

Peak	Antibody name	Immunogen/ Antigen type	Source/Strain	Phylum	Group/Cluster	Positive in this study	Supp. References
1	A139	Sonicated cells	<i>Leptospirillum ferrooxidans</i>	Nitrospirae	Fe-S oxidizers cultures		[1]
2	IVE3C_182	Intact cells	<i>Acidithiobacillus ferrooxidans</i>	Gammaproteobacteria	Fe-S oxidizers cultures	+	[1]
3	A183	Sonicated cells	<i>A. ferroxidans</i>	Gammaproteobacteria	Fe-S oxidizers cultures	+	[1]
4	A184	Intact cells	<i>A. ferroxidans</i>	Gammaproteobacteria	Fe-S oxidizers cultures	+	[1]
5	A186	Intact cells	<i>L. ferroxidans</i>	Nitrospirae	Fe-S oxidizers cultures	+	[1]
6	IVE1BF	Biofilm	<i>L. pherrifilum</i> (LPH2) fermentor	Nitrospirae	Fe-S oxidizers cultures	+	[2]
7	IVE1S100	Soluble cellular fraction (SB-S100)	<i>L. pherrifilum</i> (LPH2) fermentor	Nitrospirae	Fe-S oxidizers cultures	+	[2]
8	IVE1C2	Pellet insoluble cellular fraction	<i>L. pherrifilum</i> (LPH2) fermentor	Nitrospirae	Fe-S oxidizers cultures	+	[2]
9	IVE2S1	Supernatant	NtD ( <i>Leptospirillum ferrifilum</i> )	Nitrospirae	Fe-S oxidizers cultures	+	[2]
10	IVE3C1	Celulas Intactas	<i>Acidithiobacillus caldus</i>	Gammaproteobacteria	Fe-S oxidizers cultures	+	[2]
11	IVE3S100	Soluble cellular fraction (SB-100)	<i>A. ferroxidans</i>	Gammaproteobacteria	Fe-S oxidizers cultures	+	[2]
12	IVE4C1	Intact cells	<i>Acidithiobacillus thiooxidans</i>	Gammaproteobacteria	Fe-S oxidizers cultures	+	[2]
13	IVE4C2	Pellet S100	<i>A. thiooxidans</i>	Gammaproteobacteria	Fe-S oxidizers cultures	+	[2]
14	IVE4S100	Soluble cellular fraction (SB-100)	<i>A. thiooxidans</i>	Gammaproteobacteria	Fe-S oxidizers cultures	+	[2]
15	IVE5C1	Intact cells	<i>Acidithiobacillus albertensis</i>	Gammaproteobacteria	Fe-S oxidizers cultures	+	[2]
16	IVE6C1	Intact cells	<i>A. caldus</i>	Gammaproteobacteria	Fe-S oxidizers cultures	+	[2]
17	IVE6C2	Pellet S100	<i>A. caldus</i>	Gammaproteobacteria	Fe-S oxidizers cultures	+	[2]
18	IVE6S100	Soluble cellular fraction (SB-100)	<i>A. caldus</i>	Gammaproteobacteria	Fe-S oxidizers cultures		[2]
19	IVE7C1	Intact cells	<i>Halothiobacillus neapolitanus</i>	Gammaproteobacteria	Fe-S oxidizers cultures	+	[3]
20	IVE8C1	Intact cells	<i>Acidimicrobium ferrooxidans</i>	Actinobacteria	Fe-S oxidizers cultures	+	[3]
21	IVE9C1	Intact cells	<i>L. ferroxidans</i>	Nitrospirae	Fe-S oxidizers cultures	+	[4]
22	IA2C1	Intact cells	Green filaments from Río Tinto water	Cellular extract from environmental samples	Metal-acidic environ	+	[2]
23	IA3C1	Intact cells	Dark filaments from Río Tinto (Arroyo 3.1)	Cellular extract from environmental samples	Metal-acidic environ	+	[2]
24	IC10S1_141	First wash	Ground sediment	Extracellular substances from environmental samples	Metal-acidic environ	+	

25	IC1C1	Nycodenz	Beach lake 3,2	Cellular extract from environmental samples	Metal-acidic environ	+	[2]
26	IC3C1	Nycodenz	Source of Rio Tinto. Filaments	Cellular extract from environmental samples	Metal-acidic environ	+	[2]
