

Schedule for import of tab-separated text files into DiversityCollection

Target within DiversityCollection: Specimen

Schedule version: 1

Database version: 02.05.1964

Lines: 4 - 111

First line contains column definition: ✓

Encoding: ANSI

Language: de

Attached to table: n

Column: AccessionNumber

Description: Import scheme to attach Barcodeanalyses from BOLD to GBOL-Malaise

File columns

| | | Merge handling of table | | | Column usage | |
|------|-----------------------|--|------------------|----------------|-------------------------|--------------|
| | | Insert | Merge | Update | Attach | Decisive Key |
| Pos. | Name | Table | Column | Example | Transformed | |
| 0 | id | CollectionProject_1. | ProjectID | | | |
| 1 | GBOL_Nr_numeric | | | not imported | | |
| 2 | GBOL_Nr | CollectionSpecimen. | Accession Number | GBOL-223959611 | | |
| | | IdentificationUnitAnalysisMethodParameter_1_1_8_6. | Value | GBOL-223959611 | For all: "GBOL-Malaise" | |
| 3 | | | | not imported | | |
| 4 | | | | not imported | | |
| 5 | Feldnummer | | | not imported | | |
| 6 | Institut | | | not imported | | |
| 7 | Sammlungs_Nr | | | not imported | | |
| 8 | Ordnung | | | not imported | | |
| 9 | Familie | IdentificationUnit_1. | Taxonomic Group | | For all: "animal" | |
| 10 | Taxon_Name | | | not imported | | |
| 11 | Erstbeschreiber | | | not imported | | |
| 12 | Klammer | | | not imported | | |
| 13 | Jahr | | | not imported | | |
| 14 | _species | | | not imported | | |
| 15 | Bemerkung_Taxonomie | | | not imported | | |
| 16 | Bestimmer_Name | | | not imported | | |
| 17 | Bestimmungs_Datum | | | not imported | | |
| 18 | est_det_date | | | not imported | | |
| 19 | Gewebetyp_Menge | | | not imported | | |
| 20 | Anzahl_Individuen | | | not imported | | |
| 21 | Fixierungsmethode | | | not imported | | |
| 22 | Entwicklungsstadium | | | not imported | | |
| 23 | Sex | | | not imported | | |
| 24 | Bemerkungen_zur_Probe | | | not imported | | |
| 25 | Fundort_ID | | | not imported | | |
| 26 | Fundortbeschreibung | | | not imported | | |
| 27 | Bundesland | | | not imported | | |
| 28 | Land | | | not imported | | |
| 29 | Gemeinde_oder_Region | | | not imported | | |
| 30 | Sammel_Datum | | | not imported | | |
| 31 | est_Sammel_Datum | | | not imported | | |
| 32 | Sammelmethode | | | not imported | | |
| 33 | Breitengrad | | | not imported | | |
| 34 | GKS_R | | | not imported | | |

Description : Import scheme for GBOL-Data (attaching Barcodeanalyse from BOLD to existing CollectionSpecimens)

| | | | |
|----|--|------------------|--|
| 35 | Laengengrad | | not imported |
| 36 | GKS_H | | not imported |
| 37 | Benutzte_Methode | | not imported |
| 38 | Genauigkeitsangabe | | not imported |
| 39 | Hoehe_Tiefe_m | | not imported |
| 40 | Habitat | | not imported |
| 41 | Sammler | | not imported |
| 42 | Sammelgenehmigung_Nummer | | not imported |
| 43 | Sammelgenehmigung_Behoerde | | not imported |
| 44 | Bemerkungen_Sammelplate | | not imported |
| 45 | plate | | not imported |
| 46 | well | | |
| | IdentificationUnitAnalysisMethod_1_1_8. | MethodID | For all: "Barcode" |
| | IdentificationUnitAnalysisMethod_1_1_8. | MethodMarker | For all: "9" |
| 47 | changed | | not imported |
| 48 | changed by | | not imported |
| 49 | deleted | | not imported |
| 50 | Sammeltabelle | | not imported |
| 51 | Einsender | | not imported |
| 52 | bezahlt | | not imported |
| 53 | Kontamination | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_8_4. | ParameterID | For all: "Barcode failure" |
| | IdentificationUnitAnalysisMethodParameter_1_1_8_4. | Value | |
| | | Transformations: | <div>Filter</div> <div>Import fixed value: yes</div> <div>If column 53 ≠</div> |
| | IdentificationUnitAnalysisMethodParameter_1_1_8_5. | Value | |
| 54 | DNA-Bank | | not imported |
| 55 | Dep.Acc.No./additional notes | | not imported |
| 56 | relation notes | | not imported |
| 57 | Ext_ID | | not imported |
| 58 | Reference_ID | | not imported |
| 59 | problems | | not imported |
| 60 | External_Source | | not imported |
| 61 | external_relation | | not imported |
| 62 | rel_Uri | | not imported |
| 63 | respons_Sequenz | | |
| | IdentificationUnitAnalysis_1_1. | ResponsibleName | BIOUG |
| 64 | No_Analysis | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_1_7. | ParameterID | For all: "Barcode Sequencing Sequence_id" |
| | IdentificationUnitAnalysisMethodParameter_1_1_1_7. | Value | 6916114 |
| | IdentificationUnitAnalysisMethodParameter_1_1_2_7. | ParameterID | For all: "Barcode Sequencing Sequence_id" |
| | IdentificationUnitAnalysisMethodParameter_1_1_2_7. | Value | 6916114 |
| | IdentificationUnitAnalysisMethodParameter_1_1_3_7. | ParameterID | For all: "Barcode Sequencing Sequence_id" |
| | IdentificationUnitAnalysisMethodParameter_1_1_3_7. | Value | 6916114 |
| | IdentificationUnitAnalysisMethodParameter_1_1_4_7. | ParameterID | For all: "Barcode Sequencing Sequence_id" |
| | IdentificationUnitAnalysisMethodParameter_1_1_4_7. | Value | 6916114 |
| | IdentificationUnitAnalysisMethodParameter_1_1_5_7. | ParameterID | For all: "Barcode Sequencing Sequence_id" |

