

1 **FOR THE RECORDS**

2 **Title:**

3 Atypical effect of temperature tuning on the insertion of the catalytic iron-sulfur centre in a
4 recombinant [FeFe]-hydrogenase

5
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17
18 **Running title:**

19 Temperature effect on [FeFe]-hydrogenase expression

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23 **Tables: 1**

24

1 **Abstract**

2 The expression of recombinant [FeFe]-hydrogenases is an important step for the production of
3 large amount of these enzymes for their exploitation in biotechnology and for the
4 characterisation of the protein-metal cofactor interactions. The correct assembly of the
5 organometallic catalytic site, named H-cluster, requires a dedicated set of maturases that must
6 be co-expressed in the microbial hosts or used for *in vitro* assembly of the active enzymes. In
7 this work, the effect of the post-induction temperature on the recombinant expression of
8 CaHydA [FeFe]-hydrogenase in *E. coli* is investigated. The results show a peculiar behaviour:
9 the enzyme expression is maximum at lower temperatures (20°C), while the specific activity
10 of the purified CaHydA is higher at higher temperature (30°C), as a consequence of improved
11 protein folding and active site incorporation.

12

13 **Keywords**

14 [FeFe]-hydrogenases; Recombinant expression; Bio-hydrogen; metalloenzyme.

15

16 **50-75 words statement**

17 Post-induction temperature severely influences the recombinant expression in *E. coli* of the
18 [FeFe]-hydrogenase CaHydA, a metalloenzyme hosting the peculiar catalytic centre H-
19 cluster. The best protein yield is observed at lower temperature (20°C), while the best specific
20 activity is obtained at higher temperature (30°C), which is atypical in comparison to the usual
21 trend for recombinant holo-enzymes.

1 **Introduction**

2 [FeFe]-hydrogenases are the enzymes that reversibly catalyse the production of molecular
3 hydrogen, following the reaction $2\text{H}^+ + 2\text{e}^- \rightleftharpoons \text{H}_2$ [1]. They are widely distributed among
4 prokaryotes and eukaryotes and are essential in the energy metabolism of such organisms,
5 being usually involved in the dissipation of excess of reducing equivalents in the cell. A
6 significant biotechnological interest has been directed to their exploitation in new, clean and
7 efficient industrial processes for the production of H_2 , to be used as a valuable fuel and
8 industrial intermediate [2-7].

9 The production of [FeFe]-hydrogenases by recombinant techniques has become relevant for
10 several reasons. First of all, the recombinant techniques allow the manipulation of the protein:
11 1) by inserting tag sequences that facilitate purification [8-10], which is highly desirable given
12 the need to work under anaerobic conditions; 2) by inserting single mutations for the study of
13 target residues [11-14]; 3) by generating random mutations for the study of complex features
14 [15-19]. Moreover, recombinant expression usually grants the availability of large amount of
15 enzyme that are required for the characterisation [9,20-24] and for the development of
16 possible future applications [5,6,25]. Recombinant expression has also paved the way to study
17 the mechanisms of the insertion of the catalytic centre H-cluster in the enzyme [FeFe]-
18 hydrogenases, the so-called maturation [8,21,26-29].

19 The recombinant systems that have been developed so far are either cell-hosted or cell-free.
20 The systems that are cell-hosted are carried out in three different hosts: *Escherichia coli*
21 [8,10,20], *Clostridium acetobutylicum* [9,30] and *Shewanella oneidensis* [31]. The cell-free
22 systems are based on the *in vitro* insertion of the H-cluster into an apo-[FeFe]-hydrogenase: in
23 some cases the maturases are added [32-35], while in others the H-cluster is inserted as a
24 chemically synthesised complex [36,37].