27	IC3C3	Sonicated cells EDTA (Nycodenz)	Source of Rio Tinto. Filaments	Cellular extract from environmental samples	Metal-acidic environ	+	[2]
28	IC4C1	Nycodenz	Filaments from water dam 3,2	Cellular extract from environmental samples	Metal-acidic environ	+	[2]
29	IC4S2	Supernatant from EDTA wash	Filaments from water dam 3,3	Extracellular substances from environmental samples	Metal-acidic environ	+	[2]
30	IC6C1	Nycodenz	Red wet sediment 2cm deep (Playa 3,1 )	Cellular extract from environmental samples	Metal-acidic environ	+	[2]
31	IC7C1	Nycodenz	Dry wall 3,2 seco	Cellular extract from environmental samples	Metal-acidic environ	+	[2]
32	ID17C1	Nycodenz	Fe-S precipitate from del 3,2. Río Tinto.	Cellular extract from environmental samples	Metal-acidic environ	+	[2]
33	ID18S2	Supernatant from EDTA	Peña de Hierro (93m deep) MARTE project	Extracellular substances from environmental samples	Metal-acidic environ	+	[2]
34	ID4S2	Supernatant from EDTA wash	Baton 61a, MARTE project	Extracellular substances from environmental samples	Metal-acidic environ	+	[2]
35	IIC1C1	Whole extract	Iceland sediment	Cellular extract from environmental samples	Geothermal environment	+	
36	IVF11C1	Intact cells	<i>Micrococcus</i> , strain Eur19.1	Actinobacteria	Psychrophilic cultures	+	
37	IVF13C1	Intact cells	<i>Fronidhabitans</i> , strain AH2.2.5	Actinobacteria	Psychrophilic cultures		
38	IVF18C1	Intact cells	<i>Desulfotalea psychrophila</i>	Deltaproteobacteria	Psychrophilic cultures	+	[5]
39	IVF19C1	Intact cells	<i>Polaromonas</i> , strain Eur3.PT.21	Betaproteobacteria	Psychrophilic cultures	+	
40	IVF20C1	Intact cells	<i>Sphingobacteriu</i> , strain Eur3.AL.31	Bacteroidetes	Psychrophilic cultures		
41	IVF21C1	Intact cells	<i>Planomicrobium</i> , strain CY-C1-12	Firmicutes	Psychrophilic cultures	+	
42	IVF22C1	Intact cells	<i>Tumebacillus</i> , strain Eur1 9.5	Firmicutes	Psychrophilic cultures	+	
43	IVF23C1	Intact cells	<i>Paenibacillus</i> , strain Eur1 9.26	Firmicutes	Psychrophilic cultures	+	
44	IVF2C1	Intact cells	<i>Shewanella gelidimarina</i>	Gammaproteobacteria	Psychrophilic cultures	+	[2]
45	IVF2S100	Soluble cellular fraction (SB-100)	<i>S. gelidimarina</i>	Gammaproteobacteria	Psychrophilic cultures	+	[2]
46	IVF2S2	Supernatant from EDTA wash	<i>S. gelidimarina</i>	Gammaproteobacteria	Psychrophilic cultures	+	[2]
47	IVF31C1	Intact cells	<i>Planococcus or2</i>	Firmicutes	Psychrophilic cultures		[2]

48	IVF4C1	Intact cells	<i>Psychroserpens burtonensis</i>	Bacteroidetes	Psychrophilic cultures	+	[2]
49	IVF4S1	Supernatant	<i>P. burtonensis</i>	Bacteroidetes	Psychrophilic cultures	+	[2]
50	IVF4S2	SB del lavado de EDTA	<i>P. burtonensis</i>	Bacteroidetes	Psychrophilic cultures	+	[2]
51	IVF5C1	Intact cells	<i>Psychrobacter frigidicola</i>	Gammaproteobacteria	Psychrophilic cultures	+	[2]
52	IVF6C1	Intact cells	<i>Cryobacterium psychrophilum</i>	Actinobacteria	Psychrophilic cultures	+	[2]
