

Chapter -3

**Role of
Acidithiobacillus ferrooxidans
in
alleviating the inhibitory
effect of thiosulfate
on the growth of
acidophilic *Acidiphilium*
species isolated from
Garubathan AMD samples**

4.1 INTRODUCTION

Acidithiobacillus ferrooxidans, an acidophilic obligate chemolithoautotrophic bacterium, plays an important role in the mining and metallurgical applications. Growth of different strains of *A. ferrooxidans* is inhibited by naturally occurring compounds such as pyruvic acid, citric acid, oxaloacetic acid and glucose (Borichewski, 1967; Harrison, 1984; Frattini *et al.*, 2000). Presence of such inhibitory substances has been shown to decrease the efficiency of bioleaching (Harrison, 1984; Johnson, 1995). This inhibition could be alleviated in presence of heterotrophic acidophiles such as *Acidiphilium* sp. resulting in the enhancement of iron (pyrite) oxidation (Paiment *et al.*, 2001). Cultivation of *A. ferrooxidans* on acidic ferrous iron agar has always been difficult because of the presence of agar hydrolyzed products. Areas circumscribing point inoculations of acidophilic heterotrophs into ferrous iron agar were observed to be colonized by *A. ferrooxidans* (Harrison, 1984). Earlier authors have described a plate count procedure where acidophilic heterotrophic bacterium was used in agar medium to scavenge organic material for colony formation of *A. ferrooxidans* (Butler and Kempton, 1987). Acidophilic heterotrophs like *A. cryptum* required a very lean organic medium (oligotrophic) containing low concentrations of glucose (0.1%) and yeast extract (0.01%). Higher concentration of glucose or complex organic supplements, which are commonly used in most heterotrophic media (eutrophic), completely inhibited growth (Harrison, 1981). On the other hand, acidophilic autotrophs of the genus *Acidithiobacillus* have shown self-inhibition towards their own metabolic by-products released in the environment (Borichewski, 1967). Therefore, growth of *Acidithiobacillus* may require a relationship with acidophilic heterotrophic members of the genus *Acidiphilium* (Peccia *et al.*, 2000).

Geomicrobiological studies have shown the presence of acidophilic heterotrophic bacteria along with ferrous and/or sulfur oxidizing chemolithoautotrophs (Harrison, 1984). Few obligate acidophilic heterotrophic bacteria representing the genus *Acidiphilium*, such as *A. cryptum* (Harrison, 1981), *A. angustum*, *A. facilis* and *A. rubrum* (Lazaroff *et al.*, 1982, Jones *et al.*, 1984), *A. organovorum* (Lobos *et al.*, 1986), and *A. acidophilum* (Guay and Silver, 1975; Harrison, 1983) have been characterized. The acid-laden, sulfidic mineral-rich environments where they are characteristically found are invariably rich in ferrous and ferric ion as well as sulfur and reduced sulfur compounds.

Thiosulfate release from pyrite surface has been suggested by several authors (Luther, 1987; Rimstidt and Vaughan, 2003; Descostes *et al.*, 2004). It has been observed that thiosulfate and polythionates are the significant products of pyrite oxidation in near neutral solution (Schippers and Sand, 1999), while only low amounts of intermediate sulfur oxyanion species have been observed in acidic solution (Druschel and Borda, 2006). In an earlier study it was shown that the glucose-limited chemostat cultures of *Thiobacillus acidophilus* (re-named as *Acidiphilium acidophilum*, Hiraishi *et al.*, 1998) could oxidize thiosulfate (Pronk *et al.*, 1990). Though sulfur di-oxygenase activity has been reported in *A. acidophilum* and *A. cryptum* DSM 2389 (Rohwerdert and Sand, 2003), chemolithotrophic growth of *A. cryptum* on thiosulfate could not be demonstrated. However, there are reports on oxidation of elemental sulfur by *A. cryptum* in the presence of organic substrates (Harrison, 1983; Hallberg *et al.*, 2001).

In this study, while examining the physiological/ growth characteristics of the *Acidiphilium* sp. DKAP1.1 isolated from enrichment culture of *A. ferrooxidans* (from acid mine drainage of Garubathan, West Bengal, India), it was noticed that the *Acidiphilium* strain failed to grow in its routine culture medium in presence of thiosulfate. This observation led us to investigate as to whether growth inhibition by thiosulfate was irreversible in pure culture and the role of the chemolithoautotrophic partner, *A. ferrooxidans*, in alleviating thiosulfate mediated inhibition in mixed culture.

4.2 MATERIALS AND METHODS

4.2.1 Bacterial strains

One acidophilic heterotrophic strain DKAP1.1 and one acidophilic autotrophic strain DK6.1 were used for this study.

4.2.2 Characterization of DK6.1 and DKAP1.1

4.2.2.1 Diagnostic tests for DK6.1

Gram staining, motility, growth in different pH and temperature, autotrophic growth in different energy substrates (ferrous sulfate or elemental sulfur or thiosulfate or tetrathionate) were done according to the procedures described earlier (Chapter 1.)

4.2.2.2 Diagnostic tests for DKAP1.1

Diagnostic tests for DKAP1.1 were done as described in chapter 2.

4.2.2.3 DNA isolation

For the isolation of DNA from DK6.1 and DKAP1.1, pure culture was grown in elemental sulfur medium for two weeks and modified DSMZ 269 medium for 5-6 days respectively. Cells were harvested by centrifuging at 10,000 rpm for 10 min and frozen at -20°C. Genomic DNA was isolated and purified by the procedure described earlier (Chakraborty *et al.*, 1997), and stored in 0.1 X TE buffer or sterile double distilled water (reviewed in detail in chapter 5).

4.2.2.4 Amplification, cloning, sequencing and analysis of 16S rDNA

16S rDNA gene fragments were amplified by PCR with Taq DNA polymerase and a pair set of eubacterial universal primers 27f and 1492 (Lane, 1991). PCR products were treated with a chloroform/isoamyl alcohol mixture and cloned in pGEM-T easy vectors following the manufacturer's instruction and then transformed into *Escherichia coli* XL1 Blue. Nucleotide sequencing was performed with the ABI PRISM™ Dye Terminator Cycle Sequencing Ready Reaction kit and the reactions were analyzed on an ABI PRISM 377 DNA sequencer using SP6 and T7 promoter primers. Nucleotide sequence analysis was performed using BLAST search programs [National Center for Biotechnology Information (NCBI)] (the procedure is reviewed in detail chapter 5).

4.2.2.5 Growth of DKAP1.1 in Elemental sulfur spent medium of autotrophic DK6.1 pure culture

To test the oligotrophic nature, DKAP1.1 was grown in elemental sulfur spent medium of autotrophic DK6.1 pure culture. Three weeks old elemental sulfur culture of DK6.1 was filtered through Whatman filter paper no.1 to remove the suspended elemental sulfur particles. The filtrate was centrifuged at 10,000 rpm to exclude DK6.1 cells as pellet. The supernatant was passed through bacterial filter (pore size: 0.25 µm) and sterilized by autoclaving with flowing steam for 20 min. By this process, any traces of sulphide and sulfite that may be present in the spent elemental sulfur medium would be oxidized to sulfate. Presence of any other oxyanions in the spent medium may be ruled out

as no other oxyanions are produced during the course of elemental sulfur oxidation by *A. ferrooxidans*. Soluble form of elemental sulfur, if at all present in the spent medium, would be as low as 5µg/l (Roy and Trudinger, 1970; Steudal *et al.*, 1987). DKAP1.1 cells were grown in elemental sulfur spent medium with and without the supplementation of glucose (0.1%) and yeast extract (0.01%). Growth of the culture in different conditions was monitored by counting the viable cells in Petroff-Hausser bacterial counting chamber.