1 Given the simplicity and the technological availability of all the components, the expression
2 system for *E. coli* has been widely developed and used. In previous reports, the effect of
3 several parameters has been optimised, but the temperature was never analysed in details, as
4 most authors carried out the experiments at room temperature [8,10,20].
5 In this work, we report on the effect of the post-induction temperature on the recombinant
6 expression of *Clostridium acetobutylicum* CaHydA [FeFe]-hydrogenase in *E. coli* with a C-
7 terminal Strep-tagII.

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9

10 **Results and Discussion**

11 The effect of the post-induction temperature was assayed by SDS-PAGE (Fig. 1A) that allows
12 to observe the levels of expression of the maturases CaHydF and CaHydG, as well as western
13 blot stain against Strep-TagII (Fig. 1B) that specifically discriminates the level of CaHydA.
14 From the functional point of view, the total H₂ evolution activity was assayed on whole cells
15 by gas chromatography (Fig. 1C).

16 These results (Fig. 1A, 1B and 1C) clearly show that the amount of the maturases, the amount
17 of CaHydA and the total hydrogenase activity in whole cells reach a maximum at 20°C,
18 suggesting this temperature as the best condition.

19 To confirm the results observed in whole cells, the enzyme was anaerobically purified by
20 Strep-tagII affinity chromatography and the yield of pure protein and specific hydrogenase
21 activity were measured as previously described [18].

22 The characterisation of the purified enzyme showed that lowering the temperature results in a
23 significant increase of the pure protein yield, similarly to the observation in whole cells, but
24 also a relevant decrease in specific activity (Fig. 1D). The fact that in whole cells the total
25 activity reached a maximum at 20°C is reasonably given by the combination of a very large

1 amount of protein with low activity; on the contrary, at 30°C the amount of protein is much
2 lower, but the specific activity is higher, reaching $1880 \pm 108 \mu\text{mol H}_2/\text{min}/\text{mg}$ protein.
3 Also, it is important to consider that the purified enzyme obtained by expression at 20°C
4 formed aggregates when the concentration was increased, while the enzyme expressed either
5 at 25°C or 30°C was readily soluble and could be concentrated by ultra-filtration up to the
6 millimolar range.

7 The increase in specific activity and solubility at higher expression temperature is probably a
8 result of improved protein folding, iron sulphur clusters incorporation and maturation (*i.e.*
9 incorporation of the H-cluster catalytic centre). Even if the amount of the maturases CaHydF
10 and CaHydG is lower at 30°C, this might represent the best molar ratios between the proteins,
11 leading to optimal kinetics of the process of the metal centre assembly, and availability of the
12 cellular substrates, such as iron and tyrosine, resulting in a high proportion of holo-CaHydA.

13 Lowering the post-induction temperature is a common procedure in recombinant expression
14 of proteins in *E. coli*, as it usually leads to slower kinetics hence avoiding the formation of
15 inclusion bodies and improving recovery of the target protein [38,39]. Indeed in our case this
16 effect was observed: the protein amount was larger at lower temperatures, but it did not
17 correlate with specific activity, as this is the result of a more complex process, as discussed
18 above. Another possible tuning effect of the temperature might involve endogenous *E. coli*
19 scaffold proteins for iron-sulfur cluster biosynthesis, which must be recruited for hydrogenase
20 assembly, either affecting the H-cluster or the other FeS clusters inserted in this enzyme
21 [40,41]. For example, it was shown that the scaffold protein IscU from *Escherichia coli* has a
22 tight temperature control with a narrow range of activity [42].

