

Workshop 2020-10-26 to 2020-10-27: DiversityDescriptions as Research Data
Management System (RDMS) – 39. Diversity Workbench Workshop

Management von Forschungsdesign-Daten, Descriptor Trees und Messdaten

Janno Harjes, Gerhard Rambold
Universität Bayreuth



Workflow Segment Design

The screenshot displays the DiversityDescriptions v. 4.2.9 application window. The main area is titled 'Project name' and shows the configuration for a descriptor. The 'Descriptor' field is set to 'Project name', with 'No: 1' and 'ID: 1059003'. The 'Main attributes' tab is active, showing 'Abbrev.: Project name' and 'Details: Name of project at UBT Mycology'. The 'Notes' field is empty. The 'Natural language support' section includes 'Wording:', 'before:', and 'after:' fields. The 'Descriptor type' is set to 'Categorical'. The 'Descriptor settings' section includes 'Statistical measurement scale' (Nominal), 'State collection model' (OrSet), 'Repeatability' (Standard), and 'Availability' (Standard). The 'Measurement data' section includes 'Prefix', 'Base', 'Unit', 'Minimum' (-1,790000E+308), and 'Maximum' (1,790000E+308). The 'Molecular sequence data' section includes 'Nucleotide sequence' (selected), 'Protein sequence', 'Enable ambiguity symbols' (checked), 'Symbol length' (1), and 'Gap symbol'. A yellow banner at the bottom indicates 'Assigned to more than one project'.

DiversityDescriptions v. 4.2.9

Connection Edit Grid Query Data Administration Help

Descriptor Project name

Query results 1 - 92

- 1 - Project name
- 2 - Physical unit type
- 3 - Physical unit status
- 4 - Physical unit ID
- 5 - Physical unit origin ID
- 6 - Physical unit preceding ID
- 7 - Physical unit external ID
- 8 - Unit operational agent name
- 9 - Unit operation workflow segment
- 10 - Storage container-internal position
- 11 - Storage container ID
- 12 - Storage shelf number
- 13 - Storage rack ID
- 14 - Storage device name and ID
- 15 - Storage room name
- 16 - GBOL2TP12_environmental oper
- 17 - GBOL2TP12_host tree individual
- 18 - GBOL2TP12_host tree organ
- 19 - GBOL2TP12_environmental sampl
- 20 - GBOL2TP12_environmental sampl
- 21 - GBOL2TP12_nucleic acids extrac
- 22 - GBOL2TP12_nucleic acids extrac
- 23 - GBOL2TP12_PCR amplification
- 24 - GBOL2TP12_PCR amplicon purif
- 25 - GBOL2TP12_index PCR amplifica
- 26 - GBOL2TP12_index PCR amplicon
- 27 - GBOL2TP12_microarray name
- 28 - GBOL2TP12_microarray manufac
- 29 - GBOL2TP12_microarray develop
- 30 - GBOL2TP12_microarray format
- 31 - GBOL2TP12_microarray design i
- 32 - GBOL2TP12_microarray design i
- 33 - GBOL2TP12_microarray design p
- 34 - GBOL2TP12_microarray design p
- 35 - GBOL2TP12_microarray design p
- 36 - GBOL2TP12_microarray device i
- 37 - Halotolerant-fungi_environmental
- 38 - Halotolerant-fungi_environmental
- 39 - Halotolerant-fungi_cultivation cult
- 40 - Halotolerant-fungi_cultivation cult
- 41 - Halotolerant-fungi_cultivation cult
- 42 - Halotolerant-fungi_cultivation cult
- 43 - Halotolerant-fungi_microbial_culti
- 44 - Hoehengradient_environmental o
- 45 - Hoehengradient_soil borehole nu
- 46 - Hoehengradient_soil horizon type

order by: Sequence numb

Edit descriptors Assigned to more than one project

Workflow Segment Design

DiversityDescriptions v. 4.2.9

Connection Edit Grid Query Data Administration Help

Query results 1 - 92

- 47 - Hoehengradient_environmental sample replicate number
- 48 - Hoehengradient_environmental sampling
- 49 - Hoehengradient_nucleic acids extraction
- 50 - Hoehengradient_nucleic acids extract purification
- 51 - Hoehengradient_PCR amplification
- 52 - Hoehengradient_PCR amplicon purification
- 53 - Hoehengradient_Index PCR amplification
- 54 - Hoehengradient_Index PCR amplicon purification
- 55 - LambweValley_environmental operation location
- 56 - LambweValley_soil borehole number
- 57 - LambweValley_soil horizon type
- 58 - LambweValley_environmental sample replicate number
- 59 - LambweValley_environmental sampling
- 60 - LambweValley_nucleic acids extraction
- 61 - LambweValley_nucleic acids extract purification
- 62 - LambweValley_PCR amplification
- 63 - LambweValley_PCR amplicon purification
- 64 - LambweValley_Index PCR amplification
- 65 - LambweValley_Index PCR amplicon purification
- 66 - Microplastic_environmental operation location
- 67 - Microplastic_soil sample number
- 68 - Microplastic_environmental sampling
- 69 - Microplastic_soil sample subsample type
- 70 - Microplastic_cultivation cryoculture origin
- 71 - Microplastic_cultivation culture generation
- 72 - Microplastic_cultivation culture medium type
- 73 - Microplastic_cultivation culture medium type variation
- 74 - Microplastic_cultivation culture replicate number
- 75 - Microplastic_microbial_cultivation
- 76 - Microplastic_nucleic acids extraction
- 77 - Microplastic_nucleic acids extract purification
- 78 - Microplastic_PCR amplification
- 79 - Microplastic_PCR amplicon purification
- 80 - Microplastic_Index PCR amplification
- 81 - Microplastic_Index PCR amplicon purification
- 82 - Waldeck_environmental operation location
- 83 - Waldeck_lichen thallus individual number
- 84 - Waldeck_lichen thallus fragment
- 85 - Waldeck_environmental sample replicate number
- 86 - Waldeck_environmental sampling
- 87 - Waldeck_nucleic acids extraction
- 88 - Waldeck_nucleic acids extract purification
- 89 - Waldeck_PCR amplification
- 90 - Waldeck_PCR amplicon purification
- 91 - Waldeck_Index PCR amplification
- 92 - Waldeck_Index PCR amplicon purification

