

File columns

		Merge handling of table			Column usage		
		Insert	Merge	Update	Attach	Decisive	Key
Pos.	Name	Table	Column	Example	Transformed		
0	Beleg	CollectionSpecimen.	AccessionNumber	KR-M-0003370			
1	Gattung	IdentificationUnit_2.	LastIdentificationCache	Uromyces			
2	Art	IdentificationUnit_2.	LastIdentificationCache	psi	_psi		
			Prefix: _				
3	Wirtsgattung	IdentificationUnit_1.	LastIdentificationCache	Euphorbia			
4	Wirtsart	IdentificationUnit_1.	LastIdentificationCache	cyparissias	_cyparissias		
			Prefix: _				
5	well	IdentificationUnitAnalysisMethodParameter_2_1_2_12.	Value	h12			
		IdentificationUnitAnalysisMethodParameter_2_1_3_12.	Value	h12			
6	pcr-for-primer-name	IdentificationUnitAnalysisMethodParameter_2_1_2_2.	Value	ITS5u			
		IdentificationUnitAnalysisMethodParameter_2_1_3_2.	Value	ITS5u			
7	pcr-for-primer-seq	IdentificationUnitAnalysisMethodParameter_2_1_2_3.	Value	CAAGGTTTCTGTAGGTG			
		IdentificationUnitAnalysisMethodParameter_2_1_3_3.	Value	CAAGGTTTCTGTAGGTG			
8	pcr_rev-primer-name	IdentificationUnitAnalysisMethodParameter_2_1_2_4.	Value	ITS4u			
		IdentificationUnitAnalysisMethodParameter_2_1_3_4.	Value	ITS4u			
9	pcr-rev-primer-seq	IdentificationUnitAnalysisMethodParameter_2_1_2_5.	Value	GGCTTTTCCCTCTTCAT			
		IdentificationUnitAnalysisMethodParameter_2_1_3_5.	Value	GGCTTTTCCCTCTTCAT			
10	seq-for-primer-name	IdentificationUnitAnalysisMethodParameter_2_1_2_6.	Value	ITS5u			
11	seq-for-primer-seq	IdentificationUnitAnalysisMethodParameter_2_1_2_7.	Value	CAAGGTTTCTGTAGGTG			

12	seq_rev-primer-name	IdentificationUnitAnalysisMethodParameter_2_1_3_6.	Value	ITS4u
13	seq-rev-primer-seq	IdentificationUnitAnalysisMethodParameter_2_1_3_7.	Value	GGCTTTTCCCTCTTCAT
14	Region	IdentificationUnitAnalysis_2_1.ber	AnalysisNumber	ITS
15	Sequencing-Lab	IdentificationUnitAnalysisMethodParameter_2_1_2_8.	Value	Thines-Labor/Uni Tübingen M. Lutz
		IdentificationUnitAnalysisMethodParameter_2_1_3_8.	Value	Thines-Labor/Uni Tübingen M. Lutz
16	Edited by	not imported		
17	Annotations	not imported		
18	Problems	not imported		
19	Barcode-Sequenz	IdentificationUnitAnalysis_2_1.	AnalysisID	AGAAGGATCATTATTTAAACCAATGAGTGCACTTTA TTGTGGCTCAAACCTCTTTATCTCACCCCTTTTTTTT TGTTTGTAACACAAGTTGAATTAAGAATGTAAAAAC CTTTTATTTTAAAAATAACTTTTAACAATGGATCTCTT GGCTCTCATATCGATGAAGAACACAGTGAAATGTGA TAAGTAATGTGAATTGCAGAATTCAGTGAATCATTGA ATCTTTGAACGCACCTTGCACCTTTTGGTATTCCAAA AGGTACGCCTGTTTGAGTGTCATGAAAATCTCTCAT CAAATTAATTTTGGTGGATGTTGAGTGCTGCTGCTA TCTAGCTCACTTTAAATATATAAGTTCTTCTTTATGT TGTTGGATTGACTTGGTGAATATTTAACTTTTTTTTC ATCAAGGAAAGTAGAAATACTTGCCAACAATTTTTAA AATAAGGACTCCTAAAAATATTTATTTATAATCTTTTT TAAGACCTCAAATCAGGTGGGACTACCTGCTGAACT TAAGCATATCAATAAGCAGAGGAAAAAGAACTA
		IdentificationUnitAnalysisMethod_2_1_2.	MethodID	AGAAGGATCATTATTTAAACCAATGAGTGCACTTTA TTGTGGCTCAAACCTCTTTATCTCACCCCTTTTTTTT TGTTTGTAACACAAGTTGAATTAAGAATGTAAAAAC CTTTTATTTTAAAAATAACTTTTAACAATGGATCTCTT GGCTCTCATATCGATGAAGAACACAGTGAAATGTGA TAAGTAATGTGAATTGCAGAATTCAGTGAATCATTGA ATCTTTGAACGCACCTTGCACCTTTTGGTATTCCAAA AGGTACGCCTGTTTGAGTGTCATGAAAATCTCTCAT CAAATTAATTTTGGTGGATGTTGAGTGCTGCTGCTA TCTAGCTCACTTTAAATATATAAGTTCTTCTTTATGT TGTTGGATTGACTTGGTGAATATTTAACTTTTTTTTC ATCAAGGAAAGTAGAAATACTTGCCAACAATTTTTAA AATAAGGACTCCTAAAAATATTTATTTATAATCTTTTT TAAGACCTCAAATCAGGTGGGACTACCTGCTGAACT TAAGCATATCAATAAGCAGAGGAAAAAGAACTA
Transformation Filter				
s: Import fixed value: 16				
If column 19 ≠				
And If column 21 ≠				