23 The protocol described here, with the expression at 30°C, resulted in the highest specific
24 activity reported so far for the recombinant CaHydA. The H₂ evolution rate of 1880 ± 108
25 $\mu\text{mol H}_2/\text{min}/\text{mg}$ protein, assayed by gas chromatography with 10 mM reduced methyl

1 viologen as artificial electron donor, is in line with the specific activity of other recombinant
2 [FeFe]-hydrogenases (Table I) and in the same order of magnitude of other native [FeFe]-
3 hydrogenases from Clostridia [43-45].
4 These results may be very useful in the future to standardise the process and to simplify
5 comparison between different enzyme preparations from different laboratories. Also, the
6 effect of the temperature on specific activity of purified enzymes can contribute to explain the
7 apparent incongruences previously reported in recent mutagenesis studies [11-14,18].
8 In conclusion, the results presented here show that the post-induction temperature has a
9 relevant effect on the pure protein yield of CaHydA [FeFe]-hydrogenase and on the specific
10 activity of a properly assembled H-cluster in the purified enzyme, with reverse
11 proportionality. The maximum specific activity was observed when the post-induction
12 temperature was 30°C. Despite the lower yield of pure protein, it is clear that the solubility
13 and the higher specific activity, given by a higher proportion of holo-enzyme, are important
14 factors for the characterisation of [FeFe]-hydrogenases and for their effective exploitation in
15 future applications in biotechnology.

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17

18 **Materials and methods**

19 **Recombinant expression**

20 The plasmids pCaE2 and pCaFG encoding for CaHydA and the maturases CaHydE, CaHydF
21 and CaHydG [8] were co-transformed into *E. coli* Rosetta2(DE3). As previously described
22 [10], bacteria were aerobically grown in baffled flasks (VWR) at 37°C in terrific broth (12
23 g/L tryptone, 24 g/L yeast extract, 4 mL/L glycerol, 2.2 g/L KH₂PO₄, 9.4 g/L K₂HPO₄)
24 supplemented with 200 µg/mL carbenicillin, 50 µg/mL streptomycin, 34 µg/mL
25 chloramphenicol and 2 mM ammonium ferric citrate. When the OD₆₀₀ reached ~0.4, the

1 culture was supplemented with 2 mM cysteine, 25 mM fumarate, 0.5% w/v glucose and
2 induced with 1.5 mM IPTG.
3 Immediately after induction, the culture was split in sterile glass vials (100 mL each), sealed
4 and purged with pure argon to remove trace oxygen, allowing the expression of the active
5 enzymes. The vials were then incubated 22 hours at different temperatures ranging from 4°C
6 to 37°C.

7

8 **Protein expression analysis**

9 Total cell lisates were separated by SDS-PAGE on 10% polyacrylamide gels and stained with
10 Coomassie R350 (GE Healthcare). Western blot against Strep-TagII was performed on PVDF
11 membranes (GE Healthcare) with the Strep-Tactin HRP conjugate (IBA) and stained with
12 3,3'-diaminobenzidine (Sigma-Aldrich).

13

14 **Enzyme purification**

15 All the manipulations were carried out under strict anaerobic conditions in a glove box (Plas
16 Labs) under a hydrogen-nitrogen atmosphere. All solutions were equilibrated with the glove
17 box atmosphere and supplemented with 2-20 mM sodium dithionite before use.

18 CaHydA was purified by affinity chromatography by Strep-Tactin Superflow high capacity
19 cartridges (IBA, Goettingen, Germany) as previously described [18].

20 Purified protein yield was determined with the Bradford assay using bovine serum albumin as
21 standard (Sigma-Aldrich).

22

23 **Activity assays**

24 Hydrogenase activity (H₂ evolution) was determined at 37°C as previously described [18].

25 Briefly, reactions were set up in anaerobic 100 mM TrisHCl, 150 mM NaCl, pH 8.0 with 10

1 mM methyl viologen and 20 mM sodium dithionite. For the determination of the whole cells
2 activity 0.1% v/v Triton X-100 was also added and the reaction was started by the addition of
3 the culture. For the determination of the specific activity, the reactions were started by the
4 addition of the purified enzyme.

5 H₂ evolution was quantified by gas chromatography, using an Agilent Technologies 7890A
6 instrument equipped with purged packed inlet, Molesieve 5A column (30 m, ID 0.53 mm,
7 film 25 mm) and thermal conductivity detector; argon was used as carrier gas.

