Geomicrobiology Journal





Journal:	Geomicrobiology Journal
Manuscript ID:	UGMB-2015-0116.R1
Manuscript Type:	Special Issue
Date Submitted by the Author:	n/a
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Keywords:	biomineralization, metal reduction, bioremediation

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- 1 A novel aerobic mechanism for reductive palladium
- 2 biomineralization and recovery by Escherichia coli

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- 19 Aerobically-grown E. coli cells reduced Pd(II) via a novel mechanism using formate
- as the electron donor. This reduction was monitored in real-time using extended X-ray
- absorption fine structure. Transmission electron microscopy analysis showed that
- 22 Pd(0) nanoparticles, confirmed by X-ray diffraction, were precipitated outside the
- cells. The rate of Pd(II) reduction by E. coli mutants deficient in a range of
- 24 oxidoreductases was measured, suggesting a molybdoprotein-mediated mechanism,
- 25 distinct from the hydrogenase-mediated Pd(II) reduction previously described for

26	anaerobically-grown <i>E. coli</i> cultures. The potential implications for Pd(II) recovery
27	and bioPd catalyst fabrication are discussed.
28	
29	Keywords: palladium nanoparticles, <i>Escherichia coli</i> , biomineralization.
30	
31	Introduction
32	The microbial reduction of metals and radionuclides has attracted much interest, as it
33	can be potentially harnessed for bioremediation, metal recovery, the fabrication of
34	novel nanobiominerals and even energy generation in biobatteries (Lloyd 2003; Lloyd
35	et al. 2008; Lovley 2006;). For example, the sulfate-reducing bacterium (SRB)
36	Desulfovibrio desulfuricans has been shown to use a periplasmic hydrogenase
37	supplied with hydrogen to reduce soluble Pd(II), resulting in the precipitation of Pd(0)
38	nanoparticles in the periplasm of the cell ('bioPd'). However SRB produce H ₂ S, a
39	potent catalyst poison that must be removed before making the bioPd. Other
40	organisms capable of this metal bioreduction include the Gram-negative bacteria
41	Shewanella oneidensis (De Windt et al. 2005), Escherichia coli (Deplanche et al.
42	2010, 2014; Mabbett et al. 2006), Pseudomonas putida, Cupriavidus necator (Søbjerg
43	et al. 2009), Cupriavidus metallidurans (Gauthier et al. 2010), Paracoccus
44	denitrificans (Bunge et al. 2010), Rhodobacter sphaeroides (Redwood et al. 2008),
45	Rhodobacter capsulatus (Wood et al. 2010), and the Gram-positive bacteria Bacillus
46	sphaericus (Creamer et al. 2007), Arthrobacter oxyidans (Deplanche et al. 2014;
47	Wood et al. 2010), Micrococcus luteus (Deplanche et al. 2014), Staphylococcus sciuri
48	(Søbjerg et al. 2009) and <i>Clostridium pasteurianum</i> (Chidambaram et al. 2010). This
49	property has allowed the use of 'palladised' whole cells or processed biomineral
50	directly in industrially important reactions, often showing superior activity compared

