

Package ‘AffyCompatible’

October 16, 2020

Type Package

Title Affymetrix GeneChip software compatibility

Version 1.48.0

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Depends R (>= 2.7.0), XML (>= 2.8-1), RCurl (>= 0.8-1), methods

Imports Biostrings

Description This package provides an interface to Affymetrix chip annotation and sample attribute files. The package allows an easy way for users to download and manage local data bases of Affymetrix NetAffx annotation files. The package also provides access to GeneChip Operating System (GCOS) and GeneChip Command Console (AGCC)-compatible sample annotation files.

License Artistic-2.0

biocViews Infrastructure, Microarray, OneChannel

git_url <https://git.bioconductor.org/packages/AffyCompatible>

git_branch RELEASE_3_11

git_last_commit 6802bd1

git_last_commit_date 2020-04-27

Date/Publication 2020-10-16

R topics documented:

| | |
|--------------------------------------|----|
| AffyCompatible-package | 2 |
| accessors | 2 |
| AffyCompatible-class | 3 |
| Internal-class | 4 |
| NetAffxResource | 4 |
| NetAffxResource-class | 5 |
| readArr | 8 |
| readMage | 8 |
| readNetAffx | 9 |
| readXml | 10 |
| xclass | 11 |
| xmlValue.XMLAttributeValue | 12 |

| | |
|--|----|
| zzz-ARRCompatibility-class | 12 |
| zzz-DTTCompatibility-class | 13 |
| zzz-NetAffxCompatibility-class | 14 |

| | |
|--------------|-----------|
| Index | 16 |
|--------------|-----------|

AffyCompatible-package

Affymetrix chip annotation and sample attribute management

Description

This package manages Affymetrix NetAffx annotation files, and parses MAGE and ARR sample attribute files into R data structures.

NetAffx provides a way to manage the diverse annotation files provided by Affymetrix for their arrays.

ARR files provide attribute information about samples used in an experiment. These files are created, for instance, by the GeneChip Command Console or compliant applications, or

MAGE files also provide information about samples. Support in this package is for MAGE produced by Affymetrix GCOS compliant applications, and emphasizes sample description and processing.

Details

Package details are available in the DESCRIPTION file of the package, or with the R command `packageDescription("AffyCompatible")`

Overviews of how to use the functionality of this package are available in the package vignettes. See `vignette(package="AffyCompatible")`.

Author(s)

Written by Martin Morgan.

Maintainer: Martin Morgan <martin.t.morgan@gmail.com>

accessors

Access NetAffxResource-class content

Description

These functions exist to access the information represented in objects of `linkS4class{NetAffxResource}` and related classes.

Usage

```

annotationsFile(x)
directory(x)
password(x)
user(x)

```

Arguments

x An object for which the accessor is defined.

Value

See, e.g., `help('directory,NetAffxResource-method')` for values returned by specific methods.

Author(s)

Martin Morgan

See Also

[NetAffxResource](#)

AffyCompatible-class *Virtual classes defining the "AffyCompatibility" hierarchy*

Description

The classes `AffyCompatible`, `DTTCompatibility`, `ARRCompatibility`, and `NetAffxCompatibility` define the overall class hierarchy in this package. Information about these classes is not usually relevant to the user.

Objects from the Class

These are virtual classes: no objects may be created from them.

Methods

show signature(object = "AffyCompatible"): for each slot, provide the slot name, and then invoke `show` for the slot content.

Author(s)

Martin Morgan

Examples

```
showClass(getClass("NetAffxCompatibility",
  where=getNamespace("AffyCompatible")))
```

Internal-class *(Internal) Represent 'typed associations' from XML documents*

Description

This internal class represents 'typed associates' (vectors of identically-typed objects) that appear as vectors to the R user.

Objects from the Class

Objects are created internally, during XML document parsing.

Slots

.Data: Object of class "list" containing the association objects
.extends: Object of class "character" and length 1 describing the class each element in .Data must extend

Extends

Class "list", from data part. Class "vector", by class "list", distance 2.

Methods

[signature(x = ".TypedAssociation", i = "numeric", j = "missing"): Subset the typed association, returning a typed association with elements i.
 [signature(x = ".TypedAssociation", i = "ANY", j = "ANY"): This method is present to stop access to subsetting methods for class list; it has no user-visible functionality.
show signature(object = ".TypedAssociation"): Compactly display the typed associate, e.g., for .extends=Foo and length 5, display Foo(5).

Author(s)

Martin Morgan

NetAffxResource *Create and access a NetAffxResource database*

Description

Use this function to create and access a NetAffxResource database.

Usage

```
NetAffxResource(user = character(0),
  password = character(0),
  affxLicence = "FHCRC0607",
  directory = tempdir(), ...)
```

Arguments

| | |
|-------------|---|
| user | Length 1 character vector containing the Affymetrix user name authorized to access the NetAffx resource. |
| password | Length 1 character vector containing the Affymetrix password of the user. |
| affxLicence | Length 1 character vector containing the Affymetrix licence (typically application specific, i.e., the specified licence covers AffyCompatible). |
| directory | (Optional) length 1 character vector providing the full path to the local directory where the NetAffx database and files will be managed. Defaults to a session-specific temporary directory. |
| ... | Additional arguments, not currently used. |

Details

NetAffxResource creates a [NetAffxResource](#) object that manages a data base of NetAffx resources. The data base is located on the user file system as indicated by the directory argument. It is populated when the user requests specific files.

Users should provide a directory argument, so that the data base can be reused across sessions. This means that large files are only downloaded once.

Additional details are provided in the vignette accompanying this package.

Value

[NetAffxResource](#).

Author(s)

Martin Morgan

See Also

[NetAffxResource](#). The vignette ‘Annotations with NetAffx’.

NetAffxResource-class *Manage NetAffx annotations*

Description

This class is for accessing and retrieving NetAffx annotation files for Affymetrix chips. It requires that the user has established an account with Affymetrix. Detailed usage is described in the ‘Annotations with NetAffx’ vignette.

Objects from the Class

Objects are created by calls of the form `NetAffxResource(user="user@research.org", password="terces")`, as illustrated below and in the ‘Annotations with NetAffx’ vignette.

Slots

user: Object of class "character" containing the full user name (typically an email address) obtained when the user registers with Affymetrix.

password: Object of class "character" containing the password associated with the Affymetrix user name. The password is entered and stored in plain text, and should not be considered secure within Bioconductor.

affxLicence: Object of class "character" containing the licence, obtained from Affymetrix, for retrieving NetAffx annotations. The licence is usually software specific (e.g., for the Affy-Compatible package of Bioconductor). Users will not usually change the default value.

directory: Object of class "character", containing the local directory path where the NetAffx data base and annotation files will reside.

annotationsFile: Object of class "character" containing the file name of the NetAffx data base. Users will not usually change the default value.

affxUrl: Object of class "character" containing the URL used to download the NetAffx annotations file. Users will not usually change the default value.

Extends

Class "[NetAffxCompatibility](#)", directly. Class "[AffyCompatible](#)", by class "NetAffxCompatibility", distance 2.

Methods

[[signature(x = "NetAffxResource", i = "numeric", j = "missing")

[[signature(x = "NetAffxResource", i = "character", j = "missing"): Retrieve an object representing all NetAffx annotations available for an array. The argument i may be a numeric index or character string corresponding to an array name returned by names(x).

[[signature(x = "NetAffxResource", i = "numeric", j = "numeric")

[[signature(x = "NetAffxResource", i = "character", j = "numeric")

[[signature(x = "NetAffxResource", i = "numeric", j = "character")

[[signature(x = "NetAffxResource", i = "character", j = "character"): Retrieve an object representing a specific annotation for an array. The argument i may be a numeric index or character string corresponding to an array name returned by names(x). The argument j may be a numeric index or character string corresponding to an annotation returned by description(x[[i]]).

affxLicence signature(x = "NetAffxResource"): the character string representation of the licence used to access the NetAffx site.

affxUrl signature(x = "NetAffxResource"): the url used to access the NetAffx site.

annotationsFile signature(x = "NetAffxResource"): the file name of NetAffx data base, as stored on the local disk of the user.

directory signature(x = "NetAffxResource"): the directory in which the NetAffx data base and annotation files are stored on the local disk of the user.

names signature(x = "NetAffxResource"): the names of the arrays for which annotations are available.

password signature(x = "NetAffxResource"): The password associated with the user name obtained when the user registered with Affymetrix. The password is entered and stored as plain text, and is not secure.

readAnnotation signature(netAffxResource = "NetAffxResource", array = "numeric", annotation = "numeric", ...)

readAnnotation signature(netAffxResource = "NetAffxResource", array = "numeric", annotation = "character", ...)

readAnnotation signature(netAffxResource = "NetAffxResource", array = "character", annotation = "numeric", ...)

readAnnotation signature(netAffxResource = "NetAffxResource", array = "character", annotation = "character", ...)

readAnnotation signature(netAffxResource = "NetAffxResource", array = "missing", annotation = "AffxAnnotation", ...): Read a specific annotation file as an R object. For the first four methods, array, if present, indexes or names the array while annotation indexes or names the annotation. All methods also accept content = "logical" to indicate whether the content of the file should be returned if possible (content=TRUE, default) or only the path to the local file (content=FALSE) and update = "logical" indicating whether the file should be read from disk if available (update=FALSE, default) or retrieved from the NetAffx web site (update=TRUE). readAnnotation parses CSV, Tabular, and PSI files to data frames, and reads FASTA files to a list of description / sequence pairs (see Biostrings::readFASTA).

The ... argument is passed to corresponding read functions. CSV files are parsed using read.csv(conn, ...), Tabular files with read.delim(conn, ...), PSI files with read.delim(conn, header=FALSE) and FASTA with readFASTA(conn, ...).

show signature(object = "NetAffxResource"): compactly show the NetAffxResource object.

user signature(x = "NetAffxResource"): The user name (typically email address) obtained when the user registers with the Affymetrix web site.

Author(s)

Martin Morgan <mtmorgan@fhcrc.org>

See Also

Vignette 'Annotations with NetAffx'. Affymetrix user login site <https://www.affymetrix.com/site/login/login.affx>.

[.Rprofile](#) for ways to automatically set variables (such as password, and directory) for each session.

Examples

```
## password must be obtained by the user
rsrc <- NetAffxResource(user="mtmorgan@fhcrc.org", password=password)
head(names(rsrc))
affxDescription(rsrc[["Bovine"]])
## paradigm to display all elements of 'annotation'
sapply(affxAnnotation(rsrc[["Bovine"]]), force)
## obtain and read a single annotation
anno <- rsrc[["Porcine", "Annotations, CSV format"]]
df <- readAnnotation(rsrc, annotation=anno)
```

readArr *Read ARR sample attribute files*

Description

This functions reads the content of Affymetrix ARR sample attribute files created by GeneChip(R) Command Console. Objects are represented as R classes. Navigation and extraction of data are described in the vignette ‘Retrieving MAGE and ARR sample attributes’.

Usage

```
readArr(fls, ...)
```

Arguments

fls A character vector containing the paths to files to be read.
... Additional arguments, currently unused.

Value

An object or list of objects of class `ArrayFileSet`, depending on whether one or several file paths are provided.

Author(s)

Martin Morgan

See Also

The vignette ‘Retrieving MAGE and ARR sample attributes’ for detailed illustration of use. [xclass](#) and [readXml](#) for advanced access methods.

Examples

```
dir <- system.file("extdata", "ARR", package="AffyCompatible")
fls <- list.files(dir, full=TRUE)
readArr(fls[[1]])
readArr(fls)
```

readMage *Read DTT (MAGE) sample attribute files*

Description

This functions reads the content of Affymetrix DTT (MAGE) sample attribute files created by GCOS. Objects are represented as R classes. Navigation and extraction of data are described in the vignette ‘Retrieving MAGE and ARR sample attributes’.

Usage

```
readMage(fls, ...)
```


Arguments

`fls` A character vector containing the paths to files to be read.
`...` Additional arguments, currently unused.

Value

An object or list of objects of class `ArrayFileSet`, depending on whether one or several file paths are provided.

Author(s)

Martin Morgan

See Also

The vignette ‘Retrieving MAGE and ARR sample attributes’ for detailed illustration of use. [xclass](#) and [readXml](#) for advanced access methods.

Examples

```
dir <- system.file("extdata", "DTT", package="AffyCompatible")
fls <- list.files(dir, full=TRUE)
basename(fl)
readArr(fl[[1]])
```

readNetAffx

Update or read a NetAffx data base

Description

This function provides a way to read or update a NetAffx data base. The usual use for this is to force the data base description to be updated (by accessing the Affymetrix web site) or to retrieve the entire list of NetAffx resource data base. A more normal use case is to

Usage

```
readNetAffx(netAffxResource, update = FALSE, ...)
```

Arguments

`netAffxResource` An object of class [NetAffxResource](#) containing the path to the local data base, and permissions for access the Affymetrix NetAffx resource.
`update` An object of `logical(1)` indicating whether the data base is to be updated (TRUE) or not.
`...` Additional arguments, not currently used.

Value

An object representing the annotations found; see [NetAffxResource](#) for additional details.

Author(s)

Martin Morgan

Examples

```
## retrieve all annotations
## Not run:
example("NetAffxResource-class")
readNetAffx(rsrc)

## End(Not run)
```

| | |
|---------|--|
| readXml | <i>(Advanced) Read XML files of Affymetrix sample attributes or annotation</i> |
|---------|--|

Description

This functions reads the content of Affymetrix DTT (MAGE) or ARR sample files, or AffxNetAffx-AnnotFile array annotation files, return an internal representation of the underlying XML. The return value can be used with [xclass](#) to rapidly create R objects representing deeply nested nodes of the document. The return value can also be used with `xpathApply` in the XML package for access to character-based representation of the data.

Usage

```
readXml(f1, ...)
```

Arguments

| | |
|-----|--|
| f1 | A length-1 character vector giving the full path to the file to be parsed. |
| ... | Additional arguments, passed to <code>xmlTreeParse</code> . |

Value

An object of class `XMLInternalDocument`.

Author(s)

Martin Morgan

See Also

The vignettes ‘Retrieving MAGE and ARR sample attributes’ and ‘Annotation retrieval with NetAffxResource’ for detailed illustration of use.

[xclass](#) for instantiating objects from xpath queries.

Examples

```
mageDir <- system.file("extdata", "DTT", package="AffyCompatible")
mageFiles <- list.files(mageDir, full=TRUE)
xml <- readXml(mageFiles[[1]])
xclass(xml, "//MeasuredBioAssay")[[1]]
```

xclass *(Advanced) Construct R objects from XML document queries*

Description

This function takes an xml object, created from [readXml](#), and an xpath query referencing an element of the document, and returns an R object representing the element.

The xpath query must retrieve an element, not an attribute.

Usage

```
xclass(doc, xpathq, prefix = character(0))
```

Arguments

| | |
|--------|--|
| doc | An XMLInternalDocument retrieved using readXml . |
| xpathq | A length-1 character string representing an xpath query into the document. |
| prefix | A prefix used to map, when necessary, between the XML class name and the corresponding R class. For NetAffx-based classes, the prefix is 'Affx'; no prefix is required for DTT or ARR documents. |

Value

An R object corresponding to the class(es) referenced by the xpath query.

Author(s)

Martin Morgan

References

<http://www.w3.org/TR/xpath> provides a very useful description of xpath. Additional references are in the vignette 'Retrieving MAGE and ARR sample attributes'.

See Also

The vignettes 'Retrieving MAGE and ARR sample attributes' and 'Annotation retrieval with NetAffxResource' for detailed illustration of use.

[readXml](#) for appropriately reading XML documents.

Examples

```
example(readXml)
```

```
xmlValue.XMLAttributeValue
```

Additional functions defined in AffyCompatible

Description

This page documents additional, miscellaneous, functions defined by AffyCompatible

Usage

```
## S3 method for class 'XMLAttributeValue'
xmlValue(x, ignoreComments = FALSE, recursive = TRUE,
         encoding=XML:::CE_NATIVE, trim=FALSE)
```

Arguments

| | |
|----------------|---|
| x | An object of class XMLAttributeValue. |
| ignoreComments | Logical; ignored by the xmlValue.XMLAttributeValue method |
| recursive | ignored |
| encoding | File encoding |
| trim | Remove trailing whitespace? Ignored by this method |

Value

A character vector (usually length 1) containing the character representation of the XML attribute.

```
zzz-ARRCompatibility-class
```

(Auto-generated) classes and methods for navigating ARR sample attributes.

Description

These classes and methods facilitate navigation of the Affymetrix ARR sample attribute files. ARR files are generated by the Affymetrix AGCC (GeneChip(R) Command Console) application. Specifically, the classes are R representations of the Affymetrix classes defined in ArraySetAndTemplate-File.dtd. The 'show' methods for these classes indicate how objects are to be accessed; details are in the 'Retrieving MAGE and ARR sample attributes' vignette.

Objects from the Class

Objects are created internally, by parsing XML retrieved from ARR, and are not created directly by the user.

Slots

Objects have a variety of slots. To discover slot names and content type, see the slotNames portion of the example, below.

Extends

Class "*ARRCompatibility*", directly. Class "*AffyCompatible*", by class "*ARRCompatibility*", distance 2.

Methods

'Accessor' methods are defined on auto-generated classes to access slot content.

Accessors are apparent from displaying the object; each line of the display is of the form accessor : content. To retrieve the content of an instance object, invoke accessor(object).

Some objects contains several instances of another object. These 'typed associations' are represented by the *.TypedAssociation* class, and are navigated or subset as one would a standard R vector, e.g., using [, [[. A useful paradigm for displaying all instances is illustrated in the vignette.

Author(s)

Martin Morgan <mtmorgan@fhcrc.org>

See Also

Vignette 'Retrieving MAGE and ARR sample attributes'.

Examples

```
## defined and auto-generated classes
names(slot(getClass("ARRCompatibility",
                  where=getNamespace("AffyCompatible")),
       "subclasses"))
```

*ztt-*DTTCompatibility-class**

(Auto-generated) classes and methods for navigating DTT sample attributes.

Description

These classes and methods facilitate navigation of the Affymetrix DTT sample attribute files. DTT files are generated by the Affymetrix GCOS application. Specifically, the classes are R representations of the Affymetrix classes defined in MAGE-ML.dtd. The 'show' methods for these class indicate how objects are to be accessed; details are in the 'Retrieving MAGE and ARR sample attributes' vignette.

Objects from the Class

Objects are created internally, by parsing XML retrieved from ARR, and are not created directly by the user.

Slots

Objects have a variety of slots. To discover slot names and content type, see the slotNames portion of the example, below.

Extends

Class "DTTCompatibility", directly. Class "AffyCompatible", by class "DTTCompatibility", distance 2.

Methods

'Accessor' methods are defined on auto-generated classes to access slot content.

Accessors are apparent from displaying the object; each line of the display is of the form accessor : content. To retrieve the content of an instance object, invoke accessor(object).

Some objects contains several instances of another object. These 'typed associations' are represented by the `.TypedAssociation` class, and are navigated or subset as one would a standard R vector, e.g., using `[, [[`. A useful paradigm for displaying all instances is illustrated in the vignette.

Author(s)

Martin Morgan <mtmorgan@fhcrc.org>

See Also

Vignette 'Retrieving MAGE and ARR sample attributes'.

Examples

```
## defined and auto-generated classes
names(slot(getClass("DTTCompatibility",
                  where=getNamespace("AffyCompatible")),
       "subclasses"))
```

zzz-NetAffxCompatibility-class

(Auto-generated) classes and methods for navigating NetAffx resources

Description

These classes and methods facilitate navigation of the NetAffx resource data base. They are R representations of the Affymetrix classes defined in NetAffxAnnotFileList.dtd. The 'show' methods for these class indicate how objects are to be accessed; details are in the 'Annotations with NetAffx' vignette.

The auto-generated classes are discoverable as indicated in the example below.

Objects from the Class

Objects are created internally, by parsing XML retrieved from the local NetAffx data base or from the Affymetrix web site.

Slots

Objects have a variety of slots. To discover slot names and content type, see the slotNames portion of the example, below.

Extends

Class "[NetAffxCompatibility](#)", directly. Class "[AffyCompatible](#)", by class "NetAffxCompatibility", distance 2.

Methods

'Read' methods are defined on auto-generated classes to access slot content.

Accessors are apparent from displaying the object; each line of the display is of the form accessor: content. To retrieve the content of an instance object, invoke accessor(object).

Some objects contains several instances of another object. These 'typed associations' are represented by the [.TypedAssociation](#) class, and are navigated or subset as one would a standard R vector, e.g., using [, [[]]. A useful paradigm for displaying all instances is illustrated on the [NetAffxResource](#) page.

Author(s)

Martin Morgan <mtmorgan@fhcrc.org>

See Also

Vignette 'Annotations with NetAffx'. Affymetrix user login site <https://www.affymetrix.com/site/login/login.affx>.

[NetAffxResource](#) for creating and managing the NetAffx data base.

[NetAffxResource](#) for navigating the data base.

