

Create a CDS feature without a translation qualifier

Since version 2.1.0 of WIPO Sequence, the 'Automatically add a translation qualifier...' checkbox is selected by default. This means that by default, a translation qualifier containing the amino acid conceptual translation value as well as another amino acid sequence (if it meets the minimum length requirement) and associated protein ID will be generated automatically. If you do not want to include the translation or protein id qualifiers for one particular sequence, then follow the guide below.

Step-by-step guide

If you have not already inserted a CDS feature:

1. Uncheck the 'Automatically add a translation qualifier value' checkbox.
2. Proceed with creating the CDS feature. No translation or protein_id qualifier will be added now.
3. Consider whether adding a pseudogene or pseudo qualifier is warranted.

If you have already inserted a CDS feature:

1. Uncheck the 'Automatically add a translation qualifier value' checkbox.
2. Delete the translation qualifier.
3. Delete the protein id qualifier as well as the separate sequence which is identified by the SEQ ID included as the protein id qualifier value.
4. Consider whether adding a pseudogene or pseudo qualifier is warranted.



According to the recommendations provided in WIPO Standard ST.26, there are no mandatory qualifiers for a CDS feature. The optional qualifiers comprise:

- allele
- circular_RNA
- codon_start
- EC_number
- exception
- function
- gene
- gene_synonym
- map
- note
- number
- operon
- product
- protein_id
- pseudo
- pseudogene
- ribosomal_slippage
- standard_name
- translation
- transl_except
- transl_table
- trans_splicing

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