51	with a commercially available carbon-supported palladium catalyst. A number of
52	studies have investigated the catalytic activity of bioPd, demonstrating its use in
53	remediative reactions such as the reduction of Cr(VI) to Cr(III) (Beauregard et al.
54	2010; Mabbett et al. 2006), the dehalogenation of chlorophenol, polychlorinated
55	biphenyls, polybrominated diphenyl ethers (Baxter-Plant et al. 2003; De Windt et al.
56	2005; Harrad et al. 2007), trichloroethylene (Hennebel et al. 2009a, 2009b), and the
57	pesticide γ - hexachlorocyclohexane (Mertens et al. 2007), in 'greener' chemical
58	synthesis such as the hydrogenation of itaconic acid (Creamer et al. 2007) and 2-
59	pentyne (Bennett et al. 2010), in Heck and Suzuki reactions (Bennett et al. 2013;
60	Deplanche et al. 2014), and also in the application of bioPd as a fuel cell
61	electrocatalyst to produce electricity from hydrogen (Orozco et al. 2010; Yong et al.
62	2007). In each case where the bioPd was compared with an abiotically-produced
63	palladium catalyst (finely-divided or supported on a carbon matrix), the bioPd was
64	more active than or at least as active as the commercially available alternative.
65	
66	Production of catalytically active bioPd also was reported by an aerobically-grown
67	Serratia sp. (Beauregard et al. 2010; Deplanche et al. 2014) under which condition
68	hydrogenases are not expressed. Also, cells of <i>E. coli</i> deficient in the three major
69	hydrogenases reduced Pd(II) (albeit slowly: Deplanche et al. 2010), and showed
70	larger Pd-nanoparticles located on the outer surface of the cells. This suggested an
71	alternative mechanism of Pd(II) reduction which has not been investigated.
72	
73	$E.\ coli$ produces bioPd which is comparably active to that produced by $D.$
74	desulfuricans (Deplanche et al. 2014). This also provides a very useful model
75	organism since it is facultatively anaerobic and has well-defined molecular tools to

76	elucidate reaction mechanisms under aerobic and anaerobic conditions. The enzymes
77	potentially involved in the bioreduction of palladium by E. coli under the latter
78	conditions are the nickel-dependent hydrogenase enzymes Hyd-1, Hyd-2, and Hyd-3,
79	and the formate dehydrogenase molybdoenzymes FDH-N, and FDH-H. Another
80	molybdoenzyme, FDH-O, is expressed under both aerobic and anaerobic conditions.
81	A possible role for FDH-O is to allow bacteria to adapt rapidly to a sudden shift from
82	aerobic respiration to anaerobiosis, before FDH-N has been produced in sufficient
83	amounts to continue formate metabolism (Abaibou et al. 1995). Hyd-1, Hyd-2, FDH-
84	O, and FDH-N are membrane-bound and periplasmically oriented, whereas Hyd-3
85	and FDH-H are subunits of the formate hydrogenlyase (FHL) complex, an
86	intracellular enzyme complex that is also membrane-bound but which faces into the
87	cytoplasm. The mechanisms responsible for the formate-dependent bioreduction by
88	anaerobically-grown cultures of E. coli have been studied, showing that the
89	hydrogenase enzymes Hyd-1 and Hyd-2 are mainly responsible for Pd(II)
90	bioreduction (Deplanche et al. 2010). In a study of formate-dependent Pd(II)
91	bioreduction by Desulfovibrio fructosovorans, the deletion of the periplasmic
92	hydrogenases caused the Pd(0) nanoparticles to be relocated to the cytoplasmic
93	membrane site of the remaining hydrogenases, indicating that the periplasmic
94	hydrogenases are at least partially involved (Mikheenko et al. 2008).
95	
96	The growth yield of anaerobic cultures is lower than that of aerobic cultures, and for
97	economic production at scale a method of growth of high biomass density is required
98	When using anaerobic cultures there is also the cost of supplementing with sodium
99	fumarate and glycerol. The dual aims of this study are to establish whether <i>E. coli</i>
100	cells grown aerobically are capable of manufacturing bioPd and to identify the

enzyme(s) responsible for such metal reduction. A move away from the need for
anaerobic growth would simplify the preparation of high levels of active biomass for
catalyst production at industrial scale.

Methods

106 Bacterial growth

Starter cultures: 50 ml LB broth in a 500 ml Erlenmeyer flask was inoculated with a single isolated colony of the *E. coli* strain under investigation and incubated aerobically (37°C, shaking at 180 rpm for 18 h).

Aerobic cultures: An 11 ml starter culture was added to 99 ml LB broth in a 1 L Erlenmeyer flask. Flasks were incubated for 24 h (37°C, 180 rpm) to produce stationary phase 'resting' cells. The pH of the cells after 24 h incubation was measured to determine that organic acids had not been produced that would otherwise lower the pH considerably (Vasala et al. 2006). Oxygen saturation of a 5 ml aliquot of the broth culture was measured immediately after 24 h of incubation using an Oakton D06 Acorn Series dissolved oxygen meter.

