

研究论文

保守GTP酶ObgE的分子伴侣活性研究

薛佳莹¹, 关洪斌¹, 秦燕²

1. 山东大学威海分校海洋学院, 山东威海 264209;
2. 中国科学院生物物理研究所生物大分子国家重点实验室, 北京 100101

摘要:

分子伴侣 (molecular chaperone) 能够帮助新生多肽链或错误折叠的蛋白质形成天然构象, 但本身又不是成熟蛋白质的组成成分。蛋白质需要分子伴侣的帮助, 才能够从核糖体合成的新生肽链折叠成有生物活性的大分子。*E. coli*的ObgE蛋白是保守的GTP酶, ObgE蛋白参与信号转导、蛋白运输和细胞周期调控, 并与*E. coli*在氨基酸饥饿下的应激反应有关。本实验通过分子克隆, 将*E. coli* ObgE蛋白的基因克隆到表达载体pET-28a中, 转化到*E. coli* BL21进行蛋白表达纯化。纯化后的ObgE蛋白通过柠檬酸合成酶变复性实验、 α -葡萄糖苷酶变复性实验、牛碳酸酐酶变复性实验, 检测ObgE蛋白的分子伴侣活性, 发现ObgE具有一定的分子伴侣活性, 为该蛋白的研究应用奠定了基础。

关键词: ObgE蛋白 分子伴侣 基因克隆 蛋白纯化

Chaperone Activity of Conservative GTPase ObgE

XUE Jiaying¹, GUAN Hongbin¹, QIN Yan²

1. Marine College of Shandong University, Weihai, Shandong 264209, China;
2. National Laboratory of Biomacromolecules, Institute of Biophysics, The Chinese Academy of Sciences, Beijing 100101, China

Abstract:

Chaperone can facilitate the nascent peptide and denatured protein to fold to the native state but do not occur in these structures when the structures are performing their normal biological functions. Peptides need the help of chaperones to fold to the nature state when they are produced from the ribosomes. Protein ObgE from *E. coli* is a highly conserved GTPase. ObgE participates in cell signaling, transportation of protein and cell cycle regulation. In present work, we cloned the DNA of *E. coli* ObgE into vector pET-28a, expressed in *E. coli* BL21, and then purified the protein. After the purification of ObgE, we tested its chaperone activity by using renaturation of citrate synthase, α -glucosidase and bovine carbonic anhydrase II. We found that ObgE has chaperone activity to a certain extent. This result will have impact on the research of protein ObgE.

Keywords: ObgE Chaperone Gene clone Protein purification

收稿日期 2011-01-24 修回日期 2011-03-07 网络版发布日期

DOI: 10.3724/SP.J.1260.2011.00500

基金项目:

国家自然科学基金项目(08JM031001)

通讯作者: 秦燕, 电话: (10)64869250, E-mail: qiny@ibp.ac.cn

作者简介:

作者Email: qiny@ibp.ac.cn

参考文献:

1. Kramer G, Boehringer D, Ban N, Bukau B. The ribosome as a platform for co-translational processing, folding and targeting of newly synthesized proteins. *Nat Struct Mol Biol*, 2009, 16(6): 589~597
2. Hartl FU, Hayer-Hartl M. Converging concepts of protein folding in vitro and in vivo. *Nat Struct Mol Biol*, 2009, 16(6): 574~581