4.2.2.6 Effect of submerged layering of heterotrophic DKAP1.1 cells on the manifestation of chemoautotrophic DK6.1 colonies over modified 9K agar surface

Effect of acidophilic heterotrophs in the growth of acidophilic chemoautotroph was assessed by growing them in single and double layered modified 9K agar plates. The conventionally prepared modified 9K agar plates used for spread-plating of acidophilic chemoautotrophic cells were considered as single layer plates. To prepare double layered plates, 10 ml of molten modified 9K agar medium (45-50°C) was mixed with washed and concentrated log phase cells of acidophilic heterotroph cells ($\sim 2 \times 10^6$ cells) and poured in a Petri- plate to solidify. After solidification of the first, a second layer was made by overlaying it with 10 ml sterile molten modified 9K agar solution and allowed to solidify. Direct spread-plating of diluted *A. ferrooxidans* culture was carried out in both single and double layer plates and were incubated at 28°C. The emergence and increase in diameter of colonies with respect to the period of incubation of plates were recorded.

4.2.2.7 Growth of heterotrophic DKAP1.1 in pure and co-culture with autotrophic DK6.1 in presence of thiosulfate under laboratory condition

20 ml of modified DSMZ 269 medium containing 0.1% glucose and 0.01% yeast extract in each of the 250 ml Erlenmeyer flasks was supplemented with different concentrations of thiosulfate. 100 mM thiosulfate stock solution was prepared in distilled water and sterilized separately. The required quantity of sterile 100 mM thiosulfate solution was added to the medium to obtain the desired concentration (0.5, 0.7, 1.0, 1.5, and 2.0 mM) of thiosulfate. In the first experimental set, pre-grown DKAP1.1 cells at an initial concentration of approximately $2.07\text{-}3.3 \times 10^5$ cells/ml was added to the thiosulfate containing DSMZ 269 medium. Viability and growth of DKAP1.1 in presence of different concentration of thiosulfate was evaluated by dilution plating on DSMZ 269 agar for

determining the minimum inhibitory concentration. In another set of experiment, DKAP1.1 cells ($2.01\text{-}2.6 \times 10^5$ cells/ml) mixed with DK6.1 cells (8.8×10^6 cells/ml) were inoculated in modified DSMZ 269 broth having different concentration of thiosulfate. Survivability of DKAP1.1 cells (in presence of DK6.1) in the thiosulfate supplemented medium was assessed through dilution plating of the mixed culture aliquot at different time intervals on fresh DSMZ 269 agar. On the other hand, viability of the acidophilic chemoautotroph DK6.1 in the thiosulfate supplemented DSMZ broth was determined by periodical inoculation of 50 ml of fresh modified 9K broth with 100 μ l of the mixed culture and observation for the oxidation of ferrous iron.

4.3 RESULTS

4.3.1 Phenotypic and 16S rRNA gene characterization of the test strains

Results of Phenotypic characterization of DK6.1 and DKAP1.1 have already been presented in chapter 1 and chapter 2.

BLASTN analysis of 16S rRNA gene sequence of the isolate DK6.1 (Genbank accession no. FJ602383) revealed 99% similarity with partial 16S rRNA gene sequence of *A. ferrooxidans* LMT 4 (Fig 3.1 and Fig 3.2). Distance tree constructed using Fast minimum Evolution of BLASTN 2.2.19+ program (Zhang *et al.*, 2000) presented a robust cluster of DK6.1 sequence with partial 16S rRNA gene of *A. ferrooxidans* LMT4 (AM502931) and uncultured bacterium clone ANG Pin f8 16S rRNA gene (EU 370304).

BLASTN analysis of 16S rRNA gene sequence of the isolate DKAP1.1 (GENBANK accession no. FJ602384) produced maximum similarity (99%) with 16S rRNA gene sequence of *A. cryptum* JF-5 [being the constituent of its complete genome sequence (CP000697)] (Fig 3.3 and Fig 3.4). In the Distance tree constructed using Fast minimum Evolution of BLASTN 2.2.19+ program (Zhang *et al.*, 2000), DKAP1.1 sequence clustered strongly with 16S rRNA gene of *A. cryptum* JF-5 and 16S rRNA gene sequence of *Acidiphilium* sp. SX-F (FJ194544).

4.3.2 Growth of *A. cryptum* DKAP1.1 in Elemental sulfur spent medium of *A. ferrooxidans* DK6.1 pure culture

DKAP1.1 cells were able to grow in three weeks old elemental sulfur spent medium of *A. ferrooxidans* DK6.1, with or without supplementation of glucose and yeast

extract. However growth was not detected in fresh sterile basal salt solution of elemental sulfur medium devoid of glucose and yeast extract (Fig. 3.5). The ability of *A. cryptum* DKAP1.1 to grow in low nutrient medium such as elemental sulfur spent medium explains the oligotrophic nature of the strain.

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tcgcagctac catgcagtcg acggtacagg tcttcggatg ctgacgagtg gcggacgggt gagtaatgcg
taggaatctg tcttttagtg ggggacaacc cagggaaact tgggctaata ccgcatgagc cctgaggggg
aaagcggggg atcttcggac ctcgcgctaa gagaggagcc tacgtccgat tagctagttg gcgggtaaag
gcccaccaag gcgacgatcg gtagctggtc tgagaggagc accagccaca ctgggactga gacacggccc
agactcctac gggaggcagc agtggggaat ttttcgcaat gggggcaacc ctgacgaagc aatgcccgct
ggatgaagaa ggccttcggg ttgtaaaatc ctttcgtaga ggacgaaaag gcgggttcta atacaatctg
ctgttgacgt gaatccaaga agaagcaccg gctaactcgg tgccagcagc cgcggtaata cggggggtgc
aagcgттаат cggaatcact gggcgtaaag ggtgcgtagc ggtacgttag tctgtcgtga atccccgggc
tcaacctggg aatggcggtg gaaaccgcg cactagagta tgggagaggg tggtggaatt ccaggtgtag
cggtgaaatg cgtagagatc tggaggaaca tcagtggcga aggcggccac ctggcccact actgacgctg
aggcacgaaa gcgtggggag caaacaggat tagataacct ggtagtccac gcctaaccg atgaatacta
gatgtttggt gcctagcgtg ctgagtgctg tagctaacgc gataagtatt ccgctggga agtacggccg
caaggttaaa actcaaagga attgacgggg gcccgcaaa cgggtggagc atgtggttta atctgatgca
acgcaagaa ccttacctgg gcttgacatg tccggaattc tgacagatg cggaaagtgc cttcggggaa
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cgcaaccctt gtccttagtt gccagcggtt cggccgggca ctctagggag actgccgggt acaaacggga
ggaaggtggg gatgacgtca agtcctcatg gcctttatgt ccagggtac acagctgcta caatggcgcg
tacagagggg agccaagccg cgaggtggag cagacccag aaagcgcgct gtagttcgga ttgcagctctg
caactcgact gcataagtc ggaatcgcta gtaatcgcg atcagcatgc cgcggtgaat acgttcccgg
gccttgtaga caccggcgt cacaccatgg gagtggattg taccagaagc agctagccta acctcgggag
ggctaccac ggtacgct

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Fig. 3.1: Partial 16S rRNA gene sequence of acidophilic autotrophic strain DK6.1