Examples

```
## defined and auto-generated classes
getClass("NetAffxCompatibility", where=getNamespace("AffyCompatible"))

## slots of an auto-generated class
getSlots(getClass("AffxNetAffxAnnotFileList",
  where=getNamespace("AffyCompatible")))

## Not run:
example("NetAffxrResource-class")

## End(Not run)
```

Index

- * **classes**
 - AffyCompatible-class, 3
 - Internal-class, 4
 - NetAffxResource-class, 5
 - readArr, 8
 - readMage, 8
 - zzz-ARRCompatibility-class, 12
 - zzz-DTTCompatibility-class, 13
 - zzz-NetAffxCompatibility-class, 14
- * **database**
 - NetAffxResource, 4
 - readNetAffx, 9
- * **manip**
 - readXml, 10
 - xclass, 11
 - xmlValue.XMLAttributeValue, 12
- * **methods**
 - accessors, 2
- * **package**
 - AffyCompatible-package, 2
 - .Rprofile, 7
 - .TypedAssociation, 13–15
 - .TypedAssociation-class
 - (Internal-class), 4
 - [, .TypedAssociation, ANY, ANY-method
 - (Internal-class), 4
 - [, .TypedAssociation, numeric, missing, ANY-method
 - (Internal-class), 4
 - [, .TypedAssociation, numeric, missing-method
 - (Internal-class), 4
 - [[, NetAffxResource, character, character-method
 - (NetAffxResource-class), 5
 - [[, NetAffxResource, character, missing-method
 - (NetAffxResource-class), 5
 - [[, NetAffxResource, character, numeric-method
 - (NetAffxResource-class), 5
 - [[, NetAffxResource, numeric, character-method
 - (NetAffxResource-class), 5
 - [[, NetAffxResource, numeric, missing-method
 - (NetAffxResource-class), 5
 - [[, NetAffxResource, numeric, numeric-method
 - (NetAffxResource-class), 5
- accession (zzz-DTTCompatibility-class), 13
- accession, DatabaseEntry-method
 - (zzz-DTTCompatibility-class), 13
- accessions_assnlist
 - (zzz-DTTCompatibility-class), 13
- accessions_assnlist, BibliographicReference-method
 - (zzz-DTTCompatibility-class), 13
- Accessions_assnlist-class
 - (zzz-DTTCompatibility-class), 13
- accessionVersion
 - (zzz-DTTCompatibility-class), 13
- accessionVersion, DatabaseEntry-method
 - (zzz-DTTCompatibility-class), 13
- accessors, 2
- acpassword (NetAffxResource), 4
- action (zzz-DTTCompatibility-class), 13
- action, Audit-method
 - (zzz-DTTCompatibility-class), 13
- action_assn
 - (zzz-DTTCompatibility-class), 13
- action_assn, Treatment-method
 - (zzz-DTTCompatibility-class), 13
- Action_assn-class
 - (zzz-DTTCompatibility-class), 13
- actionMeasurement_assn
 - (zzz-DTTCompatibility-class), 13
- actionMeasurement_assn, Treatment-method
 - (zzz-DTTCompatibility-class), 13
- ActionMeasurement_assn-class
 - (zzz-DTTCompatibility-class), 13

- activityDate
 - (zzz-DTTCompatibility-class),
13
- activityDate,ProtocolApplication-method
 - (zzz-DTTCompatibility-class),
13
- address(zzz-DTTCompatibility-class), 13
- address,Organization-method
 - (zzz-DTTCompatibility-class),
13
- address,Person-method
 - (zzz-DTTCompatibility-class),
13
- adjustments_assnlist
 - (zzz-DTTCompatibility-class),
13
- adjustments_assnlist,ArrayManufactureDeviation-method(zzz-DTTCompatibility-class),
13
- Adjustments_assnlist-class
 - (zzz-DTTCompatibility-class),
13
- affiliation_assnref
 - (zzz-DTTCompatibility-class),
13
- affiliation_assnref,Person-method
 - (zzz-DTTCompatibility-class),
13
- Affiliation_assnref-class
 - (zzz-DTTCompatibility-class),
13
- affxAnnotation
 - (zzz-NetAffxCompatibility-class),
14
- affxAnnotation,AffxArray-method
 - (zzz-NetAffxCompatibility-class),
14
- AffxAnnotation-class
 - (zzz-NetAffxCompatibility-class),
14
- affxArray
 - (zzz-NetAffxCompatibility-class),
14
- affxArray,AffxNetAffxAnnotFileList-method
 - (zzz-NetAffxCompatibility-class),
14
- affxArray,Array_assnlist-method
 - (zzz-DTTCompatibility-class),
13
- AffxArray-class
 - (zzz-NetAffxCompatibility-class),
14
- affxCategory
 - (zzz-DTTCompatibility-class),
13
- affxCategory,OntologyEntry-method
 - (zzz-DTTCompatibility-class),
13
- affxComment
 - (zzz-NetAffxCompatibility-class),
14
- affxComment,AffxAnnotation-method
 - (zzz-NetAffxCompatibility-class),
14
- affxComment,PhysicalArray-method
 - (zzz-ARRCompatibility-class),
12
- AffxComment-class
 - (zzz-NetAffxCompatibility-class),
14
- affxCompression
 - (zzz-NetAffxCompatibility-class),
14
- affxCompression,AffxUrl-method
 - (zzz-NetAffxCompatibility-class),
14
- affxCrc32
 - (zzz-NetAffxCompatibility-class),
14
- affxCrc32,AffxFile-method
 - (zzz-NetAffxCompatibility-class),
14
- affxCrc32,AffxUrl-method
 - (zzz-NetAffxCompatibility-class),
14
- affxDate
 - (zzz-NetAffxCompatibility-class),
14
- affxDate,AffxFile-method
 - (zzz-NetAffxCompatibility-class),
14
- affxDate,Audit-method
 - (zzz-DTTCompatibility-class),
13
- affxDescription
 - (zzz-NetAffxCompatibility-class),
14
- affxDescription,AffxAnnotation-method
 - (zzz-NetAffxCompatibility-class),
14
- affxDescription,AffxArray-method
 - (zzz-NetAffxCompatibility-class),
14
- affxEnd(zzz-DTTCompatibility-class), 13

- affxEnd, CompositePosition-method
(zzz-DTTCompatibility-class),
13
- affxEnd, ReporterPosition-method
(zzz-DTTCompatibility-class),
13
- affxEnd, SequencePosition-method
(zzz-DTTCompatibility-class),
13
- affxFile
(zzz-NetAffxCompatibility-class),
14
- affxFile, AffxAnnotation-method
(zzz-NetAffxCompatibility-class),
14
- AffxFile-class
(zzz-NetAffxCompatibility-class),
14
- affxImage (zzz-DTTCompatibility-class),
13
- affxImage, PhysicalBioAssayData_assnlist-method
(zzz-DTTCompatibility-class),
13
- affxLicence
(zzz-NetAffxCompatibility-class),
14
- affxLicence, NetAffxResource-method
(NetAffxResource-class), 5
- affxName
(zzz-NetAffxCompatibility-class),
14
- affxName, AffxArray-method
(zzz-NetAffxCompatibility-class),
14
- affxName, AffxFile-method
(zzz-NetAffxCompatibility-class),
14
- AffxNetAffxAnnotFileList-class
(zzz-NetAffxCompatibility-class),
14
- affxOrder (zzz-DTTCompatibility-class),
13
- affxOrder, BioDataCube-method
(zzz-DTTCompatibility-class),
13
- affxOrder, Treatment-method
(zzz-DTTCompatibility-class),
13
- affxPerson
(zzz-DTTCompatibility-class),
13
- affxPerson, Contact_assnlist-method
(zzz-DTTCompatibility-class),
13
- affxPerson, Person_assnlist-method
(zzz-DTTCompatibility-class),
13
- affxRow (zzz-DTTCompatibility-class), 13
- affxRow, FeatureLocation-method
(zzz-DTTCompatibility-class),
13
- affxRow, Zone-method
(zzz-DTTCompatibility-class),
13
- affxSequence
(zzz-DTTCompatibility-class),
13
- affxSequence, BioSequence-method
(zzz-DTTCompatibility-class),
13
- affxSize
(zzz-NetAffxCompatibility-class),
14
- affxSize, AffxFile-method
(zzz-NetAffxCompatibility-class),
14
- affxSize, AffxUrl-method
(zzz-NetAffxCompatibility-class),
14
- affxStart (zzz-DTTCompatibility-class),
13
- affxStart, CompositePosition-method
(zzz-DTTCompatibility-class),
13
- affxStart, ReporterPosition-method
(zzz-DTTCompatibility-class),
13
- affxStart, SequencePosition-method
(zzz-DTTCompatibility-class),
13
- affxStatus
(zzz-NetAffxCompatibility-class),
14
- affxStatus, AffxNetAffxAnnotFileList-method
(zzz-NetAffxCompatibility-class),
14
- AffxStatus-class
(zzz-NetAffxCompatibility-class),
14
- affxText (zzz-DTTCompatibility-class),
13
- affxText, Description-method
(zzz-DTTCompatibility-class),
13

- affxText,Protocol-method
(zzz-DTTCompatibility-class),
13
- affxTitle (zzz-DTTCompatibility-class),
13
- affxTitle,BibliographicReference-method
(zzz-DTTCompatibility-class),
13
- affxTitle,Protocol-method
(zzz-DTTCompatibility-class),
13
- affxType
(zzz-NetAffxCompatibility-class),
14
- affxType,AffxAnnotation-method
(zzz-NetAffxCompatibility-class),
14
- affxType,AffxFile-method
(zzz-NetAffxCompatibility-class),
14
- affxUrl
(zzz-NetAffxCompatibility-class),
14
- affxUrl,AffxFile-method
(zzz-NetAffxCompatibility-class),
14
- affxUrl,NetAffxResource-method
(NetAffxResource-class), 5
- AffxUrl-class
(zzz-NetAffxCompatibility-class),
14
- affyBarcode
(zzz-ARRCompatibility-class),
12
- affyBarcode,PhysicalArray-method
(zzz-ARRCompatibility-class),
12
- AffyCompatible, 6, 13–15
- AffyCompatible
(AffyCompatible-package), 2
- AffyCompatible-class, 3
- AffyCompatible-package, 2
- analysisResults_assnreflist
(zzz-DTTCompatibility-class),
13
- analysisResults_assnreflist,Experiment-method
(zzz-DTTCompatibility-class),
13
- AnalysisResults_assnreflist-class
(zzz-DTTCompatibility-class),
13
- annotations_assnlist
(zzz-DTTCompatibility-class),
13
- annotations_assnlist,Description-method
(zzz-DTTCompatibility-class),
13
- annotations_assnlist,ExperimentalFactor-method
(zzz-DTTCompatibility-class),
13
- Annotations_assnlist-class
(zzz-DTTCompatibility-class),
13
- annotationsFile (accessors), 2
- annotationsFile,NetAffxResource-method
(NetAffxResource-class), 5
- Array-class
(zzz-DTTCompatibility-class),
13
- array_assnlist
(zzz-DTTCompatibility-class),
13
- array_assnlist,Array_package-method
(zzz-DTTCompatibility-class),
13
- Array_assnlist-class
(zzz-DTTCompatibility-class),
13
- array_assnref
(zzz-DTTCompatibility-class),
13
- array_assnref,BioAssayCreation-method
(zzz-DTTCompatibility-class),
13
- array_assnref,Hybridization-method
(zzz-DTTCompatibility-class),
13
- Array_assnref-class
(zzz-DTTCompatibility-class),
13
- array_package
(zzz-DTTCompatibility-class),
13
- array_package,MAGE-ML-method
(zzz-DTTCompatibility-class),
13
- Array_package-class
(zzz-DTTCompatibility-class),
13
- array_ref (zzz-DTTCompatibility-class),
13
- array_ref,Array_assnref-method
(zzz-DTTCompatibility-class),
13

- array_ref,Arrays_assnreflist-method
(zzz-DTTCompatibility-class),
13
- Array_ref-class
(zzz-DTTCompatibility-class),
13
- arrayAttribute
(zzz-ARRCompatibility-class),
12
- arrayAttribute,PhysicalArray-method
(zzz-ARRCompatibility-class),
12
- ArrayAttribute-class
(zzz-ARRCompatibility-class),
12
- arrayDesign
(zzz-DTTCompatibility-class),
13
- arrayDesign,ArrayDesign_assnlist-method
(zzz-DTTCompatibility-class),
13
- ArrayDesign-class
(zzz-DTTCompatibility-class),
13
- arrayDesign_assnlist
(zzz-DTTCompatibility-class),
13
- arrayDesign_assnlist,ArrayDesign_package-method
(zzz-DTTCompatibility-class),
13
- ArrayDesign_assnlist-class
(zzz-DTTCompatibility-class),
13
- arrayDesign_assnref
(zzz-DTTCompatibility-class),
13
- arrayDesign_assnref,Array-method
(zzz-DTTCompatibility-class),
13
- ArrayDesign_assnref-class
(zzz-DTTCompatibility-class),
13
- arrayDesign_package
(zzz-DTTCompatibility-class),
13
- arrayDesign_package,MAGE-ML-method
(zzz-DTTCompatibility-class),
13
- ArrayDesign_package-class
(zzz-DTTCompatibility-class),
13
- arrayDesign_ref
(zzz-DTTCompatibility-class),
13
- arrayDesign_ref,ArrayDesign_assnref-method
(zzz-DTTCompatibility-class),
13
- ArrayDesign_ref-class
(zzz-DTTCompatibility-class),
13
- arrayGroup
(zzz-DTTCompatibility-class),
13
- arrayGroup,ArrayGroup_assnlist-method
(zzz-DTTCompatibility-class),
13
- ArrayGroup-class
(zzz-DTTCompatibility-class),
13
- arrayGroup_assnlist
(zzz-DTTCompatibility-class),
13
- arrayGroup_assnlist,Array_package-method
(zzz-DTTCompatibility-class),
13
- ArrayGroup_assnlist-class
(zzz-DTTCompatibility-class),
13
- arrayGroup_assnref
(zzz-DTTCompatibility-class),
13
- arrayGroup_assnref,Array-method
(zzz-DTTCompatibility-class),
13
- ArrayGroup_assnref-class
(zzz-DTTCompatibility-class),
13
- arrayGroup_ref
(zzz-DTTCompatibility-class),
13
- arrayGroup_ref,ArrayGroup_assnref-method
(zzz-DTTCompatibility-class),
13
- ArrayGroup_ref-class
(zzz-DTTCompatibility-class),
13
- arrayIdentifier
(zzz-DTTCompatibility-class),
13
- arrayIdentifier,Array-method
(zzz-DTTCompatibility-class),
13
- arrayManufacture
(zzz-DTTCompatibility-class),

- 13
- arrayManufacture, ArrayManufacture_assnlist-method 12
- (zzz-DTTCompatibility-class),
- 13
- ArrayManufacture-class
- (zzz-DTTCompatibility-class),
- 13
- arrayManufacture_assnlist
- (zzz-DTTCompatibility-class),
- 13
- arrayManufacture_assnlist, Array_package-method
- (zzz-DTTCompatibility-class),
- 13
- ArrayManufacture_assnlist-class
- (zzz-DTTCompatibility-class),
- 13
- arrayManufacture_ref
- (zzz-DTTCompatibility-class),
- 13
- arrayManufacture_ref, Information_assnref-method
- (zzz-DTTCompatibility-class),
- 13
- ArrayManufacture_ref-class
- (zzz-DTTCompatibility-class),
- 13
- arrayManufactureDeviation
- (zzz-DTTCompatibility-class),
- 13
- arrayManufactureDeviation, ArrayManufactureDeviations_assnlist-method
- (zzz-DTTCompatibility-class),
- 13
- ArrayManufactureDeviation-class
- (zzz-DTTCompatibility-class),
- 13
- arrayManufactureDeviations_assnlist
- (zzz-DTTCompatibility-class),
- 13
- arrayManufactureDeviations_assnlist, Array-method
- (zzz-DTTCompatibility-class),
- 13
- ArrayManufactureDeviations_assnlist-class
- (zzz-DTTCompatibility-class),
- 13
- arrayManufacturers_assnreflist
- (zzz-DTTCompatibility-class),
- 13
- arrayManufacturers_assnreflist, ArrayManufacture-method
- (zzz-DTTCompatibility-class),
- 13
- ArrayManufacturers_assnreflist-class
- (zzz-DTTCompatibility-class),
- 13
- arrayName (zzz-ARRCompatibility-class),
- 12
- arrayName, PhysicalArray-method
- (zzz-ARRCompatibility-class),
- 12
- arrays_assnreflist
- (zzz-DTTCompatibility-class),
- 13
- arrays_assnreflist, ArrayGroup-method
- (zzz-DTTCompatibility-class),
- 13
- arrays_assnreflist, ArrayManufacture-method
- (zzz-DTTCompatibility-class),
- 13
- Arrays_assnreflist-class
- (zzz-DTTCompatibility-class),
- 13
- ArraySetFile-class
- (zzz-ARRCompatibility-class),
- 12
- arraySpacingX
- (zzz-DTTCompatibility-class),
- 13
- arraySpacingX, ArrayGroup-method
- (zzz-DTTCompatibility-class),
- 13
- arraySpacingY
- (zzz-DTTCompatibility-class),
- 13
- arraySpacingY, ArrayGroup-method
- (zzz-DTTCompatibility-class),
- 13
- arrayXOrigin
- (zzz-DTTCompatibility-class),
- 13
- arrayXOrigin, Array-method
- (zzz-DTTCompatibility-class),
- 13
- arrayYOrigin
- (zzz-DTTCompatibility-class),
- 13
- arrayYOrigin, Array-method
- (zzz-DTTCompatibility-class),
- 13
- ARRCompatibility, 13
- ARRCompatibility-class
- (AffyCompatible-class), 3
- associations_assnlist
- (zzz-DTTCompatibility-class),
- 13
- associations_assnlist, OntologyEntry-method
- (zzz-DTTCompatibility-class),

- 13
- Associations_assnlist-class
(zzz-DTTCompatibility-class),
13
- audit (zzz-DTTCompatibility-class), 13
- audit, AuditTrail_assnlist-method
(zzz-DTTCompatibility-class),
13
- Audit-class
(zzz-DTTCompatibility-class),
13
- auditAndSecurity_package
(zzz-DTTCompatibility-class),
13
- auditAndSecurity_package, MAGE-ML-method
(zzz-DTTCompatibility-class),
13
- AuditAndSecurity_package-class
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist, Array-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist, ArrayDesign-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist, ArrayGroup-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist, ArrayManufacture-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist, Audit-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist, BibliographicReference-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist, BioAssayCreation-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist, BioAssayDataCluster-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist, BioAssayDimension-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist, BioAssayMap-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist, BioAssayTreatment-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist, BioSample-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist, BioSequence-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist, BioSource-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist, Channel-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist, CompositeCompositeMap-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist, CompositeGroup-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist, CompositeSequence-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist, CompositeSequenceDimension-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist, Compound-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist, Database-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist, DerivedBioAssay-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist, DerivedBioAssayData-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist, DerivedSignal-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist, Description-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist, Error-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist, ExpectedValue-method
(zzz-DTTCompatibility-class),
13

- auditTrail_assnlist,Experiment-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,ExperimentalFactor-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,ExperimentDesign-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,FactorValue-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,Failed-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,Feature-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,FeatureDimension-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,FeatureExtraction-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,FeatureGroup-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,FeatureReporterMap-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,Fiducial-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,Hardware-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,HardwareApplication-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,Hybridization-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,Image-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,ImageAcquisition-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,LabeledExtract-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,MAGE-ML-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,ManufactureLIMS-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,ManufactureLIMSBiomaterial-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,MeasuredBioAssay-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,MeasuredBioAssayData-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,MeasuredSignal-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,Node-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,NodeContents-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,Organization-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,Parameter-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,Person-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,PhysicalArrayDesign-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,PhysicalBioAssay-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,PresentAbsent-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,Protocol-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,ProtocolApplication-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,PValue-method
(zzz-DTTCompatibility-class),
[13](#)
- auditTrail_assnlist,QuantitationTypeDimension-method
(zzz-DTTCompatibility-class),
[13](#)

- 13
- auditTrail_assnlist,QuantitationTypeMap-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist,Ratio-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist,Reporter-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist,ReporterCompositeMap-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist,ReporterDimension-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist,ReporterGroup-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist,Security-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist,SecurityGroup-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist,SeqFeature-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist,Software-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist,SoftwareApplication-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist,SpecializedQuantitationType-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist,Transformation-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist,Treatment-method
(zzz-DTTCompatibility-class),
13
- auditTrail_assnlist,Zone-method
(zzz-DTTCompatibility-class),
13
- AuditTrail_assnlist-class
(zzz-DTTCompatibility-class),
13
- authors (zzz-DTTCompatibility-class), 13
- authors,BibliographicReference-method
(zzz-DTTCompatibility-class),
13
- 13
- barcode (zzz-DTTCompatibility-class), 13
- barcode,ArrayGroup-method
(zzz-DTTCompatibility-class),
13
- basis (zzz-DTTCompatibility-class), 13
- basis,SeqFeature-method
(zzz-DTTCompatibility-class),
13
- bibliographicReference
(zzz-DTTCompatibility-class),
13
- bibliographicReference,BibliographicReferences_assnlist
(zzz-DTTCompatibility-class),
13
- BibliographicReference-class
(zzz-DTTCompatibility-class),
13
- bibliographicReferences_assnlist
(zzz-DTTCompatibility-class),
13
- bibliographicReferences_assnlist,Description-method
(zzz-DTTCompatibility-class),
13
- BibliographicReferences_assnlist-class
(zzz-DTTCompatibility-class),
13
- bioAssay_assnlist
(zzz-DTTCompatibility-class),
13
- bioAssay_assnlist,BioAssay_package-method
(zzz-DTTCompatibility-class),
13
- BioAssay_assnlist-class
(zzz-DTTCompatibility-class),
13
- bioAssay_assnref
(zzz-DTTCompatibility-class),
13
- bioAssay_assnref,BioAssayTuple-method
(zzz-DTTCompatibility-class),
13
- BioAssay_assnref-class
(zzz-DTTCompatibility-class),
13
- bioAssay_package
(zzz-DTTCompatibility-class),
13
- bioAssay_package,MAGE-ML-method
(zzz-DTTCompatibility-class),
13
- BioAssay_package-class

- (zzz-DTTCompatibility-class),
13
- BioAssay_ref-class
(zzz-DTTCompatibility-class),
13
- bioAssayCreation
(zzz-DTTCompatibility-class),
13
- bioAssayCreation,BioAssayCreation_assn-method
(zzz-DTTCompatibility-class),
13
- BioAssayCreation-class
(zzz-DTTCompatibility-class),
13
- bioAssayCreation_assn
(zzz-DTTCompatibility-class),
13
- bioAssayCreation_assn,PhysicalBioAssay-method
(zzz-DTTCompatibility-class),
13
- BioAssayCreation_assn-class
(zzz-DTTCompatibility-class),
13
- bioAssayData_assnlist
(zzz-DTTCompatibility-class),
13
- bioAssayData_assnlist,BioAssayData_package-method
(zzz-DTTCompatibility-class),
13
- BioAssayData_assnlist-class
(zzz-DTTCompatibility-class),
13
- bioAssayData_assnreflist
(zzz-DTTCompatibility-class),
13
- bioAssayData_assnreflist,Experiment-method
(zzz-DTTCompatibility-class),
13
- BioAssayData_assnreflist-class
(zzz-DTTCompatibility-class),
13
- bioAssayData_package
(zzz-DTTCompatibility-class),
13
- bioAssayData_package,MAGE-ML-method
(zzz-DTTCompatibility-class),
13
- BioAssayData_package-class
(zzz-DTTCompatibility-class),
13
- BioAssayData_ref-class
(zzz-DTTCompatibility-class),
13
- bioAssayDataCluster
(zzz-DTTCompatibility-class),
13
- bioAssayDataCluster,BioAssayDataCluster_assnlist-method
(zzz-DTTCompatibility-class),
13
- BioAssayDataCluster-class
(zzz-DTTCompatibility-class),
13
- bioAssayDataCluster_assnlist
(zzz-DTTCompatibility-class),
13
- bioAssayDataCluster_assnlist,HigherLevelAnalysis_package
(zzz-DTTCompatibility-class),
13
- BioAssayDataCluster_assnlist-class
(zzz-DTTCompatibility-class),
13
- bioAssayDataCluster_ref
(zzz-DTTCompatibility-class),
13
- bioAssayDataCluster_ref,AnalysisResults_assnreflist-method
(zzz-DTTCompatibility-class),
13
- BioAssayDataCluster_ref-class
(zzz-DTTCompatibility-class),
13
- bioAssayDataSources_assnreflist
(zzz-DTTCompatibility-class),
13
- bioAssayDataSources_assnreflist,Transformation-method
(zzz-DTTCompatibility-class),
13
- BioAssayDataSources_assnreflist-class
(zzz-DTTCompatibility-class),
13
- bioAssayDimension
(zzz-DTTCompatibility-class),
13
- bioAssayDimension,BioAssayDimension_assnlist-method
(zzz-DTTCompatibility-class),
13
- BioAssayDimension-class
(zzz-DTTCompatibility-class),
13
- bioAssayDimension_assnlist
(zzz-DTTCompatibility-class),
13
- bioAssayDimension_assnlist,BioAssayData_package-method
(zzz-DTTCompatibility-class),
13