扩展功能

本文信息

- ▶ Supporting info
- ▶ PDF(1068KB)
- ▶ [HTML全文]
- ▶ 参考文献[PDF]
- ▶ 参考文献

服务与反馈

- ▶ 把本文推荐给朋友
- ▶ 加入我的书架
- ▶ 加入引用管理器
- ▶ 引用本文
- ▶ Email Alert
- ▶ 文章反馈
- ▶ 浏览反馈信息

本文关键词相关文章

- ▶ ObgE蛋白
- ▶ 分子伴侣
- ▶ 基因克隆
- ▶ 蛋白纯化

本文作者相关文章

PubMed

3. Kadokura H, Beckwith J. Detecting folding intermediates of a protein as it passes through the bacterial translocation channel. *Cell*, 2009, 138(6): 1164~1173
4. Giglione C, Fieulaine S, Meinnel T. Cotranslational processing mechanisms: Towards a dynamic 3D model. *Trends Biochem Sci*, 2009, 34(8): 417~426
5. Bartlett AI, Radford SE. An expanding arsenal of experimental methods yields an explosion of insights into protein folding mechanisms. *Nat Struct Mol Biol*, 2009, 16(6): 582~588
6. Martinez-Hackert E, Hendrickson WA. Promiscuous substrate recognition in folding and assembly activities of the trigger factor chaperone. *Cell*, 2009, 138(5): 923~934
7. Rüdiger S, Germeroth L, Schneider-Mergener J, Bukau B. Substrate specificity of the DnaK chaperone determined by screening cellulose-bound peptide libraries. *EMBO J*, 1997, 16(7): 1501~1507
8. Weber F, Keppel F, Georgopoulos C, Hayer-Hartl MK, Hartl FU. The oligomeric structure of GroEL/GroES is required for biologically significant chaperonin function in protein folding. *Nat Struct Biol*, 1998, 5: 977~985
9. Leipe DD, Wolf YI, Koonin EV, Aravind L. Classification and evolution of P-loop GTPases and related ATPases. *J Mol Biol*, 2002, 317(1): 41~72
10. Walter S. Structure and function of the GroE chaperone. *Cell Mol Life Sci*, 2002, 59(10): 1589~1597
11. Caldas T, Laalami S, Richarme G. Chaperone properties of bacterial elongation factor EF-G and initiation factor IF2. *J Biol Chem*, 2000, 275(2): 855~860
12. Rao D, Momcilovic I, Kobayashi S, Callegari E, Ristic Z. Chaperone activity of recombinant maize chloroplast protein synthesis elongation factor, EF-Tu. *Eur J Biochem*, 2004, 271(18): 3684~3692
13. Suzuki H, Ueda T, Taguchi H, Takeuchi N. Chaperone properties of mammalian mitochondrial translation elongation factor Tu. *J Biol Chem*, 2007, 282(6): 4076~4084
14. Foti JJ, Persky NS, Ferullo DJ, Lovett ST. Chromosome segregation control by Escherichia coli ObgE GTPase. *Mol Microbiol*, 2007, 65(2): 569~581
15. Kobayashi G, Moriya S, Wada C. Deficiency of essential GTP-binding protein ObgE in Escherichia coli inhibits chromosome partition. *Mol Microbiol*, 2001, 41(5): 1037~1051
16. Jiang M, Datta K, Walker A, Strahler J, Bagamasbad P, Andrews PC, Maddock JR. The Escherichia coli GTPase CgtAE is involved in late steps of large ribosome assembly. *J Bacteriol*, 2006, 188(19): 6757~6770
17. Liu CP, Zhou JM. Trigger factor-assisted folding of bovine carbonic anhydrase II. *Biochem Biophys Res Commun*, 2004, 313(3): 509~615

本刊中的类似文章

1. 汪莉,王玉民,汪琼,郭志云,梁龙,黄培堂.致病菌III型分泌系统分子伴侣序列保守性分析及新伴侣的预测[J]. *生物物理学报*, 2005,21(2): 103-113
2. 翟春燕,刘明耀,罗剑.小鼠核因子 κ B受体活化因子配基活性区的克隆、表达及生物活性分析[J]. *生物物理学报*, 2010,26(9): 790-798

文章评论

反 馈 人	<input style="width: 95%;" type="text"/>	邮箱地址	<input style="width: 95%;" type="text"/>
反 馈 标 题	<input style="width: 95%;" type="text"/>	验证码	<input style="width: 60px;" type="text"/> 1718