

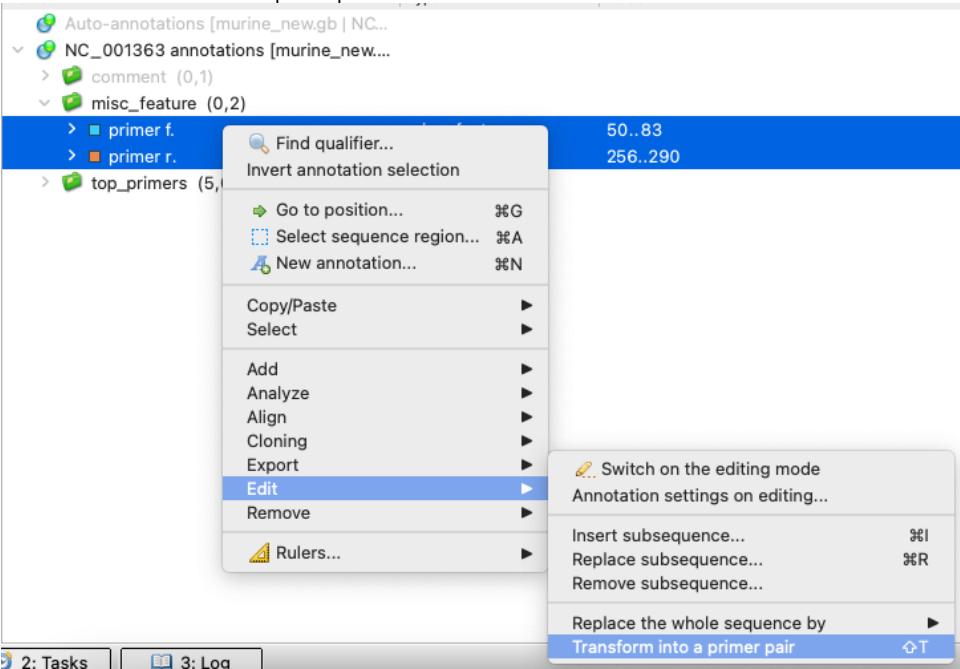
Transform into a primer pair

UGENE has certain primers format. It allows one to use some features, which are available only if this format is being recognized. Examples of these features:

- Create PCR Fragment,
- [Primer-BLAST](#).

If you have a primer pair, which doesn't fit to this format, these features will be unavailable for you. To fix this issue:

1. Select your own left and right primers.
2. Click right mouse button Edit Transform into a primer pair.



After these steps are performed, a new primer pair, fit to UGENE format, will appear:

Name	Type	Value
Auto-annotations [murine_new.gb NC...		
NC_001363 annotations [murine_new....		
comment (0,1)		
misc_feature (0,2)		
primer f.	misc_feature	50..83
primer r.	misc_feature	256..290
top_primers (6,0)		
pair 1 (0,2)		
pair 2 (0,2)		
pair 3 (0,2)		
pair 4 (0,2)		
pair 5 (0,2)		
pair 6 (0,2)		
top_primers	Primer	50..83
top_primers	Primer	complement(256..290)