DK6.1	11	CATGC-AGTCG-ACGGT-ACAGGTCTTCGGATGCTGACGAGTGGCGGACGGGTGAGTAAT	67
Aferr	27	CATGCAAGTCGAACGGTAACAGGTCTTCGGATGCTGACGAGTGGCGGACGGGTGAGTAAT	86
DK6.1	68	GCGTAGGAATCTGTCTTTTAGTGGGGGACAACCCAGGGAAACTTGGGCTAATACCGCATG	127
Aferr	87	GCGTAGGAATCTGTCTTTTAGTGGGGGACAACCCAGGGAAACTTGGGCTAATACCGCATG	146
DK6.1	128	AGCCCTGAGGGGAAAGCGGGGATCTTCGGACCTCGCGCTAAGAGAGGAGCCTACGTCC	187
Aferr	147	AGCCCTGAGGGGAAAGCGGGGATCTTCGGACCTCGCGCTAAGAGAGGAGCCTACGTCC	206
DK6.1	188	GATTAGCTAGTTGGC-GGGTAAAGGCCACCAAGGCGACGATCGGTAGCTGGTCTGAGAG	246
Aferr	207	GATTAGCTAGTTGGCGGGTAAAGGCCACCAAGGCGACGATCGGTAGCTGGTCTGAGAG	266
DK6.1	247	GACGACCAGCCACACTGGGACTGAGACACGGCCAGACTCCTACGGGAGGCAGCAGTGGG	306
Aferr	267	GACGACCAGCCACACTGGGACTGAGACACGGCCAGACTCCTACGGGAGGCAGCAGTGGG	326
DK6.1	307	GAATTTTTCGCAATGGGGGCAACCCTGACGAAGCAATGCCGCTGGATGAAGAAGGCCCTT	366
Aferr	327	GAATTTTTCGCAATGGGGGCAACCCTGACGAAGCAATGCCGCTGGATGAAGAAGGCCCTT	386
DK6.1	367	CGGGTTGTAAAGTCCTTTCGTGGAGGACGAAAAGGCGGGTCTAATACAATCTGCTGTTG	426
Aferr	387	CGGGTTGTAAAGTCCTTTCGTGGAGGACGAAAAGGCGGGTCTAATACAATCTGCTGTTG	446
DK6.1	427	ACGTGAATCCAAGAAGAAGCACCGGCTAATCCGTGCCAGCAGCCGCGGTAATACGGGGG	486
Aferr	447	ACGTGAATCCAAGAAGAAGCACCGGCTAATCCGTGCCAGCAGCCGCGGTAATACGGGGG	506

DK6.1	487	GTGCAAGCGTTAATCGGAATCACTGGGCGTAAAGGGTGCCTA-GCGGTACGTTA-GTCTG	544
Aferr	507	GTGCAAGCGTTAATCGGAATCACTGGGCGTAAAGGGTGCCTAGGCGGTACGTTAGGTCTG	566
DK6.1	545	TCGTG-AATCCCCGGGCTCAACCTGGGAATGGCGGTGAAACC CGGCGCACTAGAGTATGG	603
Aferr	567	TCGTGAAATCCCCGGGCTCAACCTGGGAATGGCGGTGAAACC CGGCGCACTAGAGTATGG	626
DK6.1	604	GAGAGGGTGGTGGAAATCCAGGTGTAGCGGTGAAATGCGTAGAGATCTGGAGGAACATCA	663
Aferr	627	GAGAGGGTGGTGGAAATCCAGGTGTAGCGGTGAAATGCGTAGAGATCTGGAGGAACATCA	686
DK6.1	664	GTGGCGAAGGCGGCCACCTGGCCCAATACTGACGCTGAGGCACGAAAGCGTGGGGAGCAA	723
Aferr	687	GTGGCGAAGGCGGCCACCTGGCCCAATACTGACGCTGAGGCACGAAAGCGTGGGGAGCAA	746
DK6.1	724	ACAGGATTAGATACCCTGGTAGTCCACGCCCTAAACGATGAATACTAGATGTTTGGTGCC	783
Aferr	747	ACAGGATTAGATACCCTGGTAGTCCACGCCCTAAACGATGAATACTAGATGTTTGGTGCC	806
DK6.1	784	TAGCGTACTGAGTGTCTGCTAGCTAACCGGATAAGTATTCGCCTGGGAAGTACGGCCGCAA	843
Aferr	807	TAGCGTACTGAGTGTCTGCTAGCTAACCGGATAAGTATTCGCCTGGGAAGTACGGCCGCAA	866
DK6.1	844	GGTTAAACTCAAAGGAATTGACGGGGGCCCGCACAAAGCGGTGGAGCATGTGGTTTAATT	903
Aferr	867	GGTTAAACTCAAAGGAATTGACGGGGGCCCGCACAAAGCGGTGGAGCATGTGGTTTAATT	926
DK6.1	904	CGATGCAACGCGAAGAACCCTTACCTGGGCTTGACATGTCCGGAATTCGCAGAGATGCGG	963
Aferr	927	CGATGCAACGCGAAGAACCCTTACCTGGGCTTGACATGTCCGGAATTCGCAGAGATGCGG	986
DK6.1	964	AAGTGCCCTTCGGGGAATCGGAACACAGGTGCTGCATGGCTGTCGTGAGCTCGTGTCTGTG	1023
Aferr	987	AAGTGCCCTTCGGGGAATCGGAACACAGGTGCTGCATGGCTGTCGTGAGCTCGTGTCTGTG	1046
DK6.1	1024	AGATGTTGGGTTAAGTCCCGCAACGAGCGCAACCCTTGTCTTAGTTGCCAGCGGTTCCG	1083
Aferr	1047	AGATGTTGGGTTAAGTCCCGCAACGAGCGCAACCCTTGTCTTAGTTGCCAGCGGTTCCG	1106
DK6.1	1084	CCGGGCACTCTAGGGGAGACTGCCGGTGACAAACCGGAGGAAGGTGGGGATGACGTCAAGT	1143
Aferr	1107	CCGGGCACTCTAGGGGAGACTGCCGGTGACAAACCGGAGGAAGGTGGGGATGACGTCAAGT	1166
DK6.1	1144	CCTCATGGCCTTTATGTCCAGGGCTACACACGTGCTACAATGGCGCGTACAGAGGGAAGC	1203
Aferr	1167	CCTCATGGCCTTTATGTCCAGGGCTACACACGTGCTACAATGGCGCGTACAGAGGGAAGC	1226
DK6.1	1204	CAAGCCGCGAGGTGGAGCAGACCCAGAAAGCGCGTCTAGTTCCGATTGCAGTCTGCAA	1263
Aferr	1227	CAAGCCGCGAGGTGGAGCAGACCCAGAAAGCGCGTCTAGTTCCGATTGCAGTCTGCAA	1286
DK6.1	1264	CTCGACTGCATGAAGTCGGAATCGCTAGTAATCGCGGATCAGCATGCCGCGGTGAATACG	1323
Aferr	1287	CTCGACTGCATGAAGTCGGAATCGCTAGTAATCGCGGATCAGCATGCCGCGGTGAATACG	1346
DK6.1	1324	TTCCCGGGCCTTGACACACCGCCCGTACACCATGGGAGTGGATTGTACCAGAAGCAGC	1383
Aferr	1347	TTCCCGGGCCTTGACACACCGCCCGTACACCATGGGAGTGGATTGTACCAGAAGCAGC	1406
DK6.1	1384	TAGCCTAACCT-CGGGAGGGCG-T-ACCACGGTA 1414	
Aferr	1407	TAGCCTAACCTTCGGGAGGGCGGTTACCACGGTA 1440	

Fig 3.2: Blastn result of 16SrRNA sequence of DK6.1 showing 99% homology with partial 16SrRNA gene sequence of *Acidithiobacillus ferrooxidans* LMT4. Vertical lines show the positions of identical bases. Aferr: *A. ferrooxidans* LMT4

4.3.3 Effect of submerged layering of *A. cryptum* DKAP1.1 cells on the manifestation of *A. ferrooxidans* DK6.1 colonies on Iron-agar

A. ferrooxidans DK6.1 cells grown in ferrous iron medium when dilution plated separately on single and double layered plates, showed marked variation with respect to

the number, emergence and robustness of colonies with time (Table 3.1). A positive effect on growth of *A. ferrooxidans* DK6.1 was observed when it was co-cultured with *A. cryptum* DKAP1.1 in double layered plates. Incorporation of heterotrophic *A. cryptum* DKAP1.1 in an underlay of ferrous-agar medium promoted growth of *A. ferrooxidans* DK6.1 colonies through enhanced ferrous iron oxidation.