order by: Sequence number

Descriptor Project name

Descriptor: Project name No: 1 ID: 1059003

Main attributes Categorical states Descriptor tree Modifier/Frequency Resource links

Descriptor tree

- Project name
- Field workflow segments
- OD - Operation design - Transformation - Mea:
- MI - Method information
- Laboratory workflow segments
- MI - Method information
- Repository workflow segments
- OD - Operation design - Transaction - Storage
- GBOL2TP12 - Chip design
- Halotolerant fungi - Workflow design
- General
- Project name
- Field workflow segments
- OD - Operation design - Transformation - Mea:
- MI - Method information
- Laboratory workflow segments
- OD - Operation design - Transformation - Mea:
- MI - Method information
- Repository workflow segments
- OD - Operation design - Transaction - Storage
- Hoehengradient - Workflow design
- General
- Project name
- Field workflow segments
- OD - Operation design - Transformation - Mea:
- MI - Method information
- Laboratory workflow segments
- MI - Method information
- Repository workflow segments
- OD - Operation design - Transaction - Storage

Recommended statistical measure

- Lower range limit (human estimate)
- Upper range limit (human estimate)
- Central or typical value (human estimate)
- Lower range limit (legacy data stat. meth. unknown)
- Upper range limit (legacy data stat. meth. unknown)
- Central or typical value (legacy data stat. meth. unknown)
- Minimum value
- Maximum value
- Mean (= average)
- Harmonic mean
- Geometric mean
- Mode
- Median
- Interquartile mean (= average)
- Variance (sample df = n-1)

Recommended modifier

Recommended frequency

Project

- Descriptor tree
- Descriptor tree node
- Descriptor

Edit descriptors - Assigned to more than one project

Workflow Segment Design

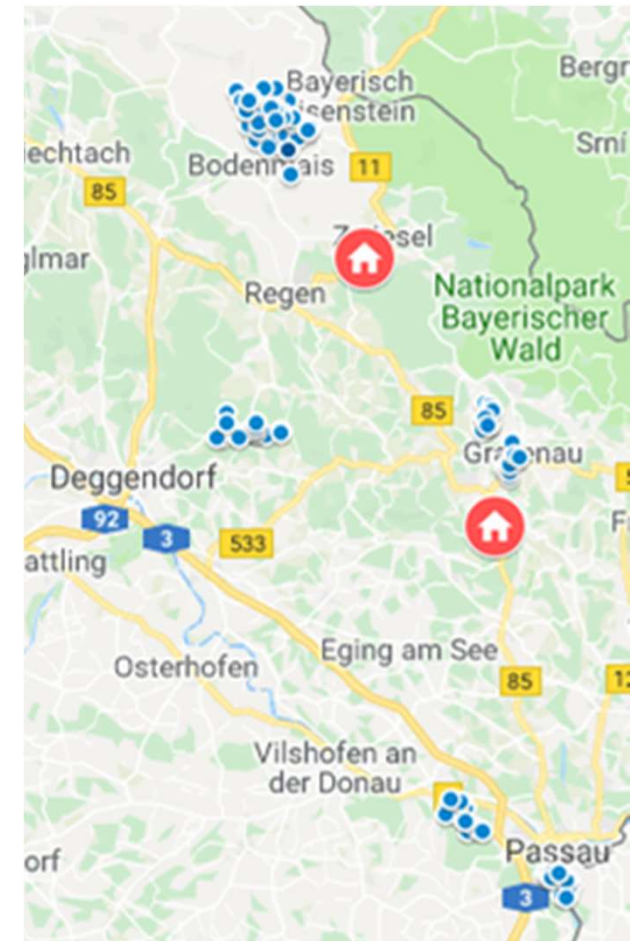
The screenshot displays the DiversityDescriptions v. 4.2.9 application window. The interface is divided into several sections:

