

Nucleotide sequence of two tRNA^{Arg}-tRNA^{Asp} tandem genes linked to duplicated *UBC* genes in *Saccharomyces cerevisiae*

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We have recently cloned the yeast genes *UBC4* and *UBC5* which are closely related in sequence and encode virtually identical ubiquitin-conjugating enzymes (1). The chromosomal location of these genes is unlinked, but interestingly, both genes are linked at their 3'-region to tRNA^{Arg}-tRNA^{Asp} tandem genes. In both cases tRNA genes are encoded on the opposite strand relative to *UBC* genes. The similarity of this gene arrangement might reflect a joint duplication of *UBC* and tRNA genes or alternatively, a *UBC* gene duplication mediated by repetitive tRNA loci.

Here we present the nucleotide sequences of both tRNA^{Arg}-tRNA^{Asp} tandem genes. The sequences given below exactly join the complementary strand of the published *UBC4* and *UBC5* sequences (1). The nucleotide sequences of tRNA^{Arg} and tRNA^{Asp} genes (underlined) correspond to the known primary structure of tRNA₃^{Arg} (2) and tRNA^{Asp} (3). In *Saccharomyces cerevisiae* tRNA^{Arg}-tRNA^{Asp} tandem genes have been identified previously (4). The sequences of the coding regions and a 10 bp spacer were found to be identical to the tRNA genes reported

here. Further similarities were restricted to nucleotides immediately preceding and following the coding sequences indicating that the tRNA genes described here represent novel copies of tRNA^{Arg}-tRNA^{Asp} tandem genes.

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| CAAAACAAA TGAATTGTCA TACAAGATA ATACTGGAA CACATCCATC TTTAGAGGAA ACATGTTTC GTTATTCAT AAAACCTTC ACAAATAGTA | 100 |
| <u>GCTCGCGTGG CGTAATGGCA ACGCGTCTGA CTTCTAATCA GAAGATTATG GGTCGACCC CCATCGTGAG TGCTTTGTTT CTTCCGTGAT AGTTTAATGG</u> | 200 |
| <u>TCAGAACATGGG CGCTTGTCGC GTGCCAGATC GGGGTTCAAT TCCCCGTGCG GGAGAATTTT TTGGCTACTC CTGCGAGTATT CTTCTGCCTC CTTAGTACAG</u> | 300 |
| TGTAATGCTC CTCAGAACATT TTGCCAAACA GAGAACCACT CATTGATCGT TGATTACATA TACTAGCTAT GGCGTTGAAC TCAAAATTT AAATAGGCTT | 400 |
| CACCCCCAGAA ATGATGTTAA TGACAATAGT AATTAACTTA CAGATAGACT AGAAAATGAAA AAGGTACGGT TAACGTTGAC GCTCCCTCGA CAAGTTAGGT | 500 |
| CAAATATGAA CCACAACTGA AATATATGCG GGATATACCC GCTTGCTGCA TGTAACTAAG CAAAAATTCT CCATACTGTT CGAGGAAAAA AATGTTCATG | 600 |
| AATGTCCTCT GAATAAATTAA AATTAAGGTT TACGATTGCA GTTGTGTTGAA AAAAAGAAAA AGTTCCTCTG TACTGTTGCC AACAAATAGTC TTATATCTTA | 700 |
| TATCAGTAAA TCTATGTT | |

tRNA^{Arg}-tRNA^{Asp} tandem genes linked to *UBC4*

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|------------------------------------------------------------------------------------------------------------------------|-----|
| TCATATTAGG CTCATTATGA GAAAAGGTTG GTGAGAAATT TAGCATATT TAAACAATCA TTTAGCTAGA TGAATCAGAA TGTTTTCCG ACAAATAGTA | 100 |
| <u>GCTCGCGTGG CGTAATGGCA ACGCGTCTGA CTTCTAATCA GAAGATTATG GGTCGACCC CCATCGTGAG TGCTTTGTTT CTTCCGTGAT AGTTTAATGG</u> | 200 |
| <u>TCAGAACATGGG CGCTTGTCGC GTGCCAGATC GGGGTTCAAT TCCCCGTGCG GGAGAATTTT TTGAAATTGA TACTAGCAGG TACAGGAGAA AGACTAAATA</u> | 300 |
| ATAGAAAGCA AGGTACGTGA TGGGAAT | |

tRNA^{Arg}-tRNA^{Asp} tandem genes linked to *UBC5*