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tcgcagctta ccatgcagtc gcacgggagc ggcaacctgt cagtggcgga cgggtgagta acgcgtagga
atctatcctt gggtagggga caaccgtggg aaactacggc taataccgca tgatccctga ggggcaaagg
cgaaagtgc ctgaggagga gectgcgtct gattaggtag ttggtggggg aaaggcctac caagcctgct
atcagtagct ggtctgagag gatgatcagc cacactggga ctgagacacg gccagactc ctacggggag
cagcagtggg gaatattgga caatgggca aagcctgatc cagcaatgcc gcgtggatga agaaggtctt
cggattgtaa agtccttttg gcggggacga tgatgacggt acccgagaa taagctccgg ctaacttcgt
gccagcagcc gcgtaatac gaagggggct agcgttgctc ggaatgactg ggcgtaaagg gcgcgtaggc
ggacggcaca gtcaggcgtg aaattcctgg gctcaacctg ggggactgct tctgagacgt gttgtcttga
gtatggaaga gggttgtgga atttccagt tagaggtgaa attcgtagat attggaaga acaccggtgg
cgaaggcggc aacctggctc attactgacg ctgaggcgcg aaagcgtggg gagcaaacag gattagatac
cctggtagtc cacgctgtaa acgatgtgtg ctggatggtg ggggtgcttag cacttcagt tctgtagctaa
cgcggtaagc acaccgctg gggagtacgg ccgcaaggtt gaaactcaa ggaattgacg ggggcccgca
caagcgtgg agcatgtggt ttaattcgaa gcaacgcgca gaaccttacc aggatttgac atggggagta
ccggtccaga gatggacttt cccgcaaggg gctctgcac aggtgctgca tggctgctgt cagctcgtgt
cgtgagatgt tgggttaagt cccgcaacga gcgcaacct cgccttcagt tgccagcatg tttgggtggg
cactctgaag gaactgccgg tgacaagccg gaggaaggtg gggatgacgt caagtctca tggcccttat
gtcctgggct acacacgtgc tacaatggcg gtgacagtgg gaagccaggt ggtgacacc agctgatctc
aaaaagcgt ctcagtcgg attgcactc gcaactcgag tgcatgaagg tggatcgct agtaatcgcg
gatcagcatg ccgggtgaa taegtcccg ggcctgtac acaccgccc tcacaccatg ggatttggtt
tgaccttaag ttggtgctgt aaccgcaag gaggcagcca

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Fig. 3.3: Partial 16SrRNA gene sequence of the acidophilic heterotroph DKAP1.1.

DKAP1.1	3	GCA-GCTT-AC-CATGC-AGTCGCACGGGCAGGGCAACCTGTCACTGGCGGACGGGTGAG	58
AcrJF-5	2012865	GCATGCTTAACACATGCAAGTCGCACGGGCAGGGCAACCTGTCACTGGCGGACGGGTGAG	2012806
DKAP1.1	59	TAACGCGTAGGAATCTATCCTTGGGTGGGGACAACCGTGGGAACTACGGCTAATACCG	118
AcrJF-5	2012805	TAACGCGTAGGAATCTATCCTTGGGTGGGGACAACCGTGGGAACTACGGCTAATACCG	2012746
DKAP1.1	119	CATGATCCCTGAGGGGCAAAGGCGAAAGTCGCCTGAGGAGGAGCCTGCGTCTGATTAGGT	178
AcrJF-5	2012745	CATGATCCCTGAGGGGCAAAGGCGAAAGTCGCCTGAGGAGGAGCCTGCGTCTGATTAGGT	2012686
DKAP1.1	179	AGTTGGTGGGGTAAAGGCTACCAAGCCTGCGATCAGTAGCTGGTCTGAGAGGATGATCA	238
AcrJF-5	2012685	AGTTGGTGGGGTAAAGGCTACCAAGCCTGCGATCAGTAGCTGGTCTGAGAGGATGATCA	2012626
DKAP1.1	239	GCCACACTGGGACTGAGACACGGCCAGACTCCTACGGGAGGCAGCAGTGGGGAATATTG	298
AcrJF-5	2012625	GCCACACTGGGACTGAGACACGGCCAGACTCCTACGGGAGGCAGCAGTGGGGAATATTG	2012566
DKAP1.1	299	GACAATGGGCGAAAGCCTGATCCAGCAATGCCCGTGGATGAAGAAGTCTTCGGATTGT	358
AcrJF-5	2012565	GACAATGGGCGAAAGCCTGATCCAGCAATGCCCGTGGATGAAGAAGTCTTCGGATTGT	2012506
DKAP1.1	359	AAAGTCCTTTTGGCGGGACGATGATGACGGTACCCGAGAAATAGCTCCGGCTAACTTC	418
AcrJF-5	2012505	AAAGTCCTTTTGGCGGGACGATGATGACGGTACCCGAGAAATAGCTCCGGCTAACTTC	2012446

DKAP1.1	419	GTGCCAGCAGCCGCGGTAATACGAAGGGGGCTAGCGTTGCTCGGAATGACTGGGCGTAAA	478
AcrJF-5	2012445	GTGCCAGCAGCCGCGGTAATACGAAGGGGGCTAGCGTTGCTCGGAATGACTGGGCGTAAA	2012386
DKAP1.1	479	GGGCGCGTAGGCGGACGGCACAGTCAGGCGTGAAATTCCTGGGCTCAACCTGGGGGACTG	538
AcrJF-5	2012385	GGGCGCGTAGGCGGACGGCACAGTCAGGCGTGAAATTCCTGGGCTCAACCT-GGGGACTG	2012327
DKAP1.1	539	CGTCTGAGACGTGTTGTCTTGAGTATGGAAGAGGGTTGTGGAATTCAGTGTAGAGGTG	598
AcrJF-5	2012326	CGTCTGAGACGTGTTGTCTTGAGTATGGAAGAGGGTTGTGGAATTCAGTGTAGAGGTG	2012267
DKAP1.1	599	AAATTCGTAGATATTGAAAGAACACCGGTGGCGAAGGCGGCAACCTGGTCCATTACTGA	658
AcrJF-5	2012266	AAATTCGTAGATATTGAAAGAACACCGGTGGCGAAGGCGGCAACCTGGTCCATTACTGA	2012207
DKAP1.1	659	CGCTGAGGCGCGAAAGCGTGGGGAGCAAACAGGATTAGATACCTGGTAGTCCACGCTGT	718
AcrJF-5	2012206	CGCTGAGGCGCGAAAGCGTGGGGAGCAAACAGGATTAGATACCTGGTAGTCCACGCTGT	2012147
DKAP1.1	719	AAACGATGTGTGCTGGATGTTGGGGTGCCTTAGCACTTCAGTGTCTAGCTAACGCGGTAA	778
AcrJF-5	2012146	AAACGATGTGTGCTGGATGTTGGGGTGCCTTAGCACTTCAGTGTCTAGCTAACGCGGTAA	2012087
DKAP1.1	779	GCACACCGCTGGGGAGTACGGCCGCAAGGTTGAAACTCAAAGGAATTGACGGGGCCCG	838
AcrJF-5	2012086	GCACACCGCTGGGGAGTACGGCCGCAAGGTTGAAACTCAAAGGAATTGACGGGGCCCG	2012027
DKAP1.1	839	CACAAGCGGTGGAGCATGTGGTTTAAATTCGAAGCAACGCGCAGAACCTTACCAGATTG	898
AcrJF-5	2012026	CACAAGCGGTGGAGCATGTGGTTTAAATTCGAAGCAACGCGCAGAACCTTACCAGATTG	2011967
DKAP1.1	899	ACATGGGGAGTACCGGTCCAGAGATGGACTTCCCGCAAGGGGCTCTCGCACAGGTGCTG	958
AcrJF-5	2011966	ACATGGGGAGTACCGGTCCAGAGATGGACTTCCCGCAAGGGGCTCCCGCACAGGTGCTG	2011907
DKAP1.1	959	CATGGCTGTGTCAGCTCGTGTGAGATGTTGGGTTAAGTCCCGCAACGAGCGCAACC	1018
AcrJF-5	2011906	CATGGCTGTGTCAGCTCGTGTGAGATGTTGGGTTAAGTCCCGCAACGAGCGCAACC	2011847
DKAP1.1	1019	CTCGCCTTCAGTTGGCAGCATGTTTGGGTGGGCACTCTGAAGGAAGTGGCGGTGACAAGC	1078
AcrJF-5	2011846	CTCGCCTTCAGTTGGCAGCATGTTTGGGTGGGCACTCTGAAGGAAGTGGCGGTGACAAGC	2011787
DKAP1.1	1079	CGGAGGAAGGTGGGGATGACGTCAAGTCCCTATGTCCTGGGCTACACACGT	1138
AcrJF-5	2011786	CGGAGGAAGGTGGGGATGACGTCAAGTCCCTATGTCCTGGGCTACACACGT	2011727
DKAP1.1	1139	GCTACAATGGCGGTGACAGTGGGAAGCCAGGTGGTGACACCGAGCTGATCTCAAAAAGCC	1198
AcrJF-5	2011726	GCTACAATGGCGGTGACAGTGGGAAGCCAGGTGGTGACACCGAGCTGATCTCAAAAAGCC	2011667
DKAP1.1	1199	GTCTCAGTTCGGATTGCACCTCTGCAACTCGAGTGCATGAAGGTGGAATCGCTAGTAATCG	1258
AcrJF-5	2011666	GTCTCAGTTCGGATTGCACCTCTGCAACTCGAGTGCATGAAGGTGGAATCGCTAGTAATCG	2011607
DKAP1.1	1259	CGGATCAGCATGCCGCGGTGAATACGTTCCCGGGCCTTGTACACACCGCCCGTACACCA	1318
AcrJF-5	2011606	CGGATCAGCATGCCGCGGTGAATACGTTCCCGGGCCTTGTACACACCGCCCGTACACCA	2011547
DKAP1.1	1319	TGGGATTTGGTTTGACCTTAAGTTGGTGCCTAACCCGCAAGG-AGGCAGCCA 1370	
AcrJF-5	2011546	TGGGATTTGGTTTGACCTTAAGTTGGTGCCTAACCCGCAAGGAGGCAGCCA 2011494	