- Left Panel:** A list of 92 query results, including items like "47 - Hoehengradient_environmental sample replicate number" and "92 - Waldeck_Index PCR amplicon purification".
- Top Bar:** Contains menu options: Connection, Edit, Grid, Query, Data, Administration, Help.
- Descriptor Panel:** Shows the "Descriptor" for "Project name" with fields for "No: 1" and "ID: 1059003".
- Descriptor Tree:** A hierarchical tree view showing the structure of the descriptor, including nodes like "Project name", "Field workflow segments", "OD - Operation design - Transformation - Mea:", "MI - Method information", "Laboratory workflow segments", "Repository workflow segments", "OD - Operation design - Transaction - Storage", "GBOL2TP12 - Chip design", "Halotolerant fungi - Workflow design", "General", and "Project name".
- Recommended statistical measure:** A list of statistical measures with checkboxes, such as "Lower range limit (human estimate)", "Upper range limit (human estimate)", "Central or typical value (human estimate)", "Lower range limit (legacy data stat. meth. unknown)", "Upper range limit (legacy data stat. meth. unknown)", "Central or typical value (legacy data stat. meth. unknown)", "Minimum value", "Maximum value", "Mean (= average)", "Harmonic mean", "Geometric mean", "Mode", "Median", "Interquartile mean (= average)", and "Variance (sample df = n-1)".
- Recommended modifier:** A section for selecting recommended modifiers.
- Recommended frequency:** A section for selecting recommended frequencies.
- Bottom Bar:** Includes a status bar with "Edit descriptors" and a yellow warning message: "Assigned to more than one project".

A yellow speech bubble with the text "Projektzuordnung" (Project assignment) is overlaid on the "Project name" node in the Descriptor tree.

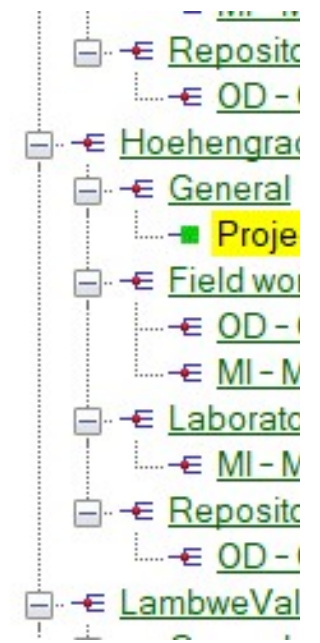
Studie ‚Hoehengradient‘

- Sampling Design hierarchy
 - Sites
 - 8 (Passau, Grafenau, and Bodenmais areas)
 - Plots
 - 144 (18 × 8)
 - Sampling holes per plot
 - 3
 - Horizons per sampling hole
 - 3
 - Replicates
 - 1 per soil horizon
 - Total number of soil samples
 - 1,296



Workflow Segment Design

42 - Halotolerant-fungi_cultivation culture medium type variation level 2
43 - Halotolerant-fungi microbial cultivation
44 - Hoehengradient_environmental operation location
45 - Hoehengradient_soil borehole number
46 - Hoehengradient_soil horizon type
47 - Hoehengradient_environmental sample replicate number
48 - Hoehengradient_environmental sampling
49 - Hoehengradient_nucleic acids extraction
50 - Hoehengradient_nucleic acids extract purification
51 - Hoehengradient_PCR amplification
52 - Hoehengradient_PCR amplicon purification
53 - Hoehengradient_Index PCR amplification
54 - Hoehengradient_Index PCR amplicon purification
55 - LambweValley_environmental operation location
56 - LambweValley_soil borehole number
57 - LambweValley_soil horizon type



Workflow Segment Design

The screenshot displays the DiversityDescriptions v. 4.2.9 application. On the left, a list of descriptors is shown, with '44 - Hohengradient_environmental operation location' selected. A yellow callout bubble points to this descriptor with the text: 'Deskriptor für die Identität der Beprobungsfläche'. The main window shows the details for descriptor 44, including a table of its instances and associated resources.

No.	Categorical state	Abbreviation	Details	Wording
1	136001	136001.x.x.x	4581304; 5439412; Rießloch; ...	
2	136001a	136001a.x.x.x	4584539; 5440597; Rießloch; ...	
3	136001b	136001b.x.x.x	4581738; 5440468; Rießloch; ...	
4	136003	136003.x.x.x	4581687; 5439428; Rießloch; ...	
5	136003a	136001a.x.x.x	4580164; 5439874; Rießloch; ...	
	136003b	136001b.x.x.x	4582332; 5439658; Rießloch; ...	
	136004	136003.x.x.x	4581307; 5439239; Rießloch; ...	
8	136004a	136003a.x.x.x	4579549; 5441485; Rießloch; ...	
9	136004b	136003b.x.x.x	4580337; 5440450; Rießloch; ...	
10	136005	136004.x.x.x	4581485; 5439448; Rießloch; ...	
11	136005a	136004a.x.x.x	4582929; 5436636; Rießloch; ...	
12	136005b	136004b.x.x.x	4579952; 5439868; Rießloch; ...	
13	136006	136005.x.x.x	4581721; 5439251; Rießloch; ...	
14	136006a	136005a.x.x.x	4581926; 5439037; Rießloch; ...	

No.	Resource name	Ranking	Role	URL	Type	Variant
1	136001		unkno...	http://www.google.com/maps/plac...		mediu...

Workflow Segment Design

The screenshot shows the DiversityDescriptions v. 4.2.9 application. The interface is divided into several sections:

- Left Panel:** A list of descriptors, with '48 - Hoehengradient_environmental sampling' selected. A callout bubble points to this list with the text: "Deskriptor für spezifische Operation".
- Top Panel:** A menu bar with options: Connection, Edit, Grid, Query, Data, Administration, Help.
- Center Panel:** A 'Descriptor' view showing a table of selected descriptors. A callout bubble points to this table with the text: "Auswahl zwischen den Elementaroperationen".
- Right Panel:** A 'Details' view for the selected descriptor, showing state resource links and a list of resources. A callout bubble points to this section with the text: "Details".
- Bottom Panel:** A 'Resources' table and a 'URL' field. A callout bubble points to the resources table with the text: "Daten-Resourcen".