53	IVF7C1	Intact cells	<i>Colwellia psychrerythraea</i>	Gammaproteobacteria	Psychrophilic cultures	+	[3]
54	IVF8C1	Intact cells	<i>Psychrobacter cryohalolentis</i>	Gammaproteobacteria	Psychrophilic cultures	+	
55	IVG1C1	Intact cells	<i>Acidocella aminolytica</i>	Alphaproteobacteria	Iron reducers	+	[3]
56	IVG2C1	Intact cells	<i>Acidiphillum</i> sp.	Alphaproteobacteria	Iron reducers	+	[3]
57	IVG3C1	Intact cells	<i>Acidobacterium capsulatum</i>	Acidobacteria	Iron reducers	+	[3]
58	IVG4C1	Intact cells	<i>Thermus scotoductus</i>	Deinococcus-Thermus	Iron reducers	+	[3]
59	IVG4C2	Pellet S100	<i>T. scotoductus</i>	Deinococcus-Thermus	Iron reducers	+	[3]
60	IVG5C1	Intact cells	<i>Sulfobacillus acidophilus</i>	Firmicutes	Iron reducers	+	[3]
61	IVH11C1	Intact cells	<i>Bacillus subtilis</i>	Firmicutes	Iron reducers		[4]
62	IVH1C1	Intact cells	<i>B. subtilis</i> (sonicated spores)	Firmicutes	Spores	+	[4]
63	IVH22C1	Intact cells	<i>Streptomyces</i> spores	Actinobacteria	Spores		[4]
64	IVH2C1	Intact cells	<i>Streptomyces</i> spores	Actinobacteria	Spores	+	
65	IVI10C1	Intact cells	<i>Desulfovibrio vulgaris</i> subsp. <i>Vulgaris</i>	Deltaproteobacteria	Mesophilic cultures	+	[2]
66	IVI11C1	Intact cells	<i>Geobacter sulfurreducens</i>	Deltaproteobacteria	Mesophilic cultures		[2]
67	IVI12C1	Intact cells	<i>Geobacter metallireducens</i>	Deltaproteobacteria	Mesophilic cultures	+	[2]
68	IVI13C1	Intact cells	<i>Thermotoga maritima</i>	Thermotogaceae	Mesophilic cultures	+	[2]
69	IVI14C1	Intact cells	<i>Verrucomicrobium spinosum</i>	Verrucomicrobia	Mesophilic cultures	+	[2]
70	IVI15C1	Intact cells	<i>Methylomicrobium capsulatum</i>	Gammaproteobacteria	Mesophilic cultures	+	[2]
71	IVI16C1	Intact cells	<i>Planctomyces limnophilus</i>	Planctomycetes	Mesophilic cultures	+	[2]
72	IVI17C1	Intact cells	<i>Hydrogenobacter thermophilus</i>	Aquificae	Mesophilic cultures	+	[2]
73	IVI19C1	Intact cells	<i>Desulfosporosinus meridiei</i>	Firmicutes	Mesophilic cultures	+	[3]
74	IVI1C1	Intact cells	<i>Pseudomonas putida</i>	Gammaproteobacteria	Mesophilic cultures	+	[2]
75	IVI20C1	Intact cells	<i>Salinibacter ruber</i> M8	Bacteroidetes	Mesophilic cultures	+	[3]

76	IVI21C1	Intact cells	<i>S. ruber</i> PR1	Bacteroidetes	Mesophilic cultures	+	[3]
77	IVI21C2	Pellet S100	<i>S. ruber</i> PR2	Bacteroidetes	Mesophilic cultures	+	[3]
78	IVI22C1	Intact cells	Isolated from Atacama	..	Mesophilic cultures		[4]
79	IVI23C1	Intact cells	Lisated mycelium from <i>Streptomyces</i>	Actinobacteria	Mesophilic cultures	+	[4]
80	IVI24C1	Intact cells	<i>Thessaracoccus lapidicapta</i>	Actinobacteria	Mesophilic cultures	+	[4]
81	IVI25C1	Intact cells	<i>Clostridium</i>	Firmicutes	Mesophilic cultures		[4]
82	IVI2C1	Intact cells	<i>Bacillus</i> spp (Rio Tinto source)	Firmicutes	Mesophilic cultures		[2]
83	IVI3C1	Intact cells	<i>Shewanella oneidensis</i>	Gammaproteobacteria	Mesophilic cultures	+	[2]
84	IVI3C2	pellet S100	<i>S. oneidensis</i>	Gammaproteobacteria	Mesophilic cultures	+	[2]