Description : Import scheme for GBOL-Data (attaching Barcodeanalyse from BOLD to existing CollectionSpecimens)

| | IdentificationUnitAnalysisMethodParameter_1_1_5_7. | Value | 6916114 | | | | | | | | | | | | | | | | | | | | | | |
|----------------|--|------------------|---|---|------|------|---|---|---|----------------|--|--|------------|----|--|---|---------|--|---|---------|--|---------|--|--|--|
| | IdentificationUnitAnalysisMethodParameter_1_1_6_7. | ParameterID | | For all: "Barcode Sequencing Sequence_id" | | | | | | | | | | | | | | | | | | | | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_6_7. | Value | 6916114 | | | | | | | | | | | | | | | | | | | | | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_7_7. | ParameterID | | For all: "Barcode Sequencing Sequence_id" | | | | | | | | | | | | | | | | | | | | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_7_7. | Value | 6916114 | | | | | | | | | | | | | | | | | | | | | | |
| 65 | Result | | | | | | | | | | | | | | | | | | | | | | | | |
| | IdentificationUnitAnalysis_1_1. | AnalysisResult | ACATTATATTTTATATTTGGG[...] | | | | | | | | | | | | | | | | | | | | | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_8_3. | Value | ACATTATATTTTATATTTGGG[...] | | | | | | | | | | | | | | | | | | | | | | |
| 66 | Date | | | | | | | | | | | | | | | | | | | | | | | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_1_4. | ParameterID | | For all: "Barcode Sequencing timestamp" | | | | | | | | | | | | | | | | | | | | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_1_4. | Value | 30.10.2014 08:21 | | | | | | | | | | | | | | | | | | | | | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_2_4. | ParameterID | | For all: "Barcode Sequencing timestamp" | | | | | | | | | | | | | | | | | | | | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_2_4. | Value | 30.10.2014 08:21 | | | | | | | | | | | | | | | | | | | | | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_3_4. | ParameterID | | For all: "Barcode Sequencing timestamp" | | | | | | | | | | | | | | | | | | | | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_3_4. | Value | 30.10.2014 08:21 | | | | | | | | | | | | | | | | | | | | | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_4_4. | ParameterID | | For all: "Barcode Sequencing timestamp" | | | | | | | | | | | | | | | | | | | | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_4_4. | Value | 30.10.2014 08:21 | | | | | | | | | | | | | | | | | | | | | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_5_4. | ParameterID | | For all: "Barcode Sequencing timestamp" | | | | | | | | | | | | | | | | | | | | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_5_4. | Value | 30.10.2014 08:21 | | | | | | | | | | | | | | | | | | | | | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_6_4. | ParameterID | | For all: "Barcode Sequencing timestamp" | | | | | | | | | | | | | | | | | | | | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_6_4. | Value | 30.10.2014 08:21 | | | | | | | | | | | | | | | | | | | | | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_7_4. | ParameterID | | For all: "Barcode Sequencing timestamp" | | | | | | | | | | | | | | | | | | | | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_7_4. | Value | 30.10.2014 08:21 | | | | | | | | | | | | | | | | | | | | | | |
| 67 | direction | | | | | | | | | | | | | | | | | | | | | | | | |
| | IdentificationUnitAnalysisMethod_1_1_1. | MethodID | R | 16 | | | | | | | | | | | | | | | | | | | | | |
| | | Transformations: | <table><tr><th>Split at</th><th>Pos.</th><th>Dir.</th></tr><tr><td>I</td><td>1</td><td>⇔</td></tr><tr><td colspan="3">Translate Into</td></tr><tr><td>F</td><td>16</td><td></td></tr><tr><td>R</td><td>16</td><td></td></tr></table> | Split at | Pos. | Dir. | I | 1 | ⇔ | Translate Into | | | F | 16 | | R | 16 | | | | | | | | |
| Split at | Pos. | Dir. | | | | | | | | | | | | | | | | | | | | | | | |
| I | 1 | ⇔ | | | | | | | | | | | | | | | | | | | | | | | |
| Translate Into | | | | | | | | | | | | | | | | | | | | | | | | | |
| F | 16 | | | | | | | | | | | | | | | | | | | | | | | | |
| R | 16 | | | | | | | | | | | | | | | | | | | | | | | | |
| | IdentificationUnitAnalysisMethod_1_1_1. | MethodMarker | | For all: "2" | | | | | | | | | | | | | | | | | | | | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_1_1. | ParameterID | | For all: "Barcode Sequencing direction" | | | | | | | | | | | | | | | | | | | | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_1_1. | Value | R reverse | | | | | | | | | | | | | | | | | | | | | | |
| | | Transformations: | <table><tr><th>Split at</th><th>Pos.</th><th>Dir.</th></tr><tr><td>I</td><td>1</td><td>⇔</td></tr><tr><td colspan="3">Translate Into</td></tr><tr><td>directions</td><td></td><td></td></tr><tr><td>F</td><td>forward</td><td></td></tr><tr><td>R</td><td>reverse</td><td></td></tr><tr><td>reverse</td><td></td><td></td></tr></table> | Split at | Pos. | Dir. | I | 1 | ⇔ | Translate Into | | | directions | | | F | forward | | R | reverse | | reverse | | | |
| Split at | Pos. | Dir. | | | | | | | | | | | | | | | | | | | | | | | |
| I | 1 | ⇔ | | | | | | | | | | | | | | | | | | | | | | | |
| Translate Into | | | | | | | | | | | | | | | | | | | | | | | | | |
| directions | | | | | | | | | | | | | | | | | | | | | | | | | |
| F | forward | | | | | | | | | | | | | | | | | | | | | | | | |
| R | reverse | | | | | | | | | | | | | | | | | | | | | | | | |
| reverse | | | | | | | | | | | | | | | | | | | | | | | | | |
| | IdentificationUnitAnalysisMethod_1_1_2. | MethodID | R | 16 | | | | | | | | | | | | | | | | | | | | | |
| | | Transformations: | <table><tr><th>Split at</th><th>Pos.</th><th>Dir.</th></tr><tr><td>I</td><td>2</td><td>⇔</td></tr><tr><td colspan="3">Translate Into</td></tr></table> | Split at | Pos. | Dir. | I | 2 | ⇔ | Translate Into | | | | | | | | | | | | | | | |
| Split at | Pos. | Dir. | | | | | | | | | | | | | | | | | | | | | | | |
| I | 2 | ⇔ | | | | | | | | | | | | | | | | | | | | | | | |
| Translate Into | | | | | | | | | | | | | | | | | | | | | | | | | |