No.	Categorical state	Abbreviation	Details
1	HG_ES protocol transformation		Soil corer (Bohrer) 0-30 cm (E...
2	HG_ES protocol measurement		Nokia Lumia 630
3	HG_ES protocol transaction		

No.	Resource name	Ranking	Role
1	modco:unit physical tran...		sk.nar...
2	modco:unit physical tran...		sk.nar...
3	modco:unit physical tran...		sk.nar...
4	modco:unit physical tran...		unkno...
5	modco:unit physical tran...		unkno...
6	Ehlerl und Partner GbR		unkno...

URL	Type	Variant
https://www.mod-co.net/wiki/modc...	applicatio...	tiny sa...

Workflow Segment Design

Art der Elementaroperation

Deskriptor zur DNA-Extraktion

Verbrauchsmittel, Geräte, Literatur

Protokoll

DiversityDescriptions v. 4.2.9

Connection Edit Grid Query Data Administration Help

Query results 1 - 92

- 21 - GBOL2TP12_nucleic acids extraction
- 22 - GBOL2TP12_nucleic acids extract purification
- 23 - GBOL2TP12_PCR amplification
- 24 - GBOL2TP12_PCR amplicon purification
- 25 - GBOL2TP12_Index PCR amplification
- 26 - GBOL2TP12_Index PCR amplicon purification
- 27 - GBOL2TP12_microarray name
- 28 - GBOL2TP12_microarray manufacturer
- 29 - GBOL2TP12_microarray development generation
- 30 - GBOL2TP12_microarray format
- 31 - GBOL2TP12_microarray design identifier internal
- 32 - GBOL2TP12_microarray design identifier external
- 33 - GBOL2TP12_microarray design probe row value
- 34 - GBOL2TP12_microarray design probe column value
- 35 - GBOL2TP12_microarray design probe ID
- 36 - GBOL2TP12_microarray device identifier
- 37 - Halotolerant-fungi_environmental operation location
- 38 - Halotolerant-fungi_environmental sampling
- 39 - Halotolerant-fungi_environmental sample replicate number
- 40 - Halotolerant-fungi_environmental sample horizon type
- 41 - Halotolerant-fungi_environmental sample replicate number
- 42 - Halotolerant-fungi_environmental sample horizon type
- 43 - Halotolerant-fungi_environmental sample replicate number
- 44 - Hoehengradient_soil borehole number
- 45 - Hoehengradient_soil horizon type
- 46 - Hoehengradient_soil horizon type
- 47 - Hoehengradient_environmental operation location
- 48 - Hoehengradient_environmental sampling
- 49 - Hoehengradient_nucleic acids extraction
- 50 - Hoehengradient_nucleic acids extract purification
- 51 - Hoehengradient_PCR amplification
- 52 - Hoehengradient_PCR amplicon purification
- 53 - Hoehengradient_Index PCR amplification
- 54 - Hoehengradient_Index PCR amplicon purification
- 55 - LambweValley_environmental operation location
- 56 - LambweValley_soil borehole number
- 57 - LambweValley_soil horizon type
- 58 - LambweValley_environmental sample replicate number
- 59 - LambweValley_environmental sampling
- 60 - LambweValley_nucleic acids extraction
- 61 - LambweValley_nucleic acids extract purification
- 62 - LambweValley_PCR amplification
- 63 - LambweValley_PCR amplicon purification
- 64 - LambweValley_Index PCR amplification
- 65 - LambweValley_Index PCR amplicon purification
- 66 - Microplastic_environmental operation location

Main workspace: No: 49 ID: 1060449

No.	Categorical s	Abbreviation	Details	Wording
1	HG_NAE protocol transformation			
2	HG_NAE protocol measurement			
3	HG_NAE protocol transaction			

State resource links

Resources

No.	Resource name	Ranking	Role
1	Dopheide, A., Xie, D., B...		unkno...
2	Soil Ex Kit NucleoSpin@...		unkno...
3	NucleoSpin@ Soil (MAC...		unkno...
4	Thermo Savant FastPre...		unkno...

URL: <https://besjournals.onlinelibrary.wiley.com/doi/full/10.1111/2041-210X.13086>

URL: <https://besjournals.onlinelibrary.wiley.com/doi/full/10.1111/2041-210X.13086>

URL: <https://www.mn-net.com/ProductsBioanalysis/DNAandRNApurification>

URL: https://www.mn-net.com/Portals/8/attachments/Redakteure_Bio/Protokoll_Hoehengradient_nucleic_acids_extraction.pdf

URL: https://web.calpoly.edu/~bio/ubl/equip_files/fastprep.pdf

order by: Sequence number

Edit descriptors

Workflow Segment Design

DiversityDescriptions v. 4.2.9

Connection Edit Grid Query Data Administration Help

Descriptor: **Hoehengradient_nucleic acids extract purification**

Descriptor: Hoehengradient_nucleic acids extract purification No: 50 ID: 1065145

Main attributes Categorical states Descriptor tree Modifier/Frequency Resource links

No.	Categorical state	Abbreviation	Details	Wording
1	HG_NAP protocol transformation			
2	HG_NAP protocol measurement		NA concentration [ng/ml]	
3	HG_NAP protocol transaction			

State resource links

Resources

No.	Resource name	Ranking	Role	URL	Type	Variant
1	Dopheide, A., Xie, D., B...		unkno...	https://besjournals.onlinelibrary.wil...	applicatio...	tiny sa...
2	NucleoSpin® Soil (MAC...		unkno...			
3	NucleoSpin® Soil (MAC...		unkno...			
4	Centrifuge 5417R/F45-3...		unkno...			