85	IVI4C1	Intact cells	<i>Burkholderia furngorum</i>	Betaproteobacteria	Mesophilic cultures	+	[2]
86	IVI5C1	Intact cells	<i>S. oneidensis</i>	Gammaproteobacteria	Mesophilic cultures	+	[2]
87	IVI5S100	Soluble cellular fraction (SB-100)	<i>S. oneidensis</i>	Gammaproteobacteria	Mesophilic cultures	+	[2]
88	IVI6C3	Sonicated cells EDTA	<i>Azotobacter vinelandii</i>	Gammaproteobacteria	Mesophilic cultures	+	[2]
89	IVI7C1	Intact cells	<i>Bacillus subtilis</i> 3610	Firmicutes	Mesophilic cultures	+	[2]
90	IVI8C1	Intact cells	<i>B. subtilis</i> 3610	Firmicutes	Mesophilic cultures	+	[2]
91	IVI9C1	Intact cells	<i>Deinococcus radiodurans</i>	Deinococcus-Thermus	Mesophilic cultures	+	[2]
92	IVJ1C1	Intact cells	<i>Haloferax mediterranei</i>	Euryarchaeota	Archaea	+	[2]
93	IVJ2C1	Intact cells	<i>Methanococoides burtonii</i>	Euryarchaeota	Archaea		[2]
94	IVJ3C1	Intact cells	<i>Thermoplasma acidophilum</i>	Euryarchaeota	Archaea	+	[2]
95	IVJ4C1	Intact cells	<i>Methanobacterium formicicum</i>	Euryarchaeota	Archaea	+	[2]
96	IVJ5C1	Intact cells	<i>Methanosarcina mazeii</i>	Euryarchaeota	Archaea	+	[2]
97	IVJ6C1	Intact cells	<i>Pyrococcus furiosus</i>	Euryarchaeota	Archaea	+	[3]
98	IVJ8C1	Intact cells	<i>Halorubrum</i> sp.	Euryarchaeota	Archaea	+	[3]
99	IVJ9C1	Intact cells	<i>Halobacterium</i> sp.	Euryarchaeota	Archaea	+	[3]
100	IVK10C1	Intact cells. Cellular extract.	<i>Leptolyngbya boryana</i>	Cyanobacteria	Cyanobacteria		[6]
101	IVK11C1	Intact cells. Cellular extract.	<i>Tolypothrix distorta</i>	Cyanobacteria	Cyanobacteria		[6]
102	IVK12C1	Intact cells. Cellular extract.	<i>Aphanizomenon aphanizomenoides</i>	Cyanobacteria	Cyanobacteria	+	[6]
103	IVK13C1	Intact cells. Cellular extract.	<i>Nostoc</i> Antártida 16	Cyanobacteria	Cyanobacteria		[6]

104	IVK14C1	Intact cells. Cellular extract.	<i>Anabaena</i> Antartica 39	Cyanobacteria	Cyanobacteria	+	[6]
105	IVK15C1	Intact cells. Cellular extract.	<i>Leptolyngbya</i> Antartica 39	Cyanobacteria	Cyanobacteria	+	[6]
106	IVK16C1	Intact cells. Cellular extract.	<i>Tolypothrix</i> 17	Cyanobacteria	Cyanobacteria		[6]
107	IVK17C1	Intact cells. Cellular extract.	<i>Plankthotrix</i>	Cyanobacteria	Cyanobacteria		[6]
108	IVK18C1	Intact cells. Cellular extract.	<i>Chroococciopsis</i> O29	Cyanobacteria	Cyanobacteria	+	[4]
109	IVK19C1	Intact cells. Cellular extract.	<i>Chroococciopsis</i> O29	Cyanobacteria	Cyanobacteria	+	
110	IVK1C1	Intact cells. Cellular extract.	<i>Nostoc</i> grown in nitrate, <i>Anabaena</i> PCC7120	Cyanobacteria	Cyanobacteria	+	[6]
111	IVK1S2	Exopolisacarides fraction	<i>Nostoc</i> grown in nitrate, <i>Anabaena</i> PCC7120	Cyanobacteria	Cyanobacteria	+	[6]
112	IVK20C1	Intact cells. Cellular extract.	<i>Chroococciopsis</i> O57	Cyanobacteria	Cyanobacteria	+	
113	IVK21C1	Intact cells. Cellular extract.	<i>Chroococciopsis</i> O57	Cyanobacteria	Cyanobacteria	+	
114	IVK22C1	Intact cells. Cellular extract.	<i>Chroococciopsis</i> 171	Cyanobacteria	Cyanobacteria		