Fig 3.4: Blast result of 16S rRNA sequence of DKAP1.1 showing 99% homology with the partial 16SrRNA sequence of *Acidiphilium cryptum* JF-5 (AcrJF-5). Vertical lines show the position of identical bases.

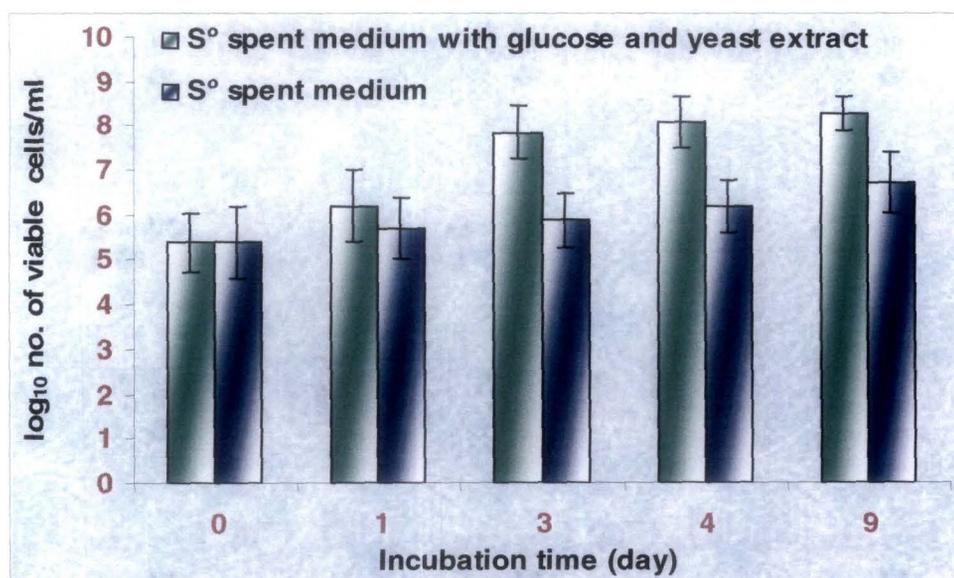


Fig 3.5: Growth of acidophilic heterotroph DKAP1.1 in spent elemental sulfur medium with or without the supplementation of glucose. Error bars represent the standard deviation in replicate samples (n =3).

Table 3.1: Comparison of the rate of emergence and increase in the diameter of *A. ferrooxidans* GBVI colonies in Fe (II) iron agar in single and double layered plates.

Single layered Fe(II)-agar plate				Double layered Fe(II)-agar plate			
Dilution of the stock	No. of colonies emerged	No. of days of incubation	Average diameter (mm)	Dilution of the stock	No. of colonies emerged	No. of days of incubation	Average diameter (mm)
10 ⁻⁷	Nil	5	na	10 ⁻⁷	0	5	nd
10 ⁻⁷	-do-	8	na	10 ⁻⁷	14	8	2.0
10 ⁻⁷	-do-	9	na	10 ⁻⁷	14	9	2.6
10 ⁻⁷	-do-	10	na	10 ⁻⁷	14	10	3.2
10 ⁻⁷	10	18	nd	10 ⁻⁷	14	12	3.4
10 ⁻⁷	10	23	0.5	10 ⁻⁷	14	13	3.6

na, not applicable

4.3.4 Inhibitory effect of thiosulfate on growth of *A. cryptum* DKAP1.1 and alleviation of inhibition in co-culture with *A. ferrooxidans* DK6.1

A decrease in viability of *A. cryptum* DKAP1.1 cells in thiosulfate supplemented modified DSMZ 269 medium was observed with the increase in the concentration of thiosulfate (Fig. 3.6). It was found that *A. cryptum* DKAP1.1 could tolerate concentrations of thiosulfate up to 0.7 mM. It was observed that the number of viable cells were higher in 48 h treated cells for a given concentration of thiosulfate as compared to 24 h treatment at the same concentration up to 0.7 mM. However at concentration above 1.0 mM, the effect of thiosulfate on the viability was much more pronounced with respect to both the concentration and time of treatment. At 1.0 mM concentration viable cells could be obtained from 24 h treated culture but no viable cells were obtained after 48 h treatment. On further increase of concentration no viable cells could be obtained from either 24 or 48 h treated cells (Fig. 3.6). On the other hand, when pure culture of *A. ferrooxidans* DK6.1 was incubated in modified DSMZ 269 medium for ≥ 24 h with or without thiosulfate, the cells lost their ability to oxidize ferrous iron when transferred to fresh 9K medium.

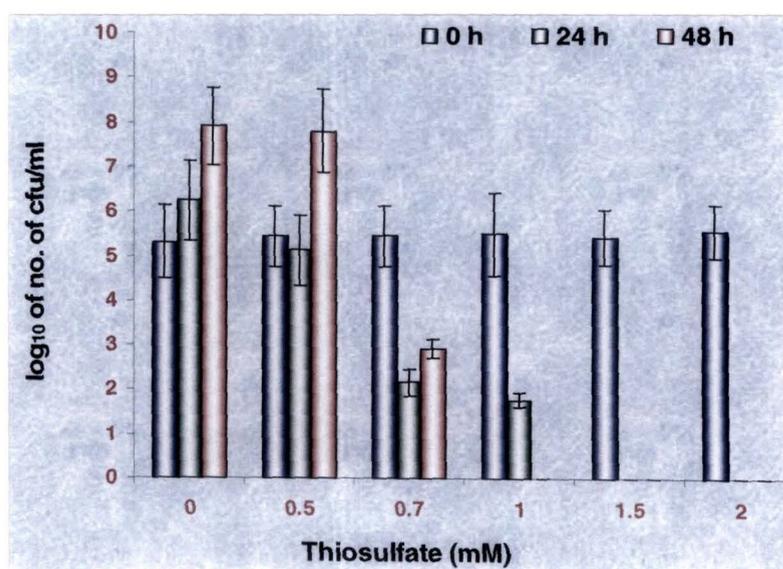


Fig 3.6: Viable cell number of pure culture of *A. cryptum* DKAP1.1 cells in DSMZ 269 medium containing different thiosulfate concentrations.