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IdentificationUnitAnalysisMethod_1_1_2.

MethodMarker

For all: "3"

IdentificationUnitAnalysisMethodParameter_1_1_2_1.

ParameterID

For all: "Barcode | Sequencing | direction"

IdentificationUnitAnalysisMethodParameter_1_1_2_1.

Value

R

Transformations:

| Split at | Pos. | Dir. |
|----------|------|------|
| I | 2 | ⇒ |

| Translate | Into |
|-----------|---------|
| F | forward |
| R | reverse |

IdentificationUnitAnalysisMethod_1_1_3.

MethodID

R

Transformations:

| Split at | Pos. | Dir. |
|----------|------|------|
| I | 3 | ⇒ |

| Translate | Into |
|-----------|------|
| F | 16 |
| R | 16 |

IdentificationUnitAnalysisMethod_1_1_3.

MethodMarker

For all: "4"

IdentificationUnitAnalysisMethodParameter_1_1_3_1.

ParameterID

For all: "Barcode | Sequencing | direction"

IdentificationUnitAnalysisMethodParameter_1_1_3_1.

Value

R

Transformations:

| Split at | Pos. | Dir. |
|----------|------|------|
| I | 3 | ⇒ |

| Translate | Into |
|-----------|---------|
| F | forward |
| R | reverse |

IdentificationUnitAnalysisMethod_1_1_4.

MethodID

R

Transformations:

| Split at | Pos. | Dir. |
|----------|------|------|
| I | 4 | ⇒ |

| Translate | Into |
|-----------|------|
| F | 16 |
| R | 16 |

IdentificationUnitAnalysisMethod_1_1_4.

MethodMarker

For all: "5"

IdentificationUnitAnalysisMethodParameter_1_1_4_1.

ParameterID

For all: "Barcode | Sequencing | direction"

IdentificationUnitAnalysisMethodParameter_1_1_4_1.

Value

R

Transformations:

| Split at | Pos. | Dir. |
|----------|------|------|
| I | 4 | ⇒ |

| Translate | Into |
|-----------|---------|
| F | forward |
| R | reverse |

IdentificationUnitAnalysisMethod_1_1_5.

MethodID

R

Transformations:

| Split at | Pos. | Dir. |
|----------|------|------|
| I | 5 | ⇒ |

| Translate | Into |
|-----------|------|
| F | 16 |
| R | 16 |

IdentificationUnitAnalysisMethod_1_1_5.

MethodMarker

For all: "6"

IdentificationUnitAnalysisMethodParameter_1_1_5_1.

ParameterID

For all: "Barcode | Sequencing | direction"

IdentificationUnitAnalysisMethodParameter_1_1_5_1.