BioAssayDimension_assnlist-class
 (zzz-DTTCompatibility-class),
 13
 (zzz-DTTCompatibility-class),
 13
 bioAssayDimension_assnref
 (zzz-DTTCompatibility-class),
 13
 bioAssayDimension_assnref,DerivedBioAssayData-method
 (zzz-DTTCompatibility-class),
 13
 bioAssayDimension_assnref,MeasuredBioAssayData-method
 (zzz-DTTCompatibility-class),
 13
 bioAssayDimension_assnref,NodeContents-method
 (zzz-DTTCompatibility-class),
 13
 BioAssayDimension_assnref-class
 (zzz-DTTCompatibility-class),
 13
 bioAssayDimension_ref
 (zzz-DTTCompatibility-class),
 13
 bioAssayDimension_ref,BioAssayDimension_assnref-method
 (zzz-DTTCompatibility-class),
 13
 BioAssayDimension_ref-class
 (zzz-DTTCompatibility-class),
 13
 bioAssayFactorValues_assnreflist
 (zzz-DTTCompatibility-class),
 13
 bioAssayFactorValues_assnreflist,DerivedBioAssay-method
 (zzz-DTTCompatibility-class),
 13
 bioAssayFactorValues_assnreflist,MeasuredBioAssay-method
 (zzz-DTTCompatibility-class),
 13
 bioAssayFactorValues_assnreflist,PhysicalBioAssay-method
 (zzz-DTTCompatibility-class),
 13
 BioAssayFactorValues_assnreflist-class
 (zzz-DTTCompatibility-class),
 13
 bioAssayMap
 (zzz-DTTCompatibility-class),
 13
 bioAssayMap,BioAssayMap_assnlist-method
 (zzz-DTTCompatibility-class),
 13
 BioAssayMap-class
 (zzz-DTTCompatibility-class),
 13
 bioAssayMap_assnlist
 (zzz-DTTCompatibility-class),
 13
 (zzz-DTTCompatibility-class),
 13
 bioAssayMap_assnlist,BioAssayData-package-method
 (zzz-DTTCompatibility-class),
 13
 BioAssayMap_assnlist-class
 (zzz-DTTCompatibility-class),
 13
 bioAssayMap_ref
 (zzz-DTTCompatibility-class),
 13
 bioAssayMap_ref,BioAssayMaps_assnreflist-method
 (zzz-DTTCompatibility-class),
 13
 bioAssayMap_ref,DerivedBioAssayMap_assnreflist-method
 (zzz-DTTCompatibility-class),
 13
 BioAssayMap_ref-class
 (zzz-DTTCompatibility-class),
 13
 bioAssayMapping
 (zzz-DTTCompatibility-class),
 13
 bioAssayMapping,BioAssayMapping_assn-method
 (zzz-DTTCompatibility-class),
 13
 BioAssayMapping-class
 (zzz-DTTCompatibility-class),
 13
 bioAssayMapping_assn
 (zzz-DTTCompatibility-class),
 13
 bioAssayMapping_assn,Transformation-method
 (zzz-DTTCompatibility-class),
 13
 BioAssayMapping_assn-class
 (zzz-DTTCompatibility-class),
 13
 bioAssayMaps_assnreflist
 (zzz-DTTCompatibility-class),
 13
 bioAssayMaps_assnreflist,BioAssayMapping-method
 (zzz-DTTCompatibility-class),
 13
 BioAssayMaps_assnreflist-class
 (zzz-DTTCompatibility-class),
 13
 bioAssayMapTarget_assnref
 (zzz-DTTCompatibility-class),
 13
 bioAssayMapTarget_assnref,BioAssayMap-method
 (zzz-DTTCompatibility-class),
 13

- 13
- BioAssayMapTarget_assnref-class
(zzz-DTTCompatibility-class),
13
- bioAssays_assnreflist
(zzz-DTTCompatibility-class),
13
- bioAssays_assnreflist,BioAssayDimension-method
(zzz-DTTCompatibility-class),
13
- bioAssays_assnreflist,Experiment-method
(zzz-DTTCompatibility-class),
13
- BioAssays_assnreflist-class
(zzz-DTTCompatibility-class),
13
- bioAssayTreatment
(zzz-DTTCompatibility-class),
13
- bioAssayTreatment,BioAssayTreatments_assnlist-method
(zzz-DTTCompatibility-class),
13
- BioAssayTreatment-class
(zzz-DTTCompatibility-class),
13
- bioAssayTreatments_assnlist
(zzz-DTTCompatibility-class),
13
- bioAssayTreatments_assnlist,PhysicalBioAssay-method
(zzz-DTTCompatibility-class),
13
- BioAssayTreatments_assnlist-class
(zzz-DTTCompatibility-class),
13
- bioAssayTuple
(zzz-DTTCompatibility-class),
13
- bioAssayTuple,BioAssayTuples_assnlist-method
(zzz-DTTCompatibility-class),
13
- BioAssayTuple-class
(zzz-DTTCompatibility-class),
13
- bioAssayTuples_assnlist
(zzz-DTTCompatibility-class),
13
- bioAssayTuples_assnlist,BioDataTuples-method
(zzz-DTTCompatibility-class),
13
- BioAssayTuples_assnlist-class
(zzz-DTTCompatibility-class),
13
- bioDataCube
(zzz-DTTCompatibility-class),
13
- bioDataCube,BioDataValues_assn-method
(zzz-DTTCompatibility-class),
13
- BioDataCube-class
(zzz-DTTCompatibility-class),
13
- bioDataTuples
(zzz-DTTCompatibility-class),
13
- bioDataTuples,BioDataValues_assn-method
(zzz-DTTCompatibility-class),
13
- BioDataTuples-class
(zzz-DTTCompatibility-class),
13
- bioDataValues_assn
(zzz-DTTCompatibility-class),
13
- bioDataValues_assn,DerivedBioAssayData-method
(zzz-DTTCompatibility-class),
13
- bioDataValues_assn,MeasuredBioAssayData-method
(zzz-DTTCompatibility-class),
13
- BioDataValues_assn-class
(zzz-DTTCompatibility-class),
13
- bioEvent_package
(zzz-DTTCompatibility-class),
13
- bioEvent_package,MAGE-ML-method
(zzz-DTTCompatibility-class),
13
- BioEvent_package-class
(zzz-DTTCompatibility-class),
13
- biologicalCharacteristics_assnreflist
(zzz-DTTCompatibility-class),
13
- biologicalCharacteristics_assnreflist,CompositeSequence
(zzz-DTTCompatibility-class),
13
- BiologicalCharacteristics_assnreflist-class
(zzz-DTTCompatibility-class),
13
- bioMaterial_assnlist
(zzz-DTTCompatibility-class),
13
- bioMaterial_assnlist,BioMaterial_package-method

- (zzz-DTTCompatibility-class),
13
- BioMaterial_assnlist-class
(zzz-DTTCompatibility-class),
13
- bioMaterial_assnref
(zzz-DTTCompatibility-class),
13
- bioMaterial_assnref,BioMaterialMeasurement-method
(zzz-DTTCompatibility-class),
13
- bioMaterial_assnref,ManufactureLIMS-method
(zzz-DTTCompatibility-class),
13
- bioMaterial_assnref,ManufactureLIMSBiomaterial-method
(zzz-DTTCompatibility-class),
13
- BioMaterial_assnref-class
(zzz-DTTCompatibility-class),
13
- bioMaterial_package
(zzz-DTTCompatibility-class),
13
- bioMaterial_package,MAGE-ML-method
(zzz-DTTCompatibility-class),
13
- BioMaterial_package-class
(zzz-DTTCompatibility-class),
13
- BioMaterial_ref-class
(zzz-DTTCompatibility-class),
13
- bioMaterialMeasurement
(zzz-DTTCompatibility-class),
13
- bioMaterialMeasurement,SourceBioMaterialMeasurements_assnlist-method
(zzz-DTTCompatibility-class),
13
- BioMaterialMeasurement-class
(zzz-DTTCompatibility-class),
13
- bioMaterialPlateCol
(zzz-DTTCompatibility-class),
13
- bioMaterialPlateCol,ManufactureLIMSBiomaterial-method
(zzz-DTTCompatibility-class),
13
- bioMaterialPlateIdentifier
(zzz-DTTCompatibility-class),
13
- bioMaterialPlateIdentifier,ManufactureLIMSBiomaterial-method
(zzz-DTTCompatibility-class),
- 13
- bioMaterialPlateRow
(zzz-DTTCompatibility-class),
13
- bioMaterialPlateRow,ManufactureLIMSBiomaterial-method
(zzz-DTTCompatibility-class),
13
- bioSample (zzz-DTTCompatibility-class),
13
- bioSample,BioMaterial_assnlist-method
(zzz-DTTCompatibility-class),
13
- BioSample-class
(zzz-DTTCompatibility-class),
- bioSample_ref
(zzz-DTTCompatibility-class),
13
- bioSample_ref,BioMaterial_assnref-method
(zzz-DTTCompatibility-class),
13
- BioSample_ref-class
(zzz-DTTCompatibility-class),
13
- bioSequence
(zzz-DTTCompatibility-class),
13
- bioSequence,BioSequence_assnlist-method
(zzz-DTTCompatibility-class),
13
- BioSequence-class
(zzz-DTTCompatibility-class),
13
- bioSequence_assnlist
(zzz-DTTCompatibility-class),
- bioSequence_assnlist,BioSequence_package-method
(zzz-DTTCompatibility-class),
13
- BioSequence_assnlist-class
(zzz-DTTCompatibility-class),
13
- bioSequence_package
(zzz-DTTCompatibility-class),
- bioSequence_package,MAGE-ML-method
(zzz-DTTCompatibility-class),
13
- BioSequence_package-class
(zzz-DTTCompatibility-class),
- bioSequence_ref

- (zzz-DTTCompatibility-class),
13
- bioSequence_ref,BiologicalCharacteristics_assnreflist-method
(zzz-DTTCompatibility-class),
13
- bioSequence_ref,ImmobilizedCharacteristics_assnreflist-method
(zzz-DTTCompatibility-class),
13
- BioSequence_ref-class
(zzz-DTTCompatibility-class),
13
- bioSource (zzz-DTTCompatibility-class),
13
- bioSource,BioMaterial_assnlist-method
(zzz-DTTCompatibility-class),
13
- BioSource-class
(zzz-DTTCompatibility-class),
13
- bioSource_ref
(zzz-DTTCompatibility-class),
13
- bioSource_ref,BioMaterial_assnref-method
(zzz-DTTCompatibility-class),
13
- BioSource_ref-class
(zzz-DTTCompatibility-class),
13
- bqs_package
(zzz-DTTCompatibility-class),
13
- bqs_package,MAGE-ML-method
(zzz-DTTCompatibility-class),
13
- BQS_package-class
(zzz-DTTCompatibility-class),
13
- category_assn
(zzz-DTTCompatibility-class),
13
- category_assn,ExperimentalFactor-method
(zzz-DTTCompatibility-class),
13
- Category_assn-class
(zzz-DTTCompatibility-class),
13
- channel (zzz-DTTCompatibility-class), 13
- channel,Channel_assnlist-method
(zzz-DTTCompatibility-class),
13
- Channel-class
(zzz-DTTCompatibility-class),
13
- channel_assnlist
(zzz-DTTCompatibility-class),
13
- channel_assnlist,BioAssay_package-method
(zzz-DTTCompatibility-class),
13
- Channel_assnlist-class
(zzz-DTTCompatibility-class),
13
- channel_assnref
(zzz-DTTCompatibility-class),
13
- channel_assnref,DerivedSignal-method
(zzz-DTTCompatibility-class),
13
- channel_assnref,Error-method
(zzz-DTTCompatibility-class),
13
- channel_assnref,ExpectedValue-method
(zzz-DTTCompatibility-class),
13
- channel_assnref,Failed-method
(zzz-DTTCompatibility-class),
13
- channel_assnref,MeasuredSignal-method
(zzz-DTTCompatibility-class),
13
- channel_assnref,PresentAbsent-method
(zzz-DTTCompatibility-class),
13
- channel_assnref,PValue-method
(zzz-DTTCompatibility-class),
13
- channel_assnref,Ratio-method
(zzz-DTTCompatibility-class),
13
- channel_assnref,SpecializedQuantitationType-method
(zzz-DTTCompatibility-class),
13
- Channel_assnref-class
(zzz-DTTCompatibility-class),
13
- channel_ref
(zzz-DTTCompatibility-class),
13
- channel_ref,Channel_assnref-method
(zzz-DTTCompatibility-class),
13
- channel_ref,Channels_assnreflist-method
(zzz-DTTCompatibility-class),
13

- Channel_ref-class
(zzz-DTTCompatibility-class),
13
- channels_assnreflist
(zzz-DTTCompatibility-class),
13
- channels_assnreflist, DerivedBioAssay-method
(zzz-DTTCompatibility-class),
13
- channels_assnreflist, Image-method
(zzz-DTTCompatibility-class),
13
- channels_assnreflist, MeasuredBioAssay-method
(zzz-DTTCompatibility-class),
13
- channels_assnreflist, PhysicalBioAssay-method
(zzz-DTTCompatibility-class),
13
- Channels_assnreflist-class
(zzz-DTTCompatibility-class),
13
- characteristics_assnlist
(zzz-DTTCompatibility-class),
13
- characteristics_assnlist, BioSample-method
(zzz-DTTCompatibility-class),
13
- characteristics_assnlist, BioSource-method
(zzz-DTTCompatibility-class),
13
- characteristics_assnlist, LabeledExtract-method
(zzz-DTTCompatibility-class),
13
- Characteristics_assnlist-class
(zzz-DTTCompatibility-class),
13
- clusterBioAssayData_assnref
(zzz-DTTCompatibility-class),
13
- clusterBioAssayData_assnref, BioAssayDataCluster-method
(zzz-DTTCompatibility-class),
13
- ClusterBioAssayData_assnref-class
(zzz-DTTCompatibility-class),
13
- column (zzz-DTTCompatibility-class), 13
- column, FeatureLocation-method
(zzz-DTTCompatibility-class),
13
- column, Zone-method
(zzz-DTTCompatibility-class),
13
- componentCompounds_assnlist
(zzz-DTTCompatibility-class),
13
- componentCompounds_assnlist, Compound-method
(zzz-DTTCompatibility-class),
13
- ComponentCompounds_assnlist-class
(zzz-DTTCompatibility-class),
13
- composite_assnref
(zzz-DTTCompatibility-class),
13
- composite_assnref, CompositePosition-method
(zzz-DTTCompatibility-class),
13
- Composite_assnref-class
(zzz-DTTCompatibility-class),
13
- compositeCompositeMap
(zzz-DTTCompatibility-class),
13
- compositeCompositeMap, CompositeCompositeMap_assnlist-method
(zzz-DTTCompatibility-class),
13
- compositeCompositeMap, DesignElementMap_assnlist-method
(zzz-DTTCompatibility-class),
13
- CompositeCompositeMap-class
(zzz-DTTCompatibility-class),
13
- compositeCompositeMap_assnlist
(zzz-DTTCompatibility-class),
13
- compositeCompositeMap_assnlist, DesignElement_package-method
(zzz-DTTCompatibility-class),
13
- CompositeCompositeMap_assnlist-class
(zzz-DTTCompatibility-class),
13
- compositeCompositeMap_ref
(zzz-DTTCompatibility-class),
13
- compositeCompositeMap_ref, CompositeCompositeMaps_assnreflist
(zzz-DTTCompatibility-class),
13
- compositeCompositeMap_ref, DesignElementMaps_assnreflist
(zzz-DTTCompatibility-class),
13
- CompositeCompositeMap_ref-class
(zzz-DTTCompatibility-class),
13
- compositeCompositeMaps_assnreflist

- (zzz-DTTCompatibility-class), 13
- compositeCompositeMaps_assnreflist, CompositeSequence-assnreflist-method (zzz-DTTCompatibility-class), 13
- CompositeCompositeMaps_assnreflist-class (zzz-DTTCompatibility-class), 13
- compositeGroup (zzz-DTTCompatibility-class), 13
- compositeGroup, CompositeGroup_assnlist-method (zzz-DTTCompatibility-class), 13
- CompositeGroup-class (zzz-DTTCompatibility-class), 13
- compositeGroup_assnlist (zzz-DTTCompatibility-class), 13
- compositeGroup_assnlist, ArrayDesign_package-method (zzz-DTTCompatibility-class), 13
- CompositeGroup_assnlist-class (zzz-DTTCompatibility-class), 13
- compositeGroup_ref (zzz-DTTCompatibility-class), 13
- compositeGroup_ref, CompositeGroups_assnreflist-method (zzz-DTTCompatibility-class), 13
- CompositeGroup_ref-class (zzz-DTTCompatibility-class), 13
- compositeGroups_assnreflist (zzz-DTTCompatibility-class), 13
- compositeGroups_assnreflist, ArrayDesign-method (zzz-DTTCompatibility-class), 13
- compositeGroups_assnreflist, PhysicalArrayDesign-method (zzz-DTTCompatibility-class), 13
- CompositeGroups_assnreflist-class (zzz-DTTCompatibility-class), 13
- compositePosition (zzz-DTTCompatibility-class), 13
- compositePosition, CompositePositionSources_assnlist-method (zzz-DTTCompatibility-class), 13
- compositePosition, Coordinate_assn-method (zzz-DTTCompatibility-class), 13
- CompositePosition-class (zzz-DTTCompatibility-class), 13
- compositePositionSources_assnlist (zzz-DTTCompatibility-class), 13
- compositePositionSources_assnlist, CompositeCompositeMap-assnreflist-method (zzz-DTTCompatibility-class), 13
- CompositePositionSources_assnlist-class (zzz-DTTCompatibility-class), 13
- compositeSequence (zzz-DTTCompatibility-class), 13
- compositeSequence, CompositeSequence_assnlist-method (zzz-DTTCompatibility-class), 13
- compositeSequence, DesignElement_assnlist-method (zzz-DTTCompatibility-class), 13
- CompositeSequence-class (zzz-DTTCompatibility-class), 13
- compositeSequence_assnlist (zzz-DTTCompatibility-class), 13
- compositeSequence_assnlist, DesignElement_package-method (zzz-DTTCompatibility-class), 13
- CompositeSequence_assnlist-class (zzz-DTTCompatibility-class), 13
- compositeSequence_assnref (zzz-DTTCompatibility-class), 13
- compositeSequence_assnref, CompositeCompositeMap-method (zzz-DTTCompatibility-class), 13
- CompositeSequence_assnref-class (zzz-DTTCompatibility-class), 13
- compositeSequence_ref (zzz-DTTCompatibility-class), 13

compositeSequence_ref, Composite_assnref-method 13
 (zzz-DTTCompatibility-class), compound_assnlist
 13 (zzz-DTTCompatibility-class),
 compositeSequence_ref, CompositeSequence_assnref-method 13
 (zzz-DTTCompatibility-class), compound_assnlist, BioMaterial_package-method
 13 (zzz-DTTCompatibility-class),
 compositeSequence_ref, CompositeSequences_assnreflist-method 13
 (zzz-DTTCompatibility-class), Compound_assnlist-class
 13 (zzz-DTTCompatibility-class),
 compositeSequence_ref, DesignElement_assnref-method 13
 (zzz-DTTCompatibility-class), compound_assnref
 13 (zzz-DTTCompatibility-class),
 CompositeSequence_ref-class 13
 (zzz-DTTCompatibility-class), compound_assnref, CompoundMeasurement-method
 13 (zzz-DTTCompatibility-class),
 compositeSequenceDimension 13
 (zzz-DTTCompatibility-class), Compound_assnref-class
 13 (zzz-DTTCompatibility-class),
 compositeSequenceDimension, DesignElementDimension_assnlist-method 13
 (zzz-DTTCompatibility-class), compound_ref
 13 (zzz-DTTCompatibility-class),
 CompositeSequenceDimension-class 13
 (zzz-DTTCompatibility-class), compound_ref, Compound_assnref-method
 13 (zzz-DTTCompatibility-class),
 compositeSequenceDimension_ref 13
 (zzz-DTTCompatibility-class), compound_ref, Labels_assnreflist-method
 13 (zzz-DTTCompatibility-class),
 compositeSequenceDimension_ref, DesignElementDimension_assnref-method 13
 (zzz-DTTCompatibility-class), Compound_ref-class
 13 (zzz-DTTCompatibility-class),
 CompositeSequenceDimension_ref-class 13
 (zzz-DTTCompatibility-class), compoundIndices_assnlist
 13 (zzz-DTTCompatibility-class),
 compositeSequences_assnreflist 13
 (zzz-DTTCompatibility-class), compoundIndices_assnlist, Compound-method
 13 (zzz-DTTCompatibility-class),
 compositeSequences_assnreflist, CompositeGroup-method 13
 (zzz-DTTCompatibility-class), CompoundIndices_assnlist-class
 13 (zzz-DTTCompatibility-class),
 compositeSequences_assnreflist, CompositeSequenceDimension-method 13
 (zzz-DTTCompatibility-class), compoundMeasurement
 13 (zzz-DTTCompatibility-class),
 CompositeSequences_assnreflist-class 13
 (zzz-DTTCompatibility-class), compoundMeasurement, ComponentCompounds_assnlist-method
 13 (zzz-DTTCompatibility-class),
 compound (zzz-DTTCompatibility-class), 13
 13 compoundMeasurement, CompoundMeasurements_assnlist-method
 compound, Compound_assnlist-method 13
 (zzz-DTTCompatibility-class), 13
 13 CompoundMeasurement-class
 Compound-class (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13

- (zzz-DTTCompatibility-class),
13
- controlledFeatures_assnreflist,Feature-method
(zzz-DTTCompatibility-class),
13
- ControlledFeatures_assnreflist-class
(zzz-DTTCompatibility-class),
13
- controlType_assn
(zzz-DTTCompatibility-class),
13
- controlType_assn,CompositeSequence-method
(zzz-DTTCompatibility-class),
13
- controlType_assn,Feature-method
(zzz-DTTCompatibility-class),
13
- controlType_assn,Reporter-method
(zzz-DTTCompatibility-class),
13
- ControlType_assn-class
(zzz-DTTCompatibility-class),
13
- coordinate_assn
(zzz-DTTCompatibility-class),
13
- coordinate_assn,SeqFeatureLocation-method
(zzz-DTTCompatibility-class),
13
- Coordinate_assn-class
(zzz-DTTCompatibility-class),
13
- createdBy (zzz-ARRCompatibility-class),
12
- createdBy,ArraySetFile-method
(zzz-ARRCompatibility-class),
12
- createdBy,PhysicalArray-method
(zzz-ARRCompatibility-class),
12
- createdBy,TemplateFile-method
(zzz-ARRCompatibility-class),
12
- createdDateTime
(zzz-ARRCompatibility-class),
12
- createdDateTime,ArraySetFile-method
(zzz-ARRCompatibility-class),
12
- createdDateTime,PhysicalArray-method
(zzz-ARRCompatibility-class),
12
- createdDateTime,TemplateFile-method
(zzz-ARRCompatibility-class),
12
- createdStep
(zzz-ARRCompatibility-class),
12
- createdStep,ArraySetFile-method
(zzz-ARRCompatibility-class),
12
- createdStep,PhysicalArray-method
(zzz-ARRCompatibility-class),
12
- database (zzz-DTTCompatibility-class),
13
- database,Database_assnlist-method
(zzz-DTTCompatibility-class),
13
- Database-class
(zzz-DTTCompatibility-class),
13
- database_assnlist
(zzz-DTTCompatibility-class),
13
- database_assnlist,Description_package-method
(zzz-DTTCompatibility-class),
13
- Database_assnlist-class
(zzz-DTTCompatibility-class),
13
- database_assnref
(zzz-DTTCompatibility-class),
13
- database_assnref,DatabaseEntry-method
(zzz-DTTCompatibility-class),
13
- Database_assnref-class
(zzz-DTTCompatibility-class),
13
- database_ref
(zzz-DTTCompatibility-class),
13
- database_ref,Database_assnref-method
(zzz-DTTCompatibility-class),
13
- Database_ref-class
(zzz-DTTCompatibility-class),
13
- databaseEntry
(zzz-DTTCompatibility-class),
13
- databaseEntry,Accessions_assnlist-method
(zzz-DTTCompatibility-class),