Reduction of Pd(II) to produce bioPd on bacteria

The aerobically grown liquid culture was divided between two 50 ml Falcon tubes and washed three times in 20 ml MOPS-NaOH (morpholinepropanesulfonic acid) buffer, 20 mM at pH7.6 after centrifugation for 20 min at 2500 g. Cell pellets were adjusted to a mass of 250 mg wet pellet weight, and resuspended in the MOPS-NaOH buffer to a volume of 1 ml. One tube of 250 mg wet weight cells was resuspended in 22.5 ml MOPS-NaOH buffer with 1 mM sodium tetrachloropalladate in a 30 ml bottle

126	sealed with a butyl rubber stopper. The bottle was incubated in the dark at 30°C for 1
127	h for the Pd(II) to biosorb to the cells (Baxter-Plant et al. 2003). 2.5 ml 10 mM
128	sodium formate was then added to the bottle to initiate bioreduction of the Pd(II).
129	
130	Use of mutants to determine electron transfer pathway to Pd(II)
131	In order to investigate the possible role of the aerobic formate dehydrogenase (FDH-O)
132	and other hydrogenase/formate dehydrogenase enzymes in the reduction of Pd(II) by
133	aerobically-grown cells of <i>E. coli</i> , the rates of reduction by six different additional
134	strains (Table 1) were compared by measuring the Pd(II) remaining in solution by
135	ICP-MS. The strains were 'palladised' as above, and rates of reduction/removal
136	compared to those in a series of controls: killed cells (MC4100), cell-free suspension,
137	and live cells (MC4100) unsupplemented with formate.
138	
139	All strains except BL21(DE3) were from the culture collection of Professor Frank
140	Sargent at the College of Life Sciences, University of Dundee. Strain BL21(DE3) was
141	obtained from Invitrogen, Paisley, UK. Strain MC4100 ΔmoaA was created by
142	disruption of the moaA gene which encodes the molybdenum cofactor biosynthesis
143	protein A, using the method of Datsenko and Wanner (2000) whereby PCR products
144	are used to disrupt the gene of choice by recombination using the plasmid-borne
145	phage λ Red recombinase.
146	
147	X-ray diffraction (XRD) analysis
148	The black precipitates were washed once in acetone and air dried, before analysis by
149	X-ray diffraction (XRD). The measurements were performed on a Bruker D8
150	Advance diffractometer, using Cu K alpha1 radiation. The samples were scanned

151	from 5-70 degrees 2theta in steps of 0.2 degrees, with a count time of 2 seconds per
152	step.
153	
154	Extended X-ray absorption fine structure (EXAFS)
155	Aliquots of the cell/Pd/formate suspension were taken at times 0 and 30 min, and 1, 3
156	and 4 h from the addition of formate, and frozen immediately in liquid nitrogen. The
157	direct reduction of Pd(II) to Pd(0) was demonstrated using EXAFS, performed at the
158	European Synchrotron Radiation Facility (ESRF), in Grenoble, France. The samples
159	were transported to the synchrotron at ESRF on dry ice, where they were thawed and
160	injected immediately into sample holders, before freezing once more in liquid
161	nitrogen and placing into the beam. X-ray absorption data were collected on beamline
162	BM29 at the Pd K-edge in the energy range 24 200 – 24 900 eV. Data were recorded
163	at low temperature (77 K) and under vacuum to reduce the thermal Debye-Waller
164	factor and prevent oxidation. A Si(III) double crystal monochromator was used,
165	calibrated with a Pd foil, and the spectra were collected in fluorescence mode using a
166	13-element solid-state detector. A reference spectrum of a palladium foil was
167	recorded in transmission mode on station 9.3 at the SRS Daresbury. The data were
168	background subtracted and the EXAFS spectra fitted in DL_Excurv
169	(http://www.cse.scitech.ac.uk/cmg/EXCURV/) using full curved wave theory
170	(Gurman et al., 1984).
171	
172	Transmission electron microscopy (TEM) and energy dispersive X-ray spectroscopy
173	(EDS)
174	Following Pd(II) reduction, cells were stored at 10°C overnight. The cell pellets were
175	then rinsed twice with deionised water, fixed in 2.5% (wt/vol) glutaraldehyde,