8

9

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1 **References.**

- 2 [1] Vignais PM, Billoud P (2007) Occurrence, classification, and biological function of
3 hydrogenases: an overview. *Chem Rev* 107:4206-4272.
- 4 [2] Levin DB, Pitt L, Love M (2004) Biohydrogen production: prospects and limitations to
5 practical application. *Int J Hydrogen Energy* 29:173-185.
- 6 [3] Hallenbeck PC (2009) Fermentative hydrogen production: Principles, progress, and
7 prognosis. *Int J Hydrogen Energy* 34:7379-7389
- 8 [4] McKinlay JB, Harwood CS (2010) Photobiological production of hydrogen gas as a
9 biofuel. *Curr Opin Biotechnol* 21:244-251.
- 10 [5] Morra S, Valetti F, Sadeghi SJ, King PW, Meyer T, Gilardi G (2011) Direct
11 electrochemistry of an [FeFe]-hydrogenase on a TiO₂ Electrode. *Chem Commun*
12 47:10566-10568.
- 13 [6] Woolerton TW, Sheard S, Chaudhary YS, Armstrong FA (2012) Enzymes and bio-
14 inspired electrocatalysts in solar fuel devices. *Energy Environ Sci* 5:7470-7490.
- 15 [7] King PW (2013) Designing interfaces of hydrogenase–nanomaterial hybrids for efficient
16 solar conversion. *Biochim Biophys Acta* 1827:949-957.
- 17 [8] King PW, Posewitz MC, Ghirardi ML, Seibert M (2006) Functional studies of [FeFe]
18 hydrogenase maturation in an *Escherichia coli* biosynthetic system. *J Bacteriol*
19 188:2163-2172.
- 20 [9] von Abendroth G, Stripp S, Silakov A, Croux C, Soucaille P, Girbal L, Happe T (2008)
21 Optimized overexpression of [FeFe] hydrogenases with high specific activity in
22 *Clostridium acetobutylicum*. *Int J Hydrogen Energy* 33:6076-6081.
- 23 [10] Yacoby I, Tegler LT, Pochekailov S, Zhang S, King PW (2012) Optimised expression
24 and purification for high-activity preparations of algal [FeFe]-hydrogenase. *PloS ONE*
25 7:e35886.

- 1 [11] Lautier T, Ezanno P, Baffert C, Fourmond V, Cournac L, Fontecilla-Camps JC,
2 Soucaille P, Bertrand P, Meynial-Salles I, Léger C (2011) The quest for a functional
3 substrate access tunnel in FeFe hydrogenase. *Faraday Discuss* 148:385-407.
- 4 [12] Knörzer P, Silakov A, Foster CE, Armstrong FA, Lubitz W, Happe T (2012) Importance
5 of the Protein Framework for Catalytic Activity of [FeFe]-Hydrogenases. *J Biol Chem*
6 286:38341-38347.
- 7 [13] Cornish J, Gärtner K, Yang H, Peters JW, Hegg WL (2011) Mechanism of Proton
8 Transfer in [FeFe]-Hydrogenase from *Clostridium pasteurianum*. *J Biol Chem*
9 286:38341-38347.
- 10 [14] Mulder DW, Ratzloff MW, Bruschi M, Greco C, Koonce E, Peters JW, King PW (2014)
11 Investigations on the Role of Proton-Coupled Electron Transfer in Hydrogen Activation
12 by [FeFe]-Hydrogenase. *J Am Chem Soc* 136:15394-15402.
- 13 [15] Nagy LE, Meuser JE, Plummer S, Seibert M, Ghirardi ML, King PW, Ahmann D,
14 Posewitz MC (2007) Application of gene-shuffling for the rapid generation of novel
15 [FeFe]-hydrogenase libraries. *Biotechnol Lett* 29:421-430.
- 16 [16] Stapleton JA, Swartz JR (2010) A Cell-Free Microtiter Plate Screen for Improved
17 [FeFe] Hydrogenases. *PLoS ONE* 5:e10554.
- 18 [17] Stapleton JA, Swartz JR (2010) Development of an In Vitro Compartmentalization
19 Screen for High-Throughput Directed Evolution of [FeFe] Hydrogenases. *PLoS ONE*
20 5:e15275.
- 21 [18] Morra S, Giraud A, Di Nardo G, King PW, Gilardi G, Valetti F (2012) Site Saturation
22 Mutagenesis Demonstrates a Central Role for Cysteine 298 as Proton Donor to the
23 Catalytic Site in CaHydA [FeFe]-Hydrogenase. *PLoS ONE* 7:e48400.
- 24 [19] Bingham S, Smith PR, Swartz JR (2012) Evolution of an [FeFe] hydrogenase with
25 decreased oxygen sensitivity. *Int J Hydrogen Energy* 37:2965-2976.