- 13
- databaseEntry, DatabaseReferences_assnlist-method (zzz-DTTCompatibility-class), 13
- databaseEntry, ExternalLIMS_assn-method (zzz-DTTCompatibility-class), 13
- databaseEntry, IdentifierLIMS_assn-method (zzz-DTTCompatibility-class), 13
- databaseEntry, OntologyReference_assn-method (zzz-DTTCompatibility-class), 13
- databaseEntry, SequenceDatabases_assnlist-method (zzz-DTTCompatibility-class), 13
- DatabaseEntry-class (zzz-DTTCompatibility-class), 13
- databaseReferences_assnlist (zzz-DTTCompatibility-class), 13
- databaseReferences_assnlist, Description-method (zzz-DTTCompatibility-class), 13
- DatabaseReferences_assnlist-class (zzz-DTTCompatibility-class), 13
- dataExternal (zzz-DTTCompatibility-class), 13
- dataExternal, DataExternal_assn-method (zzz-DTTCompatibility-class), 13
- DataExternal-class (zzz-DTTCompatibility-class), 13
- dataExternal_assn (zzz-DTTCompatibility-class), 13
- dataExternal_assn, BioDataCube-method (zzz-DTTCompatibility-class), 13
- DataExternal_assn-class (zzz-DTTCompatibility-class), 13
- dataFormat (zzz-DTTCompatibility-class), 13
- dataFormat, DataExternal-method (zzz-DTTCompatibility-class), 13
- dataFormatInfoURI (zzz-DTTCompatibility-class), 13
- dataFormatInfoURI, DataExternal-method (zzz-DTTCompatibility-class), 13
- dataInternal (zzz-DTTCompatibility-class), 13
- dataInternal, DataInternal_assn-method (zzz-DTTCompatibility-class), 13
- DataInternal-class (zzz-DTTCompatibility-class), 13
- dataInternal_assn (zzz-DTTCompatibility-class), 13
- dataInternal_assn, BioDataCube-method (zzz-DTTCompatibility-class), 13
- DataInternal_assn-class (zzz-DTTCompatibility-class), 13
- dataType_assn (zzz-DTTCompatibility-class), 13
- dataType_assn, DerivedSignal-method (zzz-DTTCompatibility-class), 13
- dataType_assn, Error-method (zzz-DTTCompatibility-class), 13
- dataType_assn, ExpectedValue-method (zzz-DTTCompatibility-class), 13
- dataType_assn, Failed-method (zzz-DTTCompatibility-class), 13
- dataType_assn, MeasuredSignal-method (zzz-DTTCompatibility-class), 13
- dataType_assn, NodeValue-method (zzz-DTTCompatibility-class), 13
- dataType_assn, Parameter-method (zzz-DTTCompatibility-class), 13
- dataType_assn, PresentAbsent-method (zzz-DTTCompatibility-class), 13
- dataType_assn, PValue-method

- (zzz-DTTCompatibility-class),
13
- dataType_assn,Ratio-method
(zzz-DTTCompatibility-class),
13
- dataType_assn,SpecializedQuantitationType-method
(zzz-DTTCompatibility-class),
13
- DataType_assn-class
(zzz-DTTCompatibility-class),
13
- datum (zzz-DTTCompatibility-class), 13
- datum,Datum_assn-method
(zzz-DTTCompatibility-class),
13
- Datum-class
(zzz-DTTCompatibility-class),
13
- datum_assn
(zzz-DTTCompatibility-class),
13
- datum_assn,QuantitationTypeTuple-method
(zzz-DTTCompatibility-class),
13
- Datum_assn-class
(zzz-DTTCompatibility-class),
13
- defaultValue
(zzz-ARRCompatibility-class),
12
- defaultValue,UserAttribute-method
(zzz-ARRCompatibility-class),
12
- defaultValue_assn
(zzz-DTTCompatibility-class),
13
- defaultValue_assn,Parameter-method
(zzz-DTTCompatibility-class),
13
- DefaultValue_assn-class
(zzz-DTTCompatibility-class),
13
- defectType_assn
(zzz-DTTCompatibility-class),
13
- defectType_assn,FeatureDefect-method
(zzz-DTTCompatibility-class),
13
- defectType_assn,ZoneDefect-method
(zzz-DTTCompatibility-class),
13
- DefectType_assn-class
(zzz-DTTCompatibility-class),
13
- deltaX (zzz-DTTCompatibility-class), 13
- deltaX,PositionDelta-method
(zzz-DTTCompatibility-class),
13
- deltaY (zzz-DTTCompatibility-class), 13
- deltaY,PositionDelta-method
(zzz-DTTCompatibility-class),
13
- derivedBioAssay
(zzz-DTTCompatibility-class),
13
- derivedBioAssay,BioAssay_assnlist-method
(zzz-DTTCompatibility-class),
13
- derivedBioAssay,DerivedBioAssay_assnlist-method
(zzz-DTTCompatibility-class),
13
- DerivedBioAssay-class
(zzz-DTTCompatibility-class),
13
- DerivedBioAssay_assnlist-class
(zzz-DTTCompatibility-class),
13
- derivedBioAssay_ref
(zzz-DTTCompatibility-class),
13
- derivedBioAssay_ref,BioAssay_assnref-method
(zzz-DTTCompatibility-class),
13
- derivedBioAssay_ref,BioAssayMapTarget_assnref-method
(zzz-DTTCompatibility-class),
13
- derivedBioAssay_ref,BioAssays_assnreflist-method
(zzz-DTTCompatibility-class),
13
- derivedBioAssay_ref,SourceBioAssays_assnreflist-method
(zzz-DTTCompatibility-class),
13
- derivedBioAssay_ref,TopLevelBioAssays_assnreflist-method
(zzz-DTTCompatibility-class),
13
- DerivedBioAssay_ref-class
(zzz-DTTCompatibility-class),
13
- derivedBioAssayData
(zzz-DTTCompatibility-class),
13
- derivedBioAssayData,BioAssayData_assnlist-method
(zzz-DTTCompatibility-class),
13

derivedBioAssayData, DerivedBioAssayData_assnlist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 13
 DerivedBioAssayData-class (zzz-DTTCompatibility-class), 13
 DerivedBioAssayData_assnlist-class (zzz-DTTCompatibility-class), 13
 derivedBioAssayData_assnreflist (zzz-DTTCompatibility-class), 13
 derivedBioAssayData_assnreflist, DerivedBioAssay-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 13
 DerivedBioAssayData_assnreflist-class (zzz-DTTCompatibility-class), 13
 derivedBioAssayData_ref (zzz-DTTCompatibility-class), 13
 derivedBioAssayData_ref, BioAssayData_assnreflist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 13
 derivedBioAssayData_ref, BioAssayDataSources_assnreflist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 13
 derivedBioAssayData_ref, ClusterBioAssayData_assnref-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 13
 derivedBioAssayData_ref, DerivedBioAssayData_assnreflist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 13
 derivedBioAssayData_ref, DerivedBioAssayDataTarget_assnref-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 13
 DerivedBioAssayData_ref-class (zzz-DTTCompatibility-class), 13
 13
 DerivedBioAssayDataTarget_assnref-class (zzz-DTTCompatibility-class), 13
 13
 derivedBioAssayMap_assnreflist (zzz-DTTCompatibility-class), 13
 13
 derivedBioAssayMap_assnreflist, DerivedBioAssay-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 13
 DerivedBioAssayMap_assnreflist-class (zzz-DTTCompatibility-class), 13
 (zzz-DTTCompatibility-class), 13
 13
 derivedSignal (zzz-DTTCompatibility-class), 13
 derivedSignal, QuantitationType_assnlist-method (zzz-DTTCompatibility-class), 13
 13
 DerivedSignal-class (zzz-DTTCompatibility-class), 13
 derivedSignal_ref (zzz-DTTCompatibility-class), 13
 derivedSignal_ref, QuantitationType_assnref-method (zzz-DTTCompatibility-class), 13
 13
 derivedSignal_ref, QuantitationTypes_assnreflist-method (zzz-DTTCompatibility-class), 13
 13
 derivedSignal_ref, SourcesQuantitationType_assnreflist-method (zzz-DTTCompatibility-class), 13
 13
 derivedSignal_ref, TargetQuantitationType_assnref-method (zzz-DTTCompatibility-class), 13
 13
 DerivedSignal_ref-class (zzz-DTTCompatibility-class), 13
 description (zzz-DTTCompatibility-class), 13
 description, Descriptions_assnlist-method (zzz-DTTCompatibility-class), 13
 description, NormalizationDescription_assn-method (zzz-DTTCompatibility-class), 13
 description, OntologyEntry-method (zzz-DTTCompatibility-class), 13
 description, QualityControlDescription_assn-method (zzz-DTTCompatibility-class), 13
 description, ReplicateDescription_assn-method (zzz-DTTCompatibility-class), 13
 Description-class (zzz-DTTCompatibility-class), 13
 description_package (zzz-DTTCompatibility-class), 13
 description_package, MAGE-ML-method (zzz-DTTCompatibility-class), 13

- 13
- Description_package-class
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,Array-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,ArrayDesign-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,ArrayGroup-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,ArrayManufacture-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,Audit-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,BibliographicReference-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,BioAssayCreation-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,BioAssayDataCluster-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,BioAssayDimension-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,BioAssayMap-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,BioAssayTreatment-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,BioSample-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,BioSequence-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,BioSource-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,Channel-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,CompositeCompositeMap-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,CompositeGroup-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,CompositeSequence-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,CompositeSequenceDimension-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,Compound-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,Database-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,DerivedBioAssay-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,DerivedBioAssayData-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,DerivedSignal-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,Description-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,Error-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,ExpectedValue-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,Experiment-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,ExperimentalFactor-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,ExperimentDesign-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,FactorValue-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,Failed-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,Feature-method

- (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, FeatureDimension-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, FeatureExtraction-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, FeatureGroup-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, FeatureReporterMap-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, Fiducial-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, Hardware-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, HardwareApplication-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, Hybridization-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, Image-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, ImageAcquisition-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, LabeledExtract-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, MAGE-ML-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, ManufactureLIMS-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, ManufactureLIMSBiomaterial-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, MeasuredBioAssay-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, MeasuredBioAssayData-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, MeasuredSignal-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, Node-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, NodeContents-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, Organization-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, Parameter-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, Person-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, PhysicalArrayDesign-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, PhysicalBioAssay-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, PresentAbsent-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, Protocol-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, ProtocolApplication-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, PValue-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, QuantitationTypeDimension-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, QuantitationTypeMap-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, Ratio-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, Reporter-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, ReporterCompositeMap-method (zzz-DTTCompatibility-class), 13
- descriptions_assnlist, ReporterDimension-method (zzz-DTTCompatibility-class), 13

- descriptions_assnlist,ReporterGroup-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,Security-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,SecurityGroup-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,SeqFeature-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,Software-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,SoftwareApplication-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,SpecializedQuantitationType-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,Transformation-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,Treatment-method
(zzz-DTTCompatibility-class),
13
- descriptions_assnlist,Zone-method
(zzz-DTTCompatibility-class),
13
- Descriptions_assnlist-class
(zzz-DTTCompatibility-class),
13
- DesignElement_assnlist-class
(zzz-DTTCompatibility-class),
13
- designElement_assnref
(zzz-DTTCompatibility-class),
13
- designElement_assnref,DesignElementTuple-method
(zzz-DTTCompatibility-class),
13
- DesignElement_assnref-class
(zzz-DTTCompatibility-class),
13
- designElement_package
(zzz-DTTCompatibility-class),
13
- designElement_package,MAGE-ML-method
(zzz-DTTCompatibility-class),
13
- DesignElement_package-class
(zzz-DTTCompatibility-class),
13
- (zzz-DTTCompatibility-class),
13
- DesignElement_ref-class
(zzz-DTTCompatibility-class),
13
- designElementDimension_assnlist
(zzz-DTTCompatibility-class),
13
- designElementDimension_assnlist,BioAssayData_package-me
(zzz-DTTCompatibility-class),
13
- DesignElementDimension_assnlist-class
(zzz-DTTCompatibility-class),
13
- designElementDimension_assnref
(zzz-DTTCompatibility-class),
13
- designElementDimension_assnref,DerivedBioAssayData-meth
(zzz-DTTCompatibility-class),
13
- designElementDimension_assnref,MeasuredBioAssayData-meth
(zzz-DTTCompatibility-class),
13
- designElementDimension_assnref,NodeContents-method
(zzz-DTTCompatibility-class),
13
- DesignElementDimension_assnref-class
(zzz-DTTCompatibility-class),
13
- DesignElementDimension_ref-class
(zzz-DTTCompatibility-class),
13
- DesignElementMap_assnlist-class
(zzz-DTTCompatibility-class),
13
- DesignElementMap_ref-class
(zzz-DTTCompatibility-class),
13
- designElementMapping
(zzz-DTTCompatibility-class),
13
- designElementMapping,DesignElementMapping_assn-method
(zzz-DTTCompatibility-class),
13
- DesignElementMapping-class
(zzz-DTTCompatibility-class),
13
- designElementMapping_assn
(zzz-DTTCompatibility-class),
13
- designElementMapping_assn,Transformation-method
(zzz-DTTCompatibility-class),
13

- 13
- DesignElementMapping_assn-class
(zzz-DTTCompatibility-class),
13
- designElementMaps_assnreflist
(zzz-DTTCompatibility-class),
13
- designElementMaps_assnreflist, DesignElementMapping-method
(zzz-DTTCompatibility-class),
13
- DesignElementMaps_assnreflist-class
(zzz-DTTCompatibility-class),
13
- designElementTuple
(zzz-DTTCompatibility-class),
13
- designElementTuple, DesignElementTuples_assnlist-method
(zzz-DTTCompatibility-class),
13
- DesignElementTuple-class
(zzz-DTTCompatibility-class),
13
- designElementTuples_assnlist
(zzz-DTTCompatibility-class),
13
- designElementTuples_assnlist, BioAssayTuple-method
(zzz-DTTCompatibility-class),
13
- DesignElementTuples_assnlist-class
(zzz-DTTCompatibility-class),
13
- designProviders_assnreflist
(zzz-DTTCompatibility-class),
13
- designProviders_assnreflist, ArrayDesign-method
(zzz-DTTCompatibility-class),
13
- designProviders_assnreflist, PhysicalArrayDesign-method
(zzz-DTTCompatibility-class),
13
- DesignProviders_assnreflist-class
(zzz-DTTCompatibility-class),
13
- directory (accessors), 2
- directory, NetAffxResource-method
(NetAffxResource-class), 5
- distanceUnit
(zzz-DTTCompatibility-class),
13
- distanceUnit, DistanceUnit_assn-method
(zzz-DTTCompatibility-class),
13
- distanceUnit, Unit_assn-method
(zzz-DTTCompatibility-class),
13
- DistanceUnit-class
(zzz-DTTCompatibility-class),
13
- distanceUnit_assn
(zzz-DTTCompatibility-class),
13
- distanceUnit_assn, ArrayGroup-method
(zzz-DTTCompatibility-class),
13
- distanceUnit_assn, FeatureGroup-method
(zzz-DTTCompatibility-class),
13
- distanceUnit_assn, Fiducial-method
(zzz-DTTCompatibility-class),
13
- distanceUnit_assn, Position-method
(zzz-DTTCompatibility-class),
13
- distanceUnit_assn, PositionDelta-method
(zzz-DTTCompatibility-class),
13
- distanceUnit_assn, Zone-method
(zzz-DTTCompatibility-class),
13
- distanceUnit_assn, ZoneGroup-method
(zzz-DTTCompatibility-class),
13
- distanceUnit_assn, ZoneLayout-method
(zzz-DTTCompatibility-class),
13
- DistanceUnit_assn-class
(zzz-DTTCompatibility-class),
13
- DTTCompatibility, 14
- DTTCompatibility-class
(AffyCompatible-class), 3
- editor (zzz-DTTCompatibility-class), 13
- editor, BibliographicReference-method
(zzz-DTTCompatibility-class),
13
- email (zzz-DTTCompatibility-class), 13
- email, Organization-method
(zzz-DTTCompatibility-class),
13
- email, Person-method
(zzz-DTTCompatibility-class),
13
- error (zzz-DTTCompatibility-class), 13

- error,ConfidenceIndicator_assnlist-method
 (zzz-DTTCompatibility-class),
 13
- error,QuantitationType_assnlist-method
 (zzz-DTTCompatibility-class),
 13
- Error-class
 (zzz-DTTCompatibility-class),
 13
- error_ref(zzz-DTTCompatibility-class),
 13
- error_ref,ConfidenceIndicators_assnreflist-method
 (zzz-DTTCompatibility-class),
 13
- error_ref,QuantitationType_assnref-method
 (zzz-DTTCompatibility-class),
 13
- error_ref,QuantitationTypes_assnreflist-method
 (zzz-DTTCompatibility-class),
 13
- error_ref,SourcesQuantitationType_assnreflist-method
 (zzz-DTTCompatibility-class),
 13
- error_ref,TargetQuantitationType_assnref-method
 (zzz-DTTCompatibility-class),
 13
- Error_ref-class
 (zzz-DTTCompatibility-class),
 13
- expectedValue
 (zzz-DTTCompatibility-class),
 13
- expectedValue,ConfidenceIndicator_assnlist-method
 (zzz-DTTCompatibility-class),
 13
- expectedValue,QuantitationType_assnlist-method
 (zzz-DTTCompatibility-class),
 13
- ExpectedValue-class
 (zzz-DTTCompatibility-class),
 13
- expectedValue_ref
 (zzz-DTTCompatibility-class),
 13
- expectedValue_ref,ConfidenceIndicators_assnreflist-method
 (zzz-DTTCompatibility-class),
 13
- expectedValue_ref,QuantitationType_assnref-method
 (zzz-DTTCompatibility-class),
 13
- expectedValue_ref,QuantitationTypes_assnreflist-method
 (zzz-DTTCompatibility-class),
 13
- expectedValue_ref,SourcesQuantitationType_assnreflist-method
 (zzz-DTTCompatibility-class),
 13
- expectedValue_ref,TargetQuantitationType_assnref-method
 (zzz-DTTCompatibility-class),
 13
- ExpectedValue_ref-class
 (zzz-DTTCompatibility-class),
 13
- experiment
 (zzz-DTTCompatibility-class),
 13
- experiment,Experiment_assnlist-method
 (zzz-DTTCompatibility-class),
 13
- Experiment-class
 (zzz-DTTCompatibility-class),
 13
- experiment_assnlist
 (zzz-DTTCompatibility-class),
 13
- experiment_assnlist,Experiment_package-method
 (zzz-DTTCompatibility-class),
 13
- Experiment_assnlist-class
 (zzz-DTTCompatibility-class),
 13
- experiment_package
 (zzz-DTTCompatibility-class),
 13
- experiment_package,MAGE-ML-method
 (zzz-DTTCompatibility-class),
 13
- Experiment_package-class
 (zzz-DTTCompatibility-class),
 13
- experimentalFactor
 (zzz-DTTCompatibility-class),
 13
- experimentalFactor,ExperimentalFactors_assnlist-method
 (zzz-DTTCompatibility-class),
 13
- ExperimentalFactor-class
 (zzz-DTTCompatibility-class),
 13
- ExperimentalFactor_assnref-class
 (zzz-DTTCompatibility-class),
 13
- experimentalFactor_ref
 (zzz-DTTCompatibility-class),
 13

- experimentalFactor_ref, ExperimentalFactor_assnref-method
(zzz-DTTCompatibility-class), 13
- ExperimentalFactor_ref-class
(zzz-DTTCompatibility-class), 13
- experimentalFactors_assnlist
(zzz-DTTCompatibility-class), 13
- experimentalFactors_assnlist, ExperimentDesign-method
(zzz-DTTCompatibility-class), 13
- ExperimentalFactors_assnlist-class
(zzz-DTTCompatibility-class), 13
- experimentDesign
(zzz-DTTCompatibility-class), 13
- experimentDesign, ExperimentDesigns_assnlist-method
(zzz-DTTCompatibility-class), 13
- ExperimentDesign-class
(zzz-DTTCompatibility-class), 13
- experimentDesigns_assnlist
(zzz-DTTCompatibility-class), 13
- experimentDesigns_assnlist, Experiment-method
(zzz-DTTCompatibility-class), 13
- ExperimentDesigns_assnlist-class
(zzz-DTTCompatibility-class), 13
- exportedFromDB
(zzz-DTTCompatibility-class), 13
- exportedFromDB, ExternalReference-method
(zzz-DTTCompatibility-class), 13
- exportedFromServer
(zzz-DTTCompatibility-class), 13
- exportedFromServer, ExternalReference-method
(zzz-DTTCompatibility-class), 13
- exportID (zzz-DTTCompatibility-class), 13
- exportID, ExternalReference-method
(zzz-DTTCompatibility-class), 13
- exportName
(zzz-DTTCompatibility-class), 13
- exportName, ExternalReference-method
(zzz-DTTCompatibility-class), 13
- externalLIMS_assn
(zzz-DTTCompatibility-class), 13
- externalLIMS_assn, Compound-method
(zzz-DTTCompatibility-class), 13
- ExternalLIMS_assn-class
(zzz-DTTCompatibility-class), 13
- externalReference
(zzz-DTTCompatibility-class), 13
- externalReference, ExternalReference_assn-method
(zzz-DTTCompatibility-class), 13
- externalReference_assn
(zzz-DTTCompatibility-class), 13
- externalReference_assn, Description-method
(zzz-DTTCompatibility-class), 13
- ExternalReference_assn-class
(zzz-DTTCompatibility-class), 13
- factorValue
(zzz-DTTCompatibility-class), 13
- factorValue, FactorValues_assnlist-method
(zzz-DTTCompatibility-class), 13
- FactorValue-class
(zzz-DTTCompatibility-class), 13
- factorValue_ref
(zzz-DTTCompatibility-class), 13
- factorValue_ref, BioAssayFactorValues_assnreflist-method
(zzz-DTTCompatibility-class), 13
- FactorValue_ref-class
(zzz-DTTCompatibility-class), 13
- factorValues_assnlist
(zzz-DTTCompatibility-class), 13
- factorValues_assnlist, ExperimentalFactor-method
(zzz-DTTCompatibility-class), 13