Value

R

Transformations:

| Split at | Pos. | Dir. |
|----------|------|------|
| I | 5 | ⇒ |

| Translate | Into |
|-----------|------|
| F | 16 |
| R | 16 |

| | | | | | | | | | | | | | | | | | |
|--|--------------|---|--|--|------|---------|---|---------|---|-----------|------|---|---------|---|---------|--|--|
| | | <table><tr><td>Translate</td><td>Into</td></tr><tr><td>F</td><td>forward</td></tr><tr><td>R</td><td>reverse</td></tr></table> | Translate | Into | F | forward | R | reverse | | | | | | | | | |
| Translate | Into | | | | | | | | | | | | | | | | |
| F | forward | | | | | | | | | | | | | | | | |
| R | reverse | | | | | | | | | | | | | | | | |
| IdentificationUnitAnalysisMethod_1_1_6. | MethodID | | R | 16 | | | | | | | | | | | | | |
| | | Transformations: | <table><tr><td>Split at</td><td>Pos.</td><td>Dir.</td></tr><tr><td>I</td><td>6</td><td>⇒</td></tr></table> <table><tr><td>Translate</td><td>Into</td></tr><tr><td>F</td><td>16</td></tr><tr><td>R</td><td>16</td></tr></table> | Split at | Pos. | Dir. | I | 6 | ⇒ | Translate | Into | F | 16 | R | 16 | | |
| Split at | Pos. | Dir. | | | | | | | | | | | | | | | |
| I | 6 | ⇒ | | | | | | | | | | | | | | | |
| Translate | Into | | | | | | | | | | | | | | | | |
| F | 16 | | | | | | | | | | | | | | | | |
| R | 16 | | | | | | | | | | | | | | | | |
| IdentificationUnitAnalysisMethod_1_1_6. | MethodMarker | | | For all: "7" | | | | | | | | | | | | | |
| IdentificationUnitAnalysisMethodParameter_1_1_6_1. | ParameterID | | | For all: "Barcode Sequencing direction" | | | | | | | | | | | | | |
| IdentificationUnitAnalysisMethodParameter_1_1_6_1. | Value | | R | | | | | | | | | | | | | | |
| | | Transformations: | <table><tr><td>Split at</td><td>Pos.</td><td>Dir.</td></tr><tr><td>I</td><td>6</td><td>⇒</td></tr></table> <table><tr><td>Translate</td><td>Into</td></tr><tr><td>F</td><td>forward</td></tr><tr><td>R</td><td>reverse</td></tr></table> | Split at | Pos. | Dir. | I | 6 | ⇒ | Translate | Into | F | forward | R | reverse | | |
| Split at | Pos. | Dir. | | | | | | | | | | | | | | | |
| I | 6 | ⇒ | | | | | | | | | | | | | | | |
| Translate | Into | | | | | | | | | | | | | | | | |
| F | forward | | | | | | | | | | | | | | | | |
| R | reverse | | | | | | | | | | | | | | | | |
| IdentificationUnitAnalysisMethod_1_1_7. | MethodID | | R | 16 | | | | | | | | | | | | | |
| | | Transformations: | <table><tr><td>Split at</td><td>Pos.</td><td>Dir.</td></tr><tr><td>I</td><td>7</td><td>⇒</td></tr></table> <table><tr><td>Translate</td><td>Into</td></tr><tr><td>F</td><td>16</td></tr><tr><td>R</td><td>16</td></tr></table> | Split at | Pos. | Dir. | I | 7 | ⇒ | Translate | Into | F | 16 | R | 16 | | |
| Split at | Pos. | Dir. | | | | | | | | | | | | | | | |
| I | 7 | ⇒ | | | | | | | | | | | | | | | |
| Translate | Into | | | | | | | | | | | | | | | | |
| F | 16 | | | | | | | | | | | | | | | | |
| R | 16 | | | | | | | | | | | | | | | | |
| IdentificationUnitAnalysisMethod_1_1_7. | MethodMarker | | | For all: "8" | | | | | | | | | | | | | |
| IdentificationUnitAnalysisMethodParameter_1_1_7_1. | ParameterID | | | For all: "Barcode Sequencing direction" | | | | | | | | | | | | | |
| IdentificationUnitAnalysisMethodParameter_1_1_7_1. | Value | | R | | | | | | | | | | | | | | |
| | | Transformations: | <table><tr><td>Split at</td><td>Pos.</td><td>Dir.</td></tr><tr><td>I</td><td>7</td><td>⇒</td></tr></table> <table><tr><td>Translate</td><td>Into</td></tr><tr><td>F</td><td>forward</td></tr><tr><td>R</td><td>reverse</td></tr></table> | Split at | Pos. | Dir. | I | 7 | ⇒ | Translate | Into | F | forward | R | reverse | | |
| Split at | Pos. | Dir. | | | | | | | | | | | | | | | |
| I | 7 | ⇒ | | | | | | | | | | | | | | | |
| Translate | Into | | | | | | | | | | | | | | | | |
| F | forward | | | | | | | | | | | | | | | | |
| R | reverse | | | | | | | | | | | | | | | | |
| seq_primers_Name | | | | | | | | | | | | | | | | | |
| IdentificationUnitAnalysisMethodParameter_1_1_1_2. | ParameterID | | | For all: "Barcode Sequencing sequencing_primer_name" | | | | | | | | | | | | | |
| IdentificationUnitAnalysisMethodParameter_1_1_1_2. | Value | C_LepFolR | | | | | | | | | | | | | | | |
| | | Transformations: | <table><tr><td>Split at</td><td>Pos.</td><td>Dir.</td></tr><tr><td>I</td><td>1</td><td>⇒</td></tr></table> | Split at | Pos. | Dir. | I | 1 | ⇒ | | | | | | | | |
| Split at | Pos. | Dir. | | | | | | | | | | | | | | | |
| I | 1 | ⇒ | | | | | | | | | | | | | | | |
| IdentificationUnitAnalysisMethodParameter_1_1_2_2. | ParameterID | C_LepFolR | | For all: "Barcode Sequencing sequencing_primer_name" | | | | | | | | | | | | | |
| IdentificationUnitAnalysisMethodParameter_1_1_2_2. | Value | C_LepFolR | | | | | | | | | | | | | | | |
| | | Transformations: | <table><tr><td>Split at</td><td>Pos.</td><td>Dir.</td></tr><tr><td>I</td><td>2</td><td>⇒</td></tr></table> | Split at | Pos. | Dir. | I | 2 | ⇒ | | | | | | | | |
| Split at | Pos. | Dir. | | | | | | | | | | | | | | | |
| I | 2 | ⇒ | | | | | | | | | | | | | | | |
| IdentificationUnitAnalysisMethodParameter_1_1_3_2. | ParameterID | | | For all: "Barcode Sequencing sequencing_primer_name" | | | | | | | | | | | | | |
| IdentificationUnitAnalysisMethodParameter_1_1_3_2. | Value | C_LepFolR | | | | | | | | | | | | | | | |
| | | Transformations: | <table><tr><td>Split at</td><td>Pos.</td><td>Dir.</td></tr><tr><td>I</td><td>3</td><td>⇒</td></tr></table> | Split at | Pos. | Dir. | I | 3 | ⇒ | | | | | | | | |
| Split at | Pos. | Dir. | | | | | | | | | | | | | | | |
| I | 3 | ⇒ | | | | | | | | | | | | | | | |
| IdentificationUnitAnalysisMethodParameter_1_1_4_2. | ParameterID | | | For all: "Barcode Sequencing sequencing_primer_name" | | | | | | | | | | | | | |
| IdentificationUnitAnalysisMethodParameter_1_1_4_2. | Value | C_LepFolR | | | | | | | | | | | | | | | |
| | | Transformations: | <table><tr><td>Split at</td><td>Pos.</td><td>Dir.</td></tr><tr><td>I</td><td></td><td></td></tr></table> | Split at | Pos. | Dir. | I | | | | | | | | | | |
| Split at | Pos. | Dir. | | | | | | | | | | | | | | | |
| I | | | | | | | | | | | | | | | | | |

