

Dataset S1. Endogenous respiration rates in heterotrophic prokaryotes

Notes to Table **S1a**:

On data collection:

Prokaryote data were compiled by searching the www.pubmedcentral.nih.gov full-text library for "bacterium" and "endogenous respiration" and subsequent analysis of the returned 570 documents (mostly papers in the Journal of Bacteriology and Journal of Applied and Environmental Microbiology, time period 1940-2006) and references therein.

On cell size and taxonomy:

Data on endogenous respiration rates (i.e. respiration rates of non-growing cells in nutrient-deprived media) in heterotrophic eukaryotes are presented. Studies of bacterial respiration **very rarely** report information on cell size, which had therefore to be retrieved from different sources. To do so, an attempt was made to assign the bacterial strains described in the metabolic sources to accepted species names, to further estimate the cell size for these species in the relevant literature. This was done using strain designations and information in the official bacterial culture collections, like ATCC (American Type Culture Collection), NCTC (National Type Culture Collection) and others.

Column "**Species (Strain)**" gives the strain designation **as given by the authors** of the respiration data paper. Column "**Valid Name**" gives the relevant valid species name for this strain, as determined from culture collections' information and/or other literature sources. Valid names follow Euzéby (1997). Cell size in the "**Mpg**" column correspond to species indicated in the "Valid Name" column. Note that this information is of approximate nature, because many respiration data come from quite old publications and it was sometimes difficult to find out the valid name of the strain used with great precision. "**Class:Order**" column contains the relevant taxonomic information for the species listed in the "Valid Name" column as given by Euzéby (1997) (<http://www.bacterio.net>).

For example, Hareland et al. (1975) reported respiration rate for *Pseudomonas acidovorans* (ATCC strain number 17455). ATCC web site (www.atcc.org) says that this strain is *Delftia acidovorans* originally deposited as *Pseudomonas acidovorans*. Cell size for *Pseudomonas acidovorans* was therefore determined from the species description of *Delftia acidovorans* given by Wen et al. (1999). "**Class:Order**" for these data was determined as given at <http://www.bacterio.net> for *Delftia acidovorans*.

Note that the taxonomic uncertainty exclusively relates to the cell size determination. Cell size information participates in the paper's results only as a crude mean for all the 173 species studied, which is unlikely biased in any significant way. Taxonomic uncertainties, if any, do not influence any of the conclusions regarding the range, mean and frequency distribution of the prokaryotic respiration rates analysed in the paper.

Abbreviations and universal conversions: DM – dry mass; WM – wet mass; N – nitrogen mass; C – carbon mass; Pr – protein mass; X/Y – X by Y mass ratio in the cell, e.g. DM/WM is the ratio of dry to wet cell mass; $1 \text{ W} = 1 \text{ J s}^{-1}$; $1 \text{ mol O}_2 = 32 \text{ g O}_2$.

Original units are the units of endogenous respiration rate measurements as given in the original publication (**Source**); **qou** is the numeric value of endogenous respiration rate in the original units. E.g., if it is “ $\mu\text{l O}_2 (5 \text{ mg DM})^{-1} (2 \text{ hr})^{-1}$ ” in the column “**Original units**” and “200” in the column “**qou**”, this means that cells amounting to 5 mg dry mass consumed 200 microliters oxygen in two hours.

qWkg is the original endogenous respiration rate **qou** converted to W (kg WM)⁻¹ (Watts per kg wet mass) using the following conversion factors: C/DM = 0.5 (Kratz & Myers 1955; Bratbak & Dundas 1984; Nagata 1986), Pr/DM = 0.5 (Gronlund & Campbell 1961; Sobek et al. 1966; Smith & Hoare 1968 (see Table S1a); Zubkov et al. 1999), N/DM = 0.1 (SI Methods, Table S12b) if not indicated otherwise, and DM/WM = 0.3 as a crude mean for all taxa applied in the analysis (SI Methods, Table S12a). Energy conversion: 1 ml O₂ = 20 J. The respiratory quotient of unity was used (1 mol CO₂ released per 1 mol O₂ consumed).

TC is ambient temperature during measurements, degrees Celsius.

q25Wkg is endogenous respiration rate converted to 25 °C using Q₁₀ = 2, $\mathbf{q25Wkg} = \mathbf{qWkg} \times 2^{(25 - \text{TC})/10}$, dimension W (kg WM)⁻¹. For each species rows are arranged in the order of increasing **q25Wkg**.

Mpg: estimated cell mass, pg (1 pg = 10⁻¹² g). In most cases it is estimated from linear dimensions (using geometric mean of the available linear size range) assuming spherical cell shape for cocci and cylindrical shape for rods. Square brackets around the **Mpg** value indicate that the cell size information was obtained from a different source than the source of endogenous respiration rate data. When converting cell volume to cell mass, cell density of 1 g ml⁻¹ was assumed.

Source: the first, unbracketed reference in this column is where the value of **qou** is taken from; references and data in square brackets refer to cell size determination. Cell size reference "BM" in brackets corresponds to Bergey's Manual of Systematic Bacteriology, 1st Edition (Holt, 1984, 1986, 1989); BM9 is Bergey's Manual of Determinative Bacteriology, 9th Edition (Holt et al. 1994). Word "genus" in brackets indicates that cell size is determined as mean for the genus. This was done for those genera where the range of minimum to maximum cell masses did not exceed a factor of ten. E.g. for an unknown Chromatium sp. (BM9 genus: rods 1-6×1.5-15 μm, which corresponds to cell mass range from 1.2 to 420 pg) cell mass was left undetermined (empty "**Mpg**" column).

Culture age: Information on culture age and the duration of respiration measurements, if available.

Comments: this column provides relevant information on culture conditions and cellular composition of the studied species, often including additional data on respiration rates that were obtained for the same strain (species) by the same group of authors.

Log₁₀-transformed values of **q25Wkg** (W (kg WM)⁻¹), minimum for each species, were used in the analyses shown in Figures 1-3 and Table 1 in the paper (a total of 173 values for n = 173 species). The corresponding rows are highlighted in [blue](#).

References within Table S1a to Tables, Figures etc. refer to the corresponding items in the original literature indicated in the **Source** column.

Table S1a. Endogenous respiration rates in heterotrophic prokaryotes.

Species (strain)	Valid name	Class: Order	Original units	MIN	gou	qWkg	TC	q25Wkg	Mpg	Source	Culture age	Comments
Acetobacter (Ch 31)	1.Acetobacter aceti	Clostridia: Clostridiales	$\mu\text{mol O}_2 (1.8 \times 45 \text{ mg WM})^{-1} (5 \text{ hr})^{-1}$	MIN	2	0.6	30	0.42	[0.75]	De Ley & Schell 1959 [BM, ellipsoid or rod-shaped 0.6-0.8x1.0-4.0 μm]	Cells incubated for 4-5 days on gelatin slants at 20 C; respiration measured for 5 hr	Cells additionally incubated for 2-3 days at 30 C in a shaking apparatus "occasionally displayed higher endogenous respiration, $13.5 \mu\text{mol O}_2 (1.8 \times 45 \text{ mg WM})^{-1} (2.5 \text{ hr})^{-1} = 8.3 \text{ W/kg}$; this respiration increased exponentially during incubation
Acholeplasma laidlawii (NCTC 10116)	2.Acholeplasma laidlawii	Mollicutes: Acholeplasmatales	$\text{nmol O}_2 (\text{mg protein})^{-1} \text{min}^{-1}$	MIN	1.2	1.4	37	0.61	[0.04]	Abu-Amero et al. 1996 [Wieslander et al. 1987, sphere diam 0.3-0.6 μm]	Cells harvested after 24-72 hr incubation	
Hydrogenomonas ruhlandii	3.Achromobacter ruhlandii	Betaproteobacteria: Burkholderiales	$\mu\text{l O}_2 (0.5 \text{ mg DM})^{-1} (2 \text{ hr})^{-1}$	MIN	9	15	30	10.61	0.2	Packer & Vishniac 1955 [rods 0.4-0.75x0.75-2.0 μm , mean 0.5x1.1 μm]	Bacteria harvested after 4-5 days' incubation on agar plates; respiration of resting cells measured for 2 h	Hydrogen-oxidizing bacterium isolated from soil
Achromobacter sp.	4.Achromobacter sp.	Betaproteobacteria: Burkholderiales	$\mu\text{mol O}_2 (100 \text{ mg DM})^{-1} (2 \text{ hr})^{-1}$	MIN	45.1	8.4	30	5.94	[0.6]	Gronlund & Campbell 1961 [Chester & Cooper 1979, 0.5-0.8x1.5-2.5 μm]	cells harvested after 20 hr growth; respiration measured for 2 hr	Classification and size determination made for the Achromobacter genus as described at www.bacterio.cict.fr . There is no such species at www.bacterio.cict.fr
Achromobacter (B8)	5.Achromobacter sp.	Betaproteobacteria: Burkholderiales	$\mu\text{l O}_2 (5 \text{ mg DM})^{-1} (2 \text{ hr})^{-1}$		200	35	30	24.75	[0.6]	Tomlinson & Campbell 1963 [Chester & Cooper 1979, 0.5-0.8x1.5-2.5 μm]	cells harvested after 20 hr growth; respiration measured for 2 hr	No drop of respiration during the first two hours Classification and size determination made for the Achromobacter genus as described at www.bacterio.cict.fr . There is no such species at www.bacterio.cict.fr
Achromobacter viscosus (ATCC 12448)	6.Achromobacter viscosus	Betaproteobacteria: Burkholderiales	$\mu\text{l O}_2 (5 \text{ mg DM})^{-1} (2 \text{ hr})^{-1}$	MIN	200	35	30	24.75	[0.6]	Tomlinson & Campbell 1963 [Chester & Cooper 1979, 0.5-0.8x1.5-2.5 μm]	cells harvested after 20 hr growth; respiration measured for 2 hr	No drop of respiration during the first two hours Classification and size determination made for the Achromobacter genus as described at www.bacterio.cict.fr . There is no such species at www.bacterio.cict.fr
Achromobacter xerosis (ATCC 14780)	7.Achromobacter xerosis	Betaproteobacteria: Burkholderiales	$\mu\text{l O}_2 (\text{mg DM})^{-1} \text{hr}^{-1}$	MIN	14	23	30	16.26	[0.5]	Jurtshuk & McQuitty 1976 [Groupé et al. 1954, 0.5x2-3 μm]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	Classification and size determination made for the Achromobacter genus as described at www.bacterio.cict.fr . There is no such species at www.bacterio.cict.fr
Hydrogenomonas facilis	8.Acidovorax facilis	Betaproteobacteria: Burkholderiales	$\mu\text{l O}_2 (\text{mg DM})^{-1} \text{hr}^{-1}$	MIN	4-11	7	30	4.95	0.3	Schatz & Bovell 1952 [rods, 0.4x2.5 μm in	Heterotrophic cultures: cells grown for 48, 21, 72, and 48 hr on lac-	Synonym Pseudomonas facilis

Strain	Reference	Phylum	Class	Order	Family	Genus	Species	Substrate	Temperature (°C)	pH	W/k	Notes					
Acinetobacter calcoaceticus (ATCC 19606)	9.Acinetobacter baumannii	Gammaproteobacteria:	Pseudomonadales					μl O ₂ (mg DM) ⁻¹ hr ⁻¹	MIN	4	12	30	8.49	[2]	Jurtshuk & McQuitty 1976 [BM, genus, rods 0.9-1.6× 1.5-2.5]	heterotrophic cultures, 0.3×2.0 in autotrophic cultures]	tate, succinate, glucose and tryptose, respectively.
Acinetobacter calcoaceticus (208)	10.Acinetobacter calcoaceticus	Gammaproteobacteria:	Pseudomonadales					μl O ₂ (mg DM) ⁻¹ hr ⁻¹	MIN	7	6.7	30	4.74	[2]	Jurtshuk & McQuitty 1976 [BM, genus, rods 0.9-1.6× 1.5-2.5]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)
Acinetobacter johnsonii (210A)	11.Acinetobacter johnsonii	Gammaproteobacteria:	Pseudomonadales					nmol O ₂ (mg protein) ⁻¹ min ⁻¹	MIN	41	46	30	32.53	[2]	van Veen et al. 1993 [BM, genus, rods 0.9-1.6× 1.5-2.5]	Cells harvested at the logarithmic phase	When starved for 12 hours, respiration decreases to "very low rates" but when glucose is added, returns back to the higher level indicating no loss of viability. This suggests that 41 W/kg is an overestimate.
Acinetobacter calcoaceticus (ATCC 31012)	12.Acinetobacter sp.	Gammaproteobacteria:	Pseudomonadales					μl O ₂ (mg DM) ⁻¹ hr ⁻¹	MIN	2.0	3.3	25	3.30	[2]	Bruheim et al. 1999 [BM, genus, rods 0.9-1.6× 1.5-2.5]	Cells grown to the early stationary phase on oil	
Acinetobacter sp.	13.Acinetobacter sp.	Gammaproteobacteria:	Pseudomonadales					μl O ₂ (4 mg DM) ⁻¹ min ⁻¹	MIN	0.25	6	30	4.24	[2]	Sparnins et al. 1974 [BM, genus, rods 0.9-1.6× 1.5-2.5]	Bacteria grown overnight to the stationary phase (Dagley & Gibson 1975)	Gram-negative, oxidase-negative coccobacillus that cannot utilize glucose was isolated from agricultural soil in St. Paul, Minnesota, USA and tentatively identified as Acinetobacter sp.
Acinetobacter sp. (4-CB1)	14.Acinetobacter sp.	Gammaproteobacteria:	Pseudomonadales					nmol O ₂ (mg protein) ⁻¹ min ⁻¹	MIN	7	8	30	5.66	[2]	Adriaens et al. 1989 [BM, genus, rods 0.9-1.6× 1.5-2.5]	Cells grown to the late exponential phase	Bacterium isolated from soil contaminated with polychlorobiphenyl
Aeromonas hydrophila (ATCC 4715)	15.Aeromonas hydrophila	Gammaproteobacteria:	Aeromonadales					μl O ₂ (mg DM) ⁻¹ hr ⁻¹	MIN	23	38	30	26.87		Jurtshuk & McQuitty 1976	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Aeromonas hydrophila (ATCC 9071)	16.Aeromonas veronii	Gammaproteobacteria:	Aeromonadales					μl O ₂ (mg DM) ⁻¹ hr ⁻¹	MIN	12	20	30	14.14		Jurtshuk & McQuitty 1976	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Agrobacterium	17.Agrobacterium	Alphaproteobacteria:						μl O ₂ (mg DM) ⁻¹ hr ⁻¹	MIN	12	20	30	14.14	[1.5]	Jurtshuk &	Cells harvested at the	

Strain	Strain	Order	Respiration rate	MIN	MAX	Q ₁₀	Q ₂₀	Q ₃₀	Q ₄₀	Q ₅₀	References	Notes
tumefaciens (ATCC 15955)	tumefaciens	Rhizobiales	DM) ⁻¹ hr ⁻¹								McQuitty 1976 [BM9, genus, rods 0.6-1.0×1.5-3.0 μm]	late-logarithmic growth phase (two thirds of the maximal growth concentration)
Hydrogenomonas eutropha (ATCC 17697)	18.Alcaligenes eutrophus	Betaproteobacteria: Burkholderiales	μmol O ₂ (3.6 mg DM) ⁻¹ hr ⁻¹	MIN	8	83	33	47.67		[0.8]	Bongers 1970 [BM, rods, 0.7×1.8-2.6 μm]	Respiration of cells withdrawn from turbidostat (steady-state growth)
Alcaligenes faecalis (ATCC 8750)	19.Alcaligenes faecalis	Betaproteobacteria: Burkholderiales	μl O ₂ (mg DM) ⁻¹ hr ⁻¹	MIN	16	27	30	19.09			Jurtshuk & McQuitty 1976	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)
Alcaligenes (strain 5)	20.Alcaligenes sp.	Betaproteobacteria: Burkholderiales	μmol O ₂ (14 mg DM) ⁻¹ (5 hr) ⁻¹	MIN	4	2.1	30	1.48			Subba-Rao & Alexander 1985	Bacteria grown for 2 days; washed; incubated in buffer for 6 to 12 hr on a rotary shaker at 30 C to reduce endogenous respiration; washed again and added to respirometer flasks (Subba-Rao & Alexander 1977)
Unnamed methylo-troph (CC495)	21.Aminobacter lissarensis	Alphaproteobacteria: Rhizobiales	nmol O ₂ (mg WM) ⁻¹ min ⁻¹	MIN	0.22	1.6	25	1.60		[0.6]	Coulter et al. 1999 [BM9, genus, rods, 0.6-1.0×1.0-3.0 μm]	Cells harvested in the late exponential phase
Amoebobacter purpureus (ML1)	22.Amoebobacter purpureus	Gammaproteobacteria: Chromatiales	nmol O ₂ (mg protein) ⁻¹ min ⁻¹	MIN	9.8	11	30	7.78		[36]	Overmann & Pfennig 1992 [Eichler & Pfennig 1988, 3.3-3.8×3.5-4.5 μm]	Respiration of cells without microscopically visible sulfure globules at oxygen concentrations of 11-67 μM; respiration rates of phototrophically (anaerobically) and chemotrophically (microaerobically) grown cells do not differ; the species displays poor if any growth in the dark.
Amoebobacter roseus (6611)	23.Amoebobacter roseus	Gammaproteobacteria: Chromatiales	nmol O ₂ (mg protein) ⁻¹ min ⁻¹	MIN	4.8	5.4	30	3.82		[5]	Overmann & Pfennig 1992 [BM9, genus, spherical cells 1.5-3 μm diam]	Respiration of cells without microscopically visible sulfure globules at oxygen concentrations of 11-67 μM; respiration rates of phototrophically (anaerobically) and chemotrophically
												Hydrogen-oxidizing bacterium, max. respiration (in the presence of H ₂) is ten times the endogenous rate
												Bacteria isolated from enrichments with benzhydrol as the sole carbon source.
												Bacterium isolated from the top 5 cm of soil in a beech wood in Northern Ireland
												Purple sulfur bacteria isolated from the chemocline of meromictic Mahoney Lake (British Columbia, Canada)
												Endogenous respiration of cells with visible sulfur globules is lower, 5.7 nmol O ₂ (mg protein) ⁻¹ min ⁻¹
												Purple sulfur bacteria
												Endogenous respiration of cells with visible sulfur globules is higher, up to 22 nmol O ₂ (mg protein) ⁻¹ min ⁻¹

