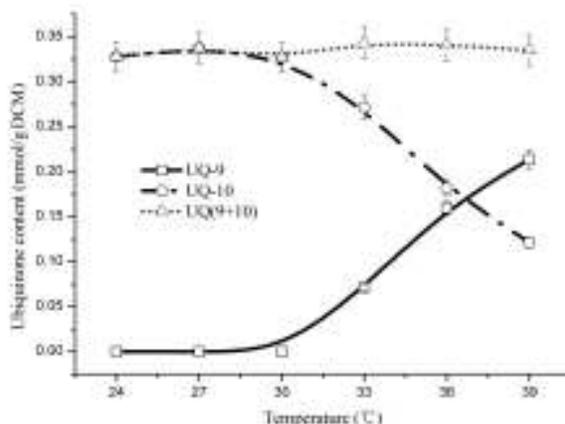
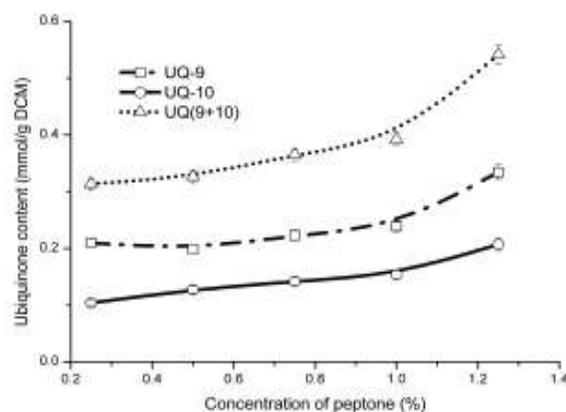Fig. 1: Temperature effect on characteristics of DPS from *R. radiobacter* in *E. coli*
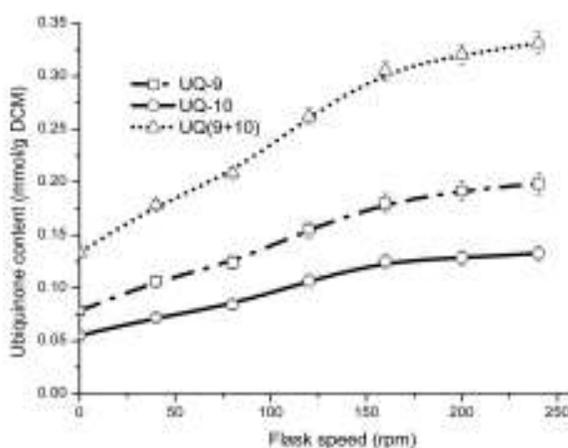

Fig. 2: DO concentration effect on characteristics of DPS from *R. radiobacter* in *E. coli*
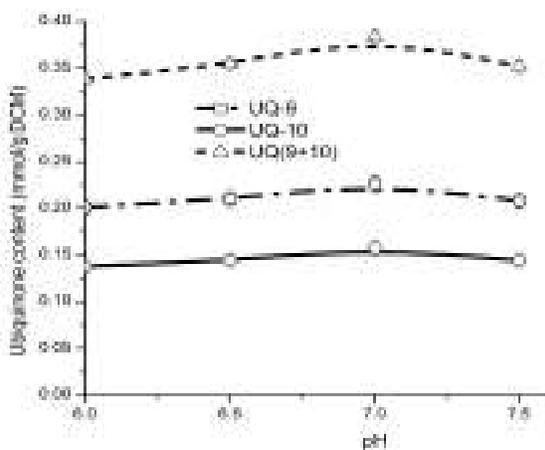

Fig. 3: pH effect on characteristics of DPS from *R. radiobacter* in *E. coli*

Both *R. radiobacter* and *E. coli* are facultative anaerobe and have two kinds of respiratory pathway. In strictly

Fig. 4: Concentration of peptone effect on characteristics of DPS from *R. radiobacter* in *E. coli*

aerobic bacterium, it compensates oxygen lack by adding the respiratory chain and leads to ubiquinone increasing. In facultative anaerobes, it can compensate the oxygen lack by anaerobic respiration without ubiquinone increasing. On the contrary, ubiquinone will decrease for aerobic respiration demand reduction.

The effects of pH and peptone concentration on the production of UQ-9 and UQ-10 at 37°C were respectively shown in Fig. 3 and 4. Results showed both pH and peptone concentration did not apparently affect the ratio of UQ-9/UQ-10. pH is an important factor for enzyme activity. In this study, culture medium's pH may not affect the pH inner cell. So the pH of culture medium seems no affect to DPS enzyme properties. Increasing the concentration of peptone can increase both UQ-9 and UQ-10 production (Fig. 4), but this factor may not affect DPS enzyme properties as pH or they have no direct relationship with intracellular enzymes' activity or stability.

CONCLUSION

In summary, our results show that higher temperature above 30°C may cause the earlier release of nonaprenyl diphosphate from DPS and reduce the production of decaprenyl diphosphate. Increasing dissolved oxygen and protein concentration in culture medium can promote total ubiquinone's production, but not the ratio of UQ-9/UQ-10. These results indicate the fermentation of UQ-10 industrial produce ought to be control at an appropriate temperature to yield higher UQ-10 ratio and increasing DO and protein supplement may help the higher yield of total ubiquinones.

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