115	IVK2C1	Intact cells. Cellular extract.	<i>Nostoc</i> grown without nitrate, <i>Anabaena</i> PCC7120	Cyanobacteria	Cyanobacteria	+	[6]
116	IVK2S2	Exopolisacarides fraction	<i>Nostoc</i> grown without nitrate, <i>Anabaena</i> PCC7120	Cyanobacteria	Cyanobacteria	+	[6]
117	IVK3C1	Intact cells. Cellular extract.	<i>Microcystis flos-aquae</i>	Cyanobacteria	Cyanobacteria		[6]
118	IVK4C1	Intact cells. Cellular extract.	<i>Microcystis novacekii</i>	Cyanobacteria	Cyanobacteria	+	[6]
119	IVK5C1	Intact cells. Cellular extract.	<i>Microcystis aeruginosa</i>	Cyanobacteria	Cyanobacteria		[6]
120	IVK6C1	Intact cells. Cellular extract.	<i>Aphanizomenon ovalisporum</i>	Cyanobacteria	Cyanobacteria		[6]
121	IVK7C1	Intact cells. Cellular extract	<i>Phormidium</i> BGU3	Cyanobacteria	Cyanobacteria	+	[6]
122	IVK8C1	Intact cells. Cellular extract	<i>Rivularia</i> sp. MU15	Cyanobacteria	Cyanobacteria	+	[6]
123	IVK9C1	Intact cells. Cellular extract	<i>Chamaesiphon</i>	Cyanobacteria	Cyanobacteria		[6]
124	IVL10C1	Intact cells. Cellular extract	<i>Dechloromarinus chlorophilus</i> , strain NSS	Gammaproteobacteria	Perchlorate reducers		
125	IVL11C1	Intact cells. Cellular extract.	<i>Dechloromonas aromatica</i> , strain RCB	Betaproteobacteria	Perchlorate reducers		
126	IVL12C1	Intact cells. Cellular extract	<i>Arcobacter</i> sp., strain CAB	Epsilonproteobacteria	Perchlorate reducers	+	[4]
127	IVL1C1	Intact cells. Cellular extract	<i>Azospira suillum</i> , strain PS	Betaproteobacteria	Perchlorate reducers		
128	IVL2C1	Intact cells. Cellular extract	<i>Magnetospirillum bellicus</i> , strain VDY	Alphaproteobacteria	Perchlorate reducers		
129	IVL3C1	Intact cells. Cellular extract	<i>Ideonella dechloratans</i>	Betaproteobacteria	Perchlorate reducers		

130	IVL4C1	Intact cells. Cellular extract	<i>Dechlorobacter hydrogenophilus</i> , strain LT-1	Betaproteobacteria	Perchlorate reducers		
131	IVL5C1	Intact cells. Cellular extract	<i>Propionivibrio militaris</i> , strain MP	Betaproteobacteria	Perchlorate reducers	+	[4]
132	IVL6C1	Intact cells. Cellular extract	<i>Dechloromonas agitata</i> , strain CKB	Betaproteobacteria	Perchlorate reducers		
133	IVL6S2	Culture supernatant with isopropanol	<i>D. agitata</i> , strain CKB	Betaproteobacteria	Perchlorate reducers		
134	IVL7C1	Intact cells. Cellular extract	<i>Magnetospirillum</i> sp., strain WD	Alphaproteobacteria	Perchlorate reducers		
135	IVL8C1	Intact cells. Cellular extract	<i>Azospira</i> sp., strain ZAP	Betaproteobacteria	Perchlorate reducers		
136	IVL9C1	Intact cells. Cellular extract	<i>Shewanella algae</i>	Gammaproteobacteria	Perchlorate reducers		
137	VIIIA1V1	Virus <i>Salinibacter ruber</i> concentrated	<i>S. ruber</i>	Bacteroidetes	Halophilic viruses		[4]
138	VIID1BF	Whole extract (Guanidinio.HCl)	Environmental sample1 (gypsum curb / sulfates)	Environmental sample	Mesophilic environment	+	