Amoebobaeter pendens (5813)	24.Amoebobaeter pendens	Gamma proteobacteria: Chromatiales	nmol O ₂ (mg protein) ⁻¹ min ⁻¹	MIN	7.6	8.5	30	6.01	[5]	Overmann & Pfennig 1992[BM9, genus, spherical cells 1.5-3 µm diam]	(microaerobically) grown cells do not differ; the species is capable of chemotrophic growth in the dark. Respiration of cells without microscopically visible sulfure globules at oxygen concentrations of 11-67 µM; respiration rates of phototrophically (anaerobically) and chemotrophically (microaerobically) grown cells do not differ.	Purple sulfur bacteria Endogenous respiration of cells with visible sulfur globules is higher, up to 35 nmol O ₂ (mg protein) ⁻¹ min ⁻¹
Spirillum (Aquaspirillum) itersonii (ATCC 12639)	25.Aquaspirillum itersonii	Beta proteobacteria: Neisseriales	µl O ₂ (mg DM) ⁻¹ hr ⁻¹	MIN	6	10	30	7.07	[0.9]	Jurtshuk & McQuitty 1976 [Krieg 1976, helical shape, Fig. 1E, Table 3, diam 0.3-0.4 µm, full length ~10 µm] Ensign 1970	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Arthrobacter tallopoietes	26.Arthrobacter crystallopoietes	Actinobacteria: Actinomycetales	µl O ₂ (mg DM) ⁻¹ hr ⁻¹	MIN	0.1	0.2	30	0.14	1.7		Cells harvested during the exponential phase of growth (48 hr for spherical cells, 4-8 hr for rods); stable endogenous respiration during 24 days of starvation at 100% viability	cell mass estimated from the dry mass data for spherical cells (0.5 mg dry mass per 10 ⁹ cells) Endogenous respiration at harvest was about 8-9 µl O ₂ (mg DM) ⁻¹ hr ⁻¹ and decreased 80-fold during the first two days of starvation Growing spherical cells contain about 40% (dry mass) of a glycogen-like polysaccharide; rods — 10% (Boylen & Ensign 1970) Boylen 1973: Bacteria of this species survived 6 months of extreme desiccation at 50% viability converting 0.0005% of their carbon per hour to carbon dioxide (≈ 10 ⁻² W/kg)
Arthrobacter globiformis (ATCC 8010)	27.Arthrobacter globiformis	Actinobacteria: Actinomycetales	µl O ₂ (mg DM) ⁻¹ hr ⁻¹	MIN	5	8.3	30	5.87	[0.5]	Jurtshuk & McQuitty 1976 [Conn & Dimmick 1947, rods 0.6-0.8x1-1.5 µm] Luscombe & Gray 1974	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Arthrobacter globiformis (NCIB)	28.Arthrobacter sp.	Actinobacteria: Actinomycetales	µl O ₂ (mg DM) ⁻¹ hr ⁻¹	MIN	0.45	0.75	25	0.75	0.2		Cells harvested from continuous cultures	Cocci survive better than rods; initial endogenous respiration was 1.74 and

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Arthrobacter sp.	29.Arthrobacter sp.	Actinobacteria: Actinomycetales	$\mu\text{l O}_2 \text{ (15 mg DM)}^{-1} \text{ (20 min)}^{-1}$	MIN	3.6	1.2	30	0.85	[1.5]	Devi et al. 1975 [BM9, genus, rods 0.8-1.2×1-8 μm ; Devi et al. 1975 indicate their strain is 2 μm in length]	Bacteria grown for 48 hr; endogenous respiration of resting cells measured for 80 min; data taken for the last 20 min	and starved for more than 2 days, steady-state viability approximately 80%. 7.33 $\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$ for cocci and rods, respectively, but “after 2 days these had both fallen to a relatively stable level of 0.45”, which was monitored for 7 days. Cell volume corresponds to the minimum dilution rate (0.01 h^{-1}); it grows up to 0.96 μm^3 at 0.3 h^{-1} . Data from Fig. 4a (induced cells), last 20 min of endogenous respiration; bacteria living in soil of Citrus plantations; rods 2 μm in length
Arthrobacter (TMP)	30.Arthrobacter sp.	Actinobacteria: Actinomycetales	$\mu\text{l O}_2 \text{ (50 mg WM)}^{-1} \text{ min}^{-1}$	MIN	0.9	6	30	4.24		Donnelly et al. 1981	Cells harvested during exponential growth	Respiration decreases with time (mean 2.3 W/kg) Isolated from soil
Arthrobacter (CA1)	31.Arthrobacter sp.	Actinobacteria: Actinomycetales	$\mu\text{mol O}_2 \text{ (5.3 mg DM)}^{-1} \text{ hr}^{-1}$	MIN	2	14	30	9.90		Ougham & Trudgill 1982	Bacteria harvested at late logarithmic phase; respiration measured for 2 hr	Bacteria isolated from field soil contaminated by aviation fuel, UK
Azotobacter agile	32.Azomonas agilis?	Gammaproteobacteria: Pseudomonadales	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$	MIN	12.6	21	26	19.59	13	Gunter & Kohn 1956	Cells harvested from 16 to 18-hr yeast agar plates	Cell mass estimated from dry mass data, Table 1, 3.8 pg DM/cell
Azorhizobium caulinodans (ORS571)	33.Azorhizobium caulinodans	Alphaproteobacteria: Rhizobiales	$\text{nmol O}_2 \text{ (mg protein)}^{-1} \text{ min}^{-1}$	MIN	34	38	30	26.87	[0.5]	Allen et al. 1991 [BM9, rods 0.5-0.6×1.5-2.5 μm]	Endogenous respiration of cells taken from continuous culture; measured for 10 min	
Azospirillum brasiliense (ATCC 29145)	34.Azospirillum brasiliense	Alphaproteobacteria: Rhodospirillales	$\mu\text{mol O}_2 \text{ (mg protein)}^{-1} \text{ min}^{-1}$	MIN	0.02	27	37	11.75	[1]	Loh et al. 1984 [BM, species image]	cells harvested during mid log-phase, starved for 4 hr at 4 C; constant respiration rate throughout the experiment (\approx 4 hr)	
Azospirillum lipoferum (ATCC 29707)	35.Azospirillum lipoferum	Alphaproteobacteria: Rhodospirillales	$\mu\text{mol O}_2 \text{ (mg protein)}^{-1} \text{ min}^{-1}$	MIN	0.03	39	37	16.98	[4]	Loh et al. 1984 [BM, species image]	cells harvested during mid log-phase, washed and starved for 4 hr at 4 C	Synonym Spirillum lipoferum (BM)
Azotobacter chroococcum (NCIB 8003)	36.Azotobacter chroococcum	Gammaproteobacteria: Pseudomonadales	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$	MIN	15	25	30	17.68	[14]	Bishop et al. 1962 [Bisset & Hale 1958, Figs. 1-3, 7, 13, diam approx. 3 μm]	Respiration measured immediately after harvesting the cells (aerobic culture) at the end of the logarithmic growth phase	Max. resp. (in the presence of glucose) was 130 $\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$
Azotobacter vinelandii (O)	37.Azotobacter vinelandii	Gammaproteobacteria: Pseudomonadales	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$	MIN	0.9	1.5	30	1.06	[0.5]	Sobek et al. 1966 [Tsai et al. 1979, Fig. 3, diam 1 μm , ATCC 12837, \approx 0.06 pgDM/cell, Fig. 1]	Respiration of glucose-grown cells harvested at 20 hr and starved for 48 hr (Table 1); viability >95% (Fig. 2)	During starvation respiration diminishes from 4.6-5.8 $\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$ (depending on growth substrate) during the first four hours and to 0.9-1.4 $\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$ between 48th and 52th hours

													Johnson et al. 1958 report that "well-washed cells of this organism possess no significant endogenous respiration"	Viability and respiration depend on the remaining store of PHB (poly-β-hydroxybutyric acid). 16-hr grown cultures were deprived of significant PHB stores and rapidly lost viability during starvation
Azotobacter vinelandii (O, ATCC 12518)	38.Azotobacter vinelandii	Gammaproteobacteria: Pseudomonadales	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$			19	30	30	21.21	[0.5]	Jurtshuk & McQuitty 1976 [Tsai et al. 1979, Fig. 3, diam 1 μm , ATCC 12837, $\approx 0.06 \text{ pgDM/cell}$, Fig. 1]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	Jurtshuk et al. 1975: Strain O cells grown to the late logarithmic phase, washed and "allowed to sit overnight at 4 C to reduce the intracellular endogenous reserve by starvation"; respiration at 30 C was 9-12 $\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1} = 15-20 \text{ W/kg}$	
Bacillus cereus (C5-25)	39.Bacillus cereus	"Bacilli": Bacillales	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$	MIN	8.5	14	37	6.09		[3.7]	Nickerson & Sherman 1952 [BM, species]	Normal (not filamentous) cells grown for 18 hr; respiration measured for about 1.5 hr; data of Table 3	N/D M=0.100-0.141	
Bacillus cereus (USDA)	40.Bacillus cereus	"Bacilli": Bacillales	$\mu\text{l O}_2 \text{ (mg N)}^{-1} \text{ hr}^{-1}$			188	31	30	21.92	[3.7]	Dietrich & Burris 1967 [BM, species]	Cells cultured for 2 weeks; respiration measured for 1 hr after 10 min equilibration	Church & Halvorson 1957 report 5 $\mu\text{l O}_2 \text{ (mg N)}^{-1} \text{ hr}^{-1} = 0.8 \text{ W/kg}$ at 30 C; heat-shocked spores stored for 4-48 months at -20 C	
Bacillus cereus	41.Bacillus cereus	"Bacilli": Bacillales	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$			56	97	30	68.59	[3.7]	Jurtshuk & McQuitty 1976 [BM, species]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)		
Bacillus firmus (ATCC 14575)	42.Bacillus firmus	"Bacilli": Bacillales	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$	MIN	8	13	30	9.19		[0.9]	Jurtshuk & McQuitty 1976 [Kanso et al. 2002, rods 0.6-0.9x1.2-4 μm]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)		
Bacillus megaterium	43.Bacillus megate-	"Bacilli": Bacillales	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$	MIN	2	3.3	30	2.33		[7]	Jurtshuk &	Cells harvested at the		

Strain	Reference	Taxonomy	Medium	Temperature (°C)	pH	Time (min)	Yield (mg)	Yield (DM)	Yield (O ₂)	Yield (O ₂ /DM)	Notes	
			DM) ⁻¹ hr ⁻¹							[[1.96]]	McQuitty 1976 [Bisset 1953, Fig. 17, rods 1.4×4.3 μm] [[Kubitschek 1969, Coulter counter]] late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Bacillus megaterium (KM)	44.Bacillus megaterium	“Bacilli”: Bacillales	μl O ₂ (mg DM) ⁻¹ hr ⁻¹	7.7	13	30	9.19	[7]	Frederick et al. 1974 [Bisset 1953, Fig. 17, rods 1.4×4.3 μm]	Flasks with washed cells equilibrated in Warburg bath for a total of 30 min; respiration measured for at least 60 min; respiration of cells starved for 2 hr (by shaking in phosphate buffer)	Obligately aerobic asporogenous strain Respiration of unstarved cells was 11 μl O ₂ (mg DM) ⁻¹ hr ⁻¹	
Bacillus megaterium (ATCC 19213)	45.Bacillus megaterium	“Bacilli”: Bacillales	natoms O (mg protein) ⁻¹ min ⁻¹	<40	25	30	17.68	[7]	Decker & Lang 1977 [Bisset 1953, Fig. 17, rods 1.4×4.3 μm]	Log-phase harvested cells (strain ATCC 19213) respired at a rate of less than 10% of 400 natoms O (mg protein) ⁻¹ min ⁻¹ = 25 W/kg; respiration measured for 1 min		
Bacillus megaterium (NCTC 9848)	46.Bacillus megaterium	“Bacilli”: Bacillales	μl O ₂ (mg DM) ⁻¹ hr ⁻¹	28	47	37	20.46	[7]	Bishop et al. 1962 [Bisset 1953, Fig. 17, rods 1.4×4.3 μm]	Cells harvested at the end of the logarithmic growth phase		
Bacillus megaterium (KM)	47.Bacillus megaterium	“Bacilli”: Bacillales	μl O ₂ (mg DM) ⁻¹ hr ⁻¹	35	58	25	58.00	[7]	Marquis 1965 [Bisset 1953, Fig. 17, rods 1.4×4.3 μm]	Cells harvested in the phase of declining growth rate; respiration measured for 90 min		
Bacillus popilliae (NRRL B-2309-P)	48.Bacillus popilliae	“Bacilli”: Bacillales	μl O ₂ (mg DM) ⁻¹ hr ⁻¹	MIN	0.5	0.8	30	0.57	[0.8]	Pepper & Costilow 1964 [Mitraka et al. 1967, Fig. 1, rods 0.6×2-4 μm]	Cells harvested in the stationary phase after 24 hr incubation	Bacterium causing "milky disease" of the Japanese beetle (Popilla japonica) larvae Similar respiration rates were measured for B. lentimorbus
Bacillus pumilus (ATCC 70)	49.Bacillus pumilus	“Bacilli”: Bacillales	μl O ₂ (mg DM) ⁻¹ hr ⁻¹	MIN	3	5	30	3.54	[0.7]	Jurtshuk & McQuitty 1976 [Hayase et al. 2004, rods 0.5-0.7×2.0-3.0 μm]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Bacillus stearothermophilus (PH24)	50.Bacillus stearothermophilus	“Bacilli”: Bacillales	μl O ₂ (14.2 mg DM) ⁻¹ (30 min) ⁻¹	MIN	35	8.2	50	1.45	[0.7]	Buswell 1975 [Montesinos et al. 1983, Coulter counter]	Cells harvested in late exponential phase (14-16 hr of growth on phenol) at 55 C and either used immediately or stored at -20 C; respiration measured for 30 min	The organism, obligate thermophile, was isolated from industrial sediment at Ravenscraig Steel Works near Motherwell, Scotland.
Bacillus subtilis (W-)	51.Bacillus subtilis	“Bacilli”: Bacillales	μl O ₂ (mg MIN)	2	3.3	30	2.33	[1.4]	Jurtshuk &	Cells harvested at the	Crook 1952 reports 4-5 μl O ₂ (mg	

Strain	Genus	Species	Phylum	Class	Order	Family	Genus	Species	Strain	Reference	Notes	
rioovorus (109J)	bacterioovorus		Bdellovibrionales						17.3	1973 [DM=10 ⁻¹³ pg/cell]	24 hr in a medium containing E. coli cells that were all lysed by the time of harvest; respiration measured for 2-4 hr Rittenberg & Shilo 1970: 13-27 nmol O ₂ (0.42 mg protein) ⁻¹ min ⁻¹ = 35-72 W/kg at 30 C	Rittenberg & Shilo 1970 report 0.42 mg protein per 10 ¹⁰ cells of strain 109 Straley et al. (1979) characterize this respiration rate as "unusually high"; Friedberg & Friedberg (1976) as "extremely high"
Pseudomonas natriegens	57.Beneckea natriegens		Gammaproteobacteria: "Vibrionales"						MIN 85 265 30 187.38 [1.5]	Cho & Eagon 1967 [Baumann et al. 1971, Figs. 12, 16, rods 0.6-1.2x1.9-3.6 µm depending on culture conditions?]	Cells harvested at the end of the logarithmic phase; respiration of resting cells measured for 45 min Oxygen uptake with all substrates is characterized as "low" (the lowest with glucose, 268 µl O ₂ (mg DM) ⁻¹ hr ⁻¹ = 450 W/kg is within the upper range of maximum specific metabolic rates in bacteria Marine bacterium with shortest known generation time (9.8 min) (Eagon 1962) Synonym Vibrio natriegens	
Bradyrhizobium japonicum (I-110)	58.Bradyrhizobium japonicum		Alphaproteobacteria: Rhizobiales						MIN 53 1.0 29 0.76 [0.7]	Frustaci et al. 1991 [BM]	Respiration of late-log phase cells	
Neisseria catarrhalis	59.Branhamella catarrhalis		Gammaproteobacteria: Pseudomonadales						MIN 1 1.7 37 0.74 [1.3]	Bishop et al. 1962 [Baumann et al. 1968, diam 1.0-1.7 µm, coccoid]	Respiration measured immediately after harvesting the cells at the end of the logarithmic growth phase	Max. resp. (in the presence of glucose) was 10 µl O ₂ (mg DM) ⁻¹ hr ⁻¹
Branhamella catarrhalis (Gp4)	60.Branhamella catarrhalis		Gammaproteobacteria: Pseudomonadales						14 23 30 16.26 [1.3]	Jurtshuk & McQuitty 1976 [Baumann et al. 1968, diam 1.0-1.7 µm, coccoid]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	Respiration of Branhamella (Neisseria) catarrhalis (ATCC 25238) and Branhamella catarrhalis (NC31) was 16 and 17 µl O ₂ (mg DM) ⁻¹ hr ⁻¹ , respectively (27-28 W/kg)
Brucella abortus (19)	61.Brucella melitensis		Alphaproteobacteria: Rhizobiales						MIN 40-82 6.5 34 3.48 [0.3]	Gerhard et al. 1950 [BM9, genus, cocci or short rods 0.5-0.7x0.6-1.5 µm]	Cells grown for 24 hr; respiration varies among equally prepared suspensions	
Burkholderia sp. (JT 1500)	62.Burkholderia sp.		Betaproteobacteria: Burkholderiales						MIN 19 21 30 14.85 0.7	Morawski et al. 1997 [rods 0.5-0.8x1.5-3.0]	Cells grown for 15-18 hr	Bacterium isolated from Miami soil
Cellvibrio gilvus	63.Cellvibrio gilvus		Gammaproteobacteria: Pseudomonadales						MIN 10 37 30 26.16 2	Hulcher & King 1958a [Hulcher & King 1958b, rods 0.75-1.5x1.5-3.75 µm]	Cells grown on cellobiose for 18 hr at room temperature	Cells in young cultures are very large (3x10 µm), 24-hr cells grown on cellulose were 0.3-0.5x0.8-1.5 µm (Hulcher & King 1958b) There is no such species at http://www.bacterio.cict.fr ; higher taxon as for Cellvibrio genus.

