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Genetic cloning and functional expression in *Escherichia coli* of an archaerhodopsin gene from *Halorubrum xinjiangense*

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Abstract Pairs of PCR primers that targeted the archae/ bacteriorhodopsin gene were used to clone the archaerhodopsin (aR) gene of Halorubrum xinjiangense strain BD-1¹, and this gene was sequenced and functionally expressed in Escherichia coli. Recombinant E. coli cells harboring the plasmid carrying this gene became slightly purple or blue depending on whether they were supplemented with all- trans retinal or 3,4-dihydroretinal, respectively, during induction with IPTG. The purple and blue membranes from the recombinant E. coli showed maximal absorption at 555 and 588 nm, respectively, which are different from maximal absorption at 568 nm of the wild-type purple membrane. Purple membranes from the recombinant E. coli and from strain BD-1^T were investigated in parallel. The *E. coli* purple membrane was fabricated into films and photoelectric responses were observed that depended on the light-on and light-off stimuli.

Keywords Archaerhodopsin · Bacteriorhodopsin · Purple membrane · *Halorubrum*

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Introduction

Bacteriorhodopsins (bR) are of interest because they are potentially useful for the fabrication of biomaterialbased devices such as artificial retinas (Frydrych et al. 2000) and optical memories (Wise et al. 2002). Khodonov et al. (1997) listed many advantages of bR as material for fabrication of devices, and those advantages could be further improved by modifications either of the protein (Wise et al. 2002; Weetall et al. 2000), or of the chromophore (Druzhko and Chamorovsky 1995; Jussila et al. 2001; Drachev et al. 1989). Modification of the chromophore has been done by the replacement of the retinal with various retinal analogs, e.g., 3,4-dihydroretinal (Khodonov et al. 1997) or 4-keto-retinal (Druzhko and Chamorovsky 1995). The procedure for such modification includes usually, bleaching of the purple membrane, and reconstitution in the presence of the retinal analog.

A bR-like protein, archaerhodopsin (aR) from the *Halorubrum* sp. was identified (Mukohata et al. 1988; Sugiyama et al. 1989; Uegaki et al. 1991). This protein shows about 56–59% homology to the bR protein from *Halobacterium* spp., and exhibits similar photochemical properties. But aR and bR molecules showed differences in their absorbance maxima, the kinetics of the photocycles, and especially in alkaline-induced red-shifted absorption. In this study, we attempted to identify the *aR* in *Halorubrum xinjiangense* strain BD-1^T, and further functionally express the gene and modify the gene product in *Escherichia coli*.

Materials and methods

Bacterial strains, plasmids, medium, and cultivation

H. xinjiangense strain BD-1^T (CGMCC $1.3527^{T} = JCM$ 12388^T) was grown at 40°C in the medium as described previously (Feng et al. 2004). *E. coli* strains (Table 1)

Table 1	Strains,	plasmids,	and	primers	used	in	this s	study
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	Characteristics or sequences	Sources (references) and notes
Bacterial/archael		
Halorubrum strain BD-1 ^T	From lab culture collection	Feng et al. (2004)
E. coli strain JM109	Cloning host	Novagen
<i>E. coli</i> strain BL21 Plasmids	Expressing host	Novagen
pGEM-T easy vector	Cloning vector	Promega
pGEMaR-1	pGEM-T carrying a fragment of aR gene from strain BD-1 ^T	This study
pGEMaR-2	pGEM-T vector carrying the entire aR gene from strain BD-1 ^T	This study
pET28a	Expression vector	Novagen
pET28aR Primers	pET28a carrying the <i>aR</i> gene of strain BD-1	This study
1	5'GAC TGG (CT)TG TTC AC(GC) AC(AG) CC 3'	Primers 1 and 2 are general primers of screen bR-like genes. Primers 3 and 4 targeting the sequence of <i>Halorubrum</i> sp. aus-1 between -11 and 795. Primers 5 and 6 targeting the sequence of strain BD-1 ^T between 1 and 777. The underlined are <i>NcoI</i> and <i>Bam</i> HI sites, respectively
2	5'A(CG)G TC(GA) A(GT) (GC) ACC AT 3'	
3	5'(A,T)(C,A,G)G A(A,T)(G,A) TGG G(T,C)A TGG AC 3'	
4	5'(C,G)GC (T,C)(A,T)G CCG ATC AGT C 3'	
5 6	5'CAT G <u>CC ATG G</u> AC CCG ATAGCG CT 3' 5'CGC <u>GGA TCC</u> GGC CAG CCG ATC AGT 3'	

