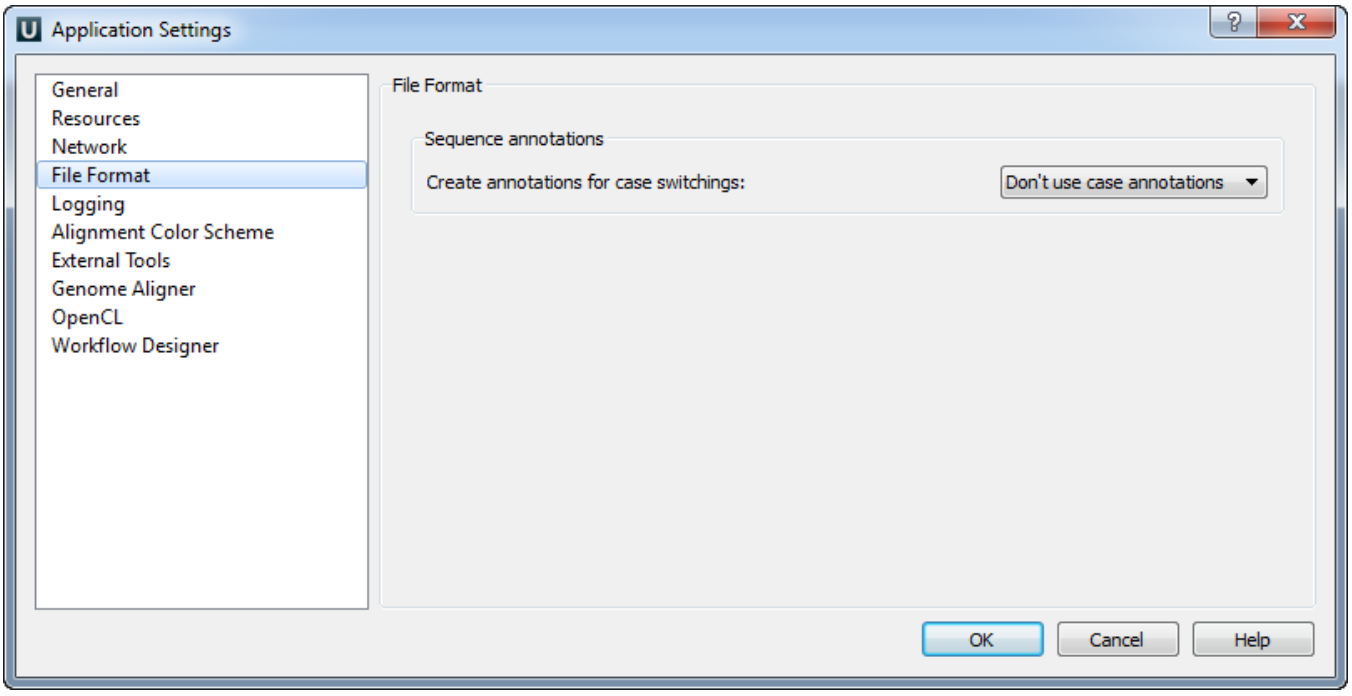
# File Format




The *Sequence Annotations* settings allows to use upper/lower case annotations during the file reading process.

Format options:

1. *Don't use case annotations* (default mode) — usual sequence reading and writing.
2. *Use lower case annotation* — sequences are read and annotations with names *lower_case* are added. When these sequences are written to file then the case becomes like original the file case (the case is saved).
3. *Use upper case annotation* — there is a similar behavior but with "upper_case" annotations.