

Known Issues and trouble-shooting

The following list of known issues have been reported to WIPO for versions of WIPO Sequence beginning at version 2.1.0. Version 1.x.x generates sequences according to outdated versions of the Standard and should not be used. Version 2.0.0 was superseded by version 2.1.0 before WIPO ST.26 entered general use and is not supported, but is known to contain some of the bugs described below. **The latest version is version 2.3.0 and this is the version we recommend that everyone uses.**

The following list of known issues have been reported to WIPO. Please only report to WIPO at wiposequence.support@wipo.int any issues which are not provided below.

Key	Description	Components	Affected Version	Resolved Version
ST26T-2893	Cannot cut and paste into or from WIPO Sequence version 2.0.0 for Mac OSX distribution	WIPO Sequence (Mac OSX only)	2.0.0	2.2.0
ST26T-2888	If there is a TAB character in the invention title recorded in an ST.25 sequence listing, during import two Invention titles are created	WIPO Sequence	2.0.0	2.1.0
ST26T-2889	Translation qualifier value is not automatically created when user manually creates it	WIPO Sequence	2.1.0	2.2.0
ST26T-2780	Terms of Use pop-up is shown to user twice in some instances	WIPO Sequence	2.1.0	-
ST26T-2882	WIPO Sequence cannot be deployed on Citrix Terminal Server	WIPO Sequence	2.1.0	-
ST26T-2873	TAB character is not detected in InventionTitle, ApplicantName, InventorName, ApplicantNameLatin, InventorNameLatin during validation process	WIPO Sequence	2.1.0	-
ST26T-2870	JSON Key rather than the value of the CDS location is shown on banner is shown on the banner when the location is too short	WIPO Sequence	2.1.0	2.2.0
ST26T-2762	XML validation does not enforce validation checks for the DOCTYPE declaration	WIPO Sequence	2.0.0	2.1.0
ST26T-2885	ncRNA qualifier added when 'n' feature key is imported from an ST.25 sequence listing	WIPO Sequence	2.1.0	2.3.0
ST26T-2872	For imported amino acid sequences from a ST.25 sequence listing, mol_type is not automatically set to 'protein'	WIPO Sequence	2.1.0	2.2.0
ST26T-2883	Wrong numbering provided during XML validation process with >5k sequences and <10k sequences on pop-up window	WIPO Sequence (Linux only)	2.1.0	-
ST26T-2876	No warning message when DNA/RNA sequence includes "n" symbols - should indicate that the default value should be assumed	WIPO Sequence	2.1.0	2.3.0
ST26T-2877	When updating manually created nucleotide sequence containing a single "n" symbol - feature location for source feature can be wrong Workaround: Uncheck "Intentionally skipped" checkbox and add the organism. The location of source feature will become correct.	WIPO Sequence	2.1.0	2.3.0
ST26T-2900	Event Viewer errors when launching WIPO Sequence	WIPO Sequence	2.1.0	-
ST26T-2902	In a project, the PDF resulting from clicking the Print button contains a wrong priority date and filing date (it reduces the day by 1).	WIPO Sequence	2.1.0	-
ST26T-2905	Non Latin characters not allowed in non English qualifier value. Error returned: "The non English free text must contain only Unicode characters, except the reserved or controlled characters"	WIPO Sequence	2.1.0	2.1.1
ST26T-2906	After ST.25 import, an error is thrown indicating that the qualifier ID is not unique Workaround: Delete the feature which includes a duplicate qualifier ID. Recreate the feature at the same location but <u>do not add</u> the qualifier before saving. Next, edit the feature to add the relevant qualifier and update the feature.	WIPO Sequence	2.1.0	2.1.2
ST26T-2907	Sporadically sequences are generated without a feature table Update: ST26T-2907: work-around	WIPO Sequence	2.1.0	2.1.2
ST26T-2910	Auto-update does not download the latest binary	WIPO Sequence	2.1.0	2.1.2
ST26T-2912	When using the organism dropdown when editing an organism, the auto-complete can be difficult to select the appropriate organism	WIPO Sequence	2.1.0	2.2.0
ST26T-2894	Importing ST.25: Newlines being stripped out of qualifier value are not being replaced with spaces	WIPO Sequence	2.1.0	2.2.0
ST26T-2922	Import message for skipped sequence indicates that the INSDSeq_division element has been set to 'PAT'	WIPO Sequence	2.1.2	2.3.0
ST26T-2913	Chinese characters are not shown when printing a sequence listing	WIPO Sequence	2.1.0	2.2.0

ST26T-2928	Missing full list of features when printing sequences from project details page	WIPO Sequence	2.1.2	2.3.0
ST26T-2917	Wrong column heading for invention title provided when printing project	WIPO Sequence	2.1.2	2.3.0
ST26T-2851	Language Dependent Qualifiers page cannot be loaded when project contains more than ~12.700 language dependent qualifiers	WIPO Sequence	2.0.0	
ST26T-2895	New person/organization pop-up: selecting 'No' can freeze the tool sporadically	WIPO Sequence	2.0.0	
ST26T-2946	Instances where SEQ ID assigned after import is not consecutive	WIPO Sequence	2.2.0	
ST26T-2944	No error is thrown if the attribute dtdVersion is not the latest version of the DTD during XML validation	WIPO Sequence	2.2.0	2.3.0
ST26T-2902	When printing a Project, output PDF file contains wrong information, including incorrect priority date	WIPO Sequence	2.2.0	2.3.0
ST26T-2932	'Synthetic construct' (with capital letter S) behavior ('Synthetic construct' is not part of the pre-defined organism names)	WIPO Sequence	2.2.0	2.3.0
ST26T-2974	Generic error on import banner shown when importing ST.26 file with a field containing more than 2 pre-defined entities	WIPO Sequence	2.2.0	
ST26T-2990	Cannot add a circular_RNA qualifier to a feature	WIPO Sequence	2.2.0	2.3.0
ST26T-2948	The Output PDF file is missing the "Inventor name:" label	WIPO Sequence	2.2.0	2.3.0
ST26T-3003	'Frequency' not being treated as a language dependent free text qualifier	WIPO Sequence	2.2.0	-
ST26T-2993	Wrong verification rule applied for circular_RNA qualifier	WIPO Sequence	2.2.0	-
ST26T-2972	No error thrown when DOCTYPE declaration include random characters at the end	WIPO Sequence	2.2.0	-
ST26T-2974	Generic error on import banner shown when importing ST.26 file with a field containing more than 2 pre-defined entities	WIPO Sequence	2.2.0	-
ST26T-2895	When pop-up for new person or organization appears, selecting 'no' can freeze the tool	WIPO Sequence	2.1.0	-
ST26T-2996	Some sequences still being generated without a feature table	WIPO Sequence	2.2.0	2.3.0
ST26T-3027	Qualifier mol_type and organism values might disappear from sequence after SL generation	WIPO Sequence	2.2.0	2.3.0
ST26T-3029	Qualifier bound_moiety is considered as mandatory for oriT feature	WIPO Sequence	2.2.0	2.3.0
ST26T-3034	Error when InventionTitle contains more than one '	WIPO Sequence	2.2.0	2.3.0
ST26T-3052	Verification rule QV_48 not always triggering	WIPO Sequence	2.3.0	
ST26T-3067	Bulk Edit only affects the first 20 sequences of a range	WIPO Sequence	2.3.0	
ST26T-3089	Cannot bulk delete more than 20 sequences at once	WIPO Sequence	2.3.0	