Description : Import scheme for GBOL-Data (attaching Barcodeanalyse from BOLD to existing CollectionSpecimens)

| | | | |
|---|-------------|-----------------------------------|--|
| IdentificationUnitAnalysisMethodParameter_1_1_5_2. | ParameterID | <div><div></div>4⇨</div> | For all: "Barcode Sequencing sequencing_primer_name" |
| | Value | C_LepFolR | |
| Transformations: <div><div>Split at</div><div>Pos.</div><div>Dir.</div><div><div></div>5⇨</div></div> | | | |
| IdentificationUnitAnalysisMethodParameter_1_1_6_2. | ParameterID | <div><div></div>6⇨</div> | For all: "Barcode Sequencing sequencing_primer_name" |
| | Value | C_LepFolR | |
| Transformations: <div><div>Split at</div><div>Pos.</div><div>Dir.</div><div><div></div>6⇨</div></div> | | | |
| IdentificationUnitAnalysisMethodParameter_1_1_7_2. | ParameterID | <div><div></div>7⇨</div> | For all: "Barcode Sequencing sequencing_primer_name" |
| | Value | C_LepFolR | |
| Transformations: <div><div>Split at</div><div>Pos.</div><div>Dir.</div><div><div></div>7⇨</div></div> | | | |
| sequencing_lab | | | |
| IdentificationUnitAnalysisMethodParameter_1_1_1_3. | ParameterID | <div><div></div>1⇨</div> | For all: "Barcode Sequencing sequencing_labor" |
| | Value | Biodiversity Institute of Ontario | |
| Transformations: <div><div>Split at</div><div>Pos.</div><div>Dir.</div><div><div></div>1⇨</div></div> | | | |
| IdentificationUnitAnalysisMethodParameter_1_1_2_3. | ParameterID | <div><div></div>2⇨</div> | For all: "Barcode Sequencing sequencing_labor" |
| | Value | Biodiversity Institute of Ontario | |
| Transformations: <div><div>Split at</div><div>Pos.</div><div>Dir.</div><div><div></div>2⇨</div></div> | | | |
| IdentificationUnitAnalysisMethodParameter_1_1_3_3. | ParameterID | <div><div></div>3⇨</div> | For all: "Barcode Sequencing sequencing_labor" |
| | Value | Biodiversity Institute of Ontario | |
| Transformations: <div><div>Split at</div><div>Pos.</div><div>Dir.</div><div><div></div>3⇨</div></div> | | | |
| IdentificationUnitAnalysisMethodParameter_1_1_4_3. | ParameterID | <div><div></div>4⇨</div> | For all: "Barcode Sequencing sequencing_labor" |
| | Value | Biodiversity Institute of Ontario | |
| Transformations: <div><div>Split at</div><div>Pos.</div><div>Dir.</div><div><div></div>4⇨</div></div> | | | |
| IdentificationUnitAnalysisMethodParameter_1_1_5_3. | ParameterID | <div><div></div>5⇨</div> | For all: "Barcode Sequencing sequencing_labor" |
| | Value | Biodiversity Institute of Ontario | |
| Transformations: <div><div>Split at</div><div>Pos.</div><div>Dir.</div><div><div></div>5⇨</div></div> | | | |
| IdentificationUnitAnalysisMethodParameter_1_1_6_3. | ParameterID | <div><div></div>6⇨</div> | For all: "Barcode Sequencing sequencing_labor" |
| | Value | Biodiversity Institute of Ontario | |
| Transformations: <div><div>Split at</div><div>Pos.</div><div>Dir.</div><div><div></div>6⇨</div></div> | | | |
| IdentificationUnitAnalysisMethodParameter_1_1_7_3. | ParameterID | <div><div></div>7⇨</div> | For all: "Barcode Sequencing sequencing_labor" |
| | Value | Biodiversity Institute of Ontario | |
| Transformations: <div><div>Split at</div><div>Pos.</div><div>Dir.</div><div><div></div>7⇨</div></div> | | | |