- FactorValues_assnlist-class
(zzz-DTTCompatibility-class),
13
- failed (zzz-DTTCompatibility-class), 13
- failed, QuantitationType_assnlist-method
(zzz-DTTCompatibility-class),
13
- Failed-class
(zzz-DTTCompatibility-class),
13
- failed_ref
(zzz-DTTCompatibility-class),
13
- failed_ref, QuantitationType_assnref-method
(zzz-DTTCompatibility-class),
13
- failed_ref, QuantitationTypes_assnreflist-method
(zzz-DTTCompatibility-class),
13
- failed_ref, SourcesQuantitationType_assnreflist-method
(zzz-DTTCompatibility-class),
13
- failed_ref, TargetQuantitationType_assnref-method
(zzz-DTTCompatibility-class),
13
- Failed_ref-class
(zzz-DTTCompatibility-class),
13
- failTypes_assnlist
(zzz-DTTCompatibility-class),
13
- failTypes_assnlist, Reporter-method
(zzz-DTTCompatibility-class),
13
- FailTypes_assnlist-class
(zzz-DTTCompatibility-class),
13
- fax (zzz-DTTCompatibility-class), 13
- fax, Organization-method
(zzz-DTTCompatibility-class),
13
- fax, Person-method
(zzz-DTTCompatibility-class),
13
- feature (zzz-DTTCompatibility-class), 13
- feature, DesignElement_assnlist-method
(zzz-DTTCompatibility-class),
13
- feature, Features_assnlist-method
(zzz-DTTCompatibility-class),
13
- Feature-class
(zzz-DTTCompatibility-class),
13
- feature_assnref
(zzz-DTTCompatibility-class),
13
- feature_assnref, FeatureDefect-method
(zzz-DTTCompatibility-class),
13
- feature_assnref, FeatureInformation-method
(zzz-DTTCompatibility-class),
13
- feature_assnref, ManufactureLIMS-method
(zzz-DTTCompatibility-class),
13
- feature_assnref, ManufactureLIMSBiomaterial-method
(zzz-DTTCompatibility-class),
13
- Feature_assnref-class
(zzz-DTTCompatibility-class),
13
- feature_ref
(zzz-DTTCompatibility-class),
13
- feature_ref, ContainedFeatures_assnreflist-method
(zzz-DTTCompatibility-class),
13
- feature_ref, ControlFeatures_assnreflist-method
(zzz-DTTCompatibility-class),
13
- feature_ref, ControlledFeatures_assnreflist-method
(zzz-DTTCompatibility-class),
13
- feature_ref, DesignElement_assnref-method
(zzz-DTTCompatibility-class),
13
- feature_ref, Feature_assnref-method
(zzz-DTTCompatibility-class),
13
- Feature_ref-class
(zzz-DTTCompatibility-class),
13
- featureDefect
(zzz-DTTCompatibility-class),
13
- featureDefect, FeatureDefects_assnlist-method
(zzz-DTTCompatibility-class),
13
- FeatureDefect-class
(zzz-DTTCompatibility-class),
13
- featureDefects_assnlist
(zzz-DTTCompatibility-class),
13

- 13
 featureDefects_assnlist, ArrayManufactureDeviation-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
- 13
 FeatureDefects_assnlist-class
 (zzz-DTTCompatibility-class), 13
- featureDimension
 (zzz-DTTCompatibility-class), 13
- featureDimension, DesignElementDimension_assnlist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
- 13
 FeatureDimension-class
 (zzz-DTTCompatibility-class), 13
- featureDimension_ref
 (zzz-DTTCompatibility-class), 13
- 13
 featureDimension_ref, DesignElementDimension_assnref-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
- 13
 FeatureDimension_ref-class
 (zzz-DTTCompatibility-class), 13
- featureExtraction
 (zzz-DTTCompatibility-class), 13
- 13
 featureExtraction, FeatureExtraction_assn-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
- 13
 FeatureExtraction-class
 (zzz-DTTCompatibility-class), 13
- featureExtraction_assn
 (zzz-DTTCompatibility-class), 13
- 13
 featureExtraction_assn, MeasuredBioAssay-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
- 13
 FeatureExtraction_assn-class
 (zzz-DTTCompatibility-class), 13
- featureGroup
 (zzz-DTTCompatibility-class), 13
- 13
 featureGroup, FeatureGroups_assnlist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
- 13
 FeatureGroup-class
 (zzz-DTTCompatibility-class), 13
- FeatureGroup_assnref-class
 (zzz-DTTCompatibility-class), 13
- featureGroup_ref
 (zzz-DTTCompatibility-class), 13
- 13
 featureGroup_ref, FeatureGroup_assnref-method
 (zzz-DTTCompatibility-class), 13
- 13
 FeatureGroup_ref-class
 (zzz-DTTCompatibility-class), 13
- featureGroups_assnlist
 (zzz-DTTCompatibility-class), 13
- 13
 featureGroups_assnlist, ArrayDesign-method
 (zzz-DTTCompatibility-class), 13
- 13
 featureGroups_assnlist, PhysicalArrayDesign-method
 (zzz-DTTCompatibility-class), 13
- 13
 FeatureGroups_assnlist-class
 (zzz-DTTCompatibility-class), 13
- featureHeight
 (zzz-DTTCompatibility-class), 13
- 13
 featureHeight, FeatureGroup-method
 (zzz-DTTCompatibility-class), 13
- 13
 featureInformation
 (zzz-DTTCompatibility-class), 13
- 13
 featureInformation, FeatureInformationSources_assnlist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
- 13
 FeatureInformation-class
 (zzz-DTTCompatibility-class), 13
- featureInformationSources_assnlist
 (zzz-DTTCompatibility-class), 13
- 13
 featureInformationSources_assnlist, FeatureReporterMap-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
- 13
 FeatureInformationSources_assnlist-class
 (zzz-DTTCompatibility-class), 13
- 13
 featureLength
 (zzz-DTTCompatibility-class), 13
- 13
 featureLength, FeatureGroup-method

- (zzz-DTTCompatibility-class),
13
- featureIMSSs_assnlist
(zzz-DTTCompatibility-class),
13
- featureIMSSs_assnlist,ArrayManufacture-method
(zzz-DTTCompatibility-class),
13
- FeatureIMSSs_assnlist-class
(zzz-DTTCompatibility-class),
13
- featureLocation
(zzz-DTTCompatibility-class),
13
- featureLocation,FeatureLocation_assn-method
(zzz-DTTCompatibility-class),
13
- FeatureLocation-class
(zzz-DTTCompatibility-class),
13
- featureLocation_assn
(zzz-DTTCompatibility-class),
13
- featureLocation_assn,Feature-method
(zzz-DTTCompatibility-class),
13
- FeatureLocation_assn-class
(zzz-DTTCompatibility-class),
13
- featureReporterMap
(zzz-DTTCompatibility-class),
13
- featureReporterMap,DesignElementMap_assnlist-method
(zzz-DTTCompatibility-class),
13
- featureReporterMap,FeatureReporterMap_assnlist-method
(zzz-DTTCompatibility-class),
13
- FeatureReporterMap-class
(zzz-DTTCompatibility-class),
13
- featureReporterMap_assnlist
(zzz-DTTCompatibility-class),
13
- featureReporterMap_assnlist,DesignElement_package-method
(zzz-DTTCompatibility-class),
13
- FeatureReporterMap_assnlist-class
(zzz-DTTCompatibility-class),
13
- featureReporterMap_ref
(zzz-DTTCompatibility-class),
13
- featureReporterMap_ref,DesignElementMaps_assnreflist-method
(zzz-DTTCompatibility-class),
13
- featureReporterMap_ref,FeatureReporterMaps_assnreflist-method
(zzz-DTTCompatibility-class),
13
- FeatureReporterMap_ref-class
(zzz-DTTCompatibility-class),
13
- featureReporterMaps_assnreflist
(zzz-DTTCompatibility-class),
13
- featureReporterMaps_assnreflist,Reporter-method
(zzz-DTTCompatibility-class),
13
- FeatureReporterMaps_assnreflist-class
(zzz-DTTCompatibility-class),
13
- features_assnlist
(zzz-DTTCompatibility-class),
13
- features_assnlist,FeatureGroup-method
(zzz-DTTCompatibility-class),
13
- Features_assnlist-class
(zzz-DTTCompatibility-class),
13
- featureShape_assn
(zzz-DTTCompatibility-class),
13
- featureShape_assn,FeatureGroup-method
(zzz-DTTCompatibility-class),
13
- FeatureShape_assn-class
(zzz-DTTCompatibility-class),
13
- featureWidth
(zzz-DTTCompatibility-class),
13
- featureWidth,FeatureGroup-method
(zzz-DTTCompatibility-class),
13
- fiducial (zzz-DTTCompatibility-class),
13
- fiducial,Fiducials_assnlist-method
(zzz-DTTCompatibility-class),
13
- Fiducial-class
(zzz-DTTCompatibility-class),
13
- fiducials_assnlist

(zzz-DTTCompatibility-class),
13

fiducials_assnlist,ArrayGroup-method
(zzz-DTTCompatibility-class),
13

Fiducials_assnlist-class
(zzz-DTTCompatibility-class),
13

fiducialType_assn
(zzz-DTTCompatibility-class),
13

fiducialType_assn,Fiducial-method
(zzz-DTTCompatibility-class),
13

FiducialType_assn-class
(zzz-DTTCompatibility-class),
13

filenameURI
(zzz-DTTCompatibility-class),
13

filenameURI,DataExternal-method
(zzz-DTTCompatibility-class),
13

firstName (zzz-DTTCompatibility-class),
13

firstName,Person-method
(zzz-DTTCompatibility-class),
13

format_assn
(zzz-DTTCompatibility-class),
13

format_assn,Image-method
(zzz-DTTCompatibility-class),
13

Format_assn-class
(zzz-DTTCompatibility-class),
13

guid (zzz-ARRCompatibility-class), 12

guid,ArraySetFile-method
(zzz-ARRCompatibility-class),
12

guid,PhysicalArray-method
(zzz-ARRCompatibility-class),
12

guid,TemplateFile-method
(zzz-ARRCompatibility-class),
12

hardware (zzz-DTTCompatibility-class),
13

hardware,Hardware_assnlist-method
(zzz-DTTCompatibility-class),

13

Hardware-class
(zzz-DTTCompatibility-class),
13

hardware_assnlist
(zzz-DTTCompatibility-class),
13

hardware_assnlist,Protocol_package-method
(zzz-DTTCompatibility-class),
13

Hardware_assnlist-class
(zzz-DTTCompatibility-class),
13

hardware_assnref
(zzz-DTTCompatibility-class),
13

hardware_assnref,HardwareApplication-method
(zzz-DTTCompatibility-class),
13

hardware_assnref,Software-method
(zzz-DTTCompatibility-class),
13

Hardware_assnref-class
(zzz-DTTCompatibility-class),
13

hardware_ref
(zzz-DTTCompatibility-class),
13

hardware_ref,Hardware_assnref-method
(zzz-DTTCompatibility-class),
13

hardware_ref,Hardwares_assnreflist-method
(zzz-DTTCompatibility-class),
13

Hardware_ref-class
(zzz-DTTCompatibility-class),
13

hardwareApplication
(zzz-DTTCompatibility-class),
13

hardwareApplication,HardwareApplications_assnlist-method
(zzz-DTTCompatibility-class),
13

HardwareApplication-class
(zzz-DTTCompatibility-class),
13

hardwareApplications_assnlist
(zzz-DTTCompatibility-class),
13

hardwareApplications_assnlist,ProtocolApplication-method
(zzz-DTTCompatibility-class),
13

- HardwareApplications_assnlist-class
(zzz-DTTCompatibility-class),
13
- hardwareManufacturers_assnreflist
(zzz-DTTCompatibility-class),
13
- hardwareManufacturers_assnreflist,Hardware-method
(zzz-DTTCompatibility-class),
13
- HardwareManufacturers_assnreflist-class
(zzz-DTTCompatibility-class),
13
- hardwares_assnreflist
(zzz-DTTCompatibility-class),
13
- hardwares_assnreflist,Protocol-method
(zzz-DTTCompatibility-class),
13
- Hardwares_assnreflist-class
(zzz-DTTCompatibility-class),
13
- higherLevelAnalysis_package
(zzz-DTTCompatibility-class),
13
- higherLevelAnalysis_package,MAGE-ML-method
(zzz-DTTCompatibility-class),
13
- HigherLevelAnalysis_package-class
(zzz-DTTCompatibility-class),
13
- hybridization
(zzz-DTTCompatibility-class),
13
- hybridization,BioAssayCreation_assn-method
(zzz-DTTCompatibility-class),
13
- Hybridization-class
(zzz-DTTCompatibility-class),
13
- identifier
(zzz-DTTCompatibility-class),
13
- identifier,Array-method
(zzz-DTTCompatibility-class),
13
- identifier,Array_ref-method
(zzz-DTTCompatibility-class),
13
- identifier,ArrayDesign-method
(zzz-DTTCompatibility-class),
13
- identifier,ArrayDesign_ref-method
(zzz-DTTCompatibility-class),
13
- identifier,ArrayGroup-method
(zzz-DTTCompatibility-class),
13
- identifier,ArrayGroup_ref-method
(zzz-DTTCompatibility-class),
13
- identifier,ArrayManufacture-method
(zzz-DTTCompatibility-class),
13
- identifier,ArrayManufacture_ref-method
(zzz-DTTCompatibility-class),
13
- identifier,BioAssay_ref-method
(zzz-DTTCompatibility-class),
13
- identifier,BioAssayCreation-method
(zzz-DTTCompatibility-class),
13
- identifier,BioAssayData_ref-method
(zzz-DTTCompatibility-class),
13
- identifier,BioAssayDataCluster-method
(zzz-DTTCompatibility-class),
13
- identifier,BioAssayDataCluster_ref-method
(zzz-DTTCompatibility-class),
13
- identifier,BioAssayDimension-method
(zzz-DTTCompatibility-class),
13
- identifier,BioAssayDimension_ref-method
(zzz-DTTCompatibility-class),
13
- identifier,BioAssayMap-method
(zzz-DTTCompatibility-class),
13
- identifier,BioAssayMap_ref-method
(zzz-DTTCompatibility-class),
13
- identifier,BioAssayTreatment-method
(zzz-DTTCompatibility-class),
13
- identifier,BioMaterial_ref-method
(zzz-DTTCompatibility-class),
13
- identifier,BioSample-method
(zzz-DTTCompatibility-class),
13
- identifier,BioSample_ref-method
(zzz-DTTCompatibility-class),
13

- 13
- identifier, BioSequence-method
(zzz-DTTCompatibility-class),
13
- identifier, BioSequence_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, BioSource-method
(zzz-DTTCompatibility-class),
13
- identifier, BioSource_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, Channel-method
(zzz-DTTCompatibility-class),
13
- identifier, Channel_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, CompositeCompositeMap-method
(zzz-DTTCompatibility-class),
13
- identifier, CompositeCompositeMap_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, CompositeGroup-method
(zzz-DTTCompatibility-class),
13
- identifier, CompositeGroup_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, CompositeSequence-method
(zzz-DTTCompatibility-class),
13
- identifier, CompositeSequence_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, CompositeSequenceDimension-method
(zzz-DTTCompatibility-class),
13
- identifier, CompositeSequenceDimension_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, Compound-method
(zzz-DTTCompatibility-class),
13
- identifier, Compound_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, ConfidenceIndicator_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, Contact_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, Database-method
(zzz-DTTCompatibility-class),
13
- identifier, Database_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, DerivedBioAssay-method
(zzz-DTTCompatibility-class),
13
- identifier, DerivedBioAssay_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, DerivedBioAssayData-method
(zzz-DTTCompatibility-class),
13
- identifier, DerivedBioAssayData_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, DerivedSignal-method
(zzz-DTTCompatibility-class),
13
- identifier, DerivedSignal_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, DesignElement_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, DesignElementDimension_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, DesignElementMap_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, Error-method
(zzz-DTTCompatibility-class),
13
- identifier, Error_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, ExpectedValue-method
(zzz-DTTCompatibility-class),
13
- identifier, ExpectedValue_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, Experiment-method
(zzz-DTTCompatibility-class),
13
- identifier, ExperimentalFactor-method

- (zzz-DTTCompatibility-class),
13
- identifier, ExperimentalFactor_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, FactorValue-method
(zzz-DTTCompatibility-class),
13
- identifier, FactorValue_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, Failed-method
(zzz-DTTCompatibility-class),
13
- identifier, Failed_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, Feature-method
(zzz-DTTCompatibility-class),
13
- identifier, Feature_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, FeatureDimension-method
(zzz-DTTCompatibility-class),
13
- identifier, FeatureDimension_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, FeatureExtraction-method
(zzz-DTTCompatibility-class),
13
- identifier, FeatureGroup-method
(zzz-DTTCompatibility-class),
13
- identifier, FeatureGroup_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, FeatureReporterMap-method
(zzz-DTTCompatibility-class),
13
- identifier, FeatureReporterMap_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, Hardware-method
(zzz-DTTCompatibility-class),
13
- identifier, Hardware_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, Hybridization-method
(zzz-DTTCompatibility-class),
13
- identifier, Image-method
(zzz-DTTCompatibility-class),
13
- identifier, Image_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, ImageAcquisition-method
(zzz-DTTCompatibility-class),
13
- identifier, LabeledExtract-method
(zzz-DTTCompatibility-class),
13
- identifier, LabeledExtract_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, MAGE-ML-method
(zzz-DTTCompatibility-class),
13
- identifier, MeasuredBioAssay-method
(zzz-DTTCompatibility-class),
13
- identifier, MeasuredBioAssay_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, MeasuredBioAssayData-method
(zzz-DTTCompatibility-class),
13
- identifier, MeasuredBioAssayData_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, MeasuredSignal-method
(zzz-DTTCompatibility-class),
13
- identifier, MeasuredSignal_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, Organization-method
(zzz-DTTCompatibility-class),
13
- identifier, Organization_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, Parameter-method
(zzz-DTTCompatibility-class),
13
- identifier, Parameter_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, Person-method
(zzz-DTTCompatibility-class),
13

- identifier, Person_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, PhysicalArrayDesign-method
(zzz-DTTCompatibility-class),
13
- identifier, PhysicalArrayDesign_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, PhysicalBioAssay-method
(zzz-DTTCompatibility-class),
13
- identifier, PhysicalBioAssay_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, PresentAbsent-method
(zzz-DTTCompatibility-class),
13
- identifier, PresentAbsent_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, Protocol-method
(zzz-DTTCompatibility-class),
13
- identifier, Protocol_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, PValue-method
(zzz-DTTCompatibility-class),
13
- identifier, PValue_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, QuantitationType_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, QuantitationTypeDimension-method
(zzz-DTTCompatibility-class),
13
- identifier, QuantitationTypeDimension_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, QuantitationTypeMap-method
(zzz-DTTCompatibility-class),
13
- identifier, QuantitationTypeMap_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, Ratio-method
(zzz-DTTCompatibility-class),
13
- identifier, Ratio_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, Reporter-method
(zzz-DTTCompatibility-class),
13
- identifier, Reporter_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, ReporterCompositeMap-method
(zzz-DTTCompatibility-class),
13
- identifier, ReporterCompositeMap_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, ReporterDimension-method
(zzz-DTTCompatibility-class),
13
- identifier, ReporterDimension_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, ReporterGroup-method
(zzz-DTTCompatibility-class),
13
- identifier, ReporterGroup_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, Security-method
(zzz-DTTCompatibility-class),
13
- identifier, Security_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, SecurityGroup-method
(zzz-DTTCompatibility-class),
13
- identifier, SecurityGroup_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, Software-method
(zzz-DTTCompatibility-class),
13
- identifier, Software_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, SpecializedQuantitationType-method
(zzz-DTTCompatibility-class),
13
- identifier, SpecializedQuantitationType_ref-method
(zzz-DTTCompatibility-class),
13
- identifier, StandardQuantitationType_ref-method
(zzz-DTTCompatibility-class),
13

- 13
- identifier,Transformation-method
(zzz-DTTCompatibility-class),
13
- identifier,Treatment-method
(zzz-DTTCompatibility-class),
13
- identifier,Zone-method
(zzz-DTTCompatibility-class),
13
- identifier,Zone_ref-method
(zzz-DTTCompatibility-class),
13
- identifierLIMS_assn
(zzz-DTTCompatibility-class),
13
- identifierLIMS_assn,ManufactureLIMS-method
(zzz-DTTCompatibility-class),
13
- identifierLIMS_assn,ManufactureLIMSBiomaterial-method
(zzz-DTTCompatibility-class),
13
- IdentifierLIMS_assn-class
(zzz-DTTCompatibility-class),
13
- Image-class
(zzz-DTTCompatibility-class),
13
- image_ref (zzz-DTTCompatibility-class),
13
- image_ref,Images_assnreflist-method
(zzz-DTTCompatibility-class),
13
- Image_ref-class
(zzz-DTTCompatibility-class),
13
- imageAcquisition
(zzz-DTTCompatibility-class),
13
- imageAcquisition,BioAssayTreatments_assnreflist-method
(zzz-DTTCompatibility-class),
13
- ImageAcquisition-class
(zzz-DTTCompatibility-class),
13
- images_assnreflist
(zzz-DTTCompatibility-class),
13
- images_assnreflist,ImageAcquisition-method
(zzz-DTTCompatibility-class),
13
- Images_assnreflist-class
(zzz-DTTCompatibility-class),
13
- immobilizedCharacteristics_assnreflist
(zzz-DTTCompatibility-class),
13
- immobilizedCharacteristics_assnreflist,Reporter-method
(zzz-DTTCompatibility-class),
13
- ImmobilizedCharacteristics_assnreflist-class
(zzz-DTTCompatibility-class),
13
- information_assnref
(zzz-DTTCompatibility-class),
13
- information_assnref,Array-method
(zzz-DTTCompatibility-class),
13
- Information_assnref-class
(zzz-DTTCompatibility-class),
13
- Internal-class, 4
- isApproximateLength
(zzz-DTTCompatibility-class),
13
- isApproximateLength,BioSequence-method
(zzz-DTTCompatibility-class),
13
- isBackground
(zzz-DTTCompatibility-class),
13
- isBackground,DerivedSignal-method
(zzz-DTTCompatibility-class),
13
- isBackground,Error-method
(zzz-DTTCompatibility-class),
13
- isBackground,ExpectedValue-method
(zzz-DTTCompatibility-class),
13
- isBackground,Failed-method
(zzz-DTTCompatibility-class),
13
- isBackground,MeasuredSignal-method
(zzz-DTTCompatibility-class),
13
- isBackground,PresentAbsent-method
(zzz-DTTCompatibility-class),
13
- isBackground,PValue-method
(zzz-DTTCompatibility-class),
13
- isBackground,Ratio-method

- (zzz-DTTCompatibility-class),
13
- isBackground, SpecializedQuantitationType-method
(zzz-DTTCompatibility-class),
13
- isCircular
(zzz-DTTCompatibility-class),
13
- isCircular, BioSequence-method
(zzz-DTTCompatibility-class),
13
- isSolvent (zzz-DTTCompatibility-class),
13
- isSolvent, Compound-method
(zzz-DTTCompatibility-class),
13
- issue (zzz-DTTCompatibility-class), 13
- issue, BibliographicReference-method
(zzz-DTTCompatibility-class),
13
- kindCV (zzz-DTTCompatibility-class), 13
- kindCV, Measurement-method
(zzz-DTTCompatibility-class),
13
- labeledExtract
(zzz-DTTCompatibility-class),
13
- labeledExtract, BioMaterial_assnlist-method
(zzz-DTTCompatibility-class),
13
- LabeledExtract-class
(zzz-DTTCompatibility-class),
13
- labeledExtract_ref
(zzz-DTTCompatibility-class),
13
- labeledExtract_ref, BioMaterial_assnref-method
(zzz-DTTCompatibility-class),
13
- LabeledExtract_ref-class
(zzz-DTTCompatibility-class),
13
- labels_assnreflist
(zzz-DTTCompatibility-class),
13
- labels_assnreflist, Channel-method
(zzz-DTTCompatibility-class),
13
- labels_assnreflist, LabeledExtract-method
(zzz-DTTCompatibility-class),
13
- Labels_assnreflist-class
(zzz-DTTCompatibility-class),
13
- lastName (zzz-DTTCompatibility-class),
13
- lastName, Person-method
(zzz-DTTCompatibility-class),
13
- length, ArrayGroup-method
(zzz-DTTCompatibility-class),
13
- length, BioSequence-method
(zzz-DTTCompatibility-class),
13
- libraryPackageName
(zzz-ARRCompatibility-class),
12
- libraryPackageName, PhysicalArray-method
(zzz-ARRCompatibility-class),
12
- list, 4
- lowerRightX
(zzz-DTTCompatibility-class),
13
- lowerRightX, Zone-method
(zzz-DTTCompatibility-class),
13
- lowerRightY
(zzz-DTTCompatibility-class),
13
- lowerRightY, Zone-method
(zzz-DTTCompatibility-class),
13
- MAGE-ML-class
(zzz-DTTCompatibility-class),
13
- make (zzz-DTTCompatibility-class), 13
- make, Hardware-method
(zzz-DTTCompatibility-class),
13
- manufactureLIMS
(zzz-DTTCompatibility-class),
13
- manufactureLIMS, FeatureLIMSS_assnlist-method
(zzz-DTTCompatibility-class),
13
- ManufactureLIMS-class
(zzz-DTTCompatibility-class),
13
- manufactureLIMSBiomaterial
(zzz-DTTCompatibility-class),
13

