

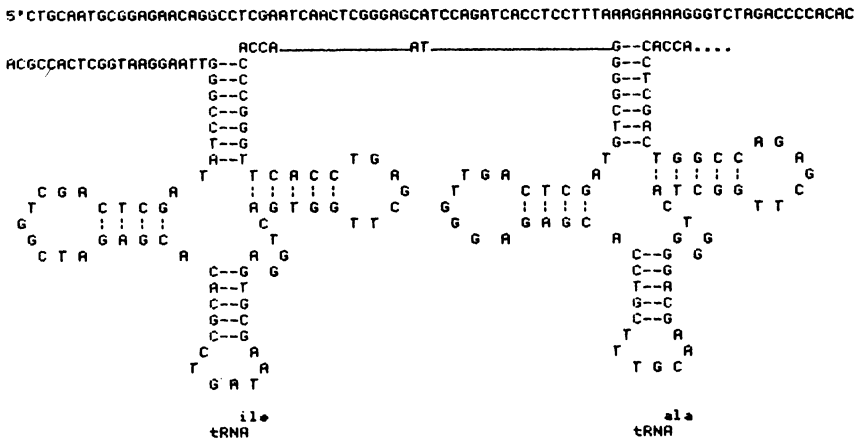
Sequence of two tRNA genes from a *Thiobacillus ferrooxidans* ribosomal operon

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**SEQUENCE ORIGIN:** Chromosomal DNA was isolated from *Thiobacillus ferrooxidans* strain A4 and a genomic library was made in pUC19. The screening was done using homologous tRNA-<sup>32</sup>P as a probe. The clone with the highest hybridization signal was mapped with restriction enzymes and subcloned in M13mp18/mp19 for sequencing analysis.



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CAGGGATGAGAGGTCCGCAAGAGCGCGGTCGCTGGGGAGCAGTAGAGAGGGGGTTTAGGGTTTAGTGTGTAGTGCTGA
ATAACCTTTAGGGATGAGSAGTAGGGTGTATTGATCACTGCCCACTCGGAGATTGAGTGGTTAGTTGTCTTTGAC
AGTTGAGGAGSAGGAGGCCATGTTTGTATCTTGGCGATGGCAGACCTCCAGGTGGATGCTTGGGGATATATGGTCAAGTGA
ATHAGGGCATACTGGTGGATGCCCTTGGCAGAGACAGGCGATGAGGACSTGGATACCTCGCAAGAGCCTC 3'
    
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**FEATURES:** No intervening sequences split coding regions. Terminal 3'-CCA sequences are gene encoded. A two-base spacer separates the tRNA gene sequences.

**COMMENTS:** Preliminary hybridization and sequence data indicate that this two tRNA genes are included in a ribosomal operon having a structural arrangement similar to those of rrnD from *E. coli* (1) or *B. subtilis* (2). Both tRNAs present high homology to those from *E. coli* and *B. subtilis* (3) (80.5% and 89.6% for tRNA<sup>Leu</sup>, and 82.9% and 84.2% for tRNA<sup>Ala</sup> respectively).

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