Chromatium sp. (Miami PBS1071)	64.Chromatium sp.		Gamma	proteobacteria: Chromatiales	nmol O ₂ (mg DM) ⁻¹ min ⁻¹	MIN	4.8	11	25	11.00		Kumazawa et al. 1983 [BM9 genus: rods 1-6×1.5-15 μm]	2-3 days' old cultures grown anaerobically in the light; respiration increased severalfold upon addition of H ₂ and fell to the endogenous level after H ₂ was exhausted	Marine purple sulfur bacterium
Chromatium vinosum (2811)	65.Chromatium vinosum		Gamma	proteobacteria: Chromatiales	nmol O ₂ (mg protein) ⁻¹ min ⁻¹	MIN	2.0	2.2	30	1.56	[1.5] [[1.21]]	Overmann & Pfennig 1992 [Montesinos et al. 1983, Coulter counter] [[Mas et al. 1985]]	Respiration of cells without microscopically visible sulfure globules at oxygen concentrations of 11-67 μM; respiration rates of phototrophically (anaerobically) and chemotrophically (microaerobically) grown cells do not differ; the species is capable of chemotrophic growth in the dark.	Purple sulfur bacteria Respiration rate increases with oxygen concentration reaching a plateau at approx. 6 μM. Endogenous respiration of cells with visible sulfur globules is higher, up to 24 nmol O ₂ (mg protein) ⁻¹ min ⁻¹
Corynebacterium diphtheriae (ATCC 11913)	66.Corynebacterium diphtheriae		Actinobacteria: Actinomycetales		μl O ₂ (mg DM) ⁻¹ hr ⁻¹	MIN	4	6.7	30	4.74		Jurtshuk & McQuitty 1976 [Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Corynebacterium sp.	67.Corynebacterium sp.		Actinobacteria: Actinomycetales		μl O ₂ (100 mg WM) ⁻¹ (120 min) ⁻¹	MIN	180	10	30	7.07		Levine & Kram-pitz 1952	Cells harvested after 2-4 days growth; respiration measured for 2 hr	A soil-isolated bacterium sometimes capable of acetone degradation
Pseudomonas acidovorans (ATCC 17455=NCIB 10013)	68.Delftia acidovorans		Beta	proteobacteria: Burkholderiales	μl O ₂ (4 mg DM) ⁻¹ min ⁻¹	MIN	0.5	13	30	9.19	[0.8]	Hareland et al. 1975 [Wen et al. 1999, rods, 0.4-0.8×2.5-4.1 μm]	Bacteria grown for 20 hr (end of logarithmic growth); respiration measured for about 10 min	Bacteria originally isolated from poultry house deep-litter
Pseudomonas B ₂ aba sp.	69.Delftia acidovorans		Beta	proteobacteria: Burkholderiales	μl O ₂ (mg DM) ⁻¹ hr ⁻¹		8	13	30	9.19	[0.8]	Kornberg & Gotto 1961 [Wen et al. 1999, rods, 0.4-0.8×2.5-4.1 μm]	Cells grown on glycollate harvested during the logarithmic phase; for succinate-grown cells 10 μl O ₂ (mg DM) ⁻¹ hr ⁻¹ (17 W/kg). Cooper & Kornberg 1964: cells grown on itaconate harvested during the logarithmic phase, 39 μl O ₂ (mg DM) ⁻¹ hr ⁻¹ = 65 W/kg; for succinate-grown	

Pseudomonas desmolytica (S449B1)	70.Delftia vorans	acido-	Betaproteobacteria: Burkholderiales	$\mu\text{l O}_2$ (0.6 mg DM) ⁻¹ (160 min) ⁻¹		14	15	30	10.61	[0.8]	Jigami et al. 1979 [Wen et al. 1999, rods, 0.4-0.8×2.5-4.1 μm]	cells 40 $\mu\text{l O}_2$ (mg DM) ⁻¹ hr ⁻¹ Cells harvested after 3 days of growth	
Pseudomonas acidovorans (ATCC 15688)	71.Delftia vorans	acido-	Betaproteobacteria: Burkholderiales	$\mu\text{l O}_2$ (mg DM) ⁻¹ hr ⁻¹		18	30	30	21.21	[0.8]	Jurtshuk & McQuitty 1976 [Wen et al. 1999, rods, 0.4-0.8×2.5-4.1 μm]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Desulfovibrio salexigens (Mast1)	72.Desulfovibrio salexigens		Deltaproteobacteria: Desulfovibrionales	nmol O ₂ (mg protein) ⁻¹ min ⁻¹	MIN	12	13	30	9.19	[1.5]	van Niel & Gottschal 1998 [BM]	Cells harvested at the end of the exponential growth phase	Organism isolated from the oxic-anoxic (top) layer of a marine microbial mat, Greece
Enterobacter aerogenes	73.Enterobacter aerogenes		Gammaproteobacteria: "Enterobacteriales"	$\mu\text{l O}_2$ (mg DM) ⁻¹ hr ⁻¹	MIN	6	10	30	7.07	[0.3]	Jurtshuk & McQuitty 1976 [Fu et al. 2003, rods 0.6×1.2 μm]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Enterobacter cloacae (17/97)	74.Enterobacter cloacae		Gammaproteobacteria: "Enterobacteriales"	nmol O ₂ (mg DM) ⁻¹ min ⁻¹	MIN	13.9	31	30	21.92	[0.9]	Majtán & Majtánová 1999 [BM9, genus, rods 0.6-1.0×1.2-3.0 μm]	Cells harvested from the exponential phase; respiration measured for 10 min	Bacterium isolated from a patient suffering from nosocomial infection.
Enterococcus cecorum (DSM 20682 ^T)	75.Enterococcus cecorum		"Bacilli": "Lactobacillales"	nmol O ₂ (mg DM) ⁻¹ min ⁻¹	MIN	1.7	3.8	30	2.69	[0.4]	Bauer et al. 2000 [BM]	Respiration of cells from aerobically glucose-grown cultures; respiration of anaerobically grown cells <1 nmol O ₂ (mg DM) ⁻¹ min ⁻¹	
Streptococcus faecalis (NCDO 581)	76.Enterococcus faecalis		"Bacilli": "Lactobacillales"	$\mu\text{l O}_2$ (mg DM) ⁻¹ hr ⁻¹	MIN	0.16	0.3	30	0.21	[0.2]	Bryan-Jones & Whittenbury 1969 [BM]	respiration of resting suspensions of cells grown aerobically with glucose	
Streptococcus faecalis (ATCC 8043)	77.Enterococcus hirae		"Bacilli": "Lactobacillales"	$\mu\text{mol O}_2$ (100 mg DM) ⁻¹ (2 hr) ⁻¹	MIN	8.0	1.5	30	1.06		Gronlund & Campbell 1961	Cells harvested after 17 hr growth; respiration measured for 2 hr	
Enterococcus hirae (ATCC 8043, ATCC 9790)	78.Enterococcus hirae		"Bacilli": "Lactobacillales"	$\mu\text{l O}_2$ (mg DM) ⁻¹ hr ⁻¹		1	1.7	30	1.20		Jurtshuk & McQuitty 1976	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Enterococcus sp. (RfL6)	79.Enterococcus sp.		"Bacilli": "Lactobacillales"	nmol O ₂ (mg DM) ⁻¹ min ⁻¹	MIN	15.1	33	30	23.33	0.8	Bauer et al. 2000 [rods, 0.6-1×1.1-3 μm]	Respiration of cells from aerobically glucose-grown cultures; respiration of anaerobically grown cells	

Escherichia coli (EMG-2 K12 Ymel)	80.	Escherichia coli	Gammaproteobacteria: "Enterobacteriales"	ngatom O (mg protein) ⁻¹ min ⁻¹	MIN	58-86	32	37	14	[0.7]	Lawford & Had-dock 1973 [Kubitschek 1969, Coulter counter, 0.33-1.46 μm ³ depending on growth rate]	15.1 nmol O ₂ (mg DM) ⁻¹ min ⁻¹ Cells were grown to the early exponential phase and starved for 2 hr by vigorous shaking at 37 C; respiration varied (depending on carbon source for growth)	
Escherichia coli (W-1485)	81.	Escherichia coli	Gammaproteobacteria: "Enterobacteriales"	μl O ₂ (mg DM) ⁻¹ hr ⁻¹		2.7	4.5	26	4.20	[0.7]	Gunter & Kohn 1956 [Kubitschek 1969, Coulter counter, 0.33-1.46 μm ³ depending on growth rate]	Cells harvested from 16 to 18-hr yeast agar plates	
Escherichia coli (O11a ₁)	82.	Escherichia coli	Gammaproteobacteria: "Enterobacteriales"	μl O ₂ (mg DM) ⁻¹ hr ⁻¹		4	6.7	30	4.74	[0.7]	Jurtshuk & McQuitty 1976 [Kubitschek 1969, Coulter counter, 0.33-1.46 μm ³ depending on growth rate]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	Respiration was 13, 7, 9, 10, 6, 8, 7, 6, and 5 μl O ₂ (mg DM) ⁻¹ hr ⁻¹ for strains B, B/r, C, Crookes, K12, K12S UTH 2593, K12S UTH 3672, and F, respectively (8.3-22 W/kg).
Escherichia coli	83.	Escherichia coli	Gammaproteobacteria: "Enterobacteriales"	μl O ₂ (mg DM) ⁻¹ hr ⁻¹		7	12	37	5.22	[0.7]	Dawes & Ribbons 1965 [Kubitschek 1969, Coulter counter, 0.33-1.46 μm ³ depending on growth rate]	Stable respiration of aerobically grown cells harvested during the exponential phase and starved aerobically for 150-180 min; no loss of viability during 12 hr of starvation	"Stationary-phase cells respire endogenously at higher rates and contain larger reserves of glycogen, which is the initial substrate oxidized [than exponential-phase cells]. Carbohydrate content of exponential-phase cells drops rapidly together with endogenous respiration during the first 100 min of starvation
Escherichia coli (B)	84.	Escherichia coli	Gammaproteobacteria: "Enterobacteriales"	μl O ₂ (10 mg DM) ⁻¹ (2 hr) ⁻¹		180	15	35	7.50	[0.7]	Sobek & Talburt 1968 [Kubitschek 1969, Coulter counter, 0.33-1.46 μm ³ depending on growth rate]	Cells grown for 20 hr at 35 C; respiration measured for 2 hr	
Escherichia coli (ATCC 4157)	85.	Escherichia coli	Gammaproteobacteria: "Enterobacteriales"	μl O ₂ (mg DM) ⁻¹ hr ⁻¹		12	20	37	8.71	[0.7]	Davis & Bateman 1960 [Kubitschek 1969, Coulter counter, 0.33-1.46 μm ³ depending on growth rate]	Cells harvested after 16.5 hr growth at 37 C; respiration measured for 2 hr	
Escherichia coli (ATCC 6894)	86.	Escherichia coli	Gammaproteobacteria: "Enterobacteriales"	μmol O ₂ (100 mg DM) ⁻¹ (2 hr) ⁻¹		79	15	30	10.61	[0.7]	Gronlund & Campbell 1961 [Kubitschek 1969, Coulter counter, 0.33-1.46 μm ³ depending on	Cells harvested after 20 hr growth; respiration measured for 2 hr	

Strain	Reference	Phylum	Class	Order	Family	Genus	Species	Strain ID	Medium	Temperature (°C)	pH	Light	Shaking	Respiration Rate (μl O ₂ (mg N) ⁻¹ hr ⁻¹)	Notes						
<i>Escherichia coli</i> (Gratia)	87.Escherichia coli	Gammaproteobacteria:	"Enterobacteriales"											105	17.5	30	12.37	[0.7]	growth rate] Dietrich & Burris 1967 [Kubitschek 1969, Coulter counter, 0.33-1.46 μm ³ depending on growth rate]	Cells cultured for 2 weeks; respiration measured for 1 hr after 10 min equilibration	Bacteria "had a lower endogenous [respiration] ... if harvested during exponential growth than after reaching the stationary phase"
<i>Escherichia coli</i>	88.Escherichia coli	Gammaproteobacteria:	"Enterobacteriales"											37	62	37	26.99	[0.7]	Bishop et al. 1962 [Kubitschek 1969, Coulter counter, 0.33-1.46 μm ³ depending on growth rate]	Respiration measured immediately after harvesting the cells (aerobic culture) at the end of the logarithmic growth phase	Anaerobically grown culture respired at 2 μl O ₂ (mg DM) ⁻¹ hr ⁻¹ = 3.3 W/kg
<i>Flavobacterium capsulatum</i> (ATCC 14666)	89.Flavobacterium capsulatum	Flavobacteria:	Flavobacteriales											38	63	30	44.55	[0.3]	Jurtshuk & McQuitty 1976 [BM9, genus, rods 0.5×1.0-3.0 μm]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	Listed in ATCC as <i>Novosphingobium capsulatum</i>
<i>Pasteurella tularensis</i> (SCHU S4, 38 A)	90.Francisella tularensis	Gammaproteobacteria:	Thiotrichales											22	3.7	37	1.61	[0.01]	Weinstein et al. 1962 [BM9, rods 0.2×0.2-0.7 μm]	Cells grown for 16 to 18 hr; respiration of strains 503, CHUR, JAP, LVS, and MAX was 60, 38, 53, 105, and 94 μl O ₂ (mg N) ⁻¹ hr ⁻¹ , respectively (6.3-17.5 W/kg)	
<i>Frankia</i> (EAN1 _{pec}) sp.	91.Frankia sp.	Actinobacteria:	Actinomycetales											242	9	25	9.00		Tisa & Ensign 1987 [BM, genus, hyphae 0.5-2.0 μm diam]	Bacteria grown for 21 days	Members of the genus <i>Frankia</i> are filamentous actinomycetes that infect roots and induce nodule formation in a variety of woody dicotyledonous plants.
<i>Haemophilus influenzae</i> (641b)	92.Haemophilus influenzae	Gammaproteobacteria:	Pasteurellales											3	0.5	37	0.22	[0.14]	Biberstein & Spencer 1962 [BM, coccobacilli or small regular rods 0.3-0.5 - 0.5-3.0 μm]	Bacteria grown for 18-24 hr at 37 C; respiration of washed cells measured for 60 min	Respiration of strains b, c, d, 525, and K 75 was 14, 6, 7, 32, and 9 μl O ₂ (mg N) ⁻¹ hr ⁻¹ , respectively (1-5.3 W/kg).
<i>Haemophilus parahaemolyticus</i> (9796)	93.Haemophilus parahaemolyticus	Gammaproteobacteria:	Pasteurellales											21	3.5	37	1.52		Biberstein & Spencer 1962	Bacteria grown for 18-24 hr at 37 C; respiration of washed cells measured for 60 min	Respiration of strain 7901 was 24 μl O ₂ (mg N) ⁻¹ hr ⁻¹ (4 W/kg).
<i>Haemophilus parainfluenzae</i> (K 98)	94.Haemophilus parainfluenzae	Gammaproteobacteria:	Pasteurellales											21	3.5	37	1.52	[0.4]	Biberstein & Spencer 1962 [Kahn 1981, Fig. 1, 0.6×1.3-1.7 μm; Kowalski et al. 1991, diam 0.75-1.25 μm]	Bacteria grown for 18-24 hr at 37 C; respiration of washed cells measured for 60 min	Respiration of strains K 8, K 17, and K 45 was 24, 32, and 41 μl O ₂ (mg N) ⁻¹ hr ⁻¹ , respectively (4-6.8 W/kg).
<i>Halobacterium salinarum</i> (1)	95.Halobacterium salinarum	Archaea: Halobacteria:	Halobacteriales											10	17	30	12.02	[3.9]	Stevenson 1966 [Mescher & Strominger 1976,	Cells grown for about 70 hr to the end of the exponential phase	White 1962: Stationary-phase cells (>15 hr old) have an insignificant endogenous respiration. Log-phase cells have a small endogenous respiratory rate. An extremely halophilic bacterium

Micrococcus denitrificans	halo-	96.Halomonas halodenitrificans	Gammaproteobacteria: Oceanospirillales	$\mu\text{l O}_2 \text{ (2 mg DM)}^{-1} \text{ (30 min)}^{-1}$	MIN	40	67	25	67.00	[0.4]	Sierra & Gibbons 1962 [Ventosa et al. 1998, rods, 0.5-0.9x0.9-1.2 μm]	Cells harvested towards the end of the logarithmic phase (about 40 hr)	Endogenous respiration and viability of starved cells remained constant at around $40 \mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$ and 100%, respectively, for 3 hr, while the amount of intracellular polyester rapidly declined. After 3 hr both endogenous respiration and viability started to decline rapidly. In polyester-poor cells this process was initiated immediately after the beginning of starvation.
Aerobacter aerogenes	aerogenes	97.Klebsiella pneumoniae? Enterobacter aerogenes?	Gammaproteobacteria: "Enterobacteriales"	$\mu\text{l O}_2 \text{ (mg N)}^{-1} \text{ hr}^{-1}$		67	11	30	7.78		Dietrich & Burris 1967	Bacteria cultured for 2 weeks in an extract from wheat plants; respiration measured for 1 hr after 10 min equilibration	Cells of this organism contain more nitrogen than other bacteria Bacteria "had a lower endogenous [respiration] ... if harvested during exponential growth than after reaching the stationary phase"
Aerobacter aerogenes	aerogenes	98.Klebsiella pneumoniae? Enterobacter aerogenes?	Gammaproteobacteria: "Enterobacteriales"	$\mu\text{mol O}_2 \text{ (100 mg DM)}^{-1} \text{ (2 hr)}^{-1}$		69.3	13	30	9.19		Gronlund & Campbell 1961	Cells harvested after 20 hr growth	
Klebsiella pneumoniae (M5a1)	pneumoniae	99.Klebsiella pneumoniae	Gammaproteobacteria: "Enterobacteriales"	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$	MIN	2	3.3	30	2.33	[0.4]	Jurtshuk & McQuitty [BM] 1976	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Klebsiella pneumoniae (ATCC 13882)	pneumoniae	100.Klebsiella pneumoniae	Gammaproteobacteria: "Enterobacteriales"	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$		4	6.7	30	4.74	[0.4]	Jurtshuk & McQuitty [BM] 1976	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Aerobacter aerogenes (NCTC 418)	aerogenes	101.Klebsiella pneumoniae	Gammaproteobacteria: "Enterobacteriales"	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$		12	20	37	8.71	[0.4]	Bishop et al. [BM] 1962	Respiration measured immediately after harvesting the cells at the end of the logarithmic growth phase	
Lactobacillus brevis (12)	brevis	102.Lactobacillus brevis	"Bacilli": "Lactobacillales"	$\mu\text{l O}_2 \text{ (12 mg DM)}^{-1} \text{ (2 hr)}^{-1}$	MIN	2.9	0.2	30	0.14	[1.6]	Walker 1959 [BM, rods 0.7-1.0x2-4]	Cells (strain 1.2) were grown for 48 hr; endogenous respiration measured for 3 hr (Fig. 3); mean respiration rate for the second and third hour was taken; during the first hour, respiration was 4 times higher	Bacterium originally isolated from New Zealand cheddar cheese Respiration decreases with starvation time
Lactobacillus casei sbsp. rhamnosus (ATCC 7469)	casei	103.Lactobacillus casei	"Bacilli": "Lactobacillales"	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$	MIN	1	1.7	30	1.20		Jurtshuk & McQuitty 1976	Cells harvested at the late-logarithmic growth phase (two	