- manufactureLIMSBiomaterial, FeatureLIMSS_assnlist-method
 (zzz-DTTCompatibility-class),
[13](#)
- ManufactureLIMSBiomaterial-class
 (zzz-DTTCompatibility-class),
[13](#)
- manufacturingDate
 (zzz-DTTCompatibility-class),
[13](#)
- manufacturingDate, ArrayManufacture-method
 (zzz-DTTCompatibility-class),
[13](#)
- massUnit (zzz-DTTCompatibility-class),
[13](#)
- massUnit, Unit_assn-method
 (zzz-DTTCompatibility-class),
[13](#)
- MassUnit-class
 (zzz-DTTCompatibility-class),
[13](#)
- masterFileGUID
 (zzz-ARRCompatibility-class),
[12](#)
- masterFileGUID, PhysicalArray-method
 (zzz-ARRCompatibility-class),
[12](#)
- masterFileName
 (zzz-ARRCompatibility-class),
[12](#)
- masterFileName, PhysicalArray-method
 (zzz-ARRCompatibility-class),
[12](#)
- materialType_assn
 (zzz-DTTCompatibility-class),
[13](#)
- materialType_assn, BioSample-method
 (zzz-DTTCompatibility-class),
[13](#)
- materialType_assn, BioSource-method
 (zzz-DTTCompatibility-class),
[13](#)
- materialType_assn, LabeledExtract-method
 (zzz-DTTCompatibility-class),
[13](#)
- MaterialType_assn-class
 (zzz-DTTCompatibility-class),
[13](#)
- measuredBioAssay
 (zzz-DTTCompatibility-class),
[13](#)
- measuredBioAssay, BioAssay_assnlist-method
 (zzz-DTTCompatibility-class),
[13](#)
- measuredBioAssay, MeasuredBioAssay_assnlist-method
 (zzz-DTTCompatibility-class),
[13](#)
- MeasuredBioAssay-class
 (zzz-DTTCompatibility-class),
[13](#)
- MeasuredBioAssay_assnlist-class
 (zzz-DTTCompatibility-class),
[13](#)
- measuredBioAssay_ref
 (zzz-DTTCompatibility-class),
[13](#)
- measuredBioAssay_ref, BioAssay_assnref-method
 (zzz-DTTCompatibility-class),
[13](#)
- measuredBioAssay_ref, BioAssays_assnreflist-method
 (zzz-DTTCompatibility-class),
[13](#)
- measuredBioAssay_ref, MeasuredBioAssayTarget_assnref-method
 (zzz-DTTCompatibility-class),
[13](#)
- measuredBioAssay_ref, SourceBioAssays_assnreflist-method
 (zzz-DTTCompatibility-class),
[13](#)
- measuredBioAssay_ref, TopLevelBioAssays_assnreflist-method
 (zzz-DTTCompatibility-class),
[13](#)
- MeasuredBioAssay_ref-class
 (zzz-DTTCompatibility-class),
[13](#)
- measuredBioAssayData
 (zzz-DTTCompatibility-class),
[13](#)
- measuredBioAssayData, BioAssayData_assnlist-method
 (zzz-DTTCompatibility-class),
[13](#)
- measuredBioAssayData, MeasuredBioAssayData_assnlist-method
 (zzz-DTTCompatibility-class),
[13](#)
- MeasuredBioAssayData-class
 (zzz-DTTCompatibility-class),
[13](#)
- MeasuredBioAssayData_assnlist-class
 (zzz-DTTCompatibility-class),
[13](#)
- measuredBioAssayData_assnreflist
 (zzz-DTTCompatibility-class),
[13](#)
- measuredBioAssayData_assnreflist, MeasuredBioAssay-method
 (zzz-DTTCompatibility-class),
[13](#)

MeasuredBioAssayData_assnreflist-class (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 measurement, ActionMeasurement_assn-method
 measuredBioAssayData_ref (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 measurement, DefaultValue_assn-method
 measuredBioAssayData_ref, BioAssayData_assnreflist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 measurement, Measurement_assn-method
 measuredBioAssayData_ref, BioAssayDataSources_assnreflist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 Measurement-class
 measuredBioAssayData_ref, ClusterBioAssayData_assnreflist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 measurement_assn
 measuredBioAssayData_ref, MeasuredBioAssayData_assnreflist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 measurement_assn, BioMaterialMeasurement-method
 MeasuredBioAssayData_ref-class (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 measurement_assn, CompoundMeasurement-method
 MeasuredBioAssayTarget_assnref-class (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 measurement_assn, FactorValue-method
 measuredSignal (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 Measurement_assn-class
 measuredSignal, QuantitationType_assnreflist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 measurement_package
 MeasuredSignal-class (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 measurement_package, MAGE-ML-method
 measuredSignal_ref (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 Measurement_package-class
 measuredSignal_ref, QuantitationType_assnreflist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 mediaCol (zzz-ARRCompatibility-class),
 mediaCol, PhysicalArray-method
 measuredSignal_ref, QuantitationTypes_assnreflist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 (zzz-ARRCompatibility-class),
 mediaFileGUID
 measuredSignal_ref, SourcesQuantitationType_assnreflist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 (zzz-ARRCompatibility-class),
 mediaFileGUID, PhysicalArray-method
 measuredSignal_ref, TargetQuantitationType_assnreflist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 (zzz-ARRCompatibility-class),
 MeasuredSignal_ref-class (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 mediaFileName
 measurement (zzz-ARRCompatibility-class),
 12

- mediaFileName,PhysicalArray-method
(zzz-ARRCompatibility-class),
[12](#)
- mediaRow (zzz-ARRCompatibility-class),
[12](#)
- mediaRow,PhysicalArray-method
(zzz-ARRCompatibility-class),
[12](#)
- mediaType (zzz-ARRCompatibility-class),
[12](#)
- mediaType,PhysicalArray-method
(zzz-ARRCompatibility-class),
[12](#)
- members_assnreflist
(zzz-DTTCompatibility-class),
[13](#)
- members_assnreflist,SecurityGroup-method
(zzz-DTTCompatibility-class),
[13](#)
- Members_assnreflist-class
(zzz-DTTCompatibility-class),
[13](#)
- midInitials
(zzz-DTTCompatibility-class),
[13](#)
- midInitials,Person-method
(zzz-DTTCompatibility-class),
[13](#)
- mismatchInformation
(zzz-DTTCompatibility-class),
[13](#)
- mismatchInformation,MismatchInformation_assnlist-method
(zzz-DTTCompatibility-class),
[13](#)
- MismatchInformation-class
(zzz-DTTCompatibility-class),
[13](#)
- mismatchInformation_assnlist
(zzz-DTTCompatibility-class),
[13](#)
- mismatchInformation_assnlist,CompositePosition-method
(zzz-DTTCompatibility-class),
[13](#)
- mismatchInformation_assnlist,FeatureInformation-method
(zzz-DTTCompatibility-class),
[13](#)
- mismatchInformation_assnlist,ReporterPosition-method
(zzz-DTTCompatibility-class),
[13](#)
- MismatchInformation_assnlist-class
(zzz-DTTCompatibility-class),
[13](#)
- model (zzz-DTTCompatibility-class), [13](#)
- model,Hardware-method
(zzz-DTTCompatibility-class),
[13](#)
- name (zzz-DTTCompatibility-class), [13](#)
- name,Array-method
(zzz-DTTCompatibility-class),
[13](#)
- name,Array_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name,ArrayAttribute-method
(zzz-ARRCompatibility-class),
[12](#)
- name,ArrayDesign-method
(zzz-DTTCompatibility-class),
[13](#)
- name,ArrayDesign_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name,ArrayGroup-method
(zzz-DTTCompatibility-class),
[13](#)
- name,ArrayGroup_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name,ArrayManufacture-method
(zzz-DTTCompatibility-class),
[13](#)
- name,ArrayManufacture_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name,BioAssay_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name,BioAssayCreation-method
(zzz-DTTCompatibility-class),
[13](#)
- name,BioAssayData_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name,BioAssayDataCluster-method
(zzz-DTTCompatibility-class),
[13](#)
- name,BioAssayDataCluster_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name,BioAssayDimension-method
(zzz-DTTCompatibility-class),
[13](#)
- name,BioAssayDimension_ref-method
(zzz-DTTCompatibility-class),
[13](#)

- [13](#)
- name, BioAssayMap-method
(zzz-DTTCompatibility-class),
[13](#)
- name, BioAssayMap_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, BioAssayTreatment-method
(zzz-DTTCompatibility-class),
[13](#)
- name, BioMaterial_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, BioSample-method
(zzz-DTTCompatibility-class),
[13](#)
- name, BioSample_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, BioSequence-method
(zzz-DTTCompatibility-class),
[13](#)
- name, BioSequence_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, BioSource-method
(zzz-DTTCompatibility-class),
[13](#)
- name, BioSource_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, Channel-method
(zzz-DTTCompatibility-class),
[13](#)
- name, Channel_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, CompositeCompositeMap-method
(zzz-DTTCompatibility-class),
[13](#)
- name, CompositeCompositeMap_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, CompositeGroup-method
(zzz-DTTCompatibility-class),
[13](#)
- name, CompositeGroup_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, CompositeSequence-method
(zzz-DTTCompatibility-class),
[13](#)
- name, CompositeSequence_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, CompositeSequenceDimension-method
(zzz-DTTCompatibility-class),
[13](#)
- name, CompositeSequenceDimension_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, Compound-method
(zzz-DTTCompatibility-class),
[13](#)
- name, Compound_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, ConfidenceIndicator_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, Contact_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, Database-method
(zzz-DTTCompatibility-class),
[13](#)
- name, Database_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, DerivedBioAssay-method
(zzz-DTTCompatibility-class),
[13](#)
- name, DerivedBioAssay_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, DerivedBioAssayData-method
(zzz-DTTCompatibility-class),
[13](#)
- name, DerivedBioAssayData_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, DerivedSignal-method
(zzz-DTTCompatibility-class),
[13](#)
- name, DerivedSignal_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, DesignElement_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, DesignElementDimension_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, DesignElementMap_ref-method

- (zzz-DTTCompatibility-class),
13
- name,Error-method
(zzz-DTTCompatibility-class),
13
- name,Error_ref-method
(zzz-DTTCompatibility-class),
13
- name,ExpectedValue-method
(zzz-DTTCompatibility-class),
13
- name,ExpectedValue_ref-method
(zzz-DTTCompatibility-class),
13
- name,Experiment-method
(zzz-DTTCompatibility-class),
13
- name,ExperimentalFactor-method
(zzz-DTTCompatibility-class),
13
- name,ExperimentalFactor_ref-method
(zzz-DTTCompatibility-class),
13
- name,FactorValue-method
(zzz-DTTCompatibility-class),
13
- name,FactorValue_ref-method
(zzz-DTTCompatibility-class),
13
- name,Failed-method
(zzz-DTTCompatibility-class),
13
- name,Failed_ref-method
(zzz-DTTCompatibility-class),
13
- name,Feature-method
(zzz-DTTCompatibility-class),
13
- name,Feature_ref-method
(zzz-DTTCompatibility-class),
13
- name,FeatureDimension-method
(zzz-DTTCompatibility-class),
13
- name,FeatureDimension_ref-method
(zzz-DTTCompatibility-class),
13
- name,FeatureExtraction-method
(zzz-DTTCompatibility-class),
13
- name,FeatureGroup-method
(zzz-DTTCompatibility-class),
13
- name,FeatureGroup_ref-method
(zzz-DTTCompatibility-class),
13
- name,FeatureReporterMap-method
(zzz-DTTCompatibility-class),
13
- name,FeatureReporterMap_ref-method
(zzz-DTTCompatibility-class),
13
- name,Hardware-method
(zzz-DTTCompatibility-class),
13
- name,Hardware_ref-method
(zzz-DTTCompatibility-class),
13
- name,Hybridization-method
(zzz-DTTCompatibility-class),
13
- name,Image-method
(zzz-DTTCompatibility-class),
13
- name,Image_ref-method
(zzz-DTTCompatibility-class),
13
- name,ImageAcquisition-method
(zzz-DTTCompatibility-class),
13
- name,LabeledExtract-method
(zzz-DTTCompatibility-class),
13
- name,LabeledExtract_ref-method
(zzz-DTTCompatibility-class),
13
- name,MAGE-ML-method
(zzz-DTTCompatibility-class),
13
- name,MeasuredBioAssay-method
(zzz-DTTCompatibility-class),
13
- name,MeasuredBioAssay_ref-method
(zzz-DTTCompatibility-class),
13
- name,MeasuredBioAssayData-method
(zzz-DTTCompatibility-class),
13
- name,MeasuredBioAssayData_ref-method
(zzz-DTTCompatibility-class),
13
- name,MeasuredSignal-method
(zzz-DTTCompatibility-class),
13

- name, MeasuredSignal_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, NameValueType-method
(zzz-DTTCompatibility-class),
[13](#)
- name, NodeValue-method
(zzz-DTTCompatibility-class),
[13](#)
- name, Organization-method
(zzz-DTTCompatibility-class),
[13](#)
- name, Organization_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, Parameter-method
(zzz-DTTCompatibility-class),
[13](#)
- name, Parameter_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, Person-method
(zzz-DTTCompatibility-class),
[13](#)
- name, Person_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, PhysicalArrayDesign-method
(zzz-DTTCompatibility-class),
[13](#)
- name, PhysicalArrayDesign_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, PhysicalBioAssay-method
(zzz-DTTCompatibility-class),
[13](#)
- name, PhysicalBioAssay_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, PresentAbsent-method
(zzz-DTTCompatibility-class),
[13](#)
- name, PresentAbsent_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, Protocol-method
(zzz-DTTCompatibility-class),
[13](#)
- name, Protocol_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, PValue-method
(zzz-DTTCompatibility-class),
[13](#)
- name, PValue_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, QuantitationType_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, QuantitationTypeDimension-method
(zzz-DTTCompatibility-class),
[13](#)
- name, QuantitationTypeDimension_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, QuantitationTypeMap-method
(zzz-DTTCompatibility-class),
[13](#)
- name, QuantitationTypeMap_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, Ratio-method
(zzz-DTTCompatibility-class),
[13](#)
- name, Ratio_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, Reporter-method
(zzz-DTTCompatibility-class),
[13](#)
- name, Reporter_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, ReporterCompositeMap-method
(zzz-DTTCompatibility-class),
[13](#)
- name, ReporterCompositeMap_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, ReporterDimension-method
(zzz-DTTCompatibility-class),
[13](#)
- name, ReporterDimension_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, ReporterGroup-method
(zzz-DTTCompatibility-class),
[13](#)
- name, ReporterGroup_ref-method
(zzz-DTTCompatibility-class),
[13](#)
- name, Security-method
(zzz-DTTCompatibility-class),
[13](#)

- 13
- name, Security_ref-method
(zzz-DTTCompatibility-class),
13
- name, SecurityGroup-method
(zzz-DTTCompatibility-class),
13
- name, SecurityGroup_ref-method
(zzz-DTTCompatibility-class),
13
- name, Software-method
(zzz-DTTCompatibility-class),
13
- name, Software_ref-method
(zzz-DTTCompatibility-class),
13
- name, SpecializedQuantitationType-method
(zzz-DTTCompatibility-class),
13
- name, SpecializedQuantitationType_ref-method
(zzz-DTTCompatibility-class),
13
- name, StandardQuantitationType_ref-method
(zzz-DTTCompatibility-class),
13
- name, Transformation-method
(zzz-DTTCompatibility-class),
13
- name, Treatment-method
(zzz-DTTCompatibility-class),
13
- name, UserAttribute-method
(zzz-ARRCompatibility-class),
12
- name, Zone-method
(zzz-DTTCompatibility-class),
13
- name, Zone_ref-method
(zzz-DTTCompatibility-class),
13
- names, AffxArray-method
(zzz-NetAffxCompatibility-class),
14
- names, NetAffxResource-method
(NetAffxResource-class), 5
- namespace (zzz-ARRCompatibility-class),
12
- namespace, UserAttribute-method
(zzz-ARRCompatibility-class),
12
- nameValueType
(zzz-DTTCompatibility-class),
13
- nameValueType, PropertySets_assnlist-method
(zzz-DTTCompatibility-class),
13
- nameValueType, QualityControlStatistics_assnlist-method
(zzz-DTTCompatibility-class),
13
- nameValueType, SummaryStatistics_assnlist-method
(zzz-DTTCompatibility-class),
13
- NameValueType-class
(zzz-DTTCompatibility-class),
13
- NetAffxCompatibility, 6, 15
- NetAffxCompatibility-class
(AffyCompatible-class), 3
- NetAffxResource, 3, 4, 5, 9, 15
- NetAffxResource-accessors (accessors), 2
- NetAffxResource-class, 5
- newSequence
(zzz-DTTCompatibility-class),
13
- newSequence, MismatchInformation-method
(zzz-DTTCompatibility-class),
13
- node (zzz-DTTCompatibility-class), 13
- node, Nodes_assnlist-method
(zzz-DTTCompatibility-class),
13
- Node-class
(zzz-DTTCompatibility-class),
13
- nodeContents
(zzz-DTTCompatibility-class),
13
- nodeContents, NodeContents_assnlist-method
(zzz-DTTCompatibility-class),
13
- NodeContents-class
(zzz-DTTCompatibility-class),
13
- nodeContents_assnlist
(zzz-DTTCompatibility-class),
13
- nodeContents_assnlist, Node-method
(zzz-DTTCompatibility-class),
13
- NodeContents_assnlist-class
(zzz-DTTCompatibility-class),
13
- nodes_assnlist
(zzz-DTTCompatibility-class),

- 13
- nodes_assnlist, BioAssayDataCluster-method
(zzz-DTTCompatibility-class),
13
- nodes_assnlist, Node-method
(zzz-DTTCompatibility-class),
13
- Nodes_assnlist-class
(zzz-DTTCompatibility-class),
13
- nodeValue (zzz-DTTCompatibility-class),
13
- nodeValue, NodeValue_assnlist-method
(zzz-DTTCompatibility-class),
13
- NodeValue-class
(zzz-DTTCompatibility-class),
13
- nodeValue_assnlist
(zzz-DTTCompatibility-class),
13
- nodeValue_assnlist, Node-method
(zzz-DTTCompatibility-class),
13
- NodeValue_assnlist-class
(zzz-DTTCompatibility-class),
13
- normalizationDescription_assn
(zzz-DTTCompatibility-class),
13
- normalizationDescription_assn, ExperimentDesign-method
(zzz-DTTCompatibility-class),
13
- NormalizationDescription_assn-class
(zzz-DTTCompatibility-class),
13
- numArrays (zzz-DTTCompatibility-class),
13
- numArrays, ArrayGroup-method
(zzz-DTTCompatibility-class),
13
- numberOfFeatures
(zzz-DTTCompatibility-class),
13
- numberOfFeatures, ArrayDesign-method
(zzz-DTTCompatibility-class),
13
- numberOfFeatures, PhysicalArrayDesign-method
(zzz-DTTCompatibility-class),
13
- numFeaturesPerCol
(zzz-DTTCompatibility-class),
13
- numFeaturesPerCol, ZoneLayout-method
(zzz-DTTCompatibility-class),
13
- numFeaturesPerRow
(zzz-DTTCompatibility-class),
13
- numFeaturesPerRow, ZoneLayout-method
(zzz-DTTCompatibility-class),
13
- ontologyEntries_assnlist
(zzz-DTTCompatibility-class),
13
- ontologyEntries_assnlist, BioSequence-method
(zzz-DTTCompatibility-class),
13
- OntologyEntries_assnlist-class
(zzz-DTTCompatibility-class),
13
- ontologyEntry
(zzz-DTTCompatibility-class),
13
- ontologyEntry, Action_assn-method
(zzz-DTTCompatibility-class),
13
- ontologyEntry, Annotations_assnlist-method
(zzz-DTTCompatibility-class),
13
- ontologyEntry, Associations_assnlist-method
(zzz-DTTCompatibility-class),
13
- ontologyEntry, Category_assn-method
(zzz-DTTCompatibility-class),
13
- ontologyEntry, Characteristics_assnlist-method
(zzz-DTTCompatibility-class),
13
- ontologyEntry, CompoundIndices_assnlist-method
(zzz-DTTCompatibility-class),
13
- ontologyEntry, ControlType_assn-method
(zzz-DTTCompatibility-class),
13
- ontologyEntry, DataType_assn-method
(zzz-DTTCompatibility-class),
13
- ontologyEntry, DefectType_assn-method
(zzz-DTTCompatibility-class),
13
- ontologyEntry, FailTypes_assnlist-method
(zzz-DTTCompatibility-class),
13

- ontologyEntry, FeatureShape_assn-method
(zzz-DTTCompatibility-class),
13
- ontologyEntry, FiducialType_assn-method
(zzz-DTTCompatibility-class),
13
- ontologyEntry, Format_assn-method
(zzz-DTTCompatibility-class),
13
- ontologyEntry, MaterialType_assn-method
(zzz-DTTCompatibility-class),
13
- ontologyEntry, OntologyEntries_assnlist-method
(zzz-DTTCompatibility-class),
13
- ontologyEntry, Parameters_assnlist-method
(zzz-DTTCompatibility-class),
13
- ontologyEntry, PolymerType_assn-method
(zzz-DTTCompatibility-class),
13
- ontologyEntry, Roles_assnlist-method
(zzz-DTTCompatibility-class),
13
- ontologyEntry, Scale_assn-method
(zzz-DTTCompatibility-class),
13
- ontologyEntry, Species_assn-method
(zzz-DTTCompatibility-class),
13
- ontologyEntry, SubstrateType_assn-method
(zzz-DTTCompatibility-class),
13
- ontologyEntry, SurfaceType_assn-method
(zzz-DTTCompatibility-class),
13
- ontologyEntry, TechnologyType_assn-method
(zzz-DTTCompatibility-class),
13
- ontologyEntry, Type_assn-method
(zzz-DTTCompatibility-class),
13
- ontologyEntry, Types_assnlist-method
(zzz-DTTCompatibility-class),
13
- ontologyEntry, Value_assn-method
(zzz-DTTCompatibility-class),
13
- ontologyEntry, WarningType_assn-method
(zzz-DTTCompatibility-class),
13
- OntologyEntry-class
(zzz-DTTCompatibility-class),
13
- ontologyReference_assn
(zzz-DTTCompatibility-class),
13
- ontologyReference_assn, OntologyEntry-method
(zzz-DTTCompatibility-class),
13
- OntologyReference_assn-class
(zzz-DTTCompatibility-class),
13
- organization
(zzz-DTTCompatibility-class),
13
- organization, Contact_assnlist-method
(zzz-DTTCompatibility-class),
13
- organization, Organization_assnlist-method
(zzz-DTTCompatibility-class),
13
- Organization-class
(zzz-DTTCompatibility-class),
13
- Organization_assnlist-class
(zzz-DTTCompatibility-class),
13
- organization_ref
(zzz-DTTCompatibility-class),
13
- organization_ref, Affiliation_assnref-method
(zzz-DTTCompatibility-class),
13
- organization_ref, ArrayManufacturers_assnreflist-method
(zzz-DTTCompatibility-class),
13
- organization_ref, Contacts_assnreflist-method
(zzz-DTTCompatibility-class),
13
- organization_ref, DesignProviders_assnreflist-method
(zzz-DTTCompatibility-class),
13
- organization_ref, HardwareManufacturers_assnreflist-method
(zzz-DTTCompatibility-class),
13
- organization_ref, Members_assnreflist-method
(zzz-DTTCompatibility-class),
13
- organization_ref, Owner_assnreflist-method
(zzz-DTTCompatibility-class),
13
- organization_ref, Parent_assnref-method
(zzz-DTTCompatibility-class),
13

