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( :10.1007/ 00253-015-6732-9)

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<sup>2</sup> D C Eg g, P S U , U P , PA 16802-4400, USA



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S	R g	S			
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B 25113	$F^{-} \lambda^{-} rph^{-1} \Delta araBAD_{AH33} lacI^{q} \Delta lacZWJ16$	E. coli G S			
	$rrnB_{T14} \Delta rhaBAD_{LD78} hsdR514$	C , U ,			
L 43	Н (PO2A) fhuA22, ДрhoA8, fadL701(T2R), relA1, glpR2(glp <sup>c</sup> ), pitA10, spoT1, glpK22( R), rrnB-2, mcrB1, creC510				

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C

A LB . T B ( 6.81 g N 2HPO4, 2.99 g KH2PO4, 0.58 g N C , 5.94 g (NH4)2SO4) g 2 g/L g 2 g/L g T 500-L E 37 C 220 . T H 7. T

## 4

B g, ..., 600 (OD<sub>600</sub>) g ..., 600 (UV-7504, ..., S g, C ). C ...

g - g (1260, Ag , S C , USA) g - g (HP -87H, B -R , H , CA) - g (RI) . A 5 M H<sub>2</sub>SO<sub>4</sub> 0.5 L/ 60. C.

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S HJ02, HJ03, HJ04, HJ05 NADP<sup>+</sup>/NADPH 60 .I NADP<sup>+</sup>/NADPH E C NADP<sup>+</sup>/NADPH B (B A S , H , CA), g D , g , M , E 2013. M g (AVOVA) **g** *P* < 0.05.

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#### glpK . . .

I HJ01 *yqhD E. coli* B 25113 g *x* T , *glpK* 

- HJ01.

- HJ01. T g g lpK mgsA HJ01 HJ02 (Fg. 3). Mg A g lpK mgsA HJ01 HJ02 (Fg. 3). Mg A g lpK lnF g lpKg L 43 g lpK lnF (P < 0.01) g mgsA g 2.5-HJ02 (P < 0.01).



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	<b>,</b> '	NAI	DPH			NADP	H/N.	ADP⁺	+	-
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g				,	CA2	24N). A	L _		<b>,</b> ,	
	,	, yq	hD 🚬		NAD	РН		35.8	% (P	< 0.01)
	NA	ADPH/	NADI	$\mathbf{p}^+$		25.1	% (P	<b>o</b> < 0.	05)	HJ02,
		(]	Г 2	2). T		gg			,	
	~	,	g	NAE	PH			5	~	1.2
			ntsG							

T

S g S NADPH (S .2013), 2g/L g  $.F_g$  4 E. coli HJ02 gg g

- g g g (CCR) (D (CCRg zwf (P < 0.01) g g -6-g (G6PDH) PP , NADPH NADPH/NADP<sup>+</sup> 1.72- 1.43- g g , , (T 2). C .(2009) S NADPH g g L , NADPH g , D.T , s s NADPH · · · g · · D
- D T, ptsG, CCRHJ02, g, HJ04. T g, ptsG- (EIICB<sup>G</sup>), G g, (G 2005). A Fg. 4, g g, T  $1.20\sigma$  /L. 69 0 % 1.20g /L, 69.0 % g HJ02. M -

- gapA . . .
- D gapA Corynebacterium glutamicum NADPH (S . 2013). H . , , gapA E. coli g

- (S . . 1997). T RNA g Ν T (2009) gapA, g g g . . . g PP . T , HJ05 E. coli д gapA . НJ04, -55 · 5 · (Fg. 4). HJ04, g gapA A
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6.9- (P < 0.01) HJ05. T zwfg 2.2- (P < 0.01). T NADPH HJ05 72.4 % g HJ02 32.3 % g HJ04 (P < 0.01) P < 0.05, ) (T 2). T NADPH/NADP<sup>+</sup> HJ05 HJ02 HJ04 (P < 0.01) P < 0.05,

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G , . 2012; C g G 2013). T (A **g** . , 2013; œ Μ E. coli**g** . - 1 g .... g , , g 2013). S G K-G D ( g . g • glpK glpD<sub>g</sub> E. coli - g · · · · · · q (M . 2013; g . . 2014; g . . 2014). I glpK g HJ01, 5.5 HJ01, glpK L 43 g . I HJ02, 1 L 43 g , , g *glpK* HJ02 -G K FBP EIIA<sup>G</sup> G K (Fg. 2). F . . . HJ02. T L 43, FBP EIIA<sup>G</sup>, g, 1971). T g-G K (F g . . 1971). g glpK HJ02 L 43 , mgsA HJ02 · · · · · ·~ • mgsA. L 43.U , (C g G HJ02 2011; J . . 2014). O (B. 2003). T L 43 22 × 22

(\_\_\_\_\_\_. 1970). I \_\_\_\_\_, \_\_\_\_ mgsA HJ02 , gg , e ( ). T g\_ mgsA g . . . .T, ... а**д** н. . . . Н. , . · · · · mgsA glpK, g mgsA В s g s g g (L . 2010; . 2014). I NADPH g . M NADPH *E. coli* g TCA (R . 1968), g (C . 2014), PP (S . 2004). P s - s NADPH RT-PCR PP g g *ZWf* g NADPH NADPH/NADP<sup>+</sup>, . T g L ' -. T g (2010) *zwf* g NADP<sup>+</sup> g pgi. T g NADPH NADPH . F , *ptsG* NADPH NADPH/NADP<sup>+</sup> . M g PTS g g I (2008)g g . I  $\Delta ptsG$  , HJ04, g g . T HJ04 a g , g ~ **g g** ~ ~ g NADPH . T ptsGg NADPH D, b g b g . . . T , gapA 12 NADPH g 6-6- g 3-g PP (K g **g** , <del>6</del>-, , 3-S 2003). S . (2013)  $\Delta gapA$ C. glutamicum g (R)-C. glutamicum g

3- gapA E. coli gapA E. coli gapA E. coli gapA g GA T gapA g GA T gapA g GA T gapA g aggapA g aggaa gaggaa

Т 973 Р.д. С -

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