IdentificationUnitAnalysisMethod thod_2_1_2.	MethodMarker	AGAAGGATCATTATTTAAACCAATGAGTGCACCTTTA TTGTGGCTCAAACCTCTTTATCTCACCCCTTTTTTTT TGTTTGTAACACAAGTTGAATTAAGAATGTAAAAAAC CTTTTATTTTGAAAATAACTTTTAACAATGGATCTCTT GGCTCTCATATCGATGAAGAACACAGTGAAATGTGA TAAGTAATGTGAATTGCAGAATTCAGTGAATCATTGA ATCTTTGAACGCACCTTGCACCTTTTGGTATTCCAAA AGGTACGCCTGTTTGAGTGTCATGAAAATCTCTCAT CAAATTAATTTTTGGTGGATGTTGAGTGCTGCTGCTA TCTAGCTCACTTTAAATATATAAGTTCTTCCTTTATGT TGTTGGATTGACTTGGTGAATATTTAACTTTTTTTTC ATCAAGGAAAGTAGAAATACTTGCCAACAATTTTTAA AATAAGGACTCCTAAAAATATTTATTTATAATCTTTTT TAAGACCTCAAATCAGGTGGGACTACCTGCTGAACT TAAGCATATCAATAAGCAGAGGAAAAGAACTA
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Transformation

Filter

S:

Import fixed value: 4

If column 19 ≠

And

If column 21 ≠

IdentificationUnitAnalysisMethod thod_2_1_3.	MethodID	AGAAGGATCATTATTTAAACCAATGAGTGCACCTTTA TTGTGGCTCAAACCTCTTTATCTCACCCCTTTTTTTT TGTTTGTAACACAAGTTGAATTAAGAATGTAAAAAAC CTTTTATTTTGAAAATAACTTTTAACAATGGATCTCTT GGCTCTCATATCGATGAAGAACACAGTGAAATGTGA TAAGTAATGTGAATTGCAGAATTCAGTGAATCATTGA ATCTTTGAACGCACCTTGCACCTTTTGGTATTCCAAA AGGTACGCCTGTTTGAGTGTCATGAAAATCTCTCAT CAAATTAATTTTTGGTGGATGTTGAGTGCTGCTGCTA TCTAGCTCACTTTAAATATATAAGTTCTTCCTTTATGT TGTTGGATTGACTTGGTGAATATTTAACTTTTTTTTC ATCAAGGAAAGTAGAAATACTTGCCAACAATTTTTAA AATAAGGACTCCTAAAAATATTTATTTATAATCTTTTT TAAGACCTCAAATCAGGTGGGACTACCTGCTGAACT TAAGCATATCAATAAGCAGAGGAAAAGAACTA
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Transformation

Filter

S:

Import fixed value: 16

If column 19 ≠

And

If column 23 ≠

IdentificationUnitAnalysisMethod thod_2_1_3.	MethodMarker	AGAAGGATCATTATTTAAACCAATGAGTGCACCTTTA TTGTGGCTCAAACCTCTTTATCTCACCCCTTTTTTTT TGTTTGTAACACAAGTTGAATTAAGAATGTAAAAAAC CTTTTATTTTGAAAATAACTTTTAACAATGGATCTCTT GGCTCTCATATCGATGAAGAACACAGTGAAATGTGA TAAGTAATGTGAATTGCAGAATTCAGTGAATCATTGA ATCTTTGAACGCACCTTGCACCTTTTGGTATTCCAAA AGGTACGCCTGTTTGAGTGTCATGAAAATCTCTCAT CAAATTAATTTTTGGTGGATGTTGAGTGCTGCTGCTA TCTAGCTCACTTTAAATATATAAGTTCTTCCTTTATGT TGTTGGATTGACTTGGTGAATATTTAACTTTTTTTTC ATCAAGGAAAGTAGAAATACTTGCCAACAATTTTTAA AATAAGGACTCCTAAAAATATTTATTTATAATCTTTTT TAAGACCTCAAATCAGGTGGGACTACCTGCTGAACT TAAGCATATCAATAAGCAGAGGAAAAGAACTA
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Transformation