Description : Import scheme for GBOL-Data (attaching Barcodeanalyse from BOLD to existing CollectionSpecimens)

| | | | |
|----|--|----------------------------|---|
| | IdentificationUnitAnalysis_1_1. | AnalysisNumber | COI-5P |
| | IdentificationUnitAnalysisMethodParameter_1_1_8_1. | Value | COI-5P |
| | 71 | genbank_accession | |
| | CollectionSpecimenRelation_2. | RelatedSpecimenURI | Prefix: http://www.ncbi.nlm.nih.gov/nuccore/ |
| | CollectionSpecimenRelation_2. | RelatedSpecimenDisplayText | |
| | CollectionSpecimenRelation_2. | RelatedSpecimenDescription | |
| 72 | image_ids | | |
| | CollectionSpecimenImage_1. | Notes | Prefix: image id: _ |
| | 73 | image_urls | |
| | CollectionSpecimenImage_1. | URI | |
| | 74 | copyright_licenses | |
| | CollectionSpecimenImage_1. | LicenseType | |
| | 75 | trace_ids | |
| | IdentificationUnitAnalysisMethodParameter_1_1_1_6. | ParameterID | For all: "Barcode Sequencing trace_id" |
| | IdentificationUnitAnalysisMethodParameter_1_1_1_6. | Value | 7126975 |
| | Transformations: Split at Pos. Dir. | | |
| | 1 ⇌ | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_2_6. | ParameterID | For all: "Barcode Sequencing trace_id" |
| | IdentificationUnitAnalysisMethodParameter_1_1_2_6. | Value | 7126975 |
| | Transformations: Split at Pos. Dir. | | |
| | 2 ⇌ | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_3_6. | ParameterID | For all: "Barcode Sequencing trace_id" |
| | IdentificationUnitAnalysisMethodParameter_1_1_3_6. | Value | 7126975 |
| | Transformations: Split at Pos. Dir. | | |
| | 3 ⇌ | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_4_6. | ParameterID | For all: "Barcode Sequencing trace_id" |
| | IdentificationUnitAnalysisMethodParameter_1_1_4_6. | Value | 7126975 |
| | Transformations: Split at Pos. Dir. | | |
| | 4 ⇌ | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_5_6. | ParameterID | For all: "Barcode Sequencing trace_id" |
| | IdentificationUnitAnalysisMethodParameter_1_1_5_6. | Value | 7126975 |
| | Transformations: Split at Pos. Dir. | | |
| | 5 ⇌ | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_6_6. | ParameterID | For all: "Barcode Sequencing trace_id" |
| | IdentificationUnitAnalysisMethodParameter_1_1_6_6. | Value | 7126975 |
| | Transformations: Split at Pos. Dir. | | |
| | 6 ⇌ | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_7_6. | ParameterID | For all: "Barcode Sequencing trace_id" |
| | IdentificationUnitAnalysisMethodParameter_1_1_7_6. | Value | 7126975 |
| | Transformations: Split at Pos. Dir. | | |
| | 7 ⇌ | | |
| 76 | trace_links | | |
| | IdentificationUnitAnalysisMethodParameter_1_1_1_5. | ParameterID | For all: "Barcode Sequencing trace_url" |
| | IdentificationUnitAnalysisMethodParameter_1_1_1_5. | Value | http://trace.boldsystems.org/traceIO/bold.org/8957035 |
| | Transformations: Split at Pos. Dir. | | |
| | 1 ⇌ | | |