Streptococcus lactis (strains 8, 9, 32)	104.Lactococcus lactis	“Bacilli”: “Lactobacillales”	$\mu\text{l O}_2$ (115 mg DM) ⁻¹ (4 hr) ⁻¹	MIN	170	0.6	30	0.42	[0.2]	Spendlove et al. 1957 [BM]	thirds of the maximal growth concentration) Cells grown for 11 hr; for 7- and 9-hr grown cells respiration was around 400 $\mu\text{l O}_2$ (115 mg DM) ⁻¹ (4 hr) ⁻¹	In 11-hr cells respiration decreases in the first 30 min of the 4-hr' measurement, then remains relatively constant; in 7-hr and 9-hr cells respiration first decreases, then starts to increase again after 2-3 hr.
Lactococcus ssp. lactis (DSM 20481 ^T)	105.Lactococcus lactis	“Bacilli”: “Lactobacillales”	nmol O ₂ (mg DM) ⁻¹ min ⁻¹		<1	2.3	30	1.63	[0.2]	Bauer et al. 2000 [BM]	Respiration of cells from aerobically and anaerobically glucose-grown cultures	
Lactococcus sp. (TmLO5)	106.Lactococcus sp.	“Bacilli”: “Lactobacillales”	nmol O ₂ (mg DM) ⁻¹ min ⁻¹	MIN	<1	2.3	30	1.63	0.8	Bauer et al. 2000 [rods, 0.6-1×1.1-3 μm]	Respiration of cells from aerobically and anaerobically glucose-grown cultures	
Legionella pneumophila (Knoxville-1, serotype 1)	107.Legionella pneumophila	Gammaproteobacteria: Legionellales	$\mu\text{l O}_2$ (mg DM) ⁻¹ min ⁻¹	MIN	0.2	20	37	8.71	[0.3]	Tesh et al. 1983 [Faulkner & Garduño 2002, rods 0.3-0.5×1.5-3.0 μm , prereplicative phase; Kowalski et al. 1999, 0.3-0.9×2 μm]	Bacteria harvested at mid- to late exponential growth phase (15-18 hr growth time) when the mass-specific oxygen consumption is highest (~70 W/kg); washed; “held at 37 C until used”; calibrated for 3 min; respiration measured when “a steady rate of endogenous respiration was established”.	
Legionella pneumophila (serogroup 1)	108.Legionella pneumophila	Gammaproteobacteria: Legionellales	$\mu\text{l O}_2$ (400 $\mu\text{g protein}$) ⁻¹ (40 min) ⁻¹		25	77	37	33.52	[0.3]	Bonach & Snyder 1983 [Faulkner & Garduño 2002, rods 0.3-0.5×1.5-3.0 μm , prereplicative phase; Kowalski et al. 1999, 0.3-0.9×2 μm]	Cells harvested at mid-log phase, shaken for 30 min, respiration measured for 40 min	
Pseudomonas AM1 (NCIB 9133)	109.Methylobacterium extorquens	Alphaproteobacteria: Rhizobiales	nmol O ₂ (mg protein) ⁻¹ min ⁻¹	MIN	5	6	30	4.24	[1]	Keevil & Anthony 1979 [Peel & Quayle 1961, rods 0.8×2.0 μm]	Cells harvested at the end of logarithmic phase; depending on growth conditions, respiration ranged from 5 to 16 orig. units; the same result obtained by O'Keeffe & Anthony 1978	
Methylomicrobium sp. (AMO 1)	110.Methylomicrobium sp.	Gammaproteobacteria: Methylococcales	nmol O ₂ (mg protein) ⁻¹ min ⁻¹	MIN	10	11	30	7.78	3	Sorokin et al. 2000 and personal communication	Cells grown with methane at pH 10 Endogenous respira-	

											with Dr. D.Yu. Sorokin (15 Nov 2006) (ovoid rods, 1-1.5×2-3 μm)	tion is usually below 10 nmol O ₂ (mg protein) ⁻¹ min ⁻¹ , except in cells grown on acetate and at high pH (10.8-11.5), when it can be 20-50 nmol O ₂ (mg protein) ⁻¹ min ⁻¹	
Methylophilus methylotrophus (NCIB 10515)	111.Methylophilus methylotrophus	Betaproteobacteria: Methylophiliales	ng-atoms O (mg DM) ⁻¹ min ⁻¹	MIN	1.4	1.6	40	0.57	[0.15]	Dawson & Jones 1981 [Jenkins et al. 1987, rods 0.3-0.6×0.8-1.5 μm]	Cells harvested from continuous culture and used within 3 hr	Obligate methylotroph using methanol as the sole source of carbon and energy	
Methylosinus trichosporium OB3b (ATCC 35070)	112.Methylosinus trichosporium	Alphaproteobacteria: Rhizobiales	nmol O ₂ (mg protein) ⁻¹ min ⁻¹	MIN	38	42	30	29.70	[1]	Lontoh et al. 1999 [Reed & Dugan 1978, Fig. 1]	Cells harvested in the exponential phase	Cell mass estimated from cell linear dimensions as shown in Fig. 1 of Reed & Dugan 1978	
Sarcina lutea	113.Micrococcus luteus	Actinobacteria: Actinomycetales	μl O ₂ (mg DM) ⁻¹ hr ⁻¹	MIN	0.7	1.2	37	0.52	[1.1]	Burleigh & Dawes 1967 [BM]	Cells were harvested after 24 hr growth on peptone; starved for 29 hr; viability 96%; respiration fell to "barely measurable" (0.3 μl O ₂ (mg DM) ⁻¹ hr ⁻¹) when starvation was prolonged to 72 hr with viability falling to 25%	Endogenous respiration decreased during starvation most rapidly in the first 5 hr of starvation (from 21.1 to 0.7 orig. units)	
											Bishop et al. 1962: respiration measured immediately after harvesting the cells at the end of the logarithmic growth phase was below detection limit (0 orig. units) (148 μl O ₂ (mg DM) ⁻¹ hr ⁻¹ in the presence of lactate)	Initial endogenous respiration depended on the time of harvesting: mid-exponential (21 hr) — 30 orig. units; onset of the stationary phase (34 hr) — 15.9; 60 hr — 6.3 orig. units; the decline of respiration is correlated with the decline of intracellular contents of free amino acids, carbohydrate etc.	
Sarcina lutea	114.Micrococcus luteus	Actinobacteria: Actinomycetales	μl O ₂ (mg DM) ⁻¹ hr ⁻¹		3.5	6	35	3.00	[1.1]	Dawes & Holms 1958 [BM]	Cells harvested after 24 hr growth at 35 C and aerated for 5 hr; respiration 3.5 μl O ₂ (mg DM) ⁻¹ hr ⁻¹ = 6 W/kg		
Micrococcus deikticus	lyso-115.Micrococcus luteus	Actinobacteria: Actinomycetales	μl O ₂ (mg DM) ⁻¹ hr ⁻¹		11	18	37	7.83	[1.1]	Davis & Bateman 1960 [BM]	Cells harvested after 16.5 hr growth at 37 C; respiration measured for 2 hr		
Sarcina flava	116.Micrococcus luteus	Actinobacteria: Actinomycetales	μl O ₂ (mg DM) ⁻¹ hr ⁻¹		9	15	30	10.61	[1.1]	Jurtshuk & McQuitty 1976 [BM]	Cells harvested at the late-logarithmic growth phase (two		

Micrococcus (S9)	sp.	117.Micrococcus sp.	Actinobacteria: Actinomycetales	$\mu\text{l O}_2 \text{ (4 mg DM)}^{-1} \text{ min}^{-1}$	MIN	0.40	10	30	7.07		Sparnins & Chapman 1976 [BM, genus, spherical cells, diam 0.5-2.0 μm]	thirds of the maximal growth concentration); respiration was 16, 9 and 30 $\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$ for Micrococcus (Sarcina) luteus, M. (S. flava) luteus and M. (lyso-deikticus) luteus ATCC 4698, respectively (27, 15 and 50 W/kg)	
Micrococcus	sp.	118.Micrococcus sp.	Actinobacteria: Actinomycetales	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$	MIN	10	17	30	12.02		Cooper et al. 1965 [BM, genus, spherical cells, diam 0.5-2.0 μm]	Cells harvested during the exponential phase	The organism was isolated from soil. Size and classification as for genus Micrococcus Cohn 1872
Moraxella osloensis		119.Moraxella osloensis	Gammaproteobacteria: Pseudomonadales	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$	MIN	8	13	30	9.19	[2.3]	Jurtshuk & McQuitty 1976 [Baumann et al. 1968, rods 0.9-1.7 \times 1.6-2.7 μm]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Mycobacterium fortuitum		120.Mycobacterium fortuitum	Actinobacteria: Actinomycetales	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$	MIN	10	17	30	12.02		Jurtshuk & McQuitty 1976	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Mycobacterium leprae		121.Mycobacterium leprae	Actinobacteria: Actinomycetales	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$	MIN	2	3.3	37	1.44		Mori et al. 1985	Cells isolated from mice	
Mycobacterium lepraemurium (Hawaiiian strain)		122.Mycobacterium lepraemurium	Actinobacteria: Actinomycetales	$\mu\text{l O}_2 \text{ (mg N)}^{-1} \text{ hr}^{-1}$	MIN	20-70	3.3	37	1.44		Gray 1952	Culture isolated from infected rats; occasionally, some suspensions respired less than 5 $\mu\text{l O}_2 \text{ (mg N)}^{-1} \text{ hr}^{-1}$	
Mycobacterium phlei (72)		123.Mycobacterium phlei	Actinobacteria: Actinomycetales	$\mu\text{mol O}_2 \text{ (mg N)}^{-1} \text{ hr}^{-1}$	MIN	4.8	14	37	6.09	[0.4]	Tepper [BM] 1968	Glucose-grown cells (strain 72) harvested from 5-day culture; respiration calculated assuming N/DM ratio of 0.08 (Tepper 1968); respiration measured for a minimum of 90 min; glycerol-grown cells had 7.3 $\mu\text{mol O}_2 \text{ (mg N)}^{-1} \text{ hr}^{-1}$ = 16 W/kg	N/DM=5.6-6.8% for glycerol-grown cells, 7.8-8.9% for glucose-grown cells

Mycobacterium phlei	124.Mycobacterium phlei	Actinobacteria: Actinomycetales	Acti-	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$		16	27	30	19.09	[0.4]	Jurtshuk & McQuitty 1976 [BM]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Mycobacterium smegmatis (607)	125.Mycobacterium smegmatis	Actinobacteria: Actinomycetales	Acti-	$\mu\text{l O}_2 \text{ (mg N)}^{-1} \text{ hr}^{-1}$	MIN	146	24	37	10.45		Forbes et al. 1962 [BM, genus, rods 0.2-0.6x3.0-3.5 μm]	Cells harvested after 22 hr incubation.	
Mycobacterium stercoris (NCTC 3820)	126.Mycobacterium smegmatis	Actinobacteria: Actinomycetales	Acti-	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$		11.9	20	30	14.14		Hunter 1953 [BM, genus, rods 0.2-0.6x3.0-3.5 μm]	Cells grown for 5-6 days; respiration of M. (butyricum) smegmatis (NCTC 337) and M. smegmatis (NCTC 523) was 13.4 and 15 $\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$, respectively	
Mycobacterium smegmatis	127.Mycobacterium smegmatis	Actinobacteria: Actinomycetales	Acti-	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$		13	22	30	15.56		Jurtshuk & McQuitty 1976 [BM, genus, rods 0.2-0.6x3.0-3.5 μm]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration); respiration was 13 $\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1} = 22 \text{ W/kg}$ at 30 C	
Mycobacterium tuberculosis var. hominis	128.Mycobacterium tuberculosis	Actinobacteria: Actinomycetales	Acti-	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ (5 hr)}^{-1}$	MIN	6.5	2.2	37	0.96	[0.2]	Engelhard et al. 1957 [BM, rods 0.2-0.5x2-4 μm]	Bacteria grown for 25 to 28 hr in batches; magnetically mixed in buffer solution for 5 hr; stored at -5 C for no more than 12 hr before analysis; respiration measured for 5 hr	Human tuberculosis bacteria
Mycobacterium tuberculosis var. hominis (H37Rv, LRv)	129.Mycobacterium tuberculosis	Actinobacteria: Actinomycetales	Acti-	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$		4.2-5.8	7	37.8	2.9	[0.2]	Segal & Bloch 1955 [BM, rods 0.2-0.5x2-4 μm]	Various growth conditions	
Mycobacterium tuberculosis var. avium	130.Mycobacterium tuberculosis	Actinobacteria: Actinomycetales	Acti-	$\mu\text{l O}_2 \text{ (10 mg WM)}^{-1} \text{ (90 min)}^{-1}$		100	37	37	16.11	[0.2]	Minami 1957 [BM, rods 0.2-0.5x2-4 μm]	Cultures 3 days old	
Mycobacterium tuberculosis var. hominis (H37Rv)	131.Mycobacterium tuberculosis	Actinobacteria: Actinomycetales	Acti-	$\mu\text{l O}_2 \text{ (mg WM)}^{-1} \text{ (hr)}^{-1}$		0.46-1.10	2.6-6	37	23.33	[0.2]	Youmans et al. 1960 [BM, rods 0.2-0.5x2-4 μm]	Various growth conditions. Similar results were obtained later by the same team (Youmans & Youmans 1962a,b)	
Myxococcus xanthus (FB)	132.Myxococcus xanthus	Deltaproteobacteria: Myxococcales		$\mu\text{l O}_2 \text{ (15 mg DM)}^{-1} \text{ (75 min)}^{-1}$	MIN	7	4.6	30	3.25	[1]	Dworkin & Niederpruem 1964 [McVittie et al. 1962, rods,	Vegetative cells harvested in the early stationary phase (30 hr); respiration meas-	No measurable respiration in microcysts

Neisseria elongata (ATCC 25295)	133.Neisseria elongata	Betaproteobacteria: Neisseriales	$\mu\text{l O}_2 (\text{mg DM})^{-1} \text{hr}^{-1}$	MIN	5	8.3	30	5.87	[0.2]	0.5×3-8 μm Jurtshuk & McQuitty 1976 [BM, short rods 0.5 μm width]	Incubated for 75 min Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Neisseria flava (ATCC 14221)	134.Neisseria flava	Betaproteobacteria: Neisseriales	$\mu\text{l O}_2 (\text{mg DM})^{-1} \text{hr}^{-1}$	MIN	7	12	30	8.49	[0.2]	Jurtshuk & McQuitty 1976 [BM, genus, cocci, diam 0.6-1.0 μm]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Neisseria gonorrhoeae	135.Neisseria gonorrhoeae	Betaproteobacteria: Neisseriales	$\text{nmol O}_2 (\text{mg protein})^{-1} \text{min}^{-1}$	MIN	0.5	0.6	37	0.26	[0.2]	Kenimer & Lapp 1978 [BM, genus, cocci, diam 0.6-1.0 μm]	Cells harvested during early stationary phase after 18-20 hours of incubation; respiration measured for 2-3 min	Cell mass estimated from linear dimensions given in BM
Neisseria mucosa	136.Neisseria mucosa	Betaproteobacteria: Neisseriales	$\mu\text{l O}_2 (\text{mg DM})^{-1} \text{hr}^{-1}$	MIN	9	15	30	10.61	[0.2]	Jurtshuk & McQuitty 1976 [BM, genus, cocci, diam 0.6-1.0 μm]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Neisseria sicca	137.Neisseria sicca	Betaproteobacteria: Neisseriales	$\mu\text{l O}_2 (\text{mg DM})^{-1} \text{hr}^{-1}$	MIN	8	13	30	9.19	[0.2]	Jurtshuk & McQuitty 1976 [BM, genus, cocci, diam 0.6-1.0 μm]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Nitrobacter agilis	138.Nitrobacter winogradskyi	Alphaproteobacteria: Rhizobiales	$\mu\text{l O}_2 (\text{mg protein})^{-1} \text{hr}^{-1}$	MIN	12	10	30	7.07	[0.24]	Smith & Hoare 1968 [Tappe et al. 1999, 0.17-0.3 μm^3]	Incubation time 20 hr	Facultative heterotroph otherwise growing on nitrite, =Nitrobacter winogradskyi (Pan 1971)
Nitrobacter agilis (ATCC 14123)	139.Nitrobacter winogradskyi	Alphaproteobacteria: Rhizobiales	$\text{ng-atoms O} (\text{mg protein})^{-1} \text{min}^{-1}$		20	11	25	11.00	[0.24]	Hollocher et al. 1982 [Tappe et al. 1999, 0.17-0.3 μm^3]	Cells grown at 30 C to the late exponential phase, stored for no more than 3 days at 0 C before measurements	Protein to dry mass ratios (Protein/DM) are 0.55, 0.47 and 0.44 for autotrophic, autotrophic plus 1mM acetate, and autotrophic plus 5mM acetate, grown cells, respectively. 1 mg WM \approx 0.1 mg protein
Nitrosomonas europaea	140.Nitrosomonas europaea	Betaproteobacteria: Nitrosomonadales	$\text{ng-atoms O} (\text{mg protein})^{-1} \text{min}^{-1}$	MIN	20	11	25	11.00	[0.6]	Hollocher et al. 1982 [Tappe et al. 1999, 0.5-0.7 μm^3]	Bacteria grown at 30 C to the late exponential phase, stored for no more than 3 days at 0 C	1 mg WM \approx 0.1 mg protein
Nocardia corallina	141.Nocardia corallina	Actinobacteria: Actinomycetales	$\mu\text{l O}_2 (\text{mg DM})^{-1} \text{hr}^{-1}$	MIN	1	1.7	30	1.20	2.1	Robertson & Batt 1973 (rods, 0.8-1.2×2-4 μm)	Cells harvested at the end of growth phase; respiration measured after 45 hours of starvation; viability more than 90%	Soil bacterium Dry mass data (Fig. 1): 4.3×10^9 cells (90% viable) make up 1.6 mg dry mass: 0.4×10^{-12} g dry mass cell $^{-1}$