The death of *A. cryptum* DKAP1.1 cells due to presence of thiosulfate in DSMZ 269 medium was prevented in co-culture with *A. ferrooxidans* DK6.1. Survival of the co-cultured DKAP1.1 cells even in presence 2.0 mM thiosulfate was noted (Fig. 3.7).

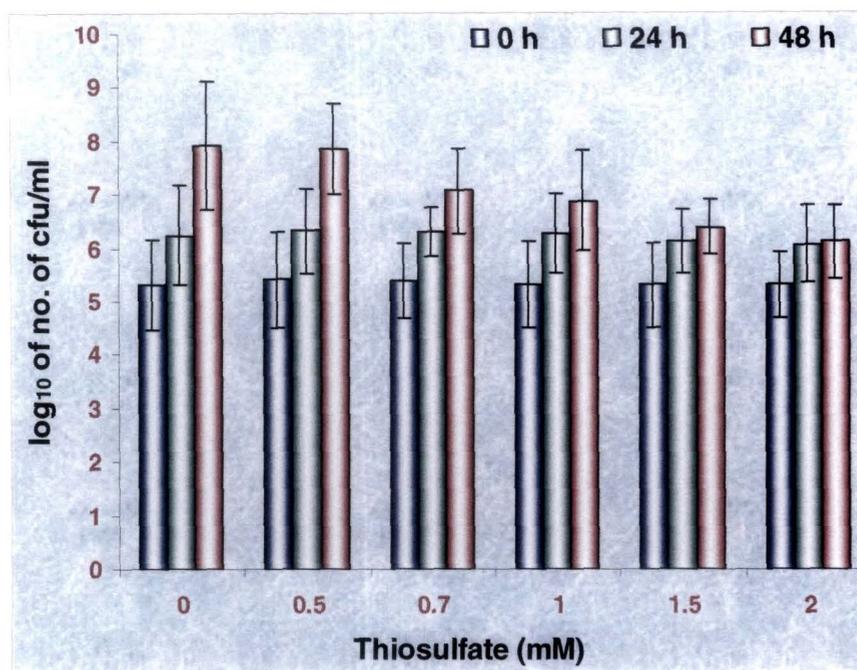


Fig 3.7: Viable cell number of *A. cryptum* DKAP 1.1 in co-culture (with *A. ferrooxidans* DK6.1) cells at different thiosulfate concentrations. Error bars represent the standard deviation on replicate

A. ferrooxidans DK6.1 cells in co-culture remained viable (qualitatively tested by their ability to oxidize ferrous iron in 9K medium) throughout the expanse of the experiment. The lethal effect of thiosulfate on *A. cryptum* DKAP1.1 in pure culture was found directly related to the concentration of thiosulfate in the medium and the lethality could be alleviated in mixed culture with *A. ferrooxidans* DK6.1.

4.4 DISCUSSION

A. ferrooxidans, unique among the acidophilic chemolithotrophic bacteria in its ability to oxidize ferrous iron and sulfidic minerals, was first isolated from acidic mine drainage (AMD) water (Temple and Colmer, 1951) and has since been isolated from a variety of acidic environments. Also, there are numerous reports of the isolation of acidophilic heterotrophs from reportedly pure cultures of *A. ferrooxidans* (Guay and

Silver, 1975; Harrison *et al.*, 1980). These heterotrophs survive during prolonged cultivation of *A. ferrooxidans* in inorganic media, probably by scavenging low concentration of organic compounds excreted by *A. ferrooxidans*. Few of the acidophilic heterotrophic bacteria like *A. cryptum* and *A. acidophilum*, have the ability to reduce ferric iron to ferrous state. In this respect it has been suggested that the removal of (toxic) organic compounds and regeneration of ferrous ions by the heterotrophs may also be advantageous to *A. ferrooxidans* (Harrison, 1984; Marchand and Silverstein, 2003).

The present work was carried out to gain a better understanding of the interactions among acidophilic chemolithoautotrophs and heterotrophs in AMD environments enriched with low amount of organic carbon substrates. In the microbial ecology of acid mine environments, members of the genus *Acidiphilium* may have considerable relations with sulfur- and iron-oxidizing bacteria. Certain conditions like acidic, mesophilic, aerobic, ferrous iron or other mineral sulfide substrate environment, which support growth of sulfur- and iron-oxidizing bacteria, are appropriate for the maintenance of some heterotrophic partners. Pure culture studies on *Acidithiobacillus thiooxidans* have shown that this bacterium excrete pyruvic and oxalo-acetic acids that are self inhibitory at 2×10^{-5} to 7×10^{-5} M (Borichevski, 1967), and therefore, growth of *A. thiooxidans* may necessitate a partnership with an acidophilic heterotroph. It has also been shown that *A. ferrooxidans* cells excrete organic compounds, such as pyruvate, glutamate, aspartate, serine, glycine, and other amino acids, which the heterotrophs can use as substrates for growth (Arkesteyn and deBont, 1980; Schnaltman and Lundgren, 1965). On the other hand, lysed cells of *A. ferrooxidans* may provide an additional source of organic matter. In each case, the quantity of obtainable organic material in autotrophically grown *A. ferrooxidans* cultures is very small. Consequently, the acidophilic heterotrophs have to be competent forager of these organic compounds. Since some of these compounds have been shown to be inhibitory to *A. ferrooxidans* (Harrison, 1984; Tabita and Lundgren, 1971), it is probable that the association is mutualistic in nature. Members of the genus *Acidiphilium* are capable of iron reduction. This biological reduction of Fe^{3+} to Fe^{2+} helps to compensate ferric iron production and thus attenuates acid production in acid mine drainage (Pronk and Johnson, 1992). The aim of this work was to determine the role of the autotroph in such an extreme habitat which was of advantage for the survival of the heterotrophs. The chemolithoautotrophic strain, *A. ferrooxidans* DK6.1, used in this study was isolated and purified from the AMD sample of Garubathan, West Bengal, India. The acidophilic

heterotrophic strain used in this study was isolated from the AMD enrichment medium (typically used for isolation of *A. ferrooxidans* strains). It was characterized as *A. cryptum* DKAP1.1. We wanted to explore the degree of obligate oligotrophy in *A. cryptum* DKAP1.1 as we have observed that it failed to grow in nutrient rich medium (eutrophic medium). The elemental sulfur spent medium of three weeks old *A. ferrooxidans* DK6.1 (which basically contained the residual basal salts and leachates/ excreta of the pure culture of the autotroph as total dissolved organic carbon) culture could support the growth of *A. cryptum* DKAP1.1. (Fig. 3.5). This growth in elemental sulfur spent medium was found to be enhanced by a factor of more than 10 in presence of glucose and yeast extract.