URL: Size: Width: Height: Duration:

order by: Sequence number

Behalten

Edit descriptors -

Query results 1 - 92

- 21 - GBOL2TP12_nucleic acids extraction
- 22 - GBOL2TP12_nucleic acids extract purification
- 23 - GBOL2TP12_PCR amplification
- 24 - GBOL2TP12_PCR amplicon purification
- 25 - GBOL2TP12_Index PCR amplification
- 26 - GBOL2TP12_Index PCR amplicon purification
- 27 - GBOL2TP12_microarray name
- 28 - GBOL2TP12_microarray manufacturer
- 29 - GBOL2TP12_microarray development generation
- 30 - GBOL2TP12_microarray format
- 31 - GBOL2TP12_microarray design identifier internal
- 32 - GBOL2TP12_microarray design identifier external
- 33 - GBOL2TP12_microarray design probe row value
- 34 - GBOL2TP12_microarray design probe column value
- 35 - GBOL2TP12_microarray design probe ID
- 36 - GBOL2TP12_microarray device identifier
- 37 - Halotolerant-fungi_environmental operation location
- 38 - Halotolerant-fungi_environmental sampling
- 39 - Halotolerant-fungi_cultivation culture generation
- 40 - Halotolerant-fungi_cultivation culture medium type
- 41 - Halotolerant-fungi_cultivation culture medium type variation level 1
- 42 - Halotolerant-fungi_cultivation culture medium type variation level 2
- 43 - Halotolerant-fungi_microbial_cultivation
- 44 - Hoehengradient_environmental operation location
- 45 - Hoehengradient_soil borehole number
- 46 - Hoehengradient_soil horizon type
- 47 - Hoehengradient_environmental sample replicate number
- 48 - Hoehengradient_environmental sampling
- 49 - Hoehengradient_nucleic acids extraction
- 50 - Hoehengradient_nucleic acids extract purification**
- 51 - Hoehengradient_PCR amplification
- 52 - Hoehengradient_PCR amplicon purification
- 53 - Hoehengradient_Index PCR amplification
- 54 - Hoehengradient_Index PCR amplicon purification
- 55 - LambweValley_environmental operation location
- 56 - LambweValley_soil borehole number
- 57 - LambweValley_soil horizon type
- 58 - LambweValley_environmental sample replicate number
- 59 - LambweValley_environmental sampling
- 60 - LambweValley_nucleic acids extraction
- 61 - LambweValley_nucleic acids extract purification
- 62 - LambweValley_PCR amplification
- 63 - LambweValley_PCR amplicon purification
- 64 - LambweValley_Index PCR amplification
- 65 - LambweValley_Index PCR amplicon purification
- 66 - Microplastic_environmental operation location

Workflow Segment Descriptions

DiversityDescriptions v. 4.2.9

Connection Edit Grid Query Data Administration Help

Query results 1 - 4177

32d6d4f3
32e0b385
32f4ba65
32f5ebeb
32fa89ad
3305f24f
33069961
330809a6
33142e69
331ca95e
333bf53a
334a7275
338317e8
33887b34
339a857e
33b7bd62

order by: Description

Query conditions

Description

Title ~

Details ~

Project = UBTMYCoperationsdesc

Description resource

Title ~

Descriptor

Categorical =

Quantitative =

Text =

Sequence =

Data status

+

Scope

Taxon ~

Geography ~

Specimen ~

Reference ~

Description: 33887b34

Project: UBTMYCoperationsdesc

Language: en

ID: 1068533

Deskriptor für Projektzuordnung

No.	Type	!	Descriptor name
1	categoryal	•	Project name
2	categoryal	•	Physical unit type
3	categoryal	•	Physical unit status
4	text	•	Physical unit ID
5	text	•	Physical unit origin ID
6	text		Physical unit preceding ID
7	text		Physical unit external ID
8	categoryal	•	Unit operational agent name
9	categoryal	•	Unit operation workflow segment
10	text		Storage container-internal positio...
11	text		Storage container ID
12	categoryal		Storage shelf number
13	text		Storage rack ID
14	categoryal		Storage device name and ID
15	categoryal		Storage room name
16	categoryal	x	GBOL2TP12_environmental oper...
17	categoryal	x	GBOL2TP12_host tree individual ...
18	categoryal	x	GBOL2TP12_host tree organ
19	quantitative	x	GBOL2TP12_environmental sam...
20	categoryal	x	GBOL2TP12_environmental sam...
21	categoryal	x	GBOL2TP12_nucleic acids extrac...
22	categoryal	x	GBOL2TP12_nucleic acids extrac...
23	categoryal	x	GBOL2TP12_PCR amplification
24	categoryal	x	GBOL2TP12_PCR amplicon purifi...
25	categoryal	x	GBOL2TP12_Index PCR amplific...
26	categoryal	x	GBOL2TP12_Index PCR amplico...

x	No.	State	Notes
<input checked="" type="checkbox"/>	1	GBOL2TP12	
<input type="checkbox"/>	2	HalotolerantFungi	
<input checked="" type="checkbox"/>	3	Hoehengradient	
<input type="checkbox"/>	4	LambweValley	
<input type="checkbox"/>	5	Microplastic	
<input type="checkbox"/>	6	SootyMoulds	
<input type="checkbox"/>	7	Waldeck	