													Respiration rate fell from 10 to 1 $\mu\text{l O}_2$ (mg DM) $^{-1}$ hr $^{-1}$ during the first 45 hr of the total 450 hr' period of starvation	Cell mass estimated from linear dimensions given by Robertson & Batt (1973), 2-4 μm (length) \times 0.8-1.2 μm (diameter), is greater, 2.4×10^{-12} g cell $^{-1}$	
													Midwinter & Batt 1953 report 1-12 $\mu\text{l O}_2$ (mg DM) $^{-1}$ hr $^{-1}$ for cells grown on different substrates for 24-96 hr at 30 C; respiration measured for about 3 hr		
Nocardia (A-6)	corallina	142.Nocardia coral-lina		Actinobacteria: Actinomycetales	Acti-	$\mu\text{l O}_2$ (4.8 mg DM) $^{-1}$ min $^{-1}$		40	4	30.3	2.77	[2.1]	Raymond et al. 1967 [Robertson & Batt 1973, rods, 0.8-1.2 \times 2-4 μm]	Cells grown for 72 hr; respiration measured for 220 min.	
Nocardia (ATCC 3308)	asteroides	143.Nocardia far-cinica		Actinobacteria: Actinomycetales	Acti-	$\mu\text{l O}_2$ (mg DM) $^{-1}$ hr $^{-1}$	MIN	5	8.3	30	5.87	[0.1]	Jurtshuk & McQuitty 1976 [Takeo & Uesaka 1975, Fig. 1, hyphae diam 0.6 μm mycellium; Kowalski et al. 1999, 1-1.25 \times 3-5 μm airborne]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Nocardia sp. (Z1)		144.Nocardia sp.		Actinobacteria: Actinomycetales	Acti-	$\mu\text{l O}_2$ (42 mg DM) $^{-1}$ hr $^{-1}$	MIN	56	2.2	30	1.56		Watson & Cain 1975 [BM9, genus, hyphae diam 0.5-1.2 μm]	Cells grown for 24 hr	Bacterium isolated from soil with by enrichment with pyridine
Nocardia sp.	erythropolis	145.Nocardia sp.		Actinobacteria: Actinomycetales	Acti-	$\mu\text{l O}_2$ (7.5 mg DM) $^{-1}$ min $^{-1}$	MIN	40	3	30	2.12		Cartwright & Cain 1959 [BM9, genus, hyphae diam 0.5-1.2 μm]	Respiration measured for 200 min	No change of respiration with time
													Cain et al. 1968: Cells harvested after 24-36 hr incubation (Smith et al. 1968); respiration measured for 90 min was 185 $\mu\text{l O}_2$ (14.3 mg DM) $^{-1}$ hr $^{-1}$ = 22 W/kg at 30 C	The organism was originally isolated from soil by enrichment with <i>p</i> -nitrobenzoate (Cain et al. 1958)	
Nonomuraea (ATCC 39727)	sp.	146.Nonomuraea sp.		Actinobacteria: Actinomycetales	Acti-	nmol O ₂ (mg protein) $^{-1}$ min $^{-1}$	MIN	1	1	25	1.00		Palese et al. 2003	Respiration of cells in the lag phase (incubated for 15 hr) prior to exponential growth	Endogenous respiration increases from lag to exponential phase from 1 to 35 nmol O ₂ (mg protein) $^{-1}$ min $^{-1}$, then starts to drop abruptly in the stationary phase (min. value measured after 160 hr incubation was 7 nmol O ₂ (mg protein) $^{-1}$ min $^{-1}$)

Micrococcus denitrificans	147.Paracoccus denitrificans	Alphaproteobacteria: Rhodobacterales	$\mu\text{l O}_2 (\text{mg DM})^{-1} \text{hr}^{-1}$	MIN	5	8.3	30	5.87	[0.16]	Kornberg & Morris 1965 [BM, spheres (0.5-0.9 μm in diam) or short rods (0.9-1.2 μm long)] Bishop et al. 1962	Cells harvested during the exponential phase	Endogenous reserves of cells were "depleted" by 4 hr aerobic shaking at 30 C in another experiment
Pasteurella pseudotuberculosis (NCTC 1101)	148.Pasteurella pseudotuberculosis	Gammaproteobacteria: Enterobacteriales	$\mu\text{l O}_2 (\text{mg DM})^{-1} \text{hr}^{-1}$	MIN	7	12	25	12.00			Respiration measured immediately after harvesting the cells at the end of the logarithmic growth phase	Max. resp. (in the presence of glucose) was 331 $\mu\text{l O}_2 (\text{mg DM})^{-1} \text{hr}^{-1}$
Pediococcus cerevisiae (ATCC 8081)	149.Pediococcus acidilactici	"Bacilli": "Lactobacillales"	$\mu\text{l O}_2 (\text{mg DM})^{-1} \text{hr}^{-1}$	MIN	2	3.3	30	2.33	[2.2]	Jurtshuk & McQuitty 1976 [BM9, genus, spheres 1.0-2.0 μm diam]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Phaeospirillum fulvum (5K, KM MGU 325)	150.Phaeospirillum fulvum	Alphaproteobacteria: Rhodospirillales	$\text{nmol O}_2 (\text{mg protein})^{-1} \text{min}^{-1}$	MIN	25.3	28	28	22.74	[2]	Berg et al. 2002 [BM]	Bacteria harvested from early exponential cultures grown photoheterotrophically	Purple bacteria
Picrophilus oshimae	151.Picrophilus oshimae	Thermoplasmata (Archaea): Thermoplasmatales	$\text{nmol O}_2 (\text{mg protein})^{-1} \text{min}^{-1}$	MIN	22.7	25	60	2.21	[1]	Van de Vossenberg et al. 1998 [Schleper et al. 1995, cocci, diam 1-1.5 μm]	Starvation for several hours	Thermoacidophilic archaeon
Proteus morganii [=Morganella morganii]	152.Proteus morganii	Gammaproteobacteria: "Enterobacteriales"	$\mu\text{l O}_2 (\text{mg DM})^{-1} \text{hr}^{-1}$	MIN	3	5	30	3.54	[0.4]	Jurtshuk & McQuitty 1976 [BM, rods, 0.6-0.7 \times 1-1.7 μm]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Proteus vulgaris	153.Proteus vulgaris	Gammaproteobacteria: "Enterobacteriales"	$\mu\text{l O}_2 (\text{mg DM})^{-1} \text{hr}^{-1}$	MIN	3	5	37	2.18	[0.4]	Bishop et al. 1962 [BM9, genus, rods 0.4-0.8 \times 1-3 μm]	Respiration measured immediately after harvesting the cells (aerobic culture) at the end of the logarithmic growth phase; endogenous respiration of anaerobic culture was below detection limit (0 orig. units)	Max. resp. (in the presence of glucose) was 87 orig. units
Proteus vulgaris	154.Proteus vulgaris	Gammaproteobacteria: "Enterobacteriales"	$\mu\text{l O}_2 (\text{mg DM})^{-1} \text{hr}^{-1}$		3	5	30	3.54	[0.4]	Jurtshuk & McQuitty 1976 [BM9, genus, rods 0.4-0.8 \times 1-3 μm]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Pseudomonas aeruginosa	155.Pseudomonas aeruginosa	Gammaproteobacteria: Pseudomonadales	$\mu\text{l O}_2 (\text{mg N})^{-1} (75 \text{ min})^{-1}$	MIN	50	6.7	30	4.74	[0.5]	Rogoff 1962 [Montesinos et al. 1983, Coulter counter; see also datra of Hou et al. 1966, 0.2-0.8 pg]	Cells isolated from soil and closely resembling P. aeruginosa were grown for 18 to 48 hr; respiration was measured for 75 min	

Pseudomonas aeruginosa (120Na)	156.Pseudomonas aeruginosa	Gammaproteobacteria: Pseudomonadales	$\mu\text{mol O}_2$ (100 mg DM) ⁻¹ (2 hr) ⁻¹	55.5	10	30	7.07	[0.5]	Gronlund & Campbell 1961 [Montesinos et al. 1983, Coulter counter; see also data of Hou et al. 1966, 0.2-0.8 pg]	Cells (strain 120Na) harvested after 20 hr growth; respiration measured for 2 hr; for strain ATCC 9027 respiration was 80 orig. units Tomlinson & Campbell 1963: cells (strain 120Na) harvested after 20 hr growth at 30 C; respiration measured for 2 hr; 120 $\mu\text{l O}_2$ (5 mg DM) ⁻¹ (2 hr) ⁻¹ = 20 W/kg Gronlund & Campbell 1963: cells (strain ATCC 9027) harvested after 20 hr growth at 30 C; respiration measured for 3 hr; hourly rates decreased from 4.75 to 4.0 to 3.19 $\mu\text{mol O}_2$ (10 mg DM) ⁻¹ hr ⁻¹ , with no loss of viability (min. rate 12 W/kg) Gronlund & Campbell 1966: cells (strain ATCC 9027) harvested after 20 hr growth at 30 C; respiration measured for 2 hr; 95 $\mu\text{l O}_2$ (5 mg DM) ⁻¹ (2 hr) ⁻¹ = 16 W/kg Warren et al. 1960 report practically the same rate, 2500 $\mu\text{l O}_2$ (100 mg DM) ⁻¹ (2 hr) ⁻¹ = 20 W/kg; the same age of culture (20 hr), strain ATCC 9027	Hou et al. 1966: Cell size of P. aeruginosa: 4.5×10 ⁸ viable cells = 29.6 ×10 ⁻⁶ g = 17.5 × 10 ⁻⁶ g protein, cells harvested after 8 hr growth: 1 cell = 0.06 ×10 ⁻¹² g dry mass = 0.2 ×10 ⁻¹² g wet mass For 24-hr phosphorus starved cells, they observed 4.18×10 ⁸ viable cells = 44.4 ×10 ⁻⁶ g = 22.8 × 10 ⁻⁶ g protein: 1 cell = 0.35 × 10 ⁻¹² g wet mass For refeed cells at 30 hr: 7.68×10 ⁸ viable cells = 182 ×10 ⁻⁶ g = 75.1 × 10 ⁻⁶ g protein: 1 cell = 0.8 × 10 ⁻¹² g wet mass Protein to DM (Protein/DM) ratio = 0.59, 0.51 and 0.41, respectively.
Pseudomonas aeruginosa (ATCC 15442)	157.Pseudomonas aeruginosa	Gammaproteobacteria: Pseudomonadales	$\mu\text{l O}_2$ (mg DM) ⁻¹ hr ⁻¹	17	28	30	19.80	[0.5]	Jurtshuk & McQuitty 1976 [Montesinos et al.	Cells harvested at the late-logarithmic growth phase (two	

Pseudomonas (pyo- ceanea) aeruginosa	158.Pseudomonas aeruginosa	Gammaproteobacteria: Pseudomonadales	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$	28	47	37	20.46	[0.5]	1983, Coulter counter; see also datra of Hou et al. 1966, 0.2-0.8 pg] Bishop et al. 1962 [Montesinos et al. 1983, Coulter counter; see also datra of Hou et al. 1966, 0.2-0.8 pg]	thirds of the maximal growth concentration) Respiration measured immediately after harvesting the cells (aerobic culture) at the end of the logarithmic growth phase; anaerobically grown culture respired at $15 \mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1} = 25 \text{ W/kg}$	
Pseudomonas aeru- ginosa (9/93 and 72/92)	159.Pseudomonas aeruginosa	Gammaproteobacteria: Pseudomonadales	$\text{nmol O}_2 \text{ (mg DM)}^{-1} \text{ min}^{-1}$	23.7	53	30	37.48	[0.5]	Majtán et al. 1995 [Montesinos et al. 1983, Coulter counter; see also datra of Hou et al. 1966, 0.2-0.8 pg]	Cells harvested at the exponential phase; respiration measured for 10 min	
Pseudomonas fluo- rescens (A.3.12)	160.Pseudomonas fluorescens	Gammaproteobacteria: Pseudomonadales	$\mu\text{l O}_2 \text{ (3 mg DM)}^{-1} \text{ hr}^{-1}$	8	4.4	28	3.57	[1]	Altekar & Rao 1963 [BM, 0.7-0.8x2.0-3.0 μm ; Gunter & Kohn 1956, 0.24 pg DM/cell = 0.8 pg/cell at 70% water content]	Cells washed after incubation for 24 hr in a growth medium on a shaker; respiration of “resting” cells measured for 4 hr; value for the last hour taken (Fig. 4)	Respiration decreases with time (mean for the 4 hrs was $\approx 8 \text{ W/kg}$)
Pseudomonas fluo- rescens (A.3.12)	161.Pseudomonas fluorescens	Gammaproteobacteria: Pseudomonadales	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$	5.9	10	26	9.33	[1]	Gunter & Kohn 1956 [BM, 0.7-0.8x2.0-3.0 μm ; Gunter & Kohn 1956, 0.24 pg DM/cell = 0.8 pg/cell at 70% water content]	Respiration of washed cell suspensions harvested from 16 to 18-hr yeast agar plates	
Pseudomonas fluo- rescens (KBI)	162.Pseudomonas fluorescens	Gammaproteobacteria: Pseudomonadales	$\mu\text{mol (10 mg DM)}^{-1} \text{ hr}^{-1}$	4	15	30	10.61	[1]	Hughes 1966 [BM, 0.7-0.8x2.0-3.0 μm ; Gunter & Kohn 1956, 0.24 pg DM/cell = 0.8 pg/cell at 70% water content]	Cells grown for 22-24 hr at 25 C; respiration measured after equilibration for 30 min at 30 C	
Pseudomonas fluo- rescens (A.3.12)	163.Pseudomonas fluorescens	Gammaproteobacteria: Pseudomonadales	$\mu\text{mol O}_2 \text{ (100 mg DM)}^{-1} \text{ (2 hr)}^{-1}$	98	18	30	12.73	[1]	Gronlund & Campbell 1961 [BM, 0.7-0.8x2.0-3.0 μm ; Gunter & Kohn 1956, 0.24 pg DM/cell = 0.8	Cells harvested after 20 hr growth; respiration measured for 2 hr at 30 C Tomlinson & Camp-	

Pseudomonas fluorescens (A.3.12)	164.Pseudomonas fluorescens	Gammaproteobacteria: Pseudomonadales	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$	20	33	30	23.33	[1]	Jacoby 1964 [BM, 0.7-0.8x2.0-3.0 μm ; Gunter & Kohn 1956, 0.24 pg DM/cell = 0.8 pg/cell at 70% water content]	pg/cell at 70% water content]	bell 1963: 190 $\mu\text{l O}_2 \text{ (5 mg DM)}^{-1} \text{ (2 hr)}^{-1} = 32 \text{ W/kg}$ for 20-hr cultures at 30 C	
Pseudomonas fluorescens (KBI)	165.Pseudomonas fluorescens	Gammaproteobacteria: Pseudomonadales	$\mu\text{l O}_2 \text{ (2 mg DM)}^{-1} \text{ (160 min)}^{-1}$	110	34	30	24.04	[1]	Kornberg 1958 [BM, 0.7-0.8x2.0-3.0 μm ; Gunter & Kohn 1956, 0.24 pg DM/cell = 0.8 pg/cell at 70% water content]	Cells grown for 10-12 hr, harvested during logarithmic phase; respiration measured for 160 min		
Pseudomonas fluorescens (ATCC 13525)	166.Pseudomonas fluorescens	Gammaproteobacteria: Pseudomonadales	$\text{nmol O}_2 \text{ (mg DM)}^{-1} \text{ (min)}^{-1}$	45.7	60	30	42.43	[1]	Eisenberg et al. 1973 [BM, 0.7-0.8x2.0-3.0 μm ; Gunter & Kohn 1956, 0.24 pg DM/cell = 0.8 pg/cell at 70% water content]	Respiration of washed cells harvested at midlog or early stationary phase		
Pseudomonas formicans	167.Pseudomonas formicans	Gammaproteobacteria: Pseudomonadales	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ (40 min)}^{-1}$	MIN	4	10	30	7.07	[7]	Sabina & Pivnick 1956 [Crawford 1954, rods 1-2x4-5 μm]	Bacteria grown for 5 days in a mixture of soluble oils; incubated for 24 hr in a nutrient broth; aerated vigorously in nutrient broth for 18 hr; harvested and washed; washed suspension "shaken for 2 to 8 hr at room temperature in an attempt to reduce the endogenous respiration"; respiration measured for 210 min (Fig. 2); value taken for the last 40 min	Bacteria isolated from used emulsifiers of industrial oil, Illinois, USA Endogenous respiration decreases with time during 2-4 hr of respiration (Figs. 2, 4)
Pseudomonas oleovorans	168.Pseudomonas oleovorans	Gammaproteobacteria: Pseudomonadales	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$	MIN	2	3.3	30	2.33	[0.16]	Sabina & Pivnick 1956 [Lee & Chandler 1941, almost coccoid, 0.5x0.8 μm]	Bacteria grown for 5 days in a mixture of soluble oils; incubated for 24 hr in a nutrient broth; aerated vigorously in nutrient broth for 18 hr; harvested and washed; washed	Bacteria isolated from used emulsifiers of industrial oil, England Endogenous respiration decreases with time during 2-4 hr of respiration (Figs. 2, 4)

Pseudomonas oleovorans	169.Pseudomonas oleovorans	Gammaproteobacteria: Pseudomonadales	nmol O ₂ (mg DM) ⁻¹ min ⁻¹	16-19	40	30	28.28	[0.16]	Peterson 1970 [Lee & Chandler 1941, almost coccoid, 0.5×0.8 μm]	suspension "shaken for 2 to 8 hr at room temperature in an attempt to reduce the endogenous respiration"; respiration measured for 210 min (Fig. 3); value taken for the last hour	Cells harvested in late-log-phase	
Pseudomonas putida (O1OC)	170.Pseudomonas putida	Gammaproteobacteria: Pseudomonadales	nmol O ₂ (25 mg DM) ⁻¹ min ⁻¹	MIN	11	1	30	0.71	[1.7]	Chapman & Ribbons 1976 [BM, 0.7-1.1×2.0-4.0 μm]	Succinate-grown cells harvested after 16-20 hr incubation in the late-logarithmic phase; respiration measured for 2 min; respiration of strains O1OC, ORC and O1 varied from 11 to 29 nmol O ₂ (25 mg DM) ⁻¹ min ⁻¹ (1-2.6 W/kg) depending on growth substrate.	The organism was isolated from orcinol enrichments Respiration measured for 1 hr (strain ORC) was 43 μl (20 mg WM) ⁻¹ hr ⁻¹ (Fig. 3) = 12 W/kg
Pseudomonas putida (arvilla (PaM1))	171.Pseudomonas putida	Gammaproteobacteria: Pseudomonadales	μl O ₂ (20 mg WM) ⁻¹ hr ⁻¹		10	2.8	30	1.98	[1.7]	Kunz & Chapman 1981 [BM, 0.7-1.1×2.0-4.0 μm]	P. putida (arvilla) mt-2 cells grown for 24 hr on toluene or 48 hr on pseudocumene; respiration 10 μl O ₂ (20 mg WM) ⁻¹ hr ⁻¹ = 2.8 W/kg	
Pseudomonas putida (biotype B)	172.Pseudomonas putida	Gammaproteobacteria: Pseudomonadales	μl O ₂ (mg DM) ⁻¹ min ⁻¹		0.27/6.1	4.4	30	3.11	[1.7]	Sebek & Barker 1968 [BM, 0.7-1.1×2.0-4.0 μm]	Cells harvested towards the end of the exponential phase, shaken for 18 hr "to reduce endogenous respiration"	
Pseudomonas putida (TM)	173.Pseudomonas putida	Gammaproteobacteria: Pseudomonadales	μl O ₂ (10 mg WM) ⁻¹ min ⁻¹		0.25	8.3	30	5.87	[1.7]	Donnelly & Dagle 1980 [BM, 0.7-1.1×2.0-4.0 μm]	Cells harvested during exponential growth	Strain isolated from rotting oak leaves
Pseudomonas fluorescens-putida	174.Pseudomonas putida	Gammaproteobacteria: Pseudomonadales	μl O ₂ (mg DM) ⁻¹ hr ⁻¹		11.2	19	37	8.27	[1.7]	Chakrabarty & Roy 1964 [BM, 0.7-1.1×2.0-4.0 μm]	Cells grown to the stationary phase (72 hr)	Respiration decreases from mid-exponential to late-exponential to stationary phase (20.3 → 17.8 → 11.2 μl O ₂ (mg DM) ⁻¹ hr ⁻¹); in cells harvested at 20 hr (mid-exponential) and starved for 0-4 hr respiration decreases from 27 (0th) to 20 (2nd) to 16.5 (3rd) to 15.6