were routinely grown at 37°C in Luria–Bertani (LB) medium. *E. coli* strains JM109 and BL21(DE3) (Novagen) and plasmids pGEM-T (Promega, WA, USA) and pET28a (Novagen) were used as hosts and vectors for gene sequencing and expression, respectively. When required, 100 mg of ampicillin per liter or 50 mg of kanamycin per liter was added to the culture medium.

DNA extraction and restriction enzyme treatment

Genomic DNA and plasmid DNA extraction, and restriction enzyme treatment on plasmid DNA and amplified DNA fragments were carried out according to Sambrook and Russell (2001).

Cloning and sequencing the aR gene from strain BD-1^T

A fragment of and the entire aR gene were PCRamplified with two pairs of primers, (Primers 1, 2 and 3, and 4, Table 1). The PCR was carried out at 30 cycles under the following conditions: denaturation for 1 min at 94°C, annealing for 1 min at 54°C, and extension for 1 min at 72°C. The amplified DNA fragments were inserted into pGEM-T easy vectors and the generated pGEMaR-1 and pGEMaR-2 were transformed into *E. coli* JM109. DNA sequences were determined by Beijing Genome Institute (Huada Corp., Beijing, China). Functional expression of aR gene from strain BD-1^T in *E. coli* in the presence of all- *trans* retinal and 3,4-dihydroretinal

The PCR-amplified *aR* gene from strain BD-1^T was ligated into pET28a at the sites of *NcoI* and *Bam*HI. The resulting plasmid, pET28aR, was transformed into *E. coli* BL21(DE3) by electroporation at the following conditions: 25 μ F, 12.5 kV/cm, and 200 Ω (ECM630, BTX, USA). Synthesis of aR proteins in recombinant *E. coli* cells harboring pET28aR were induced by the addition of 1 mM IPTG and 10 μ M all- *trans* retinal or 3,4-dihydroretinal when the culture reached OD₆₀₀ of 0.4–0.6. After further continuous cultivation for 2 h, cells were harvested by centrifugation at 10,000 g for 5 min at 4°C.

Preparation of a R membranes from strain $BD-1^T$ and recombinant *E. coli* cells

The aR membranes of strain $BD-1^{T}$ were fractionated by sucrose density gradient centrifugation, according to the method described by Oesterhelt and Stoecknius (1974). Purification of aR-membrane fractions from recombinant *E. coli* were performed in the same way except that the recombinant *E. coli* cells were broken up by sonification at 4°C (160 W, 3 s sonifying vs. 5 s break, 99 cycles) in Tris–HCl buffer (50 mM Tris–HCl, 5 mM MgCl₂, pH 8.0).

Measurement of absorption spectra

The membranes were suspended in a buffer of 50 mM Tris-HCl (pH 8.0). Before recording the absorption of spectra, membrane suspensions were light-adapted for 20 min. Absorption spectra were recorded with a Vis-UV spectrophotometer (Beckman Coulter DU800) with wavelength interval of 0.5 nm in the visible range (400–700 nm). Data were treated using origin 6.0 Software (Microcal Software Inc., Northampton, MA, USA).