State collection model: OrSet

x	ID	Status	Notes
<input checked="" type="checkbox"/>	!	To be checked	
<input type="checkbox"/>	∅	Not to be recorded	
<input type="checkbox"/>	-	Not applicable	
<input type="checkbox"/>	?	Data unavailable	
<input type="checkbox"/>	#	Not interpretable	
<input type="checkbox"/>	\$	Data withheld	
<input type="checkbox"/>	0	Missing data	

Descriptor tree: Restricted

Edit descriptions

Behalten

Workflow Segment Descriptions

DiversityDescriptions v. 4.2.9

Connection Edit Grid Query Data Administration Help

Query results 1 - 4177

32d6d4f3
32e0b385
32f4ba65
32f5ebcb
32fa89ad
3305f24f
33069961
330809a6
33142e69
331ca95e
333bf53a
334a7275
338317e8
33887b34
339a857e
33b7bd62

order by: Description

Query conditions
Description
Title ~
Details ~
Project = UBTMYCoperationsdesc

Description resource
Title ~

Descriptor
Categorical =
Quantitative =
Text =
Sequence =

Data status
+ +

Scope
Taxon ~
Geography ~
Specimen ~
Reference ~

Description 33887b34 AID: ID: 1068533

http://snb.diversityworkbench.de/Descriptions/1068533

Project UBTMYCoperationsde Language en

Description: 33887b34 AID: ID: 1068533

Main attributes Descriptor view Continuous view Resource links Sample data

No.	Type	!	Descriptor name
38	categorical		Halotolerant-fungi_environmental ...
39	categorical	x	Halotolerant-fungi_cultivation cult...
40	categorical	x	Halotolerant-fungi_cultivation cult...
41	categorical	x	Halotolerant-fungi_cultivation cult...
42	categorical	x	Halotolerant-fungi_cultivation cult...
43	categorical		Halotolerant-fungi_cultivation cult...
44	categorical	•	Halotolerant-fungi_cultivation cult...
45	quantitative	•	Halotolerant-fungi_cultivation cult...
46	categorical	•	Hoehengradient_environmental sa...
47	quantitative	•	Hoehengradient_environmental sa...
48	categorical	•	Hoehengradient_environmental sa...
49	categorical		Hoehengradient_nucleic acids ex...
50	categorical		Hoehengradient_nucleic acids ex...
51	categorical		Hoehengradient_PCR amplificati...
52	categorical		Hoehengradient_PCR amplicon ...
53	categorical		Hoehengradient_Index PCR amp...
54	categorical		Hoehengradient_Index PCR amp...
55	categorical	x	LambweValley_environmental op...
56	quantitative	x	LambweValley_soil borehole nu...
57	categorical	x	LambweValley_soil horizon type
58	quantitative	x	LambweValley_environmental sa...
59	categorical	x	LambweValley_environmental sa...
60	categorical	x	LambweValley_nucleic acids extr...
61	categorical	x	LambweValley_nucleic acids extr...
62	categorical	x	LambweValley_PCR amplification
63	categorical	x	LambweValley_PCR amplicon p...

Descriptor tree: Restricted

Category: !

Categorical states

x	No.	State	Notes
<input checked="" type="checkbox"/>	1	GBOL2TP12	
<input type="checkbox"/>	2	HalotolerantFungi	
<input checked="" type="checkbox"/>	3	Hoehengradient	
<input type="checkbox"/>	4	LambweValley	

State collection model: OrSet

Status data

x	ID	Status	Notes
<input checked="" type="checkbox"/>	!	To be checked	
<input type="checkbox"/>	∅	Not to be recorded	
<input type="checkbox"/>	-	Not applicable	
<input type="checkbox"/>	?	Data unavailable	
<input type="checkbox"/>	#	Not interpretable	
<input type="checkbox"/>	\$	Data withheld	
<input type="checkbox"/>	0	Missing data	

Behalten

Deskriptor für Umweltprobe

Edit descriptions

Workflow Segment Descriptions

The image shows a software interface with several overlapping windows. The main window displays a list of workflow segments. Three yellow arrows originate from the text '93014b. 1. 3. 1' and point to specific entries in the 'Categorical states' and 'Statistical measures' windows.

No.	Type	!	Descriptor name
39	categorical	x	Halotolerant-fungi_cultivation culture generation
40	categorical	x	Halotolerant-fungi_cultivation culture medium type
41	categorical	x	Halotolerant-fungi_cultivation culture medium type variation level 1
42	categorical	x	Halotolerant-fungi_cultivation culture medium type variation level 2
43	categorical		Halotolerant-fungi_microbial_cultivation
44	categorical	•	Hoehengradient_environmental operation location
45	quantitative	•	Hoehengradient_soil borehole number
46	categorical	•	Hoehengradient_soil horizon type
47	quantitative	•	Hoehengradient_environmental sample replicate number
48	categorical	•	Hoehengradient_environmental sampling
49	categorical		Hoehengradient_nucleic acids extraction

x	No.	State
<input type="checkbox"/>	66	93009b
<input type="checkbox"/>	67	93013
<input type="checkbox"/>	68	93013a
<input type="checkbox"/>	69	93013b
<input type="checkbox"/>	70	93014
<input type="checkbox"/>	71	93014a
<input checked="" type="checkbox"/>	72	93014b
<input type="checkbox"/>	73	94004

x	No.	Measure	V
<input checked="" type="checkbox"/>	3	[*] Central or t...	3
<input type="checkbox"/>	9	Mean (= aver...	

x	No.	State
<input type="checkbox"/>	1	HG_plant litter...
<input type="checkbox"/>	2	HG_organic l...
<input checked="" type="checkbox"/>	3	HG_mineral l...

x	No.	Measure	V
<input checked="" type="checkbox"/>	3	Central or typi...	1
<input type="checkbox"/>	9	Mean (= aver...	