- 1 [20] Kuchenreuther JM, Grady-Smith CS, Bingham AS, Gorge SJ, Cramer SP, Swartz JR
2 (2010) High-Yield Expression of Heterologous [FeFe] Hydrogenases in *Escherichia*
3 *coli*. PloS ONE 5:e15491.
- 4 [21] Mulder DW, Boyd ES, Sarma R, Lange RK, Endrizzi JA, Broderick JB, Peters JW
5 (2010) Stepwise [FeFe]-hydrogenase H-cluster assembly revealed in the structure of
6 HydA^{ΔEFG}. Nature 465:248-251.
- 7 [22] Mulder DW, Ratzloff MW, Shepard EM, Byer AS, Noone SM, Peters JW, Broderick
8 JB, King PW (2013) EPR and FTIR Analysis on the Mechanism of H₂ Activation by
9 [FeFe]-Hydrogenase HydA1 from *Chlamydomonas reinhardtii*. J Am Chem Soc
10 135:6921-6929.
- 11 [23] Myers WK, Stich TA, Suess DLM, Kuchenreuther JM, Swartz JR, Britt RD (2014) The
12 Cyanide Ligands of [FeFe] Hydrogenase: Pulse EPR Studies of ¹³C and ¹⁵N-Labeled H-
13 Cluster. J Am Chem Soc 136:12237-12240.
- 14 [24] Adamska A, Silakov A, Lambertz C, Rüdiger O, Happe T, Reijerse E, Lubitz W (2012)
15 Identification and Characterization of the “Super-Reduced” State of the H-Cluster in
16 [FeFe] Hydrogenase: A New Building Block for the Catalytic Cycle? Angew Chem Int
17 Ed 51:11458-11462.
- 18 [25] Kim S, Lu D, Park S, Wang G (2012) Production of hydrogenases as biocatalysts. Int J
19 Hydrogen Energy 37:15833-15840.
- 20 [26] Posewitz MC, King PW, Smolinski SL, Zhang Z, Seibert M, Ghirardi ML (2004)
21 Discovery of two novel radical S-adenosylmethionine proteins required for the assembly
22 of an active [Fe] hydrogenase. J Biol Chem 279:25711–25720.
- 23 [27] Kuchenreuther JM, George SJ, Grady-Smith CS, Cramer SP, Swartz JR (2011) Cell-free
24 H-cluster Synthesis and [FeFe] Hydrogenase Activation: All Five CO and CN⁻ Ligands
25 Derive from Tyrosine. PloS ONE 6:e20346.