Determination of photoelectric properties of fabricated films made from aR-membrane fractions of strain $BD-1^{T}$ and recombinant *E. coli*

Indium-tin-oxide (ITO) glass slide after negativecharged treatment was immersed into Poly(allylamine hydrochloride) (PAH) aqueous solution (2 mg/mL, pH 6.4) for 5 min, rinsed with doubly distilled water, and then dried by nitrogen flow. This modified glass slide was immersed in aR-membrane suspension (pH 9.4) for 5 min, rinsed with doubly distilled water, and then dried with nitrogen again. In this way, we obtained one bilayer of aR-membrane/PAH (M/PAH) films that were marked as (M/PAH)₁. This process was repeated six times. The ITO glass with M/PAH films was used as a working electrode and platinum wire as a counter electrode. The electrolyte solution was 0.5 M KCl, with pH 7.3. To test the photoelectric property of

Chemicals and reagents

DNA restriction enzymes, DNA ligase, and DNA polymerase were purchased from Takara or Promega. All- *trans* retinal was purchased from Sigma. The 3,4-dihydroretinal was synthesized from all- *trans* retinal by following the method of Drachev et al. (1989). PAH was purchased from Aldrich Chemicals.

The Genbank accession numbers

The Genbank accession number for the aR gene of strain BD-1^T is AY510709. Other Genbank numbers of aR or bR proteins are given in Fig. 1.

Results

Genetic cloning and characterization of aR gene from strain BD-1^T

With Primers 1 and 2 (Table 1), a DNA fragment of 388 bp was PCR-amplified from genomic DNA of strain

	Helix A	Helix B	-
Halorubrum xinjiangenseMDPIALQAGYDLLGDHalorubrum sp.aus1MDPIALTAAVGADLLGDHalorubrum sodomenseMDPIALQAGYDLLGDHalorubrum sp.aus2Haloacterium salinarumMLELLPTAVEGVS-QAQITHaloarcula argentinensisMPEHaloarcula japonicaMPEHaloarcula sp.arg2MLQSGMSTYVHaloterrigena sp.arg4MCCAALAPPMAAT	GRPETLWLGIGTLLMLIGTFYFIVKGWGVT GRPETLWLGIGTLLMLIGTFYFIVKGWGVT GRPETLWLGIGTLLMLIGTFYFIARGWGVT GRPETLWLGIGTLLMLIGTFYFIARGWGVT GRPEWIWLALGTALMGIGTLYFIVKGMGVS PGSEAIWLWLGTAGMFLGMLYFIARGWGET PGGESIFLWVGTAGMFLGMLYFIARGWSVS VGPESIWLWIGTIGMTLGTLYFVGRGRGVF	DKEAREYYSITILVPGIASAAYLSM DKEAREYYSITILVPGIASAAYLSM DKDAREYYAVTILVPGIASAAYLSM DKEAREYYAITILVPGIASAAYLAM DPDAKKFYAITTLVPAIAFTMYLAM DSRRQKFYIATILITAIAFVNYLAM DSRRQKFYIATILITAIAFVNYLAM DQRRQKFYIATIMIAAIAFVNYLSM DRKMQEFYIITIFITTIAAAMYFAM	IFFGIGLTEVQVGS- IFFGIGLTEVQVGS- IFFGIGVTEVELASG ILLGYGLTMVPFGG- IALGFGLTIVEFAG- IALGFGLTIVEFAG- IALGFGVTTIELGG- IALGFGVTEVMVGD-
Helix C Helix I	DHe	lix E	Helix F
EMLDIYYARYADWLFTTPLLLLDLALLAKVDRVSIGTLVGVDALM EMLDIYYARYADWLFTTPLLLLDLALLAKVDRVSIGTLVGVDALM EMLDIYYARYADWLFTTPLLLLDLALLAKVDRVTIGTLIGVDALM TVLDIYYARYADWLFTTPLLLDLALLAKVDRVTIGTLIGVDALM EQNPIYWARYADWLFTTPLLLDLALLVDADQGTILALVGADGIM EEHPIYWARYSDWLFTTPLLLYDLGLLAGADRNTITSLVSLDVLM EEHPIYWARYSDWLFTTPLLLYDLGLLAGADRNTIASLVSLDVLM EERAIYWARYTDWLFTTPLLLYDLALLAGADRNTIYSLVGLDVLM EALTIYWARYADWLFTTPLLLLDLSLLAGANRNTIATLIGLDVFM	IVTGLIGALSHTPLARYTWWLFST IVTGLIGALSHTPLARYTWWLFST IVTGLIGALSHTAIARYSWWLFST IVTGLIGALSKTPLARYTWWLFST IGTGLVGALTKVYSYRFVWWAIST IGTGLVATLSPGSGVLSAGAERLVWWGIST IGTGLVATLSAGSGVLSAGAERLVWWGIST IGTGALATLSAGSGVLPAGAERLVWWGIST IGTGAIAALSSTPGTRIAWWAIST	ICMIVVLYFLATSLRAAAKERGPEV ICMIVVLYFLATSLRAAAKERGPEV ICMIVVLYFLATSLRSAAKERGPEV IAFLFVLYYLLTSLRSAAKRSEEV AAMLYILYVLFFGFTSKAESMRPEV AFLIVLLYFLFSSLSGRVADLPSDT GFLLVLLYFLFSSLSGRVADLPSDT GFLLVLLYFLFSNLTDRASELSGDI GALLALLYVLVGTLSENARNRAPEV	ASTFNTLTALVL VASTFNTLTALVL VASTFNTLTALVA VASTFKVLRNVTV VRSTFKTLRNLVT VRSTFKTLRNLVT VRSTFKTLRNLVT VRSTFKTLRNLVL VASLFGRLRNLVI
Helix F Helix G	_		