Workflow Segment Descriptions

No.	Type	!	Descriptor name
39	categorical	x	Halotolerant-fungi_cultivation culture generation
40	categorical	x	Halotolerant-fungi_cultivation culture medium type
41	categorical	x	Halotolerant-fungi_cultivation culture medium type variation level 1
42	categorical	x	Halotolerant-fungi_cultivation culture medium type variation level 2
43	categorical		Halotolerant-fungi_microbial_cultivation
44	categorical	•	Hoehengradient_environmental operation location
45	quantitative	•	Hoehengradient_soil borehole number
46	categorical	•	Hoehengradient_soil horizon type
47	quantitative	•	Hoehengradient_environmental sample replicate number

x	No.	State
<input type="checkbox"/>	66	93009b
<input type="checkbox"/>	67	93013
<input type="checkbox"/>	68	93013a
<input type="checkbox"/>	69	93013b
<input type="checkbox"/>	70	
<input type="checkbox"/>	71	
<input checked="" type="checkbox"/>	72	
<input type="checkbox"/>	73	

93014b. 1. 3. 1



No.	Type	!	Descriptor name
39	categorical	x	Halotolerant-fungi_cultivation culture generation
40	categorical	x	Halotolerant-fungi_cultivation culture medium type
41	categorical	x	Halotolerant-fungi_cultivation culture medium type variation level 1
42	categorical	x	Halotolerant-fungi_cultivation culture medium type variation level 2
43	categorical		Halotolerant-fungi_microbial_cultivation
44	categorical	•	Hoehengradient_environmental operation location
45	quantitative	•	Hoehengradient_soil borehole number
46	categorical	•	Hoehengradient_soil horizon type
47	quantitative	•	Hoehengradient_environmental sample replicate number
48	categorical	•	Hoehengradient_environmental sampling
49	categorical		Hoehengradient_nucleic acids extraction

No.	Type	!	Descriptor name
39	categorical	x	Halotolerant-fungi_cultivation culture generation
40	categorical	x	Halotolerant-fungi_cultivation culture medium type
41	categorical	x	Halotolerant-fungi_cultivation culture medium type variation level 1
42	categorical	x	Halotolerant-fungi_cultivation culture medium type variation level 2
43	categorical		Halotolerant-fungi_microbial_cultivation
44	categorical	•	Hoehengradient_environmental operation location
45	quantitative	•	Hoehengradient_soil borehole number
46	categorical	•	Hoehengradient_soil horizon type
47	quantitative	•	Hoehengradient_environmental sample replicate number
48	categorical	•	Hoehengradient_environmental sampling
49	categorical		Hoehengradient_nucleic acids extraction

No.	Type	!	Descriptor name
39	categorical	x	Halotolerant-fungi_cultivation culture generation
40	categorical	x	Halotolerant-fungi_cultivation culture medium type
41	categorical	x	Halotolerant-fungi_cultivation culture medium type variation level 1
42	categorical	x	Halotolerant-fungi_cultivation culture medium type variation level 2
43	categorical		Halotolerant-fungi_microbial_cultivation
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45	quantitative	•	Hoehengradient_soil borehole number
46	categorical	•	Hoehengradient_soil horizon type
47	quantitative	•	Hoehengradient_environmental sample replicate number
48	categorical	•	Hoehengradient_environmental sampling
49	categorical		Hoehengradient_nucleic acids extraction

x	No.	Measure	V
<input checked="" type="checkbox"/>	3	Central or typi...	1
<input type="checkbox"/>	9	Mean (= aver...	

Workflow Segment Descriptions

Project: Matches: **432** Show matches 30 s 30 s

Query parameter

Add description Add scope Add descriptor Remove Save Delete

Descriptor: No.:

Categorical states & 123 abc abc

- GBOL2TP12
- HalotolerantFungi
- Hoehengradient**
- LambweValley
- Microplastic
- SootyMoulds

Data status &

- To be checked
- Not to be recorded
- Not applicable
- Data unavailable
- Not interpretable
- Data withheld
- Missing data

Descriptor: No.:

Categorical states & 123 abc abc

- HG_plant litter (mid)
- HG_organic layer (mid of 0 to 5 cm)**
- HG_mineral layer (mid of 5 to 20 cm)

Data status &

- To be checked
- Not to be recorded
- Not applicable
- Data unavailable
- Not interpretable
- Data withheld
- Missing data

Result list

all none swap View in separate window

- 0027fa56
- 002a3b81
- 00b51a51
- 01188bfd
- 01936956
- 019fabfb
- 01a5d11b
- 026f250d
- 02942fbc
- 039ec140
- 03c58821
- 071294ac
- 0820d224
- 086dda8a
- 08a387ef
- 0a8fe6d7
- 0ab96ec1
- 0abb633c
- 0ad145ed
- 0aedd672
- 0b0459a1
- 0c063adb
- 0c7636b6
- 0d1a0ea1
- 0d7a2a9e
- 0d81b21c
- 0df8e41a
- 0f9080d1
- 0fd77d96
- 1015b987
- 113cedb7
- 11d24e80
- 1211b952
- 126cb631
- 133304ee
- 1360e4c4
- 14eb5136
- 1550d1bf1