- 1 [28] Cendron L, Berto P, D'Adamo S, Vallese F, Covoni C, Posewitz MC, Giacometti GM,
2 Costantini P, Zanotti G (2011) Crystal Structure of HydF Scaffold Protein Provides
3 Insights into [FeFe]-Hydrogenase Maturation. *J Biol Chem* 286:43944-43950.
- 4 [29] Shepard EM, Mus F, Betz JN, Byer AS, Duffus BR, Peters JW, Broderick JB (2014)
5 [FeFe]-Hydrogenase Maturation. *Biochemistry* 53:4090-4104.
- 6 [30] Girbal L, von Abendroth G, Winkler M, Benton PMC, Meynial-Salles I, Croux C,
7 Peters JW, Happe T, Soucaille P (2005) Homologous and Heterologous Overexpression
8 in *Clostridium acetobutylicum* and Characterization of Purified Clostridial and Algal Fe-
9 Only Hydrogenases with High Specific Activities. *Appl Environ Microbiol* 71:2777-
10 2781.
- 11 [31] Sybirna K, Antoine T, Lindberg P, Fourmond V, Rousset M, Mèjean V, Bottin H (2008)
12 *Shewanella oneidensis*: a new and efficient system for expression and maturation of
13 heterologous [Fe-Fe] hydrogenase from *Chlamydomonas reinhardtii*. *BMC Biotechnol*
14 8:73.
- 15 [32] McGlynn SE, Ruebush SS, Naumov A, Nagy LE, Dubini A, King PW, Broderick JB,
16 Posewitz MC, Peters JW (2007) In vitro activation of [FeFe] hydrogenase: new insights
17 into hydrogenase maturation. *J Biol Inorg Chem* 12:443-447.
- 18 [33] Boyer ME, Stapleton JA, Kuchenreuther JM, Wang CW, Swartz JR (2008) Cell-Free
19 Synthesis and Maturation of [FeFe] Hydrogenases. *Biotechnol Bioeng* 99:59-67.
- 20 [34] Kuchenreuther JM, Stapleton JA, Swartz JR (2009) Tyrosine, cysteine and S-Adenosyl
21 methionine stimulate in vitro [FeFe] hydrogenase activation. *PloS ONE* 4:e7565.
- 22 [35] Kuchenreuther JM, Britt RD, Swartz JR (2012) New insights into [FeFe] hydrogenase
23 activation and maturase function. *PloS ONE* 7:e45850.

- 1 [36] Berggren G, Adamska A, Lambertz C, Simmons TR, Esselborn J, Atta M, Gambarelli S,
2 Mouesca JM, Reijerse E, Lubitz W, Happe T, Artero V, Fontecave M (2013)
3 Biomimetic assembly and activation of [FeFe]-hydrogenases. *Nature* 499:66-70.
- 4 [37] Esselborn J, Lambertz C, Adamska-Venkatesh A, Simmons T, Berggren G, Noth J,
5 Siebel J, Hemschemeier A, Artero V, Reijerse E, Fontecave M, Lubitz W, Happe T
6 (2013) Spontaneous activation of [FeFe]-hydrogenases by an inorganic [2Fe] active site
7 mimic. *Nat Chem Biol* 9 :607-609.
- 8 [38] Sørensen HP, Mortensen KK (2005) Soluble expression of recombinant proteins in the
9 cytoplasm of *Escherichia coli*. *Microb Cell Fact* 4:1.
- 10 [39] Tolia NH, Joshua-Tor L (2006) Soluble expression of recombinant proteins in the
11 cytoplasm of *Escherichia coli*. *Nat Methods* 3:55-64.
- 12 [40] Bandyopadhyay S, Chandramouli K, Johnson MK (2010) Iron-Sulphur Cluster
13 Biosynthesis. *Biochem Soc Trans* 36:1112-1119.
- 14 [41] Roche B, Aussel L, Ezraty B, Mandin P, Py B, Barras F (2013) Iron/sulfur proteins
15 biogenesis in prokaryotes: Formation, regulation and diversity. *Biochim Biophys Acta*
16 1827:455-469.
- 17 [42] Markley JL, Kim JH, Dai Z, Bothe JR, Cai K, Frederick RO, Tonelli M (2013)
18 Metamorphic protein IscU alternates conformations in the course of its role as the
19 scaffold protein for iron–sulfur cluster biosynthesis and delivery. *FEBS Letters*
20 587:1172-1179.
- 21 [43] Chen JS, Mortenson LE (1974) Purification and properties of hydrogenase from
22 *Clostridium pasteurianum* W5. *Biochim Biophys Acta* 371:283-298.
- 23 [44] Adams MWW (1990) The structure and mechanism of iron-hydrogenases. *Biochim*
24 *Biophys Acta* 1020:115-145.