Filter

S:

Import fixed value: 5

If column 19 ≠

And

If column 23 ≠

20	Rohdatenfile fwd 1	not imported
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21	Rohdatenfile fwd 2	
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IdentificationUnitAnalysisMethod thodParameter_2_1_2_9.	Value
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Transformation

S:

Split at	Pos.	Dir.
.	1	↩
Translate	Into	

				<div> <div>ab1AB1</div> <div>scfSCF</div> <div>SCFSCF</div> </div>
	IdentificationUnitAnalysisMethodParameter_2_1_2_10.	Value		
	IdentificationUnitAnalysisMethodParameter_2_1_2_11.	Value		P:\GFBio\GBOL_Rostpilze\3_Archived\Traces
		Transformations:	Filter	
			Import fixed value:	P:\GFBio\GBOL_Rostpilze\3_Archived\Traces
			If column 19 ≠	
	IdentificationUnitAnalysisMethodParameter_2_1_2_11.	Value		
22	Rohdatenfile rev 1			not imported
23	Rohdatenfile rev 2			
	IdentificationUnitAnalysisMethodParameter_2_1_3_9.	Value		
		Transformations:	Split at	Pos. Dir.
			.	1 ↩
			Translate	Into
			ab1	AB1
			scf	SCF
			SCF	SCF
	IdentificationUnitAnalysisMethodParameter_2_1_3_10.	Value		
	IdentificationUnitAnalysisMethodParameter_2_1_3_11.	Value		P:\GFBio\GBOL_Rostpilze\3_Archived\Traces
		Transformations:	Filter	
			Import fixed value:	P:\GFBio\GBOL_Rostpilze\3_Archived\Traces
			If column 19 ≠	
	IdentificationUnitAnalysisMethodParameter_2_1_3_11.	Value		
24	Withhold			
	CollectionSpecimen.	DataWithholdingReason		
25	Interne Bemerkung			not imported
26	Dateiname			not imported
27	Identification UnitID			not imported

Interface settings

Table	Table alias	Column	Value
CollectionProject	CollectionProject_1	ProjectID	940
IdentificationUnit	IdentificationUnit_1	TaxonomicGroup	plant
	IdentificationUnit_2	TaxonomicGroup	fungus
IdentificationUnitAnalysis	IdentificationUnitAnalysis_2_1	AnalysisID	161
IdentificationUnitAnalysisMethodParameter	IdentificationUnitAnalysisMethodParameter_2_1_2_1	ParameterID	73
	IdentificationUnitAnalysisMethodParameter_2_1_2_1	Value	forward
	IdentificationUnitAnalysisMethodParameter_2_1_2_2	ParameterID	63
	IdentificationUnitAnalysisMethodParameter_2_1_2_3	ParameterID	64

IdentificationUnitAnalysisMethodParameter_2_1_2_4	ParameterID	65
IdentificationUnitAnalysisMethodParameter_2_1_2_5	ParameterID	66
IdentificationUnitAnalysisMethodParameter_2_1_2_6	ParameterID	91
IdentificationUnitAnalysisMethodParameter_2_1_2_7	ParameterID	92
IdentificationUnitAnalysisMethodParameter_2_1_2_8	ParameterID	72
IdentificationUnitAnalysisMethodParameter_2_1_2_9	ParameterID	81
IdentificationUnitAnalysisMethodParameter_2_1_2_10	ParameterID	74
IdentificationUnitAnalysisMethodParameter_2_1_2_11	ParameterID	75
IdentificationUnitAnalysisMethodParameter_2_1_2_12	ParameterID	85
IdentificationUnitAnalysisMethodParameter_2_1_3_1	ParameterID	73
IdentificationUnitAnalysisMethodParameter_2_1_3_1	Value	reverse
IdentificationUnitAnalysisMethodParameter_2_1_3_2	ParameterID	63
IdentificationUnitAnalysisMethodParameter_2_1_3_3	ParameterID	64
IdentificationUnitAnalysisMethodParameter_2_1_3_4	ParameterID	65
IdentificationUnitAnalysisMethodParameter_2_1_3_5	ParameterID	66
IdentificationUnitAnalysisMethodParameter_2_1_3_6	ParameterID	91
IdentificationUnitAnalysisMethodParameter_2_1_3_7	ParameterID	92
IdentificationUnitAnalysisMethodParameter_2_1_3_8	ParameterID	72
IdentificationUnitAnalysisMethodParameter_2_1_3_9	ParameterID	81
IdentificationUnitAnalysisMethodParameter_2_1_3_10	ParameterID	74
IdentificationUnitAnalysisMethodParameter_2_1_3_11	ParameterID	75
IdentificationUnitAnalysisMethodParameter_2_1_3_12	ParameterID	85