centifuged for 3 min at 10 000 g, resuspended in 1.3 mi of 0.1 M cacodylate buffer
(pH 7) and stained in 1% osmium tetroxide in 0.1 M phosphate buffer, pH 7 (60 min)
Cells were dehydrated using an ethanol series (70, 90, 100, 100, 100% dried ethanol,
15 min each) and washed twice in propylene oxide (15 min, 9500 g). Cells were
embedded in epoxy resin and the mixture was left to polymerise (24 h; 60°C).
Sections (100-150 nm thick) were cut from the resin block, placed onto a copper grid
and viewed with a JEOL 1200CX2 TEM, accelerating voltage 80 keV. EDS was
performed on electron-dark areas, to confirm the presence of palladium.

Results

186 Palladisation of E. coli BL21(DE3)

The pH of the aerobically-grown liquid culture was between 7.7-7.9, indicating that there was not extensive production of organic acids due to overflow metabolism.

Oxygen saturation measurements showed that the liquid culture was 72% saturated following 24 h of incubation, indicating that it was not oxygen-limited. After harvesting, the cells were able to couple the reduction of Pd(II) to the oxidation of formate, indicated by the rapid formation of a black precipitate, tentatively identified as Pd(0) (Figure 1). ICP-MS analysis confirmed complete removal of Pd(II) from solution within 45 min, and the presence of crystalline Pd(0) was confirmed using XRD in this, but not in the heat-killed cells control where the cells removed substantial Pd(II) abiotically. An increase in metal biosorption by heat killed biomass as compared to live cells is well documented (Machado et al. 2009; Parameswari et al. 2009) and was attributed to loss of membrane integrity to reveal additional intracellular metal binding sites (Machado et al. 2009).

201	Extended X-ray absorption fine structure (EXAFS)
202	The nature of the Pd associated with the biomass was assessed further using X-ray
203	absorbance spectroscopy. The features in the corresponding EXAFS spectra (Figure 2)
204	are due to the wave-like nature of the photoelectron, which is released from the atom
205	with increasing energy and scattered from surrounding atoms with new waves being
206	emitted. With increasing photon energy, the interference between the waves alternates
207	between constructive and destructive, which leads to oscillations in the spectrum.
208	Examining these oscillations gives information on the number, species and distance of
209	the surrounding atoms. As seen in Figure 2, the samples taken at times 0 and 30 min,
210	which contain Pd(II), have identical EXAFS spectra. The samples taken at 60 min
211	onwards are identical to the Pd(0) foil control, which indicates that only Pd(0) was
212	present. Reduction of the Pd(II) to Pd(0) was therefore confirmed to be complete in
213	less than 30 min, as confirmed by ICP-MS analysis.
214	
215	Use of mutants to determine electron transfer pathway to Pd(II)
216	Aerobic cultures of the parental strains MC4100 and BW25113 and the strain which
217	lacked all hydrogenases (JW2682) removed Pd(II) identically with no residual Pd(II)
218	detected after 30 min (Figure 3). Removal of the hydrogenase enzymes had no effect
219	on the rate of palladium removal from solution, confirming that these hydrogenases
220	have no role in the aerobic reduction of Pd(II). The FDH-O-negative strain JW3865
221	reduced Pd(II) within 1 h, and the FDH-O/FDH-N-negative strain FTD128 within 2 h.
222	Strain MC4100 $\Delta moaA$, lacking all molybdoenzymes, reduced the palladium within 7
223	h. These results indicate the likely involvement of the FDH-O enzyme in the
224	reduction of Pd(II) by aerobically-grown E. coli using formate, although other Mo-
225	containing enzymes must also be involved given the impaired metal reduction noted