- 13
- organization_ref, Performer_assnref-method
(zzz-DTTCompatibility-class),
13
- organization_ref, Providers_assnreflist-method
(zzz-DTTCompatibility-class),
13
- organization_ref, SoftwareManufacturers_assnreflist-method
(zzz-DTTCompatibility-class),
13
- organization_ref, SourceContact_assnreflist-method
(zzz-DTTCompatibility-class),
13
- Organization_ref-class
(zzz-DTTCompatibility-class),
13
- orientationMark
(zzz-DTTCompatibility-class),
13
- orientationMark, ArrayGroup-method
(zzz-DTTCompatibility-class),
13
- orientationMarkPosition
(zzz-DTTCompatibility-class),
13
- orientationMarkPosition, ArrayGroup-method
(zzz-DTTCompatibility-class),
13
- originalProjectName
(zzz-ARRCompatibility-class),
12
- originalProjectName, ArraySetFile-method
(zzz-ARRCompatibility-class),
12
- originRelativeTo
(zzz-DTTCompatibility-class),
13
- originRelativeTo, Array-method
(zzz-DTTCompatibility-class),
13
- otherKind (zzz-DTTCompatibility-class),
13
- otherKind, Measurement-method
(zzz-DTTCompatibility-class),
13
- owner_assnreflist
(zzz-DTTCompatibility-class),
13
- owner_assnreflist, Security-method
(zzz-DTTCompatibility-class),
13
- Owner_assnreflist-class
(zzz-DTTCompatibility-class),
13
- pages (zzz-DTTCompatibility-class), 13
- pages, BibliographicReference-method
(zzz-DTTCompatibility-class),
13
- parameter (zzz-DTTCompatibility-class),
13
- parameter, ParameterTypes_assnlist-method
(zzz-DTTCompatibility-class),
13
- Parameter-class
(zzz-DTTCompatibility-class),
13
- parameter_ref
(zzz-DTTCompatibility-class),
13
- parameter_ref, ParameterType_assnref-method
(zzz-DTTCompatibility-class),
13
- Parameter_ref-class
(zzz-DTTCompatibility-class),
13
- parameters_assnlist
(zzz-DTTCompatibility-class),
13
- parameters_assnlist, BibliographicReference-method
(zzz-DTTCompatibility-class),
13
- Parameters_assnlist-class
(zzz-DTTCompatibility-class),
13
- parameterType_assnref
(zzz-DTTCompatibility-class),
13
- parameterType_assnref, ParameterValue-method
(zzz-DTTCompatibility-class),
13
- ParameterType_assnref-class
(zzz-DTTCompatibility-class),
13
- parameterTypes_assnlist
(zzz-DTTCompatibility-class),
13
- parameterTypes_assnlist, Hardware-method
(zzz-DTTCompatibility-class),
13
- parameterTypes_assnlist, Protocol-method
(zzz-DTTCompatibility-class),
13
- parameterTypes_assnlist, Software-method
(zzz-DTTCompatibility-class),

- 13
- ParameterTypes_assnlist-class
(zzz-DTTCompatibility-class),
13
- parameterValue
(zzz-DTTCompatibility-class),
13
- parameterValue,ParameterValues_assnlist-method
(zzz-DTTCompatibility-class),
13
- ParameterValue-class
(zzz-DTTCompatibility-class),
13
- parameterValues_assnlist
(zzz-DTTCompatibility-class),
13
- parameterValues_assnlist,HardwareApplication-method
(zzz-DTTCompatibility-class),
13
- parameterValues_assnlist,ProtocolApplication-method
(zzz-DTTCompatibility-class),
13
- parameterValues_assnlist,SoftwareApplication-method
(zzz-DTTCompatibility-class),
13
- ParameterValues_assnlist-class
(zzz-DTTCompatibility-class),
13
- parent_assnref
(zzz-DTTCompatibility-class),
13
- parent_assnref,Organization-method
(zzz-DTTCompatibility-class),
13
- Parent_assnref-class
(zzz-DTTCompatibility-class),
13
- password (accessors), 2
- password,NetAffxResource-method
(NetAffxResource-class), 5
- patassignmentMethod
(zzz-ARRCompatibility-class),
12
- patassignmentMethod,PhysicalArray-method
(zzz-ARRCompatibility-class),
12
- performer_assnref
(zzz-DTTCompatibility-class),
13
- performer_assnref,Audit-method
(zzz-DTTCompatibility-class),
13
- Performer_assnref-class
(zzz-DTTCompatibility-class),
13
- performers_assnreflist
(zzz-DTTCompatibility-class),
13
- performers_assnreflist,ProtocolApplication-method
(zzz-DTTCompatibility-class),
13
- Performers_assnreflist-class
(zzz-DTTCompatibility-class),
13
- Person-class
(zzz-DTTCompatibility-class),
13
- Person_assnlist-class
(zzz-DTTCompatibility-class),
13
- person_ref
(zzz-DTTCompatibility-class),
13
- person_ref,ArrayManufacturers_assnreflist-method
(zzz-DTTCompatibility-class),
13
- person_ref,Contacts_assnreflist-method
(zzz-DTTCompatibility-class),
13
- person_ref,DesignProviders_assnreflist-method
(zzz-DTTCompatibility-class),
13
- person_ref,HardwareManufacturers_assnreflist-method
(zzz-DTTCompatibility-class),
13
- person_ref,Members_assnreflist-method
(zzz-DTTCompatibility-class),
13
- person_ref,Owner_assnreflist-method
(zzz-DTTCompatibility-class),
13
- person_ref,Performer_assnref-method
(zzz-DTTCompatibility-class),
13
- person_ref,Performers_assnreflist-method
(zzz-DTTCompatibility-class),
13
- person_ref,Providers_assnreflist-method
(zzz-DTTCompatibility-class),
13
- person_ref,SoftwareManufacturers_assnreflist-method
(zzz-DTTCompatibility-class),
13
- person_ref,SourceContact_assnreflist-method

- (zzz-DTTCompatibility-class),
13
- Person_ref-class
(zzz-DTTCompatibility-class),
13
- phone (zzz-DTTCompatibility-class), 13
- phone, Organization-method
(zzz-DTTCompatibility-class),
13
- phone, Person-method
(zzz-DTTCompatibility-class),
13
- physicalArray
(zzz-ARRCompatibility-class),
12
- physicalArray, PhysicalArrays-method
(zzz-ARRCompatibility-class),
12
- PhysicalArray-class
(zzz-ARRCompatibility-class),
12
- physicalArrayDesign
(zzz-DTTCompatibility-class),
13
- physicalArrayDesign, ArrayDesign_assnlist-method
(zzz-DTTCompatibility-class),
13
- PhysicalArrayDesign-class
(zzz-DTTCompatibility-class),
13
- physicalArrayDesign_ref
(zzz-DTTCompatibility-class),
13
- physicalArrayDesign_ref, ArrayDesign_assnref-method
(zzz-DTTCompatibility-class),
13
- PhysicalArrayDesign_ref-class
(zzz-DTTCompatibility-class),
13
- physicalArrays
(zzz-ARRCompatibility-class),
12
- physicalArrays, ArraySetFile-method
(zzz-ARRCompatibility-class),
12
- PhysicalArrays-class
(zzz-ARRCompatibility-class),
12
- physicalBioAssay
(zzz-DTTCompatibility-class),
13
- physicalBioAssay, BioAssay_assnlist-method
(zzz-DTTCompatibility-class),
13
- physicalBioAssay, PhysicalBioAssay_assnlist-method
(zzz-DTTCompatibility-class),
13
- PhysicalBioAssay-class
(zzz-DTTCompatibility-class),
13
- PhysicalBioAssay_assnlist-class
(zzz-DTTCompatibility-class),
13
- PhysicalBioAssay_assnref-class
(zzz-DTTCompatibility-class),
13
- physicalBioAssay_ref
(zzz-DTTCompatibility-class),
13
- physicalBioAssay_ref, BioAssay_assnref-method
(zzz-DTTCompatibility-class),
13
- physicalBioAssay_ref, BioAssays_assnreflist-method
(zzz-DTTCompatibility-class),
13
- physicalBioAssay_ref, PhysicalBioAssay_assnref-method
(zzz-DTTCompatibility-class),
13
- physicalBioAssay_ref, PhysicalBioAssaySource_assnref-method
(zzz-DTTCompatibility-class),
13
- physicalBioAssay_ref, PhysicalBioAssayTarget_assnref-method
(zzz-DTTCompatibility-class),
13
- physicalBioAssay_ref, SourceBioAssays_assnreflist-method
(zzz-DTTCompatibility-class),
13
- physicalBioAssay_ref, Target_assnref-method
(zzz-DTTCompatibility-class),
13
- physicalBioAssay_ref, TopLevelBioAssays_assnreflist-method
(zzz-DTTCompatibility-class),
13
- PhysicalBioAssay_ref-class
(zzz-DTTCompatibility-class),
13
- physicalBioAssayData_assnlist
(zzz-DTTCompatibility-class),
13
- physicalBioAssayData_assnlist, PhysicalBioAssay-method
(zzz-DTTCompatibility-class),
13
- PhysicalBioAssayData_assnlist-class
(zzz-DTTCompatibility-class),

- 13
- physicalBioAssaySource_assnref
(zzz-DTTCompatibility-class),
13
- physicalBioAssaySource_assnref,FeatureExtraction-method
(zzz-DTTCompatibility-class),
13
- PhysicalBioAssaySource_assnref-class
(zzz-DTTCompatibility-class),
13
- PhysicalBioAssayTarget_assnref-class
(zzz-DTTCompatibility-class),
13
- polymerType_assn
(zzz-DTTCompatibility-class),
13
- polymerType_assn,BioSequence-method
(zzz-DTTCompatibility-class),
13
- PolymerType_assn-class
(zzz-DTTCompatibility-class),
13
- position (zzz-DTTCompatibility-class),
13
- position,Position_assn-method
(zzz-DTTCompatibility-class),
13
- Position-class
(zzz-DTTCompatibility-class),
13
- position_assn
(zzz-DTTCompatibility-class),
13
- position_assn,Feature-method
(zzz-DTTCompatibility-class),
13
- position_assn,Fiducial-method
(zzz-DTTCompatibility-class),
13
- Position_assn-class
(zzz-DTTCompatibility-class),
13
- positionDelta
(zzz-DTTCompatibility-class),
13
- positionDelta,PositionDelta_assn-method
(zzz-DTTCompatibility-class),
13
- PositionDelta-class
(zzz-DTTCompatibility-class),
13
- positionDelta_assn
(zzz-DTTCompatibility-class),
13
- positionDelta_assn,FeatureDefect-method
(zzz-DTTCompatibility-class),
13
- positionDelta_assn,ZoneDefect-method
(zzz-DTTCompatibility-class),
13
- PositionDelta_assn-class
(zzz-DTTCompatibility-class),
13
- presentAbsent
(zzz-DTTCompatibility-class),
13
- presentAbsent,QuantitationType_assnlist-method
(zzz-DTTCompatibility-class),
13
- PresentAbsent-class
(zzz-DTTCompatibility-class),
13
- presentAbsent_ref
(zzz-DTTCompatibility-class),
13
- presentAbsent_ref,QuantitationType_assnref-method
(zzz-DTTCompatibility-class),
13
- presentAbsent_ref,QuantitationTypes_assnreflist-method
(zzz-DTTCompatibility-class),
13
- presentAbsent_ref,SourcesQuantitationType_assnreflist-method
(zzz-DTTCompatibility-class),
13
- presentAbsent_ref,TargetQuantitationType_assnref-method
(zzz-DTTCompatibility-class),
13
- PresentAbsent_ref-class
(zzz-DTTCompatibility-class),
13
- producerTransformation_assn
(zzz-DTTCompatibility-class),
13
- producerTransformation_assn,DerivedBioAssayData-method
(zzz-DTTCompatibility-class),
13
- ProducerTransformation_assn-class
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,Array-method
(zzz-DTTCompatibility-class),

- 13
propertySets_assnlist,ArrayDesign-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,ArrayGroup-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,ArrayManufacture-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,ArrayManufactureDeviation-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,Audit-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,BibliographicReference-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,BioAssayCreation-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,BioAssayDataCluster-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,BioAssayDimension-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,BioAssayMap-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,BioAssayMapping-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,BioAssayTreatment-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,BioDataCube-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,BioDataTuples-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,BioMaterialMeasurement-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,BioSample-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,BioSequence-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,BioSource-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,Channel-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,CompositeCompositeMap-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,CompositeGroup-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,CompositePosition-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,CompositeSequence-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,CompositeSequenceDimension-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,Compound-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,CompoundMeasurement-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,ConcentrationUnit-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,Database-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,DatabaseEntry-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,DerivedBioAssay-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,DerivedBioAssayData-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,DerivedSignal-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,Description-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,DesignElementMapping-method
(zzz-DTTCompatibility-class),
13
propertySets_assnlist,DistanceUnit-method

- (zzz-DTTCompatibility-class),
13
- propertySets_assnlist,Error-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,ExpectedValue-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,Experiment-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,ExperimentalFactor-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,ExperimentDesign-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,ExternalReference-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,FactorValue-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,Failed-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,Feature-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,FeatureDefect-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,FeatureDimension-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,FeatureExtraction-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,FeatureGroup-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,FeatureInformation-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,FeatureLocation-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,FeatureReporterMap-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,Fiducial-method
(zzz-DTTCompatibility-class),
13
- 13
- propertySets_assnlist,Hardware-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,HardwareApplication-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,Hybridization-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,Image-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,ImageAcquisition-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,LabeledExtract-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,MAGE-ML-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,ManufactureLIMS-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,ManufactureLIMSBiomaterial-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,MassUnit-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,MeasuredBioAssay-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,MeasuredBioAssayData-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,MeasuredSignal-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,Measurement-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,MismatchInformation-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,NameValueType-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,Node-method
(zzz-DTTCompatibility-class),
13

- propertySets_assnlist,NodeContents-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,NodeValue-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,OntologyEntry-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,Organization-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,Parameter-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,ParameterValue-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,Person-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,PhysicalArrayDesign-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,PhysicalBioAssay-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,Position-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,PositionDelta-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,PresentAbsent-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,Protocol-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,ProtocolApplication-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,PValue-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,QuantitationTypeDimension-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,QuantitationTypeMap-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,QuantitationTypeMapping-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,QuantityUnit-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,Ratio-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,Reporter-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,ReporterCompositeMap-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,ReporterDimension-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,ReporterGroup-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,ReporterPosition-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,Security-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,SecurityGroup-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,SeqFeature-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,SeqFeatureLocation-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,SequencePosition-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,Software-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,SoftwareApplication-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,SpecializedQuantitationType-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,TemperatureUnit-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,TimeUnit-method
(zzz-DTTCompatibility-class),
13

- 13
- propertySets_assnlist,Transformation-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,Treatment-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,VolumeUnit-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,Zone-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,ZoneDefect-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,ZoneGroup-method
(zzz-DTTCompatibility-class),
13
- propertySets_assnlist,ZoneLayout-method
(zzz-DTTCompatibility-class),
13
- PropertySets_assnlist-class
(zzz-DTTCompatibility-class),
13
- protocol (zzz-DTTCompatibility-class),
13
- protocol,Protocol_assnlist-method
(zzz-DTTCompatibility-class),
13
- Protocol-class
(zzz-DTTCompatibility-class),
13
- protocol_assnlist
(zzz-DTTCompatibility-class),
13
- protocol_assnlist,Protocol_package-method
(zzz-DTTCompatibility-class),
13
- Protocol_assnlist-class
(zzz-DTTCompatibility-class),
13
- protocol_assnref
(zzz-DTTCompatibility-class),
13
- protocol_assnref,ProtocolApplication-method
(zzz-DTTCompatibility-class),
13
- Protocol_assnref-class
(zzz-DTTCompatibility-class),
13
- protocol_package
(zzz-DTTCompatibility-class),
13
- protocol_package,MAGE-ML-method
(zzz-DTTCompatibility-class),
13
- Protocol_package-class
(zzz-DTTCompatibility-class),
13
- protocol_ref
(zzz-DTTCompatibility-class),
13
- protocol_ref,Protocol_assnref-method
(zzz-DTTCompatibility-class),
13
- Protocol_ref-class
(zzz-DTTCompatibility-class),
13
- protocolApplication
(zzz-DTTCompatibility-class),
13
- protocolApplication,ProtocolApplications_assnlist-method
(zzz-DTTCompatibility-class),
13
- ProtocolApplication-class
(zzz-DTTCompatibility-class),
13
- protocolApplications_assnlist
(zzz-DTTCompatibility-class),
13
- protocolApplications_assnlist,ArrayDesign-method
(zzz-DTTCompatibility-class),
13
- protocolApplications_assnlist,ArrayManufacture-method
(zzz-DTTCompatibility-class),
13
- protocolApplications_assnlist,BioAssayCreation-method
(zzz-DTTCompatibility-class),
13
- protocolApplications_assnlist,BioAssayMap-method
(zzz-DTTCompatibility-class),
13
- protocolApplications_assnlist,BioAssayTreatment-method
(zzz-DTTCompatibility-class),
13
- protocolApplications_assnlist,CompositeCompositeMap-method
(zzz-DTTCompatibility-class),
13
- protocolApplications_assnlist,FeatureExtraction-method
(zzz-DTTCompatibility-class),
13
- protocolApplications_assnlist,FeatureReporterMap-method
(zzz-DTTCompatibility-class),

- 13
 PValue-class
 protocolApplications_assnlist,Hybridization-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 13
 pvalue_ref
 protocolApplications_assnlist,ImageAcquisition-method(zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 13
 pvalue_ref,ConfidenceIndicators_assnreflist-method
 protocolApplications_assnlist,PhysicalArrayDesign-method(zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 13
 pvalue_ref,QuantitationType_assnref-method
 protocolApplications_assnlist,QuantitationTypeMap-method(zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 13
 pvalue_ref,QuantitationTypes_assnreflist-method
 protocolApplications_assnlist,ReporterCompositeMap-method(zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 13
 pvalue_ref,SourcesQuantitationType_assnreflist-method
 protocolApplications_assnlist,Transformation-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 13
 pvalue_ref,TargetQuantitationType_assnref-method
 protocolApplications_assnlist,Treatment-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 13
 PValue_ref-class
 ProtocolApplications_assnlist-class (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 13
 13
 providers_assnreflist
 quality (zzz-DTTCompatibility-class), 13
 quality,ManufactureLIMS-method
 (zzz-DTTCompatibility-class),
 13
 providers_assnreflist,Experiment-method
 quality,ManufactureLIMSBiomaterial-method
 (zzz-DTTCompatibility-class),
 13
 13
 Providers_assnreflist-class
 qualityControlDescription_assn
 (zzz-DTTCompatibility-class),
 13
 13
 publication
 qualityControlDescription_assn,ExperimentDesign-method
 (zzz-DTTCompatibility-class),
 13
 13
 publication,BibliographicReference-method
 QualityControlDescription_assn-class
 (zzz-DTTCompatibility-class),
 13
 13
 publisher (zzz-DTTCompatibility-class),
 qualityControlStatistics_assnlist
 (zzz-DTTCompatibility-class),
 13
 13
 publisher,BibliographicReference-method
 qualityControlStatistics_assnlist,ArrayManufacture-meth
 (zzz-DTTCompatibility-class),
 13
 13
 pvalue (zzz-DTTCompatibility-class), 13
 qualityControlStatistics_assnlist,BioSample-method
 (zzz-DTTCompatibility-class),
 13
 13
 pvalue,ConfidenceIndicator_assnlist-method
 qualityControlStatistics_assnlist,BioSource-method
 (zzz-DTTCompatibility-class),
 13
 13
 pvalue,QuantitationType_assnlist-method
 (zzz-DTTCompatibility-class),
 13
 13

- qualityControlStatistics_assnlist,LabeledExtract-method (zzz-DTTCompatibility-class),
(zzz-DTTCompatibility-class), 13
- QualityControlStatistics_assnlist-class (zzz-DTTCompatibility-class), 13
- quantitationDimension_assnref (zzz-DTTCompatibility-class), 13
- quantitationDimension_assnref,NodeContents-method (zzz-DTTCompatibility-class), 13
- QuantitationDimension_assnref-class (zzz-DTTCompatibility-class), 13
- quantitationType_assnlist (zzz-DTTCompatibility-class), 13
- quantitationType_assnlist,QuantitationType_package-method (zzz-DTTCompatibility-class), 13
- QuantitationType_assnlist-class (zzz-DTTCompatibility-class), 13
- quantitationType_assnref (zzz-DTTCompatibility-class), 13
- quantitationType_assnref,QuantitationTypeTuple-method (zzz-DTTCompatibility-class), 13
- QuantitationType_assnref-class (zzz-DTTCompatibility-class), 13
- quantitationType_package (zzz-DTTCompatibility-class), 13
- quantitationType_package,MAGE-ML-method (zzz-DTTCompatibility-class), 13
- QuantitationType_package-class (zzz-DTTCompatibility-class), 13
- QuantitationType_ref-class (zzz-DTTCompatibility-class), 13
- quantitationTypeDimension (zzz-DTTCompatibility-class), 13
- quantitationTypeDimension,QuantitationTypeDimension_assnlist-method (zzz-DTTCompatibility-class), 13
- QuantitationTypeDimension-class (zzz-DTTCompatibility-class), 13
- quantitationTypeDimension_assnlist (zzz-DTTCompatibility-class), 13
- quantitationTypeDimension_assnlist,BioAssayData_package (zzz-DTTCompatibility-class), 13
- QuantitationTypeDimension_assnlist-class (zzz-DTTCompatibility-class), 13
- quantitationTypeDimension_assnref (zzz-DTTCompatibility-class), 13
- quantitationTypeDimension_assnref,DerivedBioAssayData-n (zzz-DTTCompatibility-class), 13
- quantitationTypeDimension_assnref,MeasuredBioAssayData-n (zzz-DTTCompatibility-class), 13
- QuantitationTypeDimension_assnref-class (zzz-DTTCompatibility-class), 13
- quantitationTypeDimension_ref (zzz-DTTCompatibility-class), 13
- quantitationTypeDimension_ref,QuantitationDimension_assnlist-method (zzz-DTTCompatibility-class), 13
- quantitationTypeDimension_ref,QuantitationTypeDimension (zzz-DTTCompatibility-class), 13
- QuantitationTypeDimension_ref-class (zzz-DTTCompatibility-class), 13
- quantitationTypeMap (zzz-DTTCompatibility-class), 13
- quantitationTypeMap,QuantitationTypeMap_assnlist-method (zzz-DTTCompatibility-class), 13
- QuantitationTypeMap-class (zzz-DTTCompatibility-class), 13
- quantitationTypeMap_assnlist (zzz-DTTCompatibility-class), 13
- quantitationTypeMap_assnlist,BioAssayData_package-method (zzz-DTTCompatibility-class), 13
- QuantitationTypeMap_assnlist-class (zzz-DTTCompatibility-class), 13