Description : Import scheme for GBOL-Data (attaching Barcodeanalyse from BOLD to existing CollectionSpecimens)

| | | | |
|--|-------------|---|---|
| IdentificationUnitAnalysisMethodParameter_1_1_2_5. | ParameterID | | For all: "Barcode Sequencing trace_url" |
| IdentificationUnitAnalysisMethodParameter_1_1_2_5. | Value | http://trace.boldsystems.org/traceIO/bold.org/8957035 | |
| Transformations: | | | |
| | | Split at | Pos. Dir. |
| | | | 2 ⇌ |
| IdentificationUnitAnalysisMethodParameter_1_1_3_5. | ParameterID | | For all: "Barcode Sequencing trace_url" |
| IdentificationUnitAnalysisMethodParameter_1_1_3_5. | Value | http://trace.boldsystems.org/traceIO/bold.org/8957035 | |
| Transformations: | | | |
| | | Split at | Pos. Dir. |
| | | | 3 ⇌ |
| IdentificationUnitAnalysisMethodParameter_1_1_4_5. | ParameterID | | For all: "Barcode Sequencing trace_url" |
| IdentificationUnitAnalysisMethodParameter_1_1_4_5. | Value | http://trace.boldsystems.org/traceIO/bold.org/8957035 | |
| Transformations: | | | |
| | | Split at | Pos. Dir. |
| | | | 4 ⇌ |
| IdentificationUnitAnalysisMethodParameter_1_1_5_5. | ParameterID | | For all: "Barcode Sequencing trace_url" |
| IdentificationUnitAnalysisMethodParameter_1_1_5_5. | Value | http://trace.boldsystems.org/traceIO/bold.org/8957035 | |
| Transformations: | | | |
| | | Split at | Pos. Dir. |
| | | | 5 ⇌ |
| IdentificationUnitAnalysisMethodParameter_1_1_6_5. | ParameterID | | For all: "Barcode Sequencing trace_url" |
| IdentificationUnitAnalysisMethodParameter_1_1_6_5. | Value | http://trace.boldsystems.org/traceIO/bold.org/8957035 | |
| Transformations: | | | |
| | | Split at | Pos. Dir. |
| | | | 6 ⇌ |
| IdentificationUnitAnalysisMethodParameter_1_1_7_5. | ParameterID | | For all: "Barcode Sequencing trace_url" |
| IdentificationUnitAnalysisMethodParameter_1_1_7_5. | Value | http://trace.boldsystems.org/traceIO/bold.org/8957035 | |
| Transformations: | | | |
| | | Split at | Pos. Dir. |
| | | | 7 ⇌ |

77 marker_codes

not imported

78 Ext_URI

| | | |
|-------------------------------|----------------------------|---|
| CollectionSpecimenRelation_1. | RelatedSpecimenURI | http://www.boldsystems.org/index.php/Public_RecordView?processid=GMGM B967-14 |
| CollectionSpecimenRelation_1. | RelatedSpecimenDisplayText | http://www.boldsystems.org/index.php/Public_RecordView?processid=GMGM B967-14 |
| CollectionSpecimenRelation_1. | RelatedSpecimenDescription | http://www.boldsystems.org/index.php/Public_RecordView?processid=GMGM B967-14 |

Description : Import scheme for GBOL-Data (attaching Barcodeanalyse from BOLD to existing CollectionSpecimens)