Strain	Accession	Species	Phylum	Class	Order	Family	Genus	Species	Strain	Media	Temperature	Time	Yield	Notes
Vibrio costicola (NRCC 37001)	197.Salmonella costicola	Vibrio costicola	Gammaproteobacteria: "Vibrionales"	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ min}^{-1}$	MIN	0.10	7	25	7.00	[0.4]	Kushner et al. 1983 [Huang et al. 2000, rods, 0.5x1.5-3.2 μm]	respiration measured in the dark	this species	
Salmonella typhimurium (LT2)	198.Salmonella typhimurium	Salmonella typhimurium	Gammaproteobacteria: "Enterobacteriales"	$\mu\text{l O}_2 \text{ (1.36 mg DM)}^{-1} \text{ (10 min)}^{-1}$	MIN	2	15	37	6.53	[0.66]	Hoffee & Englesberg 1962 [Montesinos et al. 1983, Coulter counter]	Cells harvested just before the stationary phase; respiration varied from 0.10 to 1.14 $\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ min}^{-1}$ (7-80 W/kg) depending on salt concentration in the medium		
Salmonella typhimurium (ATCC 6444)	199.Salmonella typhimurium	Salmonella typhimurium	Gammaproteobacteria: "Enterobacteriales"	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$		6	10	30	7.07	[0.66] [1.35]	Jurtshuk & McQuitty 1976 [Montesinos et al. 1983, Coulter counter] [Kubitschek 1969, Coulter counter]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)		
Serratia marcescens (D1)	200.Serratia marcescens	Serratia marcescens	Gammaproteobacteria: "Enterobacteriales"	$\mu\text{l O}_2 \text{ (10 mg DM)}^{-1} \text{ (3.5 hr)}^{-1}$	MIN	75	4	37	1.74	[0.4]	Blizzard & Peterson 1962 [BM, 0.5-0.8x0.9-2 μm]	Bacteria incubated at 37 C for 18-20 hr on a rotary shaker; respiration measured for 3.5 hr; it grows with time	Respiration grows (!) with time	
Chromobacter prodigiosum (NCTC 1377)	201.Serratia marcescens	Chromobacter prodigiosum	Gammaproteobacteria: "Enterobacteriales"	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$		11	18	37	7.83	[0.4]	Bishop et al. 1962 [BM, 0.5-0.8x0.9-2 μm]	Respiration measured immediately after harvesting the cells at the end of the logarithmic growth phase	Max. resp. (in the presence of glucose) was 200 $\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$	
Serratia marcescens	202.Serratia marcescens	Serratia marcescens	Gammaproteobacteria: "Enterobacteriales"	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$		8	13	30	9.19	[0.4]	Jurtshuk & McQuitty 1976 [BM, 0.5-0.8x0.9-2 μm]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)		
Serratia marcescens (8 UK)	203.Serratia marcescens	Serratia marcescens	Gammaproteobacteria: "Enterobacteriales"	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$		21	35	30	24.75	[0.4]	Davis & Bateman 1960 [BM, 0.5-0.8x0.9-2 μm]	Cells harvested after 16 hr growth		
Shigella flexneri 3 (1013)	204.Shigella flexneri	Shigella flexneri	Gammaproteobacteria: "Enterobacteriales"	$\mu\text{l O}_2 \text{ (0.4 mg N)}^{-1} \text{ hr}^{-1}$	MIN	25	10	37	4.35		Erlanson & Ruhl 1956	Bacteria grown for 18 hr at 37 C; washed; stored in a refrigerator (can be stored for 5 days with no loss of activity); cells not older than 4 days were used in the analysis; respiration measured for 2 h	Human dysentery agent	
Sphaerotilus natans (12)	205.Sphaerotilus natans	Sphaerotilus natans	Betaproteobacteria: Burkholderiales	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$	MIN	27	45	28	36.55	6.5	Stokes 1954 [1.2-1.8x3-5 μm ;	Cells grown for 16 hr at 28 C on a shaker;	Originally isolated from contaminated flowing water	

Sporosarcina ureae	206.Sporosarcina ureae	“Bacilli”: Bacillales	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$	MIN	7	12	30	8.49	[3.8]	Jurtshuk & McQuitty 1976 [BM9, genus, rods, 1-2×2-3 μm]	washed suspensions were aerated for 3-5 hr to reduce endogenous respiration, which is characterized as “rather high” perhaps “due to the large amount of fatty material stored in the cells”	Aeration for more than 5 hr “tended to destroy the oxidizing capacity of the cells”
Staphylococcus aureus	207.Staphylococcus aureus	“Bacilli”: Bacillales	$7 \mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$	MIN	9.3-15.7	16	37	7	[0.04-0.17] [[0.27]]	Ramsey 1962 [Watson et al. 1998, diam 0.41 μm for long-starved cells, 0.69 μm for exponential phase cells] [[Montesinos et al. 1983, Coulter counter]]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration) Cells grown for 12, 24, 48 and 72 hr on agar respired at 14.8, 12.0, 10.0 and 0 $\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$, respectively, at 37 C; depending on the growth medium, endogenous respiration ranged from 9.3 to 15.7 $\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1} = 16\text{-}26 \text{ W/kg}$	
Staphylococcus aureus (FDA 209P)	208.Staphylococcus aureus	“Bacilli”: Bacillales	$\mu\text{l O}_2 \text{ (14 mg DM)}^{-1} \text{ (4 hr)}^{-1}$		44	1.3	30	0.92	[0.04-0.17] [[0.27]]	Huber & Schuardt 1970 [Watson et al. 1998, diam 0.41 μm for long-starved cells, 0.69 μm for exponential phase cells] [[Montesinos et al. 1983, Coulter counter]]	Cells grown for 18 hr at 37 C with shake aeration; respiration measured for 4 hr.	
Staphylococcus aureus	209.Staphylococcus aureus	“Bacilli”: Bacillales	$\mu\text{l O}_2 \text{ (0.1-0.15 mg N)} \text{ hr}^{-1}$		2	2.2	37	0.96	[0.04-0.17] [[0.27]]	Yotis & Ekstedt 1959 [Watson et al. 1998, diam 0.41 μm for long-starved cells, 0.69 μm for exponential phase cells] [[Montesinos et al. 1983, Coulter counter]]	Cells incubated for 16 hr at 37 C; respiration measured for 6 hr; it decreases with time	Respiration decreases with starvation time from 21 $\mu\text{l O}_2 \text{ (0.1-0.15 mg N)} \text{ hr}^{-1}$ in the first hour, 11.5 (second hour), 6.5 (third, fourth hour), 2 (fifth, sixth hour), min. rate = 2.2 W/kg
Staphylococcus	210.Staphylococcus	“Bacilli”: Bacillales	$\mu\text{l O}_2 \text{ (6.02 mg)}$		9.73	2.7	37	1.18	[0.04-	Bluhm & Ordal	Cells grown for 10 hr	

aureus (MF-31)	aureus		$\text{DM}^{-1} \text{hr}^{-1}$							0.17 [[0.27]]	1969 [Watson et al. 1998, diam 0.41 μm for long-starved cells, 0.69 μm for exponential phase cells] [[Montesinos et al. 1983, Coulter counter]]	at 37 C; respiration of heat injured cells is three times lower	
Staphylococcus aureus (31-r)	211.Staphylococcus aureus	“Bacilli”: Bacillales	$\mu\text{mol O}_2 \text{ (mg DM)}^{-1} \text{hr}^{-1}$	0.19	7	37	3.05			[0.04-0.17] [[0.27]]	Krzemiński et al. 1972 [Watson et al. 1998, diam 0.41 μm for long-starved cells, 0.69 μm for exponential phase cells] [[Montesinos et al. 1983, Coulter counter]]	Cells harvested after 20 hr growth at 37 C and starved for 3 hr; respiration decreases from 0.68 to 0.43 to 0.19 $\mu\text{mol O}_2 \text{ (mg DM)}^{-1} \text{h}^{-1}$ for the 1st, 2nd and 3rd hrs at 37 C, respectively.	Respiration decreases with starvation time; it also depends on growth phase: maximum at approx. 10 hr and then decreases towards the stationary phase
Staphylococcus (albus) aureus	212.Staphylococcus aureus	“Bacilli”: Bacillales	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{hr}^{-1}$	3	5	30	3.54			[0.04-0.17] [[0.27]]	Jurtshuk & McQuitty 1976 [Watson et al. 1998, diam 0.41 μm for long-starved cells, 0.69 μm for exponential phase cells] [[Montesinos et al. 1983, Coulter counter]]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration); respiration was 3, 4 and 5 $\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{hr}^{-1}$ for strains S. (albus) aureus, S. aureus (University of Houston) and S. aureus ATCC 6538, respectively (5-8.3 W/kg), at 30 C	
Staphylococcus epidermidis (AT2)	213.Staphylococcus epidermidis	“Bacilli”: Bacillales	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{hr}^{-1}$	MIN	16	27	30	19.09		[0.5]	Jacobs & Conti 1965 [BM]	Cells grown for 8 hr at 37 C harvested at the end of the log phase	Cell mass estimated from linear dimensions given in BM
Staphylococcus albus (Micrococcus pyogenes var. albus)	214.Staphylococcus simulans	“Bacilli”: Bacillales	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{hr}^{-1}$	MIN	6	10	37	4.35			Bishop et al. 1962	Respiration measured immediately after harvesting the cells (aerobic culture) at the end of the logarithmic growth phase; endogenous respiration of anaerobic culture was 4 orig. units	Max. resp. (in the presence of lactate) was 111 orig. units
Gaffkya tetragena (ATCC 10875)	215.Staphylococcus sp.	“Bacilli”: Bacillales	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{hr}^{-1}$	MIN	10	17	30	12.02			Jurtshuk & McQuitty 1976	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration); strain ATCC 10875	

Streptococcus mastitidis (70b)	mas-	216.Streptococcus agalactiae	“Bacilli”:	“Lactobacillales”	$\mu\text{l O}_2$ (mg N) ⁻¹ (120 min) ⁻¹	MIN	10	1.7	37	0.74	[0.3]	Greisen & Gun-salus 1944 [BM]	respired at 14 $\mu\text{l O}_2$ (mg DM) ⁻¹ hr ⁻¹ = 23 W/kg Cells grown for 12 hr	N/DM \approx 0.1
Streptococcus agalactiae	aga-	217.Streptococcus agalactiae	“Bacilli”:	“Lactobacillales”	$\mu\text{l O}_2$ (mg DM) ⁻¹ hr ⁻¹		5	8.3	37	3.61	[0.3]	Mickelson 1961 [BM]	Cells harvested after 10-15 hr incubation; respiration measured for 5 hr	
Streptococcus pneumoniae (ATCC 6360)		218.Streptococcus pneumoniae	“Bacilli”:	“Lactobacillales”	$\mu\text{l O}_2$ (mg DM) ⁻¹ hr ⁻¹	MIN	1	1.7	30	1.20	[0.4]	Jurtshuk & McQuitty 1976 [Kowalski et al. 1999, diam 0.8-1 μm]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Streptococcus pyogenes (ATCC 10389)	py-	219.Streptococcus pyogenes	“Bacilli”:	“Lactobacillales”	$\mu\text{l O}_2$ (mg DM) ⁻¹ hr ⁻¹	MIN	2	3.3	30	2.33	[0.2]	Jurtshuk & McQuitty 1976 [Kowalski et al. 1999, diam 0.6-1 μm]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Streptomyces coelicolor	coeli-	220.Streptomyces coelicolor	Actinobacteria:	Actinomycetales	$\mu\text{l O}_2$ (mg DM) ⁻¹ hr ⁻¹	MIN	24	40	30	28.28		Niederpruem & Hackett 1961	Respiration of mycelium	
Streptomyces fradiae (ATCC 11903)	fradi-	221.Streptomyces fradiae	Actinobacteria:	Actinomycetales	$\mu\text{l O}_2$ (mg DM) ⁻¹ hr ⁻¹	MIN	13	22	30	15.56		Niederpruem & Hackett 1961	Respiration of mycelium	
Streptomyces griseus (3475 Waksman)	grise-	222.Streptomyces griseus	Actinobacteria:	Actinomycetales	$\mu\text{l O}_2$ (mg N) ⁻¹ hr ⁻¹	MIN	105	18	30	12.73		Gilmour et al. 1955	Cells grown for 24 hr	
Streptomyces griseus (ATCC 10137)	grise-	223.Streptomyces griseus	Actinobacteria:	Actinomycetales	$\mu\text{l O}_2$ (mg DM) ⁻¹ hr ⁻¹		12	20	30	14.14		Jurtshuk & McQuitty 1976	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Actinomyces longispororuber	longi-	224.Streptomyces longispororuber	Actinobacteria:	Actinomycetales	$\mu\text{l O}_2$ (mg DM) ⁻¹ hr ⁻¹	MIN	2	3.3	30	2.33		Feofilova et al. 1966	Respiration of 24-hr-old mycelium	Endogenous respiration was minimal (around 2 $\mu\text{l O}_2$ (mg DM) ⁻¹ hr ⁻¹) in the beginning of growth before the exponential phase (lag phase); at maximal growth rate it increased up to 14 $\mu\text{l O}_2$ (mg DM) ⁻¹ hr ⁻¹ (48 hr) and then started to decline gradually to 7-11 $\mu\text{l O}_2$ (mg DM) ⁻¹ hr ⁻¹ at 72 hr and 4-8 $\mu\text{l O}_2$ (mg DM) ⁻¹ hr ⁻¹ at 96 hr
Streptomyces olivaceus (NRRL B-1125)	oliva-	225.Streptomyces olivaceus	Actinobacteria:	Actinomycetales	$\mu\text{mol O}_2$ (mg N) ⁻¹ hr ⁻¹	MIN	54	9	37	3.92		Maitra & Roy 1961	Cells harvested at the end of growth at 24 hr; respiration measured for 60 min; pH 5.5; at pH 7.2 endogenous	

Sulfolobus acidocalcadareus	226.Sulfolobus acidocalcadareus	Archaea: Thermoprotei (Crenarchaeota): Sulfolobales	nmol O ₂ (mg protein) ⁻¹ min ⁻¹	MIN	13.5	15	60	1.33	[0.4]	Schäfer 1996 [Takayanagi et al. 1996, lobed cells, diam 0.8-1.0 µm]	respiration was 201.1 orig. units Steady-state respiration on endogenous substrate	
Contagious equine metritis bacterium (E-CMO)	227.Taylorella equigenitalis	Betaproteobacteria: Burkholderiales	pmol O ₂ (mg protein) ⁻¹ min ⁻¹	MIN	82	0.1	30	0.07	[0.4]	Lindmark et al. 1982 [BM9, genus, 0.7×0.7-1.8 µm]	Cells (English E-CMO strain) harvested from late log phase after 24 hr	Contagious equine metritis bacterium
Ferrobacillus ferrooxidans (Thiobacillus ferrooxidans)	228.Thiobacillus ferrooxidans	Betaproteobacteria: Hydrogenophilales	µmol O ₂ (5.6 mg protein) ⁻¹ hr ⁻¹	MIN	0.12	0.5	25	0.50	[0.25]	Silver 1970 [Kelly & Wood 2000, rods 0.4×2.0 µm]	respiration of cells during 60 min of substrate deprivation	
Thiobacillus intermedius	229.Thiobacillus intermedius	Betaproteobacteria: Hydrogenophilales	µmol O ₂ (mg protein) ⁻¹ hr ⁻¹	MIN	0.6	11	30	7.78	[0.4]	London & Rittenberg 1966 [BM9, genus, 0.5-1.0-4.0 µm]	Cells were grown in the presence of glucose to the stationary phase; respiration of cells grown without glucose was "nil"	Facultative autotroph oxidizing thiosulfate
Thiobacillus thiooxidans	230.Thiobacillus thiooxidans	Betaproteobacteria: Hydrogenophilales	µl O ₂ (mg N) ⁻¹ hr ⁻¹	MIN	4-10	1.5	28	1.22	[0.25]	Vogler 1942b [Kelly & Wood 2000, rods 0.4×2.0 µm]	young cultures (as cited by Newburgh 1954) Vogler 1942a: late cultures respire at 10-40 µl O ₂ (mg N) ⁻¹ hr ⁻¹	Autotroph growing on sulfur; in the presence of sulfur respiration increases by 20-100 times
Thiocapsa roseopersicina (M1)	231.Thiocapsa roseopersicina	Gammaproteobacteria: Chromatiales	nmol O ₂ (mg protein) ⁻¹ min ⁻¹	MIN	5.0	5.6	30	3.96	[1]	Overmann & Pfennig 1992 [Montesinos et al. 1983, Coulter counter]	Respiration of cells without microscopically visible sulfure globules at oxygen concentrations of 11-67 µM; respiration rates of phototrophically (anaerobically) and chemotrophically (microaerobically) grown cells do not differ; the species is capable of chemotrophic growth in the dark.	Purple sulfur bacteria Endogenous respiration of cells with visible sulfur globules is higher, up to 15 nmol O ₂ (mg protein) ⁻¹ min ⁻¹
Thiocystis violacea (2711)	232.Thiocystis violacea	Gammaproteobacteria: Chromatiales	nmol O ₂ (mg protein) ⁻¹ min ⁻¹	MIN	2.2	2.5	30	1.77	[11]	Overmann & Pfennig 1992 [BM9, genus, spherical or ovoid, 2.5-3.0 µm diam]	Respiration of cells without microscopically visible sulfure globules at oxygen concentrations of 11-67 µM; respiration rates of phototrophically (anaerobically) and chemotrophically (microaerobically) grown cells do not	Purple sulfur bacteria Endogenous respiration of cells with visible sulfur globules is higher, up to 30 orig. units