VLWTAYPILWIIGTEG-AGVVGLGIETLLFMVLDVTAKVGFGFILLRSRAILGDTEAPEPSAGAEASAAD-(J05165) VLWTAYPILWIIGTEG-AGVVGLGIETLLFMVLDVTAKVGFGFILLRSRAILGDTEAPEPSAGADVSAAD-(D50848) VLWTAYPILWIVGTEG-AGVVGLGIETLAFMVLDVTAKVGFGFVLLRSRAILGETEAPEPSAGADASAAD-(S56354) VLWSAYPVVWLIGSEG-AGIVPLNIETLLFMVLDVSAKVGKGLILLRSRAIFGEAEAPEPSAGDGAAATSD (D11058) VVWLVYPVWWLIGTEG-IGLVGIGIETAGFMVIDLTAKVGFGIILLRSHGVLDG-AAETTGTGATPADD--(D31880) VVWLVYPVWWLIGTEG-LGLVGIGIETAGFMVIDLTAKVGFGIILLRSHGVLDG-AAETTGAGATATAD--(AB029320) VLWLVYPVLWLVGTEG-LGLVGLPIETAAFMVLDLTAKIGFGIILLQSHAVLD--EGQTASEGAAVAD---(S76743)(AB009620) ALWFLYPVVWILGTEGTFGILPLYWETAAFMVLDLSAKVGFGVILLQSRSVLER-VATPTAAPT-

Fig. 1 The alignment of nine archaerhodopsin and bacteriorhodopsin proteins that function as proton pump. The names of the bacterial species that the archaerhodopsin and bacteriorhodopsin isolated are given at the *beginning* and Genbank accession numbers of these proteins are provided at the *end*. All conserved amino acid residues are shaded in *gray*

BD-1^T. This fragment showed 93% identity to a part of the aR gene of Halorubrum aus-1, indicating that an aR gene existed in strain BD-1^T. A second pair of primers (Primers 3 and 4, Table 1) according to the aR gene sequence of Halorubrum aus-1 was synthesized and used for amplification of the entire aR gene from strain BD- 1^{T} . The amplified entire *aR* gene of strain BD- 1^{T} was 777 bp in length, and encoded a protein of 258 amino acids. When this protein was aligned to other known aR or bR proteins, all the amino acid residues that were previously revealed to be essential for proton transport and linkage to retinal were completely conserved (Fig. 1). The aR protein from strain $BD-1^{T}$ showed high identity (86-95%) to the aRs of Halorubrum aus-1, aus-2, and Halorubrum sodomense, and showed relatively low identities to the bacteriorhodopsin (55%) of Halobacterium salinarum(Dunn et al. 1981), to the cruxrhodopsins of *Haloarcula* spp. (38–47%, Yatsunami et al. 1997; Kitajima et al. 1996; Tateno et al. 1994; Otomo et al. 1992), and to the archaerhodopsin of Haloterrigena (49%, Ihara et al. 1999).