Interface settings

| Table | Table alias | Column | Value | |
|---|---|--------------------------------|-----------------|----------------------------|
| CollectionProject | CollectionProject_1 | ProjectID | 29995 | GBOL-Malaise |
| CollectionSpecimenRelation | CollectionSpecimenRelation_1 | RelationType | Public database | |
| | CollectionSpecimenRelation_1 | RelatedSpecimen Description | BOLD Process ID | |
| | CollectionSpecimenRelation_1 | IsInternalRelationCache | 0 | |
| | CollectionSpecimenRelation_2 | RelationType | public database | |
| | CollectionSpecimenRelation_2 | RelatedSpecimen Description | GenBank Acc. No | |
| | CollectionSpecimenRelation_2 | IsInternalRelationCache | 0 | |
| IdentificationUnit | IdentificationUnit_1 | TaxonomicGroup | animal | |
| IdentificationUnitAnalysis | IdentificationUnitAnalysis_1_1 | AnalysisID | 161 | Barcode_Analyse161 |
| IdentificationUnitAnalysisMethod | IdentificationUnitAnalysisMethod_1_1_1 | MethodMarker | 2 | |
| | IdentificationUnitAnalysisMethod_1_1_2 | MethodMarker | 3 | |
| | IdentificationUnitAnalysisMethod_1_1_3 | MethodMarker | 4 | |
| | IdentificationUnitAnalysisMethod_1_1_4 | MethodMarker | 5 | |
| | IdentificationUnitAnalysisMethod_1_1_5 | MethodMarker | 6 | |
| | IdentificationUnitAnalysisMethod_1_1_6 | MethodMarker | 7 | |
| | IdentificationUnitAnalysisMethod_1_1_7 | MethodMarker | 8 | |
| | IdentificationUnitAnalysisMethod_1_1_8 | MethodID | 12 | Barcode |
| | IdentificationUnitAnalysisMethod_1_1_8 | MethodMarker | 9 | |
| IdentificationUnitAnalysisMethodParameter | IdentificationUnitAnalysisMethodParameter_1_1_1_1 | ParameterID | 73 | direction |
| | IdentificationUnitAnalysisMethodParameter_1_1_1_2 | ParameterID | 91 | sequencing_primer_name |
| | IdentificationUnitAnalysisMethodParameter_1_1_1_3 | ParameterID | 72 | sequencing_primer_sequence |
| | IdentificationUnitAnalysisMethodParameter_1_1_1_4 | ParameterID | 71 | sequencing_timestamp |
| | IdentificationUnitAnalysisMethodParameter_1_1_1_5 | ParameterID | 75 | trace_file_url |
| | IdentificationUnitAnalysisMethodParameter_1_1_1_6 | ParameterID | 83 | trace_id |
| | IdentificationUnitAnalysisMethodParameter_1_1_1_7 | ParameterID | 84 | sequence_id |
| | IdentificationUnitAnalysisMethodParameter_1_1_2_1 | ParameterID | 73 | direction |
| | IdentificationUnitAnalysisMethodParameter_1_1_2_2 | ParameterID | 91 | sequencing_primer_name |
| | IdentificationUnitAnalysisMethodParameter_1_1_2_3 | ParameterID | 72 | sequencing_primer_sequence |
| | IdentificationUnitAnalysisMethodParameter_1_1_2_4 | ParameterID | 71 | sequencing_timestamp |
| | IdentificationUnitAnalysisMethodParameter_1_1_2_5 | ParameterID | 75 | trace_file_url |
| | IdentificationUnitAnalysisMethodParameter_1_1_2_6 | ParameterID | 83 | trace_id |
| | IdentificationUnitAnalysisMethodParameter_1_1_2_7 | ParameterID | 84 | sequence_id |
| | IdentificationUnitAnalysisMethodParameter_1_1_3_1 | ParameterID | 73 | direction |
| | IdentificationUnitAnalysisMethodParameter_1_1_3_2 | ParameterID | 91 | sequencing_primer_name |
| | IdentificationUnitAnalysisMethodParameter_1_1_3_3 | ParameterID | 72 | sequencing_primer_sequence |
| | IdentificationUnitAnalysisMethodParameter_1_1_3_4 | ParameterID | 71 | sequencing_timestamp |
| | IdentificationUnitAnalysisMethodParameter_1_1_3_5 | ParameterID | 75 | trace_file_url |
| | IdentificationUnitAnalysisMethodParameter_1_1_3_6 | ParameterID | 83 | trace_id |

Description : Import scheme for GBOL-Data (attaching Barcodeanalyse from BOLD to existing CollectionSpecimens)

| | | | |
|---|-------------|----|----------------------------|
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 84 | sequence_id |
| _1_3_7 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 73 | direction |
| _1_4_1 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 91 | sequencing_primer_name |
| _1_4_2 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 72 | sequencing_primer_sequence |
| _1_4_3 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 71 | sequencing_timestamp |
| _1_4_4 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 75 | trace_file_url |
| _1_4_5 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 83 | trace_id |
| _1_4_6 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 84 | sequence_id |
| _1_4_7 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 73 | direction |
| _1_5_1 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 91 | sequencing_primer_name |
| _1_5_2 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 72 | sequencing_primer_sequence |
| _1_5_3 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 71 | sequencing_timestamp |
| _1_5_4 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 75 | trace_file_url |
| _1_5_5 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 83 | trace_id |
| _1_5_6 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 84 | sequence_id |
| _1_5_7 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 73 | direction |
| _1_6_1 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 91 | sequencing_primer_name |
| _1_6_2 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 72 | sequencing_primer_sequence |
| _1_6_3 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 71 | sequencing_timestamp |
| _1_6_4 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 75 | trace_file_url |
| _1_6_5 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 83 | trace_id |
| _1_6_6 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 84 | sequence_id |
| _1_6_7 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 73 | direction |
| _1_7_1 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 91 | sequencing_primer_name |
| _1_7_2 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 72 | sequencing_primer_sequence |
| _1_7_3 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 71 | sequencing_timestamp |
| _1_7_4 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 75 | trace_file_url |
| _1_7_5 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 83 | trace_id |
| _1_7_6 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 84 | sequence_id |
| _1_7_7 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 62 | region |
| _1_8_1 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 90 | sequence |
| _1_8_3 | | | |
| IdentificationUnitAnalysisMethodParameter_1 | ParameterID | 28 | failure |
| _1_8_4 | | | |

Description : Import scheme for GBOL-Data (attaching Barcodeanalyse from BOLD to existing CollectionSpecimens)

| | | |
|---|--------------|----------------|
| IdentificationUnitAnalysisMethodParameter_1 ParameterID _1_8_5 | 29 | failure_detail |
| IdentificationUnitAnalysisMethodParameter_1 ParameterID _1_8_6 | 27 | project |
| IdentificationUnitAnalysisMethodParameter_1 Value _1_8_6 | GBOL-Malaise | |