139	VD2BF	Whole extract	Biofilm from Mansimongs mines (SA7 + SA9) South Africa	Biofilm from environmental samples	Mines	+	
140	ASB	ATP synthase, subunit B/Purified recombinant polipeptide	<i>Archaeoglobus fulgidus</i>	Euryarchaeota	Proteins and peptides	+	[3]
141	ASF1	ATP synthase F1, subunit alpha/Purified recombinant polipeptide	<i>Thermotoga maritima</i>	Thermotogaceae	Proteins and peptides	+	[3]
142	cld	Alpha-chlorite dismutase	<i>Dechloromonas agitata</i> strain CKB	Betaproteobacteria	Proteins and peptides	+	[7]
143	NRA	Nitrate reductase, alpha subunit/Purified recombinant polipeptide	<i>Geobacter metallireducens</i>	Deltaproteobacteria	Proteins and peptides	+	[3]
144	Prot_PCR	Perchlorate reductase	<i>Dechloromonas aromatica</i>	Betaproteobacteria	Proteins and peptides	+	[4]
145	Prot_ABCtransporter	ABC transporter for nitrogenase (purified recombinant polypeptide)	<i>T. scotoductus</i>	Gammaproteobacteria	Proteins and peptides	+	[5]
146	Prot_ApsA_RB11754	Adenylylsulfate reductase alpha subunit	<i>Desulfovibrio desulfuricans</i>	Deltaproteobacteria	Proteins and peptides	+	[3]
147	Prot_DsrA_RB11365	Sulfite reductase, dissimilatory-type subunit alpha	<i>A. fulgidus</i>	Euryarchaeota	Proteins and peptides	+	[3]
148	Prot_DsrB_RB11368	Sulfite reductase, dissimilatory-type subunit beta	<i>A. fulgidus</i>	Euryarchaeota	Proteins and peptides	+	[3]
149	Prot_EFG_RB11359	Elongation factor G	<i>T. maritima</i>	Thermotogaceae	Proteins and peptides	+	[3]
150	Prot_ICDH_RB11756	Peroxisomal isocitrate dehydrogenase	<i>Arabidopsis thaliana</i>	Eukaryota (Viridiplantae)	Proteins and peptides	+	[3]
151	Prot_Lasparaginase	L-Asparaginase (purified)	<i>Escherichia coli</i> (Batch culture)	Gammaproteobacteria	Proteins and peptides	+	BIODESIGN (K59171R)

152	Prot_LBP+BSA	Lipopolysaccharide (in BSA)	<i>Pseudomonas sp.</i>	Gammaproteobacteria	Proteins and peptides	+	[3]
153	Prot_NADH_RB11364	NADH-quinone oxidoreductase subunit G	<i>Pseudomonas putida</i>	Gammaproteobacteria	Proteins and peptides	+	[3]
154	PCR A/B	Perchlorate reductase subunit alpha/beta	<i>Dechloromonas aromatica</i>	Betaproteobacteria	Proteins and peptides	+	[4]
155	Prot_FeReTs_983	Iron reductase/Purified recombinant polipeptide	<i>T.scotoductus</i>	Gammaproteobacteria	Proteins and peptides	+	[3]
156	Prot-BaFER	Bacterioferritin	<i>D. desulfuricans</i>	Deltaproteobacteria	Proteins and peptides	+	[5]
157	Prot-LR1	ABC transporter (from South Africa mines)	<i>T. scotoductus</i>	Gammaproteobacteria	Proteins and peptides	+	[5]
158	Prot-Pfu-DPS	DNA protection during starvation protein	<i>P. furiosus</i>	Archaea	Proteins and peptides	+	[5]
159	Prot-Pfu-FER	Ferritin	<i>P. furiosus</i>	Archaea	Proteins and peptides	+	[5]
160	Prot-Sso-DPS	DNA protection during starvation protein	<i>Sulfolobus solfataricus</i>	Archaea	Proteins and peptides	+	[5]
161	RbcL	Ribulose biphosphate carboxylase large chain	<i>A. thaliana</i>	Eukaryota (Viridiplantae)	Proteins and peptides	+	[3]