1 [45] Peters JW, Lanzilotta WN, Lemon BJ, Seefeldt LC (1998) X-ray Crystal Structure of
2 the Fe-Only Hydrogenase (CpI) from *Clostridium pasteurianum* to 1.8 Angstrom
3 Resolution. *Science* 282:1853-1858.

4

1 **Table and Caption.**

Type	Host	Maturases	Enzyme	Specific activity ($\mu\text{mol H}_2/\text{min}/\text{mg}$)	Yield (mg/L)	Ref.
Cell- Hosted	<i>E. coli</i> Rosetta2(DE3)	Ca	CaHydA	1880±108 GC 10 mM MV pH 8.0	1.2	This work
	<i>E. coli</i> Rosetta2(DE3)	Ca	Fd-CrHydA1	1000 GC 10 mM MV pH 8.0	5	[10]
	<i>E. coli</i> BL21(DE3) <i>ΔiscR</i>	So	CrHydA1	641±88 GC 5 mM MV pH 6.8	30±11	[20]
	<i>S. oneidensis</i>	endog.	CrHydA1	740±56 Electrode 5 mM MV pH 6.7	0.5	[31]
	<i>C. acetobutylicum</i>	endog.	CaHydA	1750* GC MV pH 6.8	0.8	[9]
		endog.	CrHydA1	625* GC MV pH 6.8	1	
	<i>E. coli</i> BL21(DE3)	Ca	CaHydA	75.2 GC 5 mM MV pH 7-8	NR	[8]
		Ca	CrHydA1	150 GC 5 mM MV pH 7-8	0.8-1.0	
Cell- Free	-	-	CpI	2037±616 GC 10 mM MV pH 6.8	NR	[37]
	-	Ca	CrHydA1	700-800 GC 10 mM MV pH 6.8	NR	[36]
	-	So	CpI	~700** Spect. MV	NR	[35]
	-	Ca	CsHydA	~2.5 GC 10 mM MV pH 7.5	NR	[32]

2

1 Table I. Comparison of the specific activity and yield of CaHydA with other recombinant
2 [FeFe]-hydrogenases. Ca) *Clostridium acetobutylicum*. Cr) *Chlamydomonas reinhardtii*. So)
3 *Shewanella oneidensis*. Cs) *Clostridium saccharobutylicum*. endog.) endogenous maturases.
4 Fd) ferredoxin. Without other specification, specific activity is reported as H₂ evolution rate.
5 *) V_{max}. **) H₂ oxidation rate. The methodology used is also indicated: GC) Gas
6 chromatography. Spect.) Spectrophotometric assay. MV) methyl viologen as artificial redox
7 partner. The pH of the assay is also specified. Protein yield is reported as mg pure protein
8 obtained per litre of culture. NR) not reported.
9

1 **Figure Caption**

2

3 Figure 1. (2-column fitting)

4 Effect of the post-induction temperature on the recombinant expression of CaHydA. A)

5 Coomassie stained SDS-PAGE of whole cells lisates; bands at the molecular weight of

6 CaHydF (46 kDa) and CaHydG (53 kDa) are marked. NI = Not induced. B) Western blot

7 against Strep-tagII; a band at the molecular weight of CaHydA (65 kDa) can be identified. C)

8 Total hydrogenase activity of whole cells. D) Specific activity of purified CaHydA

9 (continuous line, filled squares) and yield of pure protein (dashed line, open squares).

10