Thiorhodovibrio winogradskyi (SSP1)	233.Thiorhodovibrio winogradskyi	Gammaproteobacteria: Chromatiales	nmol O ₂ (mg protein) ⁻¹ min ⁻¹	MIN	5.9	6.6	30	4.67		Overmann & Pfennig 1992	differ; the species is capable of chemotrophic growth in the dark. Respiration of cells without microscopically visible sulfure globules at oxygen concentrations of 11-67 μM; respiration rates of phototrophically (anaerobically) and chemotrophically (microaerobically) grown cells do not differ; the species is capable of chemotrophic growth in the dark.	Purple sulfur bacteria isolated from the littoral sediment of meromictic Mahoney Lake (British Columbia, Canada) Maximum respiration in the presence of substrate (H ₂ S) is 264 orig. units
Pseudomonas butanovora (ATCC 43655)	234.Unidentified bacterium	Gammaproteobacteria: Pseudomonadales	nmol O ₂ (mg protein) ⁻¹ min ⁻¹	MIN	10-25	11	30	7.78	[0.6]	Vangnai et al. 2002 and personal communication with Dr. Luis A. Sayavedra-Soto (7 Nov 2006) [BM, species, rods, 0.6-0.8×1.1-2.4 μm]	Bacteria grown to the stationary phase (35-40 hr); kept at 25 C for at least 1 hr to lower endogenous respiration With a cell suspension kept at room temperature, the cells would show less endogenous respiration as time passes, down to 1-5 nmol O ₂ (mg protein) ⁻¹ min ⁻¹	Classification made for the Pseudomonas genus as described at www.bacterio.cict.fr. There is no such species at www.bacterio.cict.fr; strain ATCC 43655 is an "unidentified bacterium"
Unknown sp. (A-50)	235.Unknown	Unknown	μl O ₂ (16 mg DM) ⁻¹ min ⁻¹	MIN	32	6.7	30	4.74		Trudinger 1967	Cells grown for 16 hr at 28 C; respiration measured for 30 min	"Where necessary to reduce endogenous respiration, the bacteria were shaken at 30 C for 1 to 2 hr"
Unknown sp. (C-3)	236.Unknown	Unknown	μl O ₂ (3.1 mg DM) ⁻¹ min ⁻¹	MIN	28	30	30	21.21		Trudinger 1967	Cells grown for 16 hr at 28 C; respiration measured for 30 min	"Where necessary to reduce endogenous respiration, the bacteria were shaken at 30 C for 1 to 2 hr"
Achromobacter hartlebii (NCIB 8129)	237.Unnamed rhizobiaceae	Alphaproteobacteria: Rhizobiales	μl O ₂ (mg DM) ⁻¹ hr ⁻¹	MIN	25	42	25	42.00		Bishop et al. 1962	Respiration measured immediately after harvesting the cells at	The organism was isolated from percolation units containing garden soil and elemental sulfur. Max. resp. (in the presence of lactate) was 220 μl O ₂ (mg DM) ⁻¹ hr ⁻¹

Vibrio parahaemolyticus (biotype alginolyticus 156-70)	238.Vibrio alginolyticus	Gammaproteobacteria: "Vibrionales"	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$	MIN	2	3.3	30	2.33	[0.6]	Jurtshuk & McQuitty 1976[BM9, genus, rods 0.5-0.8×1.4-2.6 μm]	the end of the logarithmic growth phase Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Vibrio fischeri (MAC 401)	239.Vibrio fischeri	Gammaproteobacteria: "Vibrionales"	$\text{n atoms O (mg DM)}^{-1} \text{ min}^{-1}$	MIN	20	22	20	31.11	[0.6]	Droniuk et al. 1987 [Allen & Baumann 1971, Fig. 16, 0.7×1.5 μm]	Respiration at 100 mM Na^+	Marine bacterium
Vibrio (metschnikovii) cholerae biotype proteus (ATCC 7708)	240.Vibrio metschnikovii	Gammaproteobacteria: "Vibrionales"	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$	MIN	2	3.3	30	2.33	[0.6]	Jurtshuk & McQuitty 1976[BM9, genus, rods 0.5-0.8×1.4-2.6 μm]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration); respiration 1 orig. unit both studied strains, FC1011 and SAK3	
Vibrio parahaemolyticus (FC1011, SAK3)	241.Vibrio parahaemolyticus	Gammaproteobacteria: "Vibrionales"	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$	MIN	1	1.7	30	1.20	[0.6]	Jurtshuk & McQuitty 1976 [BM9, genus, rods 0.5-0.8×1.4-2.6 μm]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration); respiration 1 orig. unit both studied strains, FC1011 and SAK3	
Vibrio sp. (Ant-300)	242.Vibrio sp.	Gammaproteobacteria: "Vibrionales"	% cellular carbon hr^{-1}	MIN	0.0071	0.11	5	0.44	0.14	Novitsky & Morita 1977 [Novitsky & Morita 1976, Figs. 3b,c, cells starved for several weeks, cocci, diam 0.6-0.7 μm ; unstarved cells, rods, 1×2-4 $\mu\text{m} = 2 \mu\text{m}^3$]	Stable respiration of cells starved for several weeks; viability 50-100% judged by plate counts; during the first week of starvation, respiration is reduced by over 99%	Cell size estimated from linear dimensions of starved cells as shown in Fig. 3b of Novitsky & Morita 1976 During starvation, cells first increase in numbers at the expense of internal genetic material (nuclear bodies); as testified by Amy and Morita 1983 for 16 other marine bacterial isolates, this property of Ant-300 is not unique
Vitreoscilla stercoraria (ATCC 15218)	243.Vitreoscilla stercoraria	Betaproteobacteria: Neisseriales	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$	MIN	21	35	30	24.75		Jurtshuk & McQuitty 1976 [BM9, genus, filaments 1-3 μm diam]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Xanthomonas phaseoli (ATCC 9563)	244.Xanthomonas axonopodis	Gammaproteobacteria: Xanthomonadales	$\mu\text{l O}_2 \text{ (mg DM)}^{-1} \text{ hr}^{-1}$	MIN	22	37	30	26.16	[0.25]	Jurtshuk & McQuitty 1976 [BM9, genus, rods 0.4-0.7×0.7-1.8 μm]	Cells harvested at the late-logarithmic growth phase (two thirds of the maximal growth concentration)	
Pasteurella pestis (virulent Alexander strain)	245.Yersinia pestis	Gammaproteobacteria: Enterobacteriales	$\mu\text{l O}_2 \text{ (mg N)}^{-1} \text{ hr}^{-1}$	MIN	29	4.8	28	3.90	[0.55]	Wessman & Miller 1966 [Kowalski et al.	Cells (virulent Alexander strain) harvested after 24 to 30 hr	Respiration decreased by approx. 1.5-fold during the first 1-2 hr of starvation and then stabilized for 72 hr

1999, 0.5-1×1-2 growth; stable respira-
μm] tion after 1.5 hr of
starvation

Notes on additional data not included into Table **S1a**:

- 1) *Listeria monocytogenes* (Friedman & Alm 1962) resting cells from cultures grown for 16 hr had no measurable endogenous respiration. The lowest reported values were 17.7 and 8.1 $\mu\text{l O}_2 (\text{mg N})^{-1} \text{hr}^{-1}$ for growth in the presence of pyruvate. The same result was obtained by Welch et al. 1979.
- 2) Data needing verification (can be unrealistic): Gaudy et al. 1963, *E.coli* 500 mg DM/l consumed 10 mg O_2 in 7 hr (Fig. 3) = 0.003 W/kg at 25 C.
- 3) Goldshmidt & Wiss 1966: "Since the high endogenous respiration of 24-hr *Azotobacter* cultures can be reduced markedly by aerating in saline, washed vegetative cells were shaken for 4 hr before exposure to the EDTA-Tris system."
- 4) *Neisseria meningitidis* (Yu & deVoe 1980): washed whole cells were devoid of detectable endogenous respiration; Mallavia & Weiss 1970 obtained the same result (typical rates in the presence of substrates were about 5 $\mu\text{mol O}_2 (\text{mg protein}) \text{hr}^{-1} = 93 \text{ W/kg}$)
- 5) Data needing verification (can be unrealistic): Mårdén et al. 1985 studied the decline of endogenous respiration of marine bacteria during several days' starvation and observed values of the order of $0.5 \times 10^{-10} \text{ mg O}_2 (\mu\text{m}^3)^{-1} \text{hr}^{-1} \sim 200 \text{ W/kg}$. Since this value is in the upper range of respiration values in the presences of substrates (!) (Makarieva et al. 2005), there should be some error in the reported respiration units. Moreover, in the inlet to Fig. 2 showing respiration rate per biosurface the units are again similar (μm^{-3}) instead of (μm^{-2}) again indicating an inconsistency. Finally, Morton et al. 1994 characterize these rates as "low but detectable", which could hardly be plausible if they were indeed in the vicinity of several hundred W/kg. In the related work by Kjelleberg et al. 1982 it is stated that after five days of starvation a marine *Vibrio* respired at a rate of not less than 9 ng atoms $\text{O}_2 (10^9 \text{ viable cells})^{-1} \text{min}^{-1}$ and cell volume was 0.4 μm^3 . This gives a mass-specific rate of 90 W/kg. Even taking a conservative estimate of energy content of the living matter of $4 \times 10^6 \text{ J/kg}$, we conclude that the bacterium should have eaten itself about ten times in five days, had it possessed such a high respiration rate. The data of Mårdén et al. 1985 and Kjelleberg et al. 1982 were not included into the present analysis.
- 6) Acidophilic bacterium PW2 (Goulbourne et al. 1986) starved at pH 3 or 4 progressively lost both respiration and viability, with no stabilization. In the presence of Mg^+ ions half-life of cells was 72 hr and respiration dropped from 84 nmol $\text{O}_2 (\text{mg protein})^{-1} \text{min}^{-1} = 94 \text{ W/kg}$ to virtually zero.
- 7) Data needing verification (can be unrealistic): *Azospirillum brasiliense* Sp7 and *Azospirillum lipoferum* Sp59b respired endogeneously at 0.73 and 0.98 $\mu\text{mol O}_2 (\text{mg protein})^{-1} \text{min}^{-1}$, respectively (820 and 1100 W/kg) (Martinez-Drets et al. 1984)

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Table **S1b**. Numeric values used in the analyses presented in Figures 1-3 and Table 1 in the paper (after Table S1a). Log is decimal logarithm of the corresponding variable.

	Valid name	Class: Order	MIN	qWkg	LogqWkg	TC	q25Wkg	Logq25Wkg	Mpg	LogMpg
1.	<i>Acetobacter aceti</i>	Clostridia: Clostridiales	MIN	0.6	-0.222	30	0.42	-0.377	0.75	-0.125
2.	<i>Acholeplasma laidlawii</i>	Mollicutes: Acholeplasmatales	MIN	1.4	0.146	37	0.61	-0.215	0.04	-1.398
3.	<i>Achromobacter ruhlandii</i>	Betaproteobacteria: Burkholderiales	MIN	15	1.176	30	10.61	1.026	0.2	-0.699
4.	<i>Achromobacter</i> sp.	Betaproteobacteria: Burkholderiales	MIN	8.4	0.924	30	5.94	0.774	0.6	-0.222
5.	<i>Achromobacter viscosus</i>	Betaproteobacteria: Burkholderiales	MIN	35	1.544	30	24.75	1.394	0.6	-0.222
6.	<i>Achromobacter xerosis</i>	Betaproteobacteria: Burkholderiales	MIN	23	1.362	30	16.26	1.211	0.5	-0.301
7.	<i>Acidovorax facilis</i>	Betaproteobacteria: Burkholderiales	MIN	7	0.845	30	4.95	0.695	0.3	-0.523
8.	<i>Acinetobacter baumannii</i>	Gammaproteobacteria: Pseudomonadales	MIN	12	1.079	30	8.49	0.929	2	0.301
9.	<i>Acinetobacter calcoaceticus</i>	Gammaproteobacteria: Pseudomonadales	MIN	6.7	0.826	30	4.74	0.676	2	0.301
10.	<i>Acinetobacter johnsonii</i>	Gammaproteobacteria: Pseudomonadales	MIN	46	1.663	30	32.53	1.512	2	0.301
11.	<i>Acinetobacter</i> sp.	Gammaproteobacteria: Pseudomonadales	MIN	3.3	0.519	25	3.30	0.519	2	0.301
12.	<i>Acinetobacter</i> sp.	Gammaproteobacteria: Pseudomonadales	MIN	6	0.778	30	4.24	0.627	2	0.301
13.	<i>Acinetobacter</i> sp.	Gammaproteobacteria: Pseudomonadales	MIN	8	0.903	30	5.66	0.753	2	0.301
14.	<i>Aeromonas hydrophila</i>	Gammaproteobacteria: Aeromonadales	MIN	38	1.580	30	26.87	1.429		
15.	<i>Aeromonas veronii</i>	Gammaproteobacteria: Aeromonadales	MIN	20	1.301	30	14.14	1.150		
16.	<i>Agrobacterium tumefaciens</i>	Alphaproteobacteria: Rhizobiales	MIN	20	1.301	30	14.14	1.150	1.5	0.176
17.	<i>Alcaligenes eutrophus</i>	Betaproteobacteria: Burkholderiales	MIN	83	1.919	33	47.67	1.678	0.8	-0.097
18.	<i>Alcaligenes faecalis</i>	Betaproteobacteria: Burkholderiales	MIN	27	1.431	30	19.09	1.281		
19.	<i>Alcaligenes</i> sp.	Betaproteobacteria: Burkholderiales	MIN	2.1	0.322	30	1.48	0.170		
20.	<i>Aminobacter lissarensis</i>	Alphaproteobacteria: Rhizobiales	MIN	1.6	0.204	25	1.60	0.204	0.6	-0.222
21.	<i>Amoebobacter purpureus</i>	Gammaproteobacteria: Chromatiales	MIN	11	1.041	30	7.78	0.891	36	1.556
22.	<i>Amoebobacter roseus</i>	Gammaproteobacteria: Chromatiales	MIN	5.4	0.732	30	3.82	0.582	5	0.699
23.	<i>Amoebobaeter pendens</i>	Gammaproteobacteria: Chromatiales	MIN	8.5	0.929	30	6.01	0.779	5	0.699
24.	<i>Aquaspirillum itersonii</i>	Betaproteobacteria: Neisseriales	MIN	10	1.000	30	7.07	0.849	0.9	-0.046
25.	<i>Arthrobacter crystallopoietes</i>	Actinobacteria: Actinomycetales	MIN	0.2	-0.699	30	0.14	-0.854	1.7	0.230
26.	<i>Arthrobacter globiformis</i>	Actinobacteria: Actinomycetales	MIN	8.3	0.919	30	5.87	0.769	0.5	-0.301
27.	<i>Arthrobacter</i> sp.	Actinobacteria: Actinomycetales	MIN	1.2	0.079	30	0.85	-0.071	1.5	0.176
28.	<i>Arthrobacter</i> sp.	Actinobacteria: Actinomycetales	MIN	0.75	-0.125	25	0.75	-0.125	0.2	-0.699
29.	<i>Arthrobacter</i> sp.	Actinobacteria: Actinomycetales	MIN	6	0.778	30	4.24	0.627		
30.	<i>Arthrobacter</i> sp.	Actinobacteria: Actinomycetales	MIN	14	1.146	30	9.90	0.996		