Workflow Segment Descriptions

DiversityDescriptions

Project:

Descriptor tree:

From descriptor: to descriptor:

Separate quantitative measures

Display description table

Description: **0027fa56**

Descriptor:

Description title	44 Hoehengradient - Workflow design Field workflow segments OD - Operation design - Transformation - Measurement Hoehengradient_environmental operation location	45 Hoehengradient - Workflow design Field workflow segments OD - Operation design - Transformation - Measurement Hoehengradient_soil borehole number	46 Hoehengradient - Workflow design Field workflow segments OD - Operation design - Transformation - Measurement Hoehengradient_soil horizon type	47 Hoehengradient - Workflow design Field workflow segments OD - Operation design - Transformation - Measurement Hoehengradient_environmental sample replicate number
32e0b385	140	3	2	*
331ca95e	66	1	2	*
33db1592	126	1	2	*
34739fdd	48	1	2	*
367d73e5	84	2	2	*
3681e530	18	2	2	*
375062cc	126	3	2	*
384e3f0	100	3	2	*
39d00512	9	1	2	*
3aa8b207	100	1	2	*
3ce743e7	38	3	2	*
3e07ba8f	29	1	2	*
3e660b29	83	3	2	*
3f76e6e	142	3	2	*

Cancel

OK

Workflow Segment Measurements

The screenshot shows the DiversityDescriptions v. 4.2.9 application window. The main interface is titled 'Description' and shows details for descriptor '1e28d61b'. The 'Descriptor view' tab is active, displaying a table of descriptors. A yellow callout bubble labeled 'Deskriptoransicht' points to this tab. Another yellow callout bubble labeled 'Messkampagne' points to the row for 'Unit measurement [num]' (No. 21). The interface includes a left sidebar with a list of descriptors, a 'Query conditions' section, and a 'Statistical measures' table on the right.

No.	Type	!	Descriptor name
1	categorical	•	Project name
2	categorical	•	Unit elementary type
2	categorical	•	Unit physical type
4	text	•	Unit origin sample type
5	text	•	Unit physical ID
6	text	•	Unit digital URI
7	text	•	Unit taxon name
8	text	•	Unit taxon name ID
9	text	•	Unit taxon identification agent name
10	text		Unit taxon identification date
11	text		Unit physical generation date
12	categorical	•	Unit physical generation agent name
13	categorical	•	Unit measurement agent name
14	text		Unit measurement gene locus sequence data
15	text		Unit measurement gene locus sequence data
16	text		Unit measurement gene locus sequence data
17	text		Unit measurement gene locus sequence data
18	text		Unit measurement gene locus sequence data
19	text		Unit measurement gene locus sequence data
20	text		Unit measurement gene locus sequence data
21	quantitative		Unit measurement [num]
22	text		Unit measurement [txt]

x	No.	Measure	Value	Notes
<input checked="" type="checkbox"/>	6	Central or typical value (legacy da...		

x	ID	Status	Notes
<input checked="" type="checkbox"/>	!	To be checked	
<input type="checkbox"/>	∅	Not to be recorded	
<input type="checkbox"/>	-	Not applicable	
<input type="checkbox"/>	?	Data unavailable	
<input type="checkbox"/>	#	Not interpretable	
<input type="checkbox"/>	\$	Data withheld	
<input type="checkbox"/>	0	Missing data	

Workflow Segment Measurements

Messreihen

The screenshot shows a software interface for managing workflow segment measurements. The main window is titled 'Description' and contains several sections:

- Left Sidebar:** A list of IDs, with '14bb1c54' selected.
- Top Menu:** Connection, Edit, Grid, Query, Data, Administration, Help.
- Main Description Area:**
 - Description:** 14bb1c54
 - Main attributes:** Descriptor view, Continuous view, Resource links, Sample data.
 - Sampling events:** A list of events including 'Colony-growth_2018-12-04_2019-01-10', 'Colony-growth_2019-01-15_2019-02-15', and 'Colony-growth_2019-02-19_2019-03-13'.
 - Event data:** Details for 'Colony-growth_2018-12-04_2019-01-10' with ID 1088526. Fields include (Start) date (04.12.2018), End date, Geo. area, Lat. (NS), Long. (EW), Geo. datum, and Geo. notes.
- Sampling units:** A table with 10 units, showing a descriptor tree from 0 to 62.
- Data Table:** A table with columns for Sampling unit ID, Unit physical generation agent name, Unit measurement agent name, Unit measurement gene locus sequence data, Unit measurement gene locus BLAST result, Unit measurement date, Unit measurement time, Unit measurement location coordinates, Unit measurement type, Unit measurement measurement Unit, Unit measurement [µm], and Unit measurement [xt]. The table contains data for units 2989 through 2998.
- Bottom Panel:** Query conditions, Description resource, Descriptor, Data status, and Scope (Taxon, Geography, Specimen).

Messwerte

Reference

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Article Contents

FAIR digital objects in environmental and life sciences should comprise workflow operation design data and method information for repeatability of study setups and reproducibility of results


Janno Harjes, Anton Link, Tanja Weibulat, Dagmar Triebel, Gerhard Rambold 

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