with the Δ*moa* mutant. Controls containing no biomass showed no abiotic reduction of Pd(II) using formate (Figure 1B), although a brown precipitate was seen in the noformate control. The X-ray powder diffraction pattern did not show the presence of any peaks characteristic of Pd(0) in this precipitate, indicating that it was probably amorphous and non-crystalline. Time zero on Figure 3 is the point at which formate was added, following 1 h of incubation to allow biosorption of the Pd(II) to the cells; hence the abiotic Pd(II) removal by killed cells (Figure 1) was apparent at the time of formate addition with no evidence for further Pd(II) reduction.

Transmission electron microscopy (TEM)

TEM images of thin sections of cells showed that with all strains the reduced palladium was precipitated predominantly in the extracellular matrix of the cultures (Figure 4), although it appears that the nanoparticles may be associated with the outer membrane of the cells. Energy dispersive X-ray spectroscopy (EDS) confirmed the presence of palladium in these precipitates.

Discussion

The results from this study demonstrate that it is possible for aerobically-grown cultures of *E. coli* to reduce Pd(II) enzymatically, with no need to remove oxygen from the experimental system during the bioreduction step. Autoclaved control experiments indicate that Pd(II) bioreduction in these cultures is enzymatic, with reduction of palladium not occurring in the absence of viable cells irrespective of the length of incubation. The major enzymes shown to be involved include the formate dehydrogenases FDH-O and FDH-N, although bioreduction still occurs in strains without these enzymes albeit at a much lower rate. Other molybdoenzymes must

therefore be involved. The strain that lacked all molybdoenzymes did however still reduce the palladium, although this took 7 h, compared with less than 30 min by the wild-type strains. Hydrogenases, implicated as the dominant Pd(II) reductases in other experimental systems grown under anaerobic conditions (Deplanche et al. 2010; Mikheenko et al. 2008), are not expressed in aerobically grown cultures, and their lack of involvement was evident as the strain lacking hydrogenase enzymes reduced palladium at the same rate as the wild-type strains in this study.

Furthermore, whichever biological system is responsible for the aerobic bioreduction of Pd(II), there seems to be little impact on the site of Pd(0) deposition. The location of the bioreduced Pd(0) in our experiments is almost always extracellular, although often associated with the outer membrane of the cells. This is particularly the case with the MC4100 Δ*moaA* strain (which lacks all molybdoenzymes), in which the majority of the Pd(0) nanoparticles are closely associated with the outer membrane (Figure 4E). One conclusion that may be drawn from this is that whilst cells that lack the formate dehydrogenases are still capable of reducing Pd(II), when all of these enzymes are missing a cellular component associated with the outer membrane may be responsible. Furthermore, this formate oxidation activity is much weaker than that seen with the strains containing formate dehydrogenases, where Pd(II) reduction is more rapid. It is possible however that following the initial enzymatic reduction of a small percentage of the Pd(II), the Pd(0) nanoparticles formed may themselves be responsible for catalysing the reduction of the remainder of the Pd(II) (Yong et al. 2002), which would mean that only a minor, initial biological input is required.