31. <i>Azomonas agilis</i> ?	Gammaproteobacteria: Pseudomonadales	MIN	21	1.322	26	19.59	1.292	13	1.114
32. <i>Azorhizobium caulinodans</i>	Alphaproteobacteria: Rhizobiales	MIN	38	1.580	30	26.87	1.429	0.5	-0.301
33. <i>Azospirillum brasiliense</i>	Alphaproteobacteria: Rhodospirillales	MIN	27	1.431	37	11.75	1.070	1	0.000
34. <i>Azospirillum lipoferum</i>	Alphaproteobacteria: Rhodospirillales	MIN	39	1.591	37	16.98	1.230	4	0.602
35. <i>Azotobacter chroococcum</i>	Gammaproteobacteria: Pseudomonadales	MIN	25	1.398	30	17.68	1.247	14	1.146
36. <i>Azotobacter vinelandii</i>	Gammaproteobacteria: Pseudomonadales	MIN	1.5	0.176	30	1.06	0.025	0.5	-0.301
37. <i>Bacillus cereus</i>	"Bacilli": Bacillales	MIN	14	1.146	37	6.09	0.785	3.7	0.568
38. <i>Bacillus firmus</i>	"Bacilli": Bacillales	MIN	13	1.114	30	9.19	0.963	0.9	-0.046
39. <i>Bacillus megaterium</i>	"Bacilli": Bacillales	MIN	3.3	0.519	30	2.33	0.367	7	0.845
40. <i>Bacillus popilliae</i>	"Bacilli": Bacillales	MIN	0.8	-0.097	30	0.57	-0.244	0.8	-0.097
41. <i>Bacillus pumilus</i>	"Bacilli": Bacillales	MIN	5	0.699	30	3.54	0.549	0.7	-0.155
42. <i>Bacillus stearothermophilus</i>	"Bacilli": Bacillales	MIN	8.2	0.914	50	1.45	0.161	0.7	-0.155
43. <i>Bacillus subtilis</i>	"Bacilli": Bacillales	MIN	3.3	0.519	30	2.33	0.367	1.4	0.146
44. <i>Bdellovibrio bacteriovorus</i>	Deltaproteobacteria: Bdellovibrionales	MIN	25	1.398	30	17.68	1.247	0.3	-0.523
45. <i>Beneckea natriegens</i>	Gammaproteobacteria: "Vibrionales"	MIN	265	2.423	30	187.38	2.273	1.5	0.176
46. <i>Bradyrhizobium japonicum</i>	Alphaproteobacteria: Rhizobiales	MIN	1.0	0.000	29	0.76	-0.119	0.7	-0.155
47. <i>Branhamella catarrhalis</i>	Gammaproteobacteria: Pseudomonadales	MIN	1.7	0.230	37	0.74	-0.131	1.3	0.114
48. <i>Brucella melitensis</i>	Alphaproteobacteria: Rhizobiales	MIN	6.5	0.813	34	3.48	0.542	0.3	-0.523
49. <i>Burkholderia</i> sp.	Betaproteobacteria: Burkholderiales	MIN	21	1.322	30	14.85	1.172	0.7	-0.155
50. <i>Cellvibrio gilvus</i>	Gammaproteobacteria: Pseudomonadales	MIN	37	1.568	30	26.16	1.418	2	0.301
51. <i>Chromatium</i> sp.	Gammaproteobacteria: Chromatiales	MIN	11	1.041	25	11.00	1.041		
52. <i>Chromatium vinosum</i>	Gammaproteobacteria: Chromatiales	MIN	2.2	0.342	30	1.56	0.193	1.5	0.176
53. <i>Corynebacterium diphtheriae</i>	Actinobacteria: Actinomycetales	MIN	6.7	0.826	30	4.74	0.676		
54. <i>Corynebacterium</i> sp.	Actinobacteria: Actinomycetales	MIN	10	1.000	30	7.07	0.849		
55. <i>Delftia acidovorans</i>	Betaproteobacteria: Burkholderiales	MIN	13	1.114	30	9.19	0.963	0.8	-0.097
56. <i>Desulfovibrio salexigens</i>	Deltaproteobacteria: Desulfovibrionales	MIN	13	1.114	30	9.19	0.963	1.5	0.176
57. <i>Enterobacter aerogenes</i>	Gammaproteobacteria: "Enterobacteriales"	MIN	10	1.000	30	7.07	0.849	0.3	-0.523
58. <i>Enterobacter cloacae</i>	Gammaproteobacteria: "Enterobacteriales"	MIN	31	1.491	30	21.92	1.341	0.9	-0.046
59. <i>Enterococcus cecorum</i>	"Bacilli": "Lactobacillales"	MIN	3.8	0.580	30	2.69	0.430	0.4	-0.398
60. <i>Enterococcus faecalis</i>	"Bacilli": "Lactobacillales"	MIN	0.3	-0.523	30	0.21	-0.678	0.2	-0.699
61. <i>Enterococcus hirae</i>	"Bacilli": "Lactobacillales"	MIN	1.5	0.176	30	1.06	0.025		
62. <i>Enterococcus</i> sp.	"Bacilli": "Lactobacillales"	MIN	33	1.519	30	23.33	1.368	0.8	-0.097
63. <i>Escherichia coli</i>	Gammaproteobacteria: "Enterobacteriales"	MIN	32	1.505	37	14	1.146	0.7	-0.155
64. <i>Flavobacterium capsulatum</i>	Flavobacteria: Flavobacteriales	MIN	63	1.799	30	44.55	1.649	0.3	-0.523
65. <i>Francisella tularensis</i>	Gammaproteobacteria: Thiotrichales	MIN	3.7	0.568	37	1.61	0.207	0.01	-2.000
66. <i>Frankia</i> sp.	Actinobacteria: Actinomycetales	MIN	9	0.954	25	9.00	0.954		
67. <i>Haemophilus influenzae</i>	Gammaproteobacteria: Pasteurellales	MIN	0.5	-0.301	37	0.22	-0.658	0.14	-0.854
68. <i>Haemophilus parahaemolyticus</i>	Gammaproteobacteria: Pasteurellales	MIN	3.5	0.544	37	1.52	0.182		
69. <i>Haemophilus parainfluenzae</i>	Gammaproteobacteria: Pasteurellales	MIN	3.5	0.544	37	1.52	0.182	0.4	-0.398
70. <i>Halobacterium salinarum</i>	Archaea: Halobacteria: Halobacteriales	MIN	17	1.230	30	12.02	1.080	3.9	0.591
71. <i>Halomonas halodenitrificans</i>	Gammaproteobacteria: Oceanospirillales	MIN	67	1.826	25	67.00	1.826	0.4	-0.398
72. <i>Klebsiella pneumoniae</i>	Gammaproteobacteria: "Enterobacteriales"	MIN	3.3	0.519	30	2.33	0.367	0.4	-0.398
73. <i>Lactobacillus brevis</i>	"Bacilli": "Lactobacillales"	MIN	0.2	-0.699	30	0.14	-0.854	1.6	0.204
74. <i>Lactobacillus casei</i>	"Bacilli": "Lactobacillales"	MIN	1.7	0.230	30	1.20	0.079		
75. <i>Lactococcus lactis</i>	"Bacilli": "Lactobacillales"	MIN	0.6	-0.222	30	0.42	-0.377	0.2	-0.699
76. <i>Lactococcus</i> sp.	"Bacilli": "Lactobacillales"	MIN	2.3	0.362	30	1.63	0.212	0.8	-0.097

77. Legionella pneumophila	Gammaproteobacteria: Legionellales	MIN	20	1.301	37	8.71	0.940	0.3	-0.523
78. Methylobacterium extorquens	Alphaproteobacteria: Rhizobiales	MIN	6	0.778	30	4.24	0.627	1	0.000
79. Methylococcobium sp.	Gammaproteobacteria: Methylococcales	MIN	11	1.041	30	7.78	0.891	3	0.477
80. Methylophilus methylotrophus	Betaproteobacteria: Methylophiliales	MIN	1.6	0.204	40	0.57	-0.244	0.15	-0.824
81. Methylosinus trichosporium	Alphaproteobacteria: Rhizobiales	MIN	42	1.623	30	29.70	1.473	1	0.000
82. Micrococcus luteus	Actinobacteria: Actinomycetales	MIN	1.2	0.079	37	0.52	-0.284	1.1	0.041
83. Micrococcus sp.	Actinobacteria: Actinomycetales	MIN	10	1.000	30	7.07	0.849		
84. Micrococcus sp.	Actinobacteria: Actinomycetales	MIN	17	1.230	30	12.02	1.080		
85. Moraxella osloensis	Gammaproteobacteria: Pseudomonadales	MIN	13	1.114	30	9.19	0.963	2.3	0.362
86. Mycobacterium fortuitum	Actinobacteria: Actinomycetales	MIN	17	1.230	30	12.02	1.080		
87. Mycobacterium leprae	Actinobacteria: Actinomycetales	MIN	3.3	0.519	37	1.44	0.158		
88. Mycobacterium lepraemurium	Actinobacteria: Actinomycetales	MIN	3.3	0.519	37	1.44	0.158		
89. Mycobacterium phlei	Actinobacteria: Actinomycetales	MIN	14	1.146	37	6.09	0.785	0.4	-0.398
90. Mycobacterium smegmatis	Actinobacteria: Actinomycetales	MIN	24	1.380	37	10.45	1.019		
91. Mycobacterium tuberculosis	Actinobacteria: Actinomycetales	MIN	2.2	0.342	37	0.96	-0.018	0.2	-0.699
92. Myxococcus xanthus	Deltaproteobacteria: Myxococcales	MIN	4.6	0.663	30	3.25	0.512	1	0.000
93. Neisseria elongata	Betaproteobacteria: Neisseriales	MIN	8.3	0.919	30	5.87	0.769	0.2	-0.699
94. Neisseria flava	Betaproteobacteria: Neisseriales	MIN	12	1.079	30	8.49	0.929	0.2	-0.699
95. Neisseria gonorrhoeae	Betaproteobacteria: Neisseriales	MIN	0.6	-0.222	37	0.26	-0.585	0.2	-0.699
96. Neisseria mucosa	Betaproteobacteria: Neisseriales	MIN	15	1.176	30	10.61	1.026	0.2	-0.699
97. Neisseria sicca	Betaproteobacteria: Neisseriales	MIN	13	1.114	30	9.19	0.963	0.2	-0.699
98. Nitrobacter winogradskyi	Alphaproteobacteria: Rhizobiales	MIN	10	1.000	30	7.07	0.849	0.24	-0.620
99. Nitrosomonas europaea	Betaproteobacteria: Nitrosomonadales	MIN	11	1.041	25	11.00	1.041	0.6	-0.222
100. Nocardia corallina	Actinobacteria: Actinomycetales	MIN	1.7	0.230	30	1.20	0.079	2.1	0.322
101. Nocardia farcinica	Actinobacteria: Actinomycetales	MIN	8.3	0.919	30	5.87	0.769	0.1	-1.000
102. Nocardia sp.	Actinobacteria: Actinomycetales	MIN	2.2	0.342	30	1.56	0.193		
103. Nocardia sp.	Actinobacteria: Actinomycetales	MIN	3	0.477	30	2.12	0.326		
104. Nonomuraea sp.	Actinobacteria: Actinomycetales	MIN	1	0.000	25	1.00	0.000		
105. Paracoccus denitrificans	Alphaproteobacteria: Rhodobacterales	MIN	8.3	0.919	30	5.87	0.769	0.16	-0.796
106. Pasteurella pseudotuberculosis	Gammaproteobacteria: Enterobacteriales	MIN	12	1.079	25	12.00	1.079		
107. Pediococcus acidilactici	"Bacilli": "Lactobacillales"	MIN	3.3	0.519	30	2.33	0.367	2.2	0.342
108. Phaeospirillum fulvum	Alphaproteobacteria: Rhodospirillales	MIN	28	1.447	28	22.74	1.357	2	0.301
109. Picophilus oshimae	Thermoplasmata (Archaea): Thermoplasmatales	MIN	25	1.398	60	2.21	0.344	1	0.000
110. Proteus morgani	Gammaproteobacteria: "Enterobacteriales"	MIN	5	0.699	30	3.54	0.549	0.4	-0.398
111. Proteus vulgaris	Gammaproteobacteria: "Enterobacteriales"	MIN	5	0.699	37	2.18	0.338	0.4	-0.398
112. Pseudomonas aeruginosa	Gammaproteobacteria: Pseudomonadales	MIN	6.7	0.826	30	4.74	0.676	0.5	-0.301
113. Pseudomonas fluorescens	Gammaproteobacteria: Pseudomonadales	MIN	4.4	0.643	28	3.57	0.553	1	0.000
114. Pseudomonas formicans	Gammaproteobacteria: Pseudomonadales	MIN	10	1.000	30	7.07	0.849	7	0.845
115. Pseudomonas oleovorans	Gammaproteobacteria: Pseudomonadales	MIN	3.3	0.519	30	2.33	0.367	0.16	-0.796
116. Pseudomonas putida	Gammaproteobacteria: Pseudomonadales	MIN	1	0.000	30	0.71	-0.149	1.7	0.230
117. Pseudomonas saccharophila	Gammaproteobacteria: Pseudomonadales	MIN	33	1.519	30	23.33	1.368		
118. Pseudomonas sp.	Gammaproteobacteria: Pseudomonadales	MIN	2.2	0.342	30	1.56	0.193		
119. Pseudomonas sp.	Gammaproteobacteria: Pseudomonadales	MIN	8	0.903	30	5.66	0.753		
120. Pseudomonas sp.	Gammaproteobacteria: Pseudomonadales	MIN	3.3	0.519	25	3.30	0.519		
121. Pseudomonas sp.	Gammaproteobacteria: Pseudomonadales	MIN	8.3	0.919	30	5.87	0.769		
122. Pseudomonas sp.	Gammaproteobacteria: Pseudomonadales	MIN	15	1.176	30	10.61	1.026		

123. <i>Pseudomonas</i> sp.	Gammaproteobacteria: Pseudomonadales	MIN	18	1.255	30	12.73	1.105		
124. <i>Pseudomonas</i> sp.	Gammaproteobacteria: Pseudomonadales	MIN	35	1.544	20	49.50	1.695		
125. <i>Pseudomonas</i> sp.	Gammaproteobacteria: Pseudomonadales	MIN	17	1.230	30	12.02	1.080		
126. <i>Pseudonocardia nitrificans</i>	Actinobacteria: Actinomycetes	MIN	2.5	0.398	30	1.77	0.248		
127. <i>Rhizobium japonicum</i>	Alphaproteobacteria: Rhizobiales	MIN	16	1.204	23	18.38	1.264	0.7	-0.155
128. <i>Rhizobium leguminosarum</i>	Alphaproteobacteria: Rhizobiales	MIN	9.5	0.978	30	6.72	0.827	0.6	-0.222
129. <i>Rhizobium meliloti</i>	Alphaproteobacteria: Rhizobiales	MIN	12	1.079	30	8.49	0.929	0.7	-0.155
130. <i>Rhodobacter sphaeroides</i>	Alphaproteobacteria: Rhodobacterales	MIN	17	1.230	28	13.81	1.140	0.8	-0.097
131. <i>Rhodococcus rhodochrous</i>	Actinobacteria: Actinomycetales	MIN	17	1.230	30	12.02	1.080		
132. <i>Rhodococcus</i> sp.	Actinobacteria: Actinomycetales	MIN	2.2	0.342	25	2.20	0.342		
133. <i>Rhodococcus</i> sp.	Actinobacteria: Actinomycetales	MIN	7	0.845	30	4.95	0.695		
134. <i>Rhodococcus</i> sp.	Actinobacteria: Actinomycetales	MIN	7.4	0.869	25	7.40	0.869		
135. <i>Rhodospirillum rubrum</i>	Alphaproteobacteria: Rhodospirillales	MIN	3.8	0.580	28	3.09	0.490	9	0.954
136. <i>Sallinivibrio costicola</i>	Gammaproteobacteria: "Vibrionales"	MIN	7	0.845	25	7.00	0.845	0.4	-0.398
137. <i>Salmonella typhimurium</i>	Gammaproteobacteria: "Enterobacteriales"	MIN	15	1.176	37	6.53	0.815	0.66	-0.180
138. <i>Serratia marcescens</i>	Gammaproteobacteria: "Enterobacteriales"	MIN	4	0.602	37	1.74	0.241	0.4	-0.398
139. <i>Shigella flexneri</i>	Gammaproteobacteria: "Enterobacteriales"	MIN	10	1.000	37	4.35	0.638		
140. <i>Sphaerotilus natans</i>	Betaproteobacteria: Burkholderiales	MIN	45	1.653	28	36.55	1.563	6.5	0.813
141. <i>Sporosarcina ureae</i>	"Bacilli": Bacillales	MIN	12	1.079	30	8.49	0.929	3.8	0.580
142. <i>Staphylococcus aureus</i>	"Bacilli": Bacillales	MIN	16	1.204	37	7	0.845	0.27	-0.569
143. <i>Staphylococcus epidermidis</i>	"Bacilli": Bacillales	MIN	27	1.431	30	19.09	1.281	0.5	-0.301
144. <i>Staphylococcus simulans</i>	"Bacilli": Bacillales	MIN	10	1.000	37	4.35	0.638		
145. <i>Staphylococcus</i> sp.	"Bacilli": Bacillales	MIN	17	1.230	30	12.02	1.080		
146. <i>Streptococcus agalactiae</i>	"Bacilli": "Lactobacillales"	MIN	1.7	0.230	37	0.74	-0.131	0.3	-0.523
147. <i>Streptococcus pneumoniae</i>	"Bacilli": "Lactobacillales"	MIN	1.7	0.230	30	1.20	0.079	0.4	-0.398
148. <i>Streptococcus pyogenes</i>	"Bacilli": "Lactobacillales"	MIN	3.3	0.519	30	2.33	0.367	0.2	-0.699
149. <i>Streptomyces coelicolor</i>	Actinobacteria: Actinomycetales	MIN	40	1.602	30	28.28	1.451		
150. <i>Streptomyces fradiae</i>	Actinobacteria: Actinomycetales	MIN	22	1.342	30	15.56	1.192		
151. <i>Streptomyces griseus</i>	Actinobacteria: Actinomycetales	MIN	18	1.255	30	12.73	1.105		
152. <i>Streptomyces longispororuber</i>	Actinobacteria: Actinomycetales	MIN	3.3	0.519	30	2.33	0.367		
153. <i>Streptomyces olivaceus</i>	Actinobacteria: Actinomycetales	MIN	9	0.954	37	3.92	0.593		
154. <i>Sulfolobus acidocaldareus</i>	Archaea: Thermoprotei (Crenarchaeota): Sulfolobales	MIN	15	1.176	60	1.33	0.124	0.4	-0.398
155. <i>Taylorella equigenitalis</i>	Betaproteobacteria: Burkholderiales	MIN	0.1	-1.000	30	0.07	-1.155	0.4	-0.398
156. <i>Thiobacillus ferrooxidans</i>	Betaproteobacteria: Hydrogenophilales	MIN	0.5	-0.301	25	0.50	-0.301	0.25	-0.602
157. <i>Thiobacillus intermedius</i>	Betaproteobacteria: Hydrogenophilales	MIN	11	1.041	30	7.78	0.891	0.4	-0.398
158. <i>Thiobacillus thiooxidans</i>	Betaproteobacteria: Hydrogenophilales	MIN	1.5	0.176	28	1.22	0.086	0.25	-0.602
159. <i>Thiocapsa roseopersicina</i>	Gammaproteobacteria: Chromatiales	MIN	5.6	0.748	30	3.96	0.598	1	0.000
160. <i>Thiocystis violacea</i>	Gammaproteobacteria: Chromatiales	MIN	2.5	0.398	30	1.77	0.248	11	1.041
161. <i>Thiorhodovibrio winogradskyi</i>	Gammaproteobacteria: Chromatiales	MIN	6.6	0.820	30	4.67	0.669		
162. Unidentified bacterium	Gammaproteobacteria: Pseudomonadales	MIN	11	1.041	30	7.78	0.891	0.6	-0.222
163. Unknown	Unknown	MIN	6.7	0.826	30	4.74	0.676		
164. Unknown	Unknown	MIN	30	1.477	30	21.21	1.327		
165. Unnamed rhizobiaceae	Alphaproteobacteria: Rhizobiales	MIN	42	1.623	25	42.00	1.623		
166. <i>Vibrio alginolyticus</i>	Gammaproteobacteria: "Vibrionales"	MIN	3.3	0.519	30	2.33	0.367	0.6	-0.222
167. <i>Vibrio fischeri</i>	Gammaproteobacteria: "Vibrionales"	MIN	22	1.342	20	31.11	1.493	0.6	-0.222
168. <i>Vibrio metschnikovii</i>	Gammaproteobacteria: "Vibrionales"	MIN	3.3	0.519	30	2.33	0.367	0.6	-0.222

169. <i>Vibrio parahaemolyticus</i>	Gammaproteobacteria: "Vibrionales"	MIN	1.7	0.230	30	1.20	0.079	0.6	-0.222
170. <i>Vibrio</i> sp.	Gammaproteobacteria: "Vibrionales"	MIN	0.11	-0.959	5	0.44	-0.357	0.14	-0.854
171. <i>Vitreoscilla stercoraria</i>	Betaproteobacteria: Neisseriales	MIN	35	1.544	30	24.75	1.394		
172. <i>Xanthomonas axonopodis</i>	Gammaproteobacteria: Xanthomonadales	MIN	37	1.568	30	26.16	1.418	0.25	-0.602
173. <i>Yersinia pestis</i>	Gammaproteobacteria: Enterobacteriales	MIN	4.8	0.681	28	3.90	0.591	0.55	-0.260