It is well accepted that thiosulfate is central in much of the thinking concerning pyrite oxidation pathways (Druschel and Borda, 2006). Thiosulfate was found to be an important product during short term (< 24 h) experiment of pyrite oxidation at pH 5- 9.5 (Goldhaber and Rey, 1977; Goldhaber, 1983). The rate of tetrathionate formation in pyrite-surface-catalyzed oxidation of thiosulfate was found to be of first order with respect to the pyrite surface concentration and fractional order with respect to thiosulfate concentration (Yong and Schoonen, 1995). Tetrathionate accumulation to the extent of 1.3% (after 24 h of incubation) was observed in chemical pyrite oxidation with 10 mM FeCl₃ at pH 1.9 at 28°C (Schippers and Sand, 1999). Conversely, at neutral pH, the accumulation of thiosulfate (approximately 0.5 mM) was evident in pyrite oxidation (Schippers *et al.*, 1996). However, pyrite oxidation attending the formation of sedimentary roll-type uranium deposits does not occur in very low pH range (Goldhaber and Rey, 1977) and intermediate sulfur compounds like sulfite, thiosulfate, and tetrathionate offer suitable substrates for moderately acidophilic thiobacilli. The role of thiosulfate as an initial soluble intermediate in the pathway of pyrite oxidation has been postulated by different authors (Luther, 1987; Rimstidt and Vaughan, 2003). Earlier authors have suggested that pyrite oxidation generates thiosulfate, which reacts to tetrathionate in acidic solutions containing ferric iron. The acid-insoluble metal sulfides FeS₂, MoS₂, and WS₂ are chemically attacked by iron (III) hexahydrate ions, generating thiosulfate, which is oxidized via tetrathionate, disulfane-monosulfonic acid, and trithionate to sulfuric acid (Schippers and Sand, 1999). The predominant metal sulfide dissolving microorganisms are extremely acidophilic bacteria that are able to oxidize either inorganic sulfur compounds and/or Fe²⁺ ions. Although autotrophic Fe²⁺ oxidizing microorganisms are believed to play a more important role in acidic mineral rich environments, heterotrophic microbes have routinely been

cultured and detected in these extremely substrate limited environmental niches (Peccia *et al.*, 2000). *A. ferrooxidans* thrives on the oxidation of reduced sulfur compounds and in addition, is able to oxidize H_2 , $HCOOH$, Fe^{2+} ions and other metal ions. *Leptospirillum ferrooxidans* and *L. ferriphilum* can grow only by aerobically oxidizing Fe^{2+} ions. Bacteria of the genus *Acidiphilum* display most diverse metabolic flexibility. Addition of thiosulfate profoundly affected the growth of cultures of acidophilic bacteria typical of acid mine drainage (AMD) sites: the iron and sulfur oxidizing autotrophic bacteria *A. ferrooxidans*, and a common heterotrophic strain *A. cryptum*. Growth of *A. cryptum* DKAP1.1 on DSMZ 269 medium was significantly inhibited in the presence of 1.0 mM thiosulfate in spite of favorable cultural conditions. The actual mechanism of this inhibition is not known. However, the lethal effect of the dual presence of tetrathionate and thiosulfate on *Escherichia coli* has been reported (Palumbo and Alford, 1970). The maximum lethal effect on *E. coli* occurred at a 3 to 1 molar ratio of thiosulfate-tetrathionate in growth medium. Tetrathionate is known to react with free sulfhydryl groups of enzyme and to cause their inactivation (Parker and Allison, 1969). Thiosulfate can also react with sulfhydryl groups (Postgate, 1963). It was therefore suggested that thiosulfate-tetrathionate mixture in growth medium of *E. coli* interfered with the synthesis, the activity, or both, of sulfur-containing enzymes or cell wall and membrane components (Palumbo and Alford, 1970). Thiosulfate is unstable in acidic solutions. Inhibition of growth of heterotrophic *A. cryptum* DKAP1.1 in presence of thiosulfate in acidic growth medium could be due to similar effect as was suggested for *E. coli*.

When *A. ferrooxidans* DK6.1 cells that were inoculated at the 1.0 g/l glucose condition in DSMZ 269 medium (with or without thiosulfate) were transferred to a glucose-free 9K ferrous medium after 24 h of incubation, no iron oxidation was observed (even after a month incubation) suggesting that incubation in presence of dissolved organic carbon including glucose had an acute, and perhaps irreversibly toxic, effect on the autotroph. Interestingly, in co-culture both the autotroph and the heterotroph survived in thiosulfate containing DSMZ 269 medium. In the mixed-culture flask that experienced viability of *A. cryptum* DKAP1.1, it is likely that consumption of reduced sulfur species by autotrophic bacteria relieved the inhibitory stress of thiosulfate or its oxidized products in acidic DSMZ 269 medium, thereby allowing *A. cryptum* DK6.1 to grow. On the other hand, following *A. cryptum* DKAP1.1 growth due to heterotrophic metabolism, it was likely that a significant fraction of the total organic carbon including glucose was not

present at inhibiting high concentration to affect the viability of *A. ferrooxidans* DK6.1. Inhibition of growth of *A. ferrooxidans* in the presence of 1.0 g/l glucose and alleviation of the glucose toxicity by an acidophilic heterotroph in co-culture was demonstrated by earlier authors (Marchand and Silverstein, 2003). In this chapter we have also demonstrated a positive growth effect for *A. ferrooxidans* DK6.1 caused by co-culturing in ferrous iron agar medium in the presence of *A. cryptum* DKAP1.1. It was shown by earlier authors that heterotrophic cells can enhance iron oxidation by *A. ferrooxidans* in aerobic condition, even if the heterotrophs are not actively metabolizing organic carbon (Marchand and Silverstein, 2003). The presence of heterotrophic *A. cryptum* DKAP1.1 culture in the agar plate was probably responsible for lowering the bio-available total organic carbon (which resulted from acid hydrolysis of agar) to a level that permitted enhancement of iron oxidation by *A. ferrooxidans* DK6.1. Acidophilic heterotrophic bacterium has been used by earlier authors to scavenge organic material from normal agar in order to facilitate colony formation of *A. ferrooxidans* (Butler and Kempton, 1987). This work has provided additional evidences in support of the synergism between obligately autotrophic *A. ferrooxidans* and obligately oligotrophic heterotroph *A. cryptum* perceived to occur in natural ecosystem.

4.5 CONCLUSION

1. Increase in the robustness of colony formation of *A. ferrooxidans* in ferrous iron agar plate in deliberate presence of *Acidiphilium cryptum* was observed. This suggested that the heterotroph plays an important role in the manifestation of colonies in ferrous iron agar plate.
2. The ability of *A. cryptum* to grow in elemental sulfur spent medium showed the oligotrophic nature of the organism and also provided the best possible explanation for the co-existence of *A. cryptum* with *A. ferrooxidans* culture.
3. Thiosulfate at the level of ≥ 0.7 mM was found to be toxic to the strains of *Acidiphilium cryptum*. However, this toxicity could be alleviated in presence of *A. ferrooxidans* which provides the additional support to the synergism that exists between *A. ferrooxidans* and *A. cryptum*.

4.6 REFERENCE

- Arkesteyn, G. J. M. W. and deBont, J. A. M. (1980). *Thiobacillus asidophilus*: a study of its presence in *Thiobacillus ferrooxidans* cultures. *Can. J. Microbiol.* **26**: 1057-1065.
- Borichewski, R. M. (1967). Keto acids as growth-limiting factors in autotrophic bacteria that share their habitat. *Annu. Rev. Microbiol.* **38**: 265-292.
- Butler, B. J. and Kempton, A. G. (1987). Growth of *Thiobacillus ferrooxidans* on solid media containing heterotrophic bacteria. *J. Indust. Microbiol.* **2**: 41-45.
- Chakraborty, R., Deb, C., Lohia, A. and Roy, P. (1997). Cloning and characterization of a high-copy-number novel insertion sequence from chemolithotrophic *Thiobacillus ferrooxidans*. *Plasmid* **38**(2): 129-134.
- Descostes, M, Vitorage, P. and Beaucaire, C. (2004). Pyrite dissolution in acidic media. *Geochim. Cosmochim. Acta.* **68** (22): 4559-4569.
- Druschel, G. and Borda, M. (2006). Comment on "pyrite dissolution in acidic media" by M. Descostes, P. Vitorage, and C. Beaucaire. *Geochim. Cosmochim. Acta.* **70**: 5246-5250.
- Frattini, C. J., Leduc, L. G. and Ferroni, G. D. (2000). Strain variability and the effects of organic compounds on the growth of the chemolithotrophic bacterium *Thiobacillus ferrooxidans*. *Antonie van Leeuwenhoek.* **77**: 57-64.
- Goldhaber, M. B. (1983). Experimental study of metastable sulfur oxyanion formation during pyrite oxidation at pH 6-9 and 30 degrees C. *American Journal of Science.* **283**: 193-217.
- Goldhaber, M. B. and Rey, R. L. (1977). Experimental study of pyrite oxidation at pH 5-9.5; Implications for formation of Roll-type Uranium deposits. *AAPG Bull.* **61**: DOI: 10.1306/C1EA44C8-16C9-11D7-8645000102C1865D.
- Guay, R. and Silver, M. (1975). *Thiobacillus acidophilus* sp. nov.; isolation and some physiological characteristics. *Can. J. Microbiol.* **21**: 281-288.
- Hallberg, K. B., Thomson, H. E. C., Boeselt, I. and Johnson, D. B. (2001). Aerobic and anaerobic sulfur metabolism by acidophilic bacteria. pp. 423-431. *In Biohydrometallurgy: Fundamentals, Technology and Sustainable Development Process Metallurgy 2001*. Edited by V. S. T. Ciminelli and O. Garcia Jr. Amsterdam: Elsevier.
- Harrison, A. P. Jr. (1981). *Acidiphilium cryptum* gen. nov., sp. nov., heterotrophic bacterium from acidic mineral environments. *Int. J. Syst. Bacteriol.* **31**: 327-332.
- Harrison, A. P. Jr. (1983). Genomic and physiological comparisons between heterotrophic thiobacilli and *Acidiphilium cryptum*, *Thiobacillus versutus* sp. nov., and *Thiobacillus acidophilus* nom. rev. *Int. J. Syst. Bacteriol.* **33**: 211-217.