Although the formate dehydrogenase enzyme systems implicated in Pd(II)
bioreduction by $E.\ coli$ are periplasmic, the majority of the reduced $Pd(0)$ precipitates
outside the cell. It is possible that an electron shuttle system exists similar to that
found in Shewanella oneidensis (von Canstein et al. 2008) that is as yet undiscovered
in $E.\ coli.$ It is also possible that the first $Pd(0)$ nanoparticles to form breach the outer
membrane, and themselves form an electron conduit for further Pd(II) reduction
outside the cell. The pH of these experiments is also higher than others where Pd(0)
nanoparticles accumulated in the periplasm (Redwood et al. 2008), which could
indicate the higher biosorption of cationic metal to the outer membrane and
extracellular polymeric substances, which are then not able to enter the periplasm.
The influence of a higher pH in the location of the Pd(0) may be confirmed by the
observation that $Pd(0)$ nanoparticles were located on the cell surface of D .
desulfuricans when bioreduction of Pd(II) was performed at pH 7 (Yong et al. 2002).
In conclusion, this study has demonstrated the presence of a novel biological
mechanism responsible for the bioreduction of Pd(II) in aerobically-grown cultures of
E. coli, catalysed mainly by molybdenum-containing enzyme systems. Subsequent
studies will investigate the catalytic activity and selectivity of the Pd(0) nanoparticles
produced under aerobic conditions in a range of industrially important reactions. If
active, this new form of bioPd has the advantage over that produced by anaerobic
culture as it is easier to produce at high yield, from increased biomass levels
associated with aerobic growth. There is also no requirement for additional processing
steps to remove H ₂ S (produced by SRB systems), and the use of formate instead of
hydrogen gas means that the procedure is less hazardous and more controllable. The
advantages of this more scalable method of synthesis would need to be considered

against any alterations in activity/selectivity of the resulting catalyst (versus synthetic and other bioPds), using a cost-benefit analysis. Importantly, identification of the specific enzymatic process(es) involved in the biomanufacture of bioPd is the first step towards application of the tools of synthetic biology for 'designer catalyst' production for specific applications.

In a geomicrobiological context, this study shows that aerobic cells of *E. coli* restrict the deposition of Pd(0) to locations outside the cell. However in both *D. desulfuricans* (grown anaerobically) and *Bacillus benzeovorans* (grown aerobically) intracellular depositions of small Pd-nanoparticles were observed at the expense of both hydrogen and formate (JB Omajali, IP Mikheenko, ML Merroun, J Wood and LE Macaskie, in press) and, notably, were also seen in *E. coli* grown anaerobically (LE Macaskie, A Williams, R Priestley and J Courtney, unpublished). This raises questions about potential biochemical 'trafficking' pathways of Pd(II), the possibility of Pd(II) efflux by aerobic (but not anaerobic) cells and, following from that, the possibility of biogeochemical cycling of this element.

Acknowledgement

The authors would like to thank Dr Victoria Coker, Prof Richard Pattrick, and Dr John Charnock of the University of Manchester for assistance with EXAFS.

Funding

322 This study was supported by a grant from the Biotechnology and Biological Sciences

Research Council (BBSRC).

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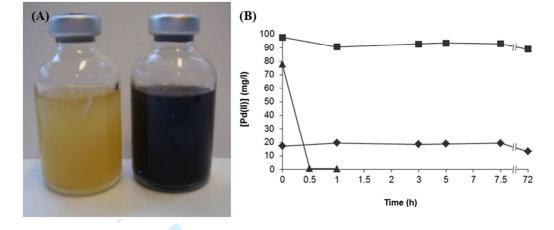
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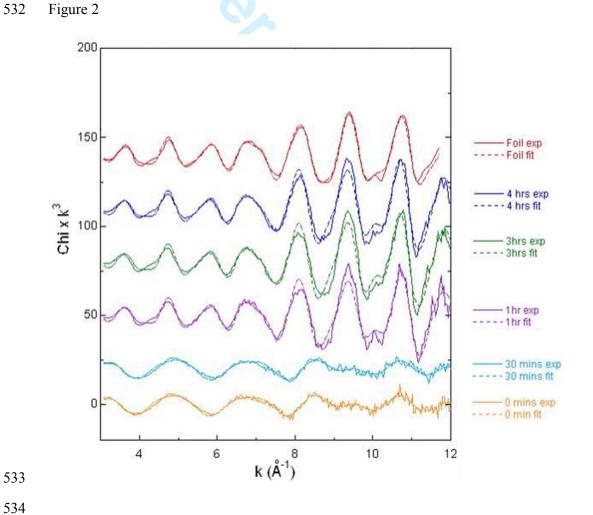
496 palladium (II) using formate as the electron donor.