162	HtpG	Heat shock proteinG/Purified recombinant polipeptide	<i>Nostoc PCC73102</i>	Cyanobacteria	Proteins and peptides		[3]
163	HupS	Ni-Fe membrane hydrogenase S chain/purified recombinant polipeptide	<i>L. ferrooxidans</i>	Nitrospirae	Proteins and peptides		[3]
164	McrB	Methyl CoM reductase I, B subunit	<i>Methanococcoides burtonii</i>	Euryarchaeota	Proteins and peptides		[3]
165	NifD	Nitrogenase protein alpha chain	<i>G. metallireducens</i>	Deltaproteobacteria	Proteins and peptides		[3]
166	NirS	Nitrite reductase	<i>Pseudomonas aeruginosa</i>	Gammaproteobacteria	Proteins and peptides		[3]
167	NOR1	Nitrite oxidoreductase Beta subunit	<i>Nitrobacter hamburgensis</i>	Alphaproteobacteria	Proteins and peptides		[3]
168	CspA	Cold shock protein A	<i>P. putida</i>	Gammaproteobacteria	Proteins and peptides		[3]

**Table S1.** List of antibodies printed in LDChip168 for this study. Please note that numbers in the first column (peak) correspond to the peak numbers in Fig. 2 and the positive sign (+) in the seventh column indicates those antibodies detected in one or more samples of this study.

### Supplementary references

1. Parro, V.; Rodríguez-Manfredi, J.A.; Briones, C.; Compostizo, C.; Herrero, P.L.; Vez, E.; Sebastián, E.; Moreno-Paz, M.; García-Villadangos, M.; Fernández-Calvo, P., et al. Instrument development to search for biomarkers on mars: Terrestrial acidophile, iron-powered chemolithoautotrophic communities as model systems. *Planetary and Space Science* 2005, 53, 729-737, doi:<https://doi.org/10.1016/j.pss.2005.02.003>.
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Ab Name/ID	Ab* (labelled) concentration (dilution factor from original [2 mg mL <sup>-1</sup> ])	Ag limit detection (cells/ml)
Bou. I	1/900	1/1000*
Bou. II	1/800	1/1*
MSIs	1/800	1/100*
SMs	1/800	1/100*
IS. SS1	1/800	1/100*
MASE-BB-1	1/800	10 <sup>3</sup>
MASE-IM-5	1/900	10 <sup>3</sup>
MASE-IM-4	1/800	10 <sup>4</sup>
MASE-SM-3	1/800	10 <sup>4</sup>
MASE-SM-2	1/800	10 <sup>4</sup>
MASE-SM-1	1/900	10 <sup>3</sup>
MASE-IM-7	1/800	10 <sup>3</sup>
MASE-LG-2	1/900	10 <sup>3</sup>
ET2	1/900	10 <sup>4</sup>
MASE-Glacier-SS3	1/800	10 <sup>4</sup>

**Table S2.** Data from assays of new antibodies performed in the core of MASE project and printed as part of the MAS-Chips. First column refers to the working dilutions of antibodies ranged from 1/800 to 1/900 on the basis of the original concentration which is raised to 2 mg mL<sup>-1</sup> after the purification of each one with Protein A (see materials and methods, section 2.3). The second columns shows the lower detection limit of antibodies that was established between 10<sup>4</sup> and 10<sup>3</sup> cells mL<sup>-1</sup> in the antigenic sample (\*it could not be established for the environmental samples in terms of number of cells mL<sup>-1</sup> but was approximated by serial dilutions of the extracted antigen used to generate the corresponding Ab).