- Harrison, A. P. Jr. (1984). The acidophilic thiobacilli and other acidophilic bacteria that share their habitat. *Annu. Rev. Microbiol.* **38**: 265-292.
- Harrison, A. P. Jr., Bruce, W., Jarvis, T. and Johnson, J. L. (1980). Heterotrophic Bacteria from Cultures of Autotrophic *Thiobacillus ferrooxidans*: Relationships as Studied by Means of Deoxyribonucleic Acid Homology. *J. Bact.* **143**: 448-454.
- Hiraishi, A., Nagashima, K. V. P., Matura K., Shimada, K., Takaichi, K., Wokao, N., and Katayama, Y.(1998). Phylogeny and photosynthetic features of *Thiobacillus acidophilus* and related acidophilic bacteria: its transfer to the genus *Acidiphilium* as *Acidiphilium acidophilum* comb. nov. *Int. J. Syst. Bacteriol.* **48**: 1389-1398.
- Johnson, D.B. (1995). Selective solid media for isolating and enumerating acidophilic bacteria. *J. Microbiol. Method.* **23**: 205-218.
- Jones, J. G., Davison, W. and Gardener, S. (1984). Iron reduction by bacteria: range of organisms involved and metals reduced. *FEMS. Microbiol. Lett.* **21**: 133-136.
- Lane, D. J. (1991). 16S/23S rRNA sequencing. pp. 115-175. *In Nucleic Acid Techniques in Bacterial Systematics.* (ed.) E. Stackebrandt & M. Goodfellow. Chichester: Wiley.
- Lazaroff, N., Sigal, W. and Wasserman, A. (1982). Iron oxidation and precipitation of ferric hydroxysulfates by resting *Thiobacillus ferrooxidans* cells. *Appl. Environ. Microbiol.* **43**: 924-938.
- Lobos, J., Chisolm, T. E., Bopp, L. H. and Holmes, D. S. (1986). *Acidiphilium organovororum* sp. nov., an acidophilic heterotroph isolated from a *Thiobacillus ferrooxidans* culture. *Int. J. Syst. Bacteriol.* **36**: 39-144.
- Luther, G. W. III. (1987). Pyrite oxidation and reduction: molecular orbital theory considerations. *Geochim. Cosmochim. Acta.* **51**: 3193-3199.
- Marchand, E. A. and Silverstein, J. (2003). The Role of enhanced heterotrophic bacterial growth on iron oxidation by *Acidithiobacillus ferrooxidans*. *Geomicrobiol. J.* **20**: 231-244.
- Paiment, A., Leduc, L. G. and Ferroni, G. D. (2001). The effect of the facultative chemolithotrophic bacterium *Thiobacillus acidophilus* on the leaching of low-grade Cu-Ni sulfide ore by *Thiobacillus ferrooxidans*. *Geomicrobiol. J.* **18**(2): 157-165.
- Palumbo, S. A. and Alford, J. H. (1970). Inhibitory action of tetrathionate enrichment broth. *Appl. Microbiol.* **20**: 970-976.
- Parker, D. J. and Allison, W. S. (1969). The mechanism of inactivation of glyceraldehydes 3-phosphate dehydrogenase by tetrathionate, *o*-iodosobenzoate, and iodine monochloride. *J. Biol. Chem.* **244**: 180-189.
- Peccia, J., Marchand, E. A., Silverstein, J. and Hernandez, M. (2000). Development and application of small-subunit rRNA probes for assessment of selected *Thiobacillus*

- species and members of the genus *Acidiphilium*. *Appl. Environ. Microbiol.* **66**(7): 3065-3072.
- Postgate, J. R. (1963). The examination of sulfur auxotrophs: a warning. *J. Gen. Microbiol.* **30**: 481-484.
- Pronk, J. T. and Johnson, D. B. (1992). Oxidation and reduction of iron by acidophilic bacteria. *Geomicrobiol. J.* **10**: 153-171
- Pronk, J. T., Meesters, P. J. W., van Dijken, J. P., Bos, P. and Kuenen, J. G. (1990). Heterotrophic growth of *Thiobacillus acidophilus* in batch and chemostat cultures. *Arch. Microbiol.* **153**: 392-398.
- Rimstidt, J. D., and Vaughan, D. J. (2003). Pyrite oxidation: a state-of-the-art assessment of the reaction mechanism. *Geochim. Cosmochim. Acta.* **67**(5): 873-880.
- Rohwerdert, T., and Sand, W. (2003). The sulfane sulfur of persulfides is the actual substrate of the sulfur-oxidizing enzymes from *Acidithiobacillus* and *Acidiphilium* spp. *Microbiol.* **149**: 1699-1709.
- Roy, A. B. and Trudinger, P. A. (1970). The biochemistry of inorganic compounds of sulfur. University Press, Cambridge, pp.400.
- Schippers, A, Jozsa, P. and Sand, W. (1996). Sulfur chemistry in bacterial leaching of pyrite. *Appl. Environ. Microbiol.* **62**: 3424-3431.
- Schippers, A. and Sand, W. (1999). Bacterial leaching of metal sulfides proceeds by two indirect mechanisms via thiosulfate or via polysulfides and sulfur. *Appl. Environ. Microbiol.* **65**: 319-321.
- Schnaltman, C. and Lundgren, D. G. (1965). Organic compounds in the spent medium of *Ferrobacillus ferrooxidans*. *Can. J. Microbiol.* **11**: 23-27.
- Studel, R., Holdt, G., Gobel, T. and Hazeu, W. (1987). Chromatographic separation of higher polythionates $S_nO_6^{2-}$ ($n=3...22$) and their detection in cultures of *Thiobacillus ferrooxidans*; molecular composition of bacterial sulfur excretions. *Angew. Chem. Int.* **26**: 151-153.
- Tabita, R. and Lundgren, D. G. (1971). Heterotrophic metabolism of chemolithotroph *Thiobacillus ferrooxidans*. *J. Bacteriol.* **108**: 334-342.
- Temple, K. L. and Colmer, A. R. (1951). The autotrophic oxidation of iron by a new bacterium: *Thiobacillus ferrooxidans*. *J. Bacteriol.* **62**: 605-611.
- Yong, X. and Schoonen, M. A. A. (1995). The stability of thiosulfate in the presence of pyrite in low-temperature aqueous solutions. *Geochim. Cosmochim. Acta.* **59**: 4605-4622(18).
- Zhang, Z., Schwartz, S., Wagner, L., and Miller, W. (2000). A greedy algorithm for aligning DNA sequences. *J. Comput. Biol.* **7**(1-2): 203-214.