Strain	Genotype	Phenotype	Reference
BL21(DE3)	F2 ompT gal dcm lon	Wild type strain	(Studier and
	$hsdS_B(r_B^- m_B^-) \lambda(DE3$	commonly used for	Moffatt 1986)
	[lacI lacUV5-T7 gene 1	recombinant protein	
	ind1 sam7 nin5])	expression.	
MC4100	F- ΔlacU169 araD139	Parental strain for	(Casadaban
	rpsL150 relA1 ptsF rbs	FTD128 and	and Cohen
	flbB5301	$\Delta moaA$.	1979)
BW25113	$lacl^q rrbB_{T14} \Delta lacZ_{WJ16}$	Parental strain for	(Datsenko and
	hsd R514 $\Delta araBAD_{AH33}$	JW2682 and	Wanner 2000)
	$\Delta rhaBAD_{ ext{LD78}}$	JW3865.	
FTD128	As MC4100, with in-	FDH-O & FDH-N	(Luke et al.
	frame deletion in the	negative.	2008)
	fdhE gene.		
JW2682	As BW25113, with in-	Deficient in all	(Baba et al.
	frame deletion of the	hydrogenases.	2006)
	hypF gene.		
JW3865	As BW25113, with in-	FDH-O negative.	(Baba et al.
	frame deletion of the		2006)
	fdoG gene.		
MC4100 ΔmoaA	As MC4100, disruption	Deficient in all	This study.
	of the moaA gene.	molybdoenzymes	5.

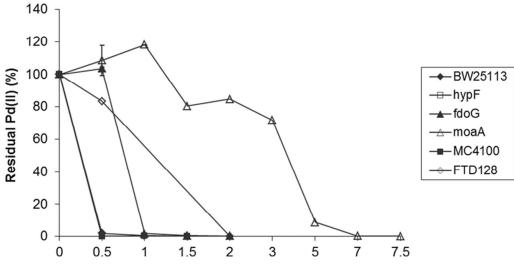
503	Fig. 1. (A) Complete reduction of Pd(II) to Pd(0) by an aerobically-grown culture of
504	E. coli. Both bottles contain cells resuspended in 20 mM MOPS buffer at pH7.6, and
505	1 mM sodium tetrachloropalladate (total volume 25 ml). This image was taken 45 mir
506	after the addition of formate to the bottle on the right. (B) Reduction by E. coli
507	MC4100 and by controls showing no abiotic reduction of Pd(II). Controls used were
508	killed (autoclaved) cells and cell-free suspension. Soluble Pd(II) in the supernatant
509	was measured using ICP-MS. ▲ = MC4100; ■ = no cells; ♦ = killed cells.
510	
511	Fig. 2. EXAFS data showing the presence of Pd(II) at 0 and 30 min (bottom two
512	traces), and Pd(0) at 1, 3 and 4 h (ascending series). The top trace is palladium foil.
513	
514	Fig. 3. Pd(II) reduction by six different strains of E. coli, using formate as the electron
515	donor. Soluble Pd(II) in the supernatant was measured using ICP-MS. ♦ = BW25113;
516	\Box = JW2682; Δ = JW3865; Δ = MC4100 Δ <i>moaA</i> ; ■ = MC4100; ◊ = FTD128. Data
517	points for BW25113, JW2682 and JW3865 are mean values of triplicates, with
518	standard error shown.
519	
520	Fig. 4. TEM of thin sections of aerobically grown cells showing extracellular
521	palladium; (A) MC4100, inset BL21; (B) BW25113, inset BL21 (no Pd); (C)
522	FTD128; (D) JW2682; (E) MC4100 $\Delta moaA$; (F) JW3865. Scale bar (A) = 100 nm;
523	(B)-(F) = 500 nm; insets = $1 \mu m$.
524	
525	
526	
527	











Time from addition of formate (h)



























552 Figure 4