Expression of *aR* gene and absorption spectra of aR-membranes from *E. coli* cells

The entire *aR* gene was PCR-amplified with primers 5 and 6 (Table 1) from strain BD-1^T and was ligated to pET28a. The resulting plasmid, pET28aR, was electroporated into *E. coli* cells. Depending on the presence of all- *trans* retinal or 3,4-dihydroretinal, recombinant *E. coli* cells that harbored pET28aR became purple (with all- *trans* retinal) or grayish blue (with 3,4-dihydroretinal) during cultivation in LB broth and induction with IPTG. The membrane fractions were isolated from the recombinant *E. coli* cells and their absorption spectra were determined (Fig. 2). The maximal absorption of the purple membrane obtained with all- *trans* retinal was at 555 nm and of the blue membrane with 3,4-dihydroretinal was 588 nm (Fig. 2), which were different from



Fig. 2 Absorption spectra of purple (*solid curve*) and blue (*broken curve*) membranes from recombinant *E. coli*

the maximal absorption at 568 nm of purple membrane from halophilic archaea (Fig. 2 and also Lukashev et al. 1994). The difference in maximal absorption of purple membrane from recombinant *E. coli* and from *Halorubrum* sp. was attributed to the monomer state of archaerhodopsin in *E. coli* and the trimer state in wild membrane, as revealed by Corcelli et al. (2002).

Photoelectric response of film made from wild and recombinant purple membranes

To demonstrate that the aR proteins synthesized in recombinant *E. coli* are active for proton transport, the purple membranes from both wild strain BD-1 and recombinant *E. coli* were used to fabricate thin films (Chu et al. 2003). Upon illumination, the film generated electric currents due to proton movement across the film. Figure 3 shows the photoelectric response profiles of the films that were made of purple membranes from either strain BD-1^T (Fig. 3a) or recombinant *E. coli* (Fig. 3b). Positive and anodic responses were corresponding to light-on and light-off photocurrents that were caused by proton release and uptake, respectively. This provided further evidence that the aR proteins in the recombinant *E. coli* membrane were correctly folded and functionally active.



Fig. 3 Photoelectric response of films made from purple membranes of strain BD-1^T (**a**) and from recombinant *E. coli* (**b**)

Discussion

Heterologous expression of bR proteins in E. coli had been previously studied, e.g., Dunn et al. (1987) reported the synthesis and purification of retinal-free bacteriorhodopsin and Hohenfeld et al. (1999) reported the purification of histidine-tagged bR from recombinant E. coli. But to obtain active aR protein or functional membrane, it was necessary to refold the bR protein, in the presence of retinal, and reconstitute with polar lipid to form purple membrane (Dunn et al. 1987; Hohenfeld et al. 1999). Active expression of proteorhodopsin gene from uncultured proteobacteria and phoborhodopsin (a photosensory protein) gene from Natronobacterium *pharaonis*(NCIMB 2191) was reported (Béjà et al. 2000: Shimono et al. 1998). In this study, we had succeeded in the construction of an E. coli system that produces in one-step the active aR and purple membrane. Moreover, with this system, one aR with all- trans retinal and one aR analog with 3,4-dihydroretinal were synthesized in E. coli. Purple and blue membranes were obtained. We believe that preparation of other aR analogs with different retinal analogs, such as 3-hydroxyretinal and 4-ketoretinal, with this system is possible.

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