- 13
- quantitationTypeMap_ref (zzz-DTTCompatibility-class), 13
- quantitationTypeMap_ref, QuantitationTypeMaps_assnreflist-method (zzz-DTTCompatibility-class), 13
- QuantitationTypeMap_ref-class (zzz-DTTCompatibility-class), 13
- quantitationTypeMapping (zzz-DTTCompatibility-class), 13
- quantitationTypeMapping, QuantitationTypeMapping_assnreflist-method (zzz-DTTCompatibility-class), 13
- QuantitationTypeMapping-class (zzz-DTTCompatibility-class), 13
- quantitationTypeMapping_assn (zzz-DTTCompatibility-class), 13
- quantitationTypeMapping_assn, Transformation-method (zzz-DTTCompatibility-class), 13
- QuantitationTypeMapping_assn-class (zzz-DTTCompatibility-class), 13
- quantitationTypeMaps_assnreflist (zzz-DTTCompatibility-class), 13
- quantitationTypeMaps_assnreflist, DerivedSignal-method (zzz-DTTCompatibility-class), 13
- quantitationTypeMaps_assnreflist, Error-method (zzz-DTTCompatibility-class), 13
- quantitationTypeMaps_assnreflist, ExpectedValue-method (zzz-DTTCompatibility-class), 13
- quantitationTypeMaps_assnreflist, Failed-method (zzz-DTTCompatibility-class), 13
- quantitationTypeMaps_assnreflist, MeasuredSignal-method (zzz-DTTCompatibility-class), 13
- quantitationTypeMaps_assnreflist, PresentAbsent-method (zzz-DTTCompatibility-class), 13
- quantitationTypeMaps_assnreflist, PValue-method (zzz-DTTCompatibility-class), 13
- quantitationTypeMaps_assnreflist, QuantitationTypeMapping-assnreflist-method (zzz-DTTCompatibility-class), 13
- quantitationTypeMaps_assnreflist, Ratio-method (zzz-DTTCompatibility-class), 13
- quantitationTypeMaps_assnreflist, SpecializedQuantitationTypeMapping-assnreflist-method (zzz-DTTCompatibility-class), 13
- QuantitationTypeMaps_assnreflist-class (zzz-DTTCompatibility-class), 13
- quantitationTypes_assnreflist (zzz-DTTCompatibility-class), 13
- quantitationTypes_assnreflist, QuantitationTypeDimensionality-assnreflist-method (zzz-DTTCompatibility-class), 13
- QuantitationTypes_assnreflist-class (zzz-DTTCompatibility-class), 13
- quantitationTypeTuple (zzz-DTTCompatibility-class), 13
- quantitationTypeTuple, QuantitationTypeTuples_assnreflist-method (zzz-DTTCompatibility-class), 13
- QuantitationTypeTuple-class (zzz-DTTCompatibility-class), 13
- quantitationTypeTuples_assnreflist (zzz-DTTCompatibility-class), 13
- quantitationTypeTuples_assnreflist, DesignElementTuple-method (zzz-DTTCompatibility-class), 13
- QuantitationTypeTuples_assnreflist-class (zzz-DTTCompatibility-class), 13
- quantityUnit (zzz-DTTCompatibility-class), 13
- quantityUnit, Unit_assnreflist-method (zzz-DTTCompatibility-class), 13
- QuantityUnit-class (zzz-DTTCompatibility-class), 13
- ratio (zzz-DTTCompatibility-class), 13
- ratio, QuantitationType_assnreflist-method (zzz-DTTCompatibility-class), 13

- Ratio-class (zzz-DTTCompatibility-class), 13
- ratio_ref (zzz-DTTCompatibility-class), 13
- ratio_ref, QuantitationType_assnref-method (zzz-DTTCompatibility-class), 13
- ratio_ref, QuantitationTypes_assnreflist-method (zzz-DTTCompatibility-class), 13
- ratio_ref, SourcesQuantitationType_assnreflist-method (zzz-DTTCompatibility-class), 13
- ratio_ref, TargetQuantitationType_assnref-method (zzz-DTTCompatibility-class), 13
- Ratio_ref-class (zzz-DTTCompatibility-class), 13
- readAnnotation (NetAffxResource-class), 5
- readAnnotation, NetAffxResource, character, character-methods (NetAffxResource-class), 5
- readAnnotation, NetAffxResource, character, numeric-methods (NetAffxResource-class), 5
- readAnnotation, NetAffxResource, missing, AffxAnnotation-method (NetAffxResource-class), 5
- readAnnotation, NetAffxResource, numeric, character-methods (NetAffxResource-class), 5
- readAnnotation, NetAffxResource, numeric, numeric-methods (NetAffxResource-class), 5
- readArr, 8
- readMage, 8
- readNetAffx, 9
- readXml, 8, 9, 10, 11
- regions_assnlist (zzz-DTTCompatibility-class), 13
- regions_assnlist, SeqFeature-method (zzz-DTTCompatibility-class), 13
- Regions_assnlist-class (zzz-DTTCompatibility-class), 13
- releaseDate (zzz-DTTCompatibility-class), 13
- releaseDate, SoftwareApplication-method (zzz-DTTCompatibility-class), 13
- replacedLength (zzz-DTTCompatibility-class), 13
- replacedLength, MismatchInformation-method (zzz-DTTCompatibility-class), 13
- replicateDescription_assn (zzz-DTTCompatibility-class), 13
- replicateDescription_assn, ExperimentDesign-method (zzz-DTTCompatibility-class), 13
- Reporter-class (zzz-DTTCompatibility-class), 13
- Reporter, DesignElement_assnlist-method (zzz-DTTCompatibility-class), 13
- Reporter, Reporter_assnlist-method (zzz-DTTCompatibility-class), 13
- Reporter-methods (zzz-DTTCompatibility-class), 13
- Reporter_assnlist (zzz-DTTCompatibility-class), 13
- Reporter_assnlist, DesignElement_package-method (zzz-DTTCompatibility-class), 13
- Reporter_assnlist-class (zzz-DTTCompatibility-class), 13
- Reporter_assnref (zzz-DTTCompatibility-class), 13
- Reporter_assnref, FeatureReporterMap-method (zzz-DTTCompatibility-class), 13
- Reporter_assnref, ReporterPosition-method (zzz-DTTCompatibility-class), 13
- Reporter_assnref-class (zzz-DTTCompatibility-class), 13
- Reporter_ref (zzz-DTTCompatibility-class), 13
- Reporter_ref, DesignElement_assnref-method (zzz-DTTCompatibility-class), 13

- reporter_ref,Reporter_assnref-method (zzz-DTTCompatibility-class), 13
- reporter_ref,Reporters_assnreflist-method (zzz-DTTCompatibility-class), 13
- Reporter_ref-class (zzz-DTTCompatibility-class), 13
- reporterCompositeMap (zzz-DTTCompatibility-class), 13
- reporterCompositeMap,DesignElementMap_assnlist-method(zzz-DTTCompatibility-class), 13
- reporterCompositeMap,ReporterCompositeMap_assnlist-method(zzz-DTTCompatibility-class), 13
- ReporterCompositeMap-class (zzz-DTTCompatibility-class), 13
- reporterCompositeMap_assnlist (zzz-DTTCompatibility-class), 13
- reporterCompositeMap_assnlist,DesignElement_package-method (zzz-DTTCompatibility-class), 13
- ReporterCompositeMap_assnlist-class (zzz-DTTCompatibility-class), 13
- reporterCompositeMap_ref (zzz-DTTCompatibility-class), 13
- reporterCompositeMap_ref,DesignElementMaps_assnreflist(zzz-DTTCompatibility-class), 13
- reporterCompositeMap_ref,ReporterCompositeMaps_assnreflist(zzz-DTTCompatibility-class), 13
- ReporterCompositeMap_ref-class (zzz-DTTCompatibility-class), 13
- reporterCompositeMaps_assnreflist (zzz-DTTCompatibility-class), 13
- reporterCompositeMaps_assnreflist,CompositeSequence-method (zzz-DTTCompatibility-class), 13
- ReporterCompositeMaps_assnreflist-class (zzz-DTTCompatibility-class), 13
- reporterDimension (zzz-DTTCompatibility-class), 13
- reporterDimension,DesignElementDimension_assnlist-method (zzz-DTTCompatibility-class), 13
- ReporterDimension-class (zzz-DTTCompatibility-class), 13
- reporterDimension_ref (zzz-DTTCompatibility-class), 13
- reporterDimension_ref,DesignElementDimension_assnref-method(zzz-DTTCompatibility-class), 13
- ReporterDimension_ref-class (zzz-DTTCompatibility-class), 13
- reporterGroup (zzz-DTTCompatibility-class), 13
- reporterGroup,ReporterGroup_assnlist-method (zzz-DTTCompatibility-class), 13
- ReporterGroup-class (zzz-DTTCompatibility-class), 13
- reporterGroup_assnlist (zzz-DTTCompatibility-class), 13
- reporterGroup_assnlist,ArrayDesign_package-method (zzz-DTTCompatibility-class), 13
- ReporterGroup_assnlist-class (zzz-DTTCompatibility-class), 13
- reporterGroup_ref (zzz-DTTCompatibility-class), 13
- reporterGroup_ref,ReporterGroups_assnreflist-method (zzz-DTTCompatibility-class), 13
- ReporterGroup_ref-class (zzz-DTTCompatibility-class), 13
- reporterGroups_assnreflist (zzz-DTTCompatibility-class), 13
- reporterGroups_assnreflist,ArrayDesign-method (zzz-DTTCompatibility-class), 13
- reporterGroups_assnreflist,PhysicalArrayDesign-method (zzz-DTTCompatibility-class), 13

- 13
- ReporterGroups_assnreflist-class
(zzz-DTTCompatibility-class),
13
- reporterPosition
(zzz-DTTCompatibility-class),
13
- reporterPosition,Coordinate_assn-method
(zzz-DTTCompatibility-class),
13
- reporterPosition,ReporterPositionSources_assnlist-method
(zzz-DTTCompatibility-class),
13
- ReporterPosition-class
(zzz-DTTCompatibility-class),
13
- reporterPositionSources_assnlist
(zzz-DTTCompatibility-class),
13
- reporterPositionSources_assnlist,ReporterCompositeMap-method
(zzz-DTTCompatibility-class),
13
- ReporterPositionSources_assnlist-class
(zzz-DTTCompatibility-class),
13
- reporters_assnreflist
(zzz-DTTCompatibility-class),
13
- reporters_assnreflist,ReporterDimension-method
(zzz-DTTCompatibility-class),
13
- reporters_assnreflist,ReporterGroup-method
(zzz-DTTCompatibility-class),
13
- Reporters_assnreflist-class
(zzz-DTTCompatibility-class),
13
- required(zzz-ARRCompatibility-class),
12
- required,UserAttribute-method
(zzz-ARRCompatibility-class),
12
- roles_assnlist
(zzz-DTTCompatibility-class),
13
- roles_assnlist,Organization-method
(zzz-DTTCompatibility-class),
13
- roles_assnlist,Person-method
(zzz-DTTCompatibility-class),
13
- Roles_assnlist-class
(zzz-DTTCompatibility-class),
13
- scale_assn
(zzz-DTTCompatibility-class),
13
- scale_assn,DerivedSignal-method
(zzz-DTTCompatibility-class),
13
- scale_assn,Error-method
(zzz-DTTCompatibility-class),
13
- scale_assn,ExpectedValue-method
(zzz-DTTCompatibility-class),
13
- scale_assn,Failed-method
(zzz-DTTCompatibility-class),
13
- scale_assn,MeasuredSignal-method
(zzz-DTTCompatibility-class),
13
- scale_assn,NodeValue-method
(zzz-DTTCompatibility-class),
13
- scale_assn,PresentAbsent-method
(zzz-DTTCompatibility-class),
13
- scale_assn,PValue-method
(zzz-DTTCompatibility-class),
13
- scale_assn,Ratio-method
(zzz-DTTCompatibility-class),
13
- scale_assn,SpecializedQuantitationType-method
(zzz-DTTCompatibility-class),
13
- Scale_assn-class
(zzz-DTTCompatibility-class),
13
- security(zzz-DTTCompatibility-class),
13
- security,Security_assnlist-method
(zzz-DTTCompatibility-class),
13
- Security-class
(zzz-DTTCompatibility-class),
13
- security_assnlist
(zzz-DTTCompatibility-class),
13
- security_assnlist,AuditAndSecurity_package-method
(zzz-DTTCompatibility-class),
13

- Security_assnlist-class
(zzz-DTTCompatibility-class),
13
- security_assnref
(zzz-DTTCompatibility-class),
13
- security_assnref,Array-method
(zzz-DTTCompatibility-class),
13
- security_assnref,ArrayDesign-method
(zzz-DTTCompatibility-class),
13
- security_assnref,ArrayGroup-method
(zzz-DTTCompatibility-class),
13
- security_assnref,ArrayManufacture-method
(zzz-DTTCompatibility-class),
13
- security_assnref,Audit-method
(zzz-DTTCompatibility-class),
13
- security_assnref,BibliographicReference-method
(zzz-DTTCompatibility-class),
13
- security_assnref,BioAssayCreation-method
(zzz-DTTCompatibility-class),
13
- security_assnref,BioAssayDataCluster-method
(zzz-DTTCompatibility-class),
13
- security_assnref,BioAssayDimension-method
(zzz-DTTCompatibility-class),
13
- security_assnref,BioAssayMap-method
(zzz-DTTCompatibility-class),
13
- security_assnref,BioAssayTreatment-method
(zzz-DTTCompatibility-class),
13
- security_assnref,BioSample-method
(zzz-DTTCompatibility-class),
13
- security_assnref,BioSequence-method
(zzz-DTTCompatibility-class),
13
- security_assnref,BioSource-method
(zzz-DTTCompatibility-class),
13
- security_assnref,Channel-method
(zzz-DTTCompatibility-class),
13
- security_assnref,CompositeCompositeMap-method
(zzz-DTTCompatibility-class),
13
- security_assnref,CompositeGroup-method
(zzz-DTTCompatibility-class),
13
- security_assnref,CompositeSequence-method
(zzz-DTTCompatibility-class),
13
- security_assnref,CompositeSequenceDimension-method
(zzz-DTTCompatibility-class),
13
- security_assnref,Compound-method
(zzz-DTTCompatibility-class),
13
- security_assnref,Database-method
(zzz-DTTCompatibility-class),
13
- security_assnref,DerivedBioAssay-method
(zzz-DTTCompatibility-class),
13
- security_assnref,DerivedBioAssayData-method
(zzz-DTTCompatibility-class),
13
- security_assnref,DerivedSignal-method
(zzz-DTTCompatibility-class),
13
- security_assnref,Description-method
(zzz-DTTCompatibility-class),
13
- security_assnref,Error-method
(zzz-DTTCompatibility-class),
13
- security_assnref,ExpectedValue-method
(zzz-DTTCompatibility-class),
13
- security_assnref,Experiment-method
(zzz-DTTCompatibility-class),
13
- security_assnref,ExperimentalFactor-method
(zzz-DTTCompatibility-class),
13
- security_assnref,ExperimentDesign-method
(zzz-DTTCompatibility-class),
13
- security_assnref,FactorValue-method
(zzz-DTTCompatibility-class),
13
- security_assnref,Failed-method
(zzz-DTTCompatibility-class),
13
- security_assnref,Feature-method
(zzz-DTTCompatibility-class),

- 13
- security_assnref, FeatureDimension-method
(zzz-DTTCompatibility-class),
13
- security_assnref, FeatureExtraction-method
(zzz-DTTCompatibility-class),
13
- security_assnref, FeatureGroup-method
(zzz-DTTCompatibility-class),
13
- security_assnref, FeatureReporterMap-method
(zzz-DTTCompatibility-class),
13
- security_assnref, Fiducial-method
(zzz-DTTCompatibility-class),
13
- security_assnref, Hardware-method
(zzz-DTTCompatibility-class),
13
- security_assnref, HardwareApplication-method
(zzz-DTTCompatibility-class),
13
- security_assnref, Hybridization-method
(zzz-DTTCompatibility-class),
13
- security_assnref, Image-method
(zzz-DTTCompatibility-class),
13
- security_assnref, ImageAcquisition-method
(zzz-DTTCompatibility-class),
13
- security_assnref, LabeledExtract-method
(zzz-DTTCompatibility-class),
13
- security_assnref, MAGE-ML-method
(zzz-DTTCompatibility-class),
13
- security_assnref, ManufactureLIMS-method
(zzz-DTTCompatibility-class),
13
- security_assnref, ManufactureLIMSBiomaterial-method
(zzz-DTTCompatibility-class),
13
- security_assnref, MeasuredBioAssay-method
(zzz-DTTCompatibility-class),
13
- security_assnref, MeasuredBioAssayData-method
(zzz-DTTCompatibility-class),
13
- security_assnref, MeasuredSignal-method
(zzz-DTTCompatibility-class),
13
- security_assnref, Node-method
(zzz-DTTCompatibility-class),
13
- security_assnref, NodeContents-method
(zzz-DTTCompatibility-class),
13
- security_assnref, Organization-method
(zzz-DTTCompatibility-class),
13
- security_assnref, Parameter-method
(zzz-DTTCompatibility-class),
13
- security_assnref, Person-method
(zzz-DTTCompatibility-class),
13
- security_assnref, PhysicalArrayDesign-method
(zzz-DTTCompatibility-class),
13
- security_assnref, PhysicalBioAssay-method
(zzz-DTTCompatibility-class),
13
- security_assnref, PresentAbsent-method
(zzz-DTTCompatibility-class),
13
- security_assnref, Protocol-method
(zzz-DTTCompatibility-class),
13
- security_assnref, ProtocolApplication-method
(zzz-DTTCompatibility-class),
13
- security_assnref, PValue-method
(zzz-DTTCompatibility-class),
13
- security_assnref, QuantitationTypeDimension-method
(zzz-DTTCompatibility-class),
13
- security_assnref, QuantitationTypeMap-method
(zzz-DTTCompatibility-class),
13
- security_assnref, Ratio-method
(zzz-DTTCompatibility-class),
13
- security_assnref, Reporter-method
(zzz-DTTCompatibility-class),
13
- security_assnref, ReporterCompositeMap-method
(zzz-DTTCompatibility-class),
13
- security_assnref, ReporterDimension-method
(zzz-DTTCompatibility-class),
13
- security_assnref, ReporterGroup-method

- (zzz-DTTCompatibility-class),
13
- security_assnref, Security-method
(zzz-DTTCompatibility-class),
13
- security_assnref, SecurityGroup-method
(zzz-DTTCompatibility-class),
13
- security_assnref, SeqFeature-method
(zzz-DTTCompatibility-class),
13
- security_assnref, Software-method
(zzz-DTTCompatibility-class),
13
- security_assnref, SoftwareApplication-method
(zzz-DTTCompatibility-class),
13
- security_assnref, SpecializedQuantitationType-method
(zzz-DTTCompatibility-class),
13
- security_assnref, Transformation-method
(zzz-DTTCompatibility-class),
13
- security_assnref, Treatment-method
(zzz-DTTCompatibility-class),
13
- security_assnref, Zone-method
(zzz-DTTCompatibility-class),
13
- Security_assnref-class
(zzz-DTTCompatibility-class),
13
- security_ref
(zzz-DTTCompatibility-class),
13
- security_ref, Security_assnref-method
(zzz-DTTCompatibility-class),
13
- Security_ref-class
(zzz-DTTCompatibility-class),
13
- securityGroup
(zzz-DTTCompatibility-class),
13
- securityGroup, SecurityGroup_assnlist-method
(zzz-DTTCompatibility-class),
13
- SecurityGroup-class
(zzz-DTTCompatibility-class),
13
- securityGroup_assnlist
(zzz-DTTCompatibility-class),
13
- securityGroup_assnlist, AuditAndSecurity_package-method
(zzz-DTTCompatibility-class),
13
- SecurityGroup_assnlist-class
(zzz-DTTCompatibility-class),
13
- securityGroup_ref
(zzz-DTTCompatibility-class),
13
- securityGroup_ref, SecurityGroups_assnreflist-method
(zzz-DTTCompatibility-class),
13
- SecurityGroup_ref-class
(zzz-DTTCompatibility-class),
13
- securityGroups_assnreflist
(zzz-DTTCompatibility-class),
13
- securityGroups_assnreflist, Security-method
(zzz-DTTCompatibility-class),
13
- SecurityGroups_assnreflist-class
(zzz-DTTCompatibility-class),
13
- seqFeature
(zzz-DTTCompatibility-class),
13
- seqFeature, SeqFeatures_assnlist-method
(zzz-DTTCompatibility-class),
13
- SeqFeature-class
(zzz-DTTCompatibility-class),
13
- seqFeatureLocation
(zzz-DTTCompatibility-class),
13
- seqFeatureLocation, Regions_assnlist-method
(zzz-DTTCompatibility-class),
13
- seqFeatureLocation, Subregions_assnlist-method
(zzz-DTTCompatibility-class),
13
- SeqFeatureLocation-class
(zzz-DTTCompatibility-class),
13
- seqFeatures_assnlist
(zzz-DTTCompatibility-class),
13
- seqFeatures_assnlist, BioSequence-method
(zzz-DTTCompatibility-class),
13

- SeqFeatures_assnlist-class
(zzz-DTTCompatibility-class),
13
- sequenceDatabases_assnlist
(zzz-DTTCompatibility-class),
13
- sequenceDatabases_assnlist,BioSequence-method
(zzz-DTTCompatibility-class),
13
- SequenceDatabases_assnlist-class
(zzz-DTTCompatibility-class),
13
- sequencePosition
(zzz-DTTCompatibility-class),
13
- sequencePosition,Coordinate_assn-method
(zzz-DTTCompatibility-class),
13
- SequencePosition-class
(zzz-DTTCompatibility-class),
13
- serialNumber
(zzz-DTTCompatibility-class),
13
- serialNumber,HardwareApplication-method
(zzz-DTTCompatibility-class),
13
- show,.TypedAssociation-method
(Internal-class), 4
- show,AffyCompatible-method
(AffyCompatible-class), 3
- show,NetAffxResource-method
(NetAffxResource-class), 5
- software(zzz-DTTCompatibility-class),
13
- software,Software_assnlist-method
(zzz-DTTCompatibility-class),
13
- Software-class
(zzz-DTTCompatibility-class),
13
- software_assnlist
(zzz-DTTCompatibility-class),
13
- software_assnlist,Protocol_package-method
(zzz-DTTCompatibility-class),
13
- Software_assnlist-class
(zzz-DTTCompatibility-class),
13
- software_assnref
(zzz-DTTCompatibility-class),
13
- software_assnref,SoftwareApplication-method
(zzz-DTTCompatibility-class),
13
- Software_assnref-class
(zzz-DTTCompatibility-class),
13
- software_ref
(zzz-DTTCompatibility-class),
13
- software_ref,Software_assnref-method
(zzz-DTTCompatibility-class),
13
- software_ref,Softwares_assnreflist-method
(zzz-DTTCompatibility-class),
13
- Software_ref-class
(zzz-DTTCompatibility-class),
13
- softwareApplication
(zzz-DTTCompatibility-class),
13
- softwareApplication,SoftwareApplications_assnlist-method
(zzz-DTTCompatibility-class),
13
- SoftwareApplication-class
(zzz-DTTCompatibility-class),
13
- softwareApplications_assnlist
(zzz-DTTCompatibility-class),
13
- softwareApplications_assnlist,ProtocolApplication-method
(zzz-DTTCompatibility-class),
13
- SoftwareApplications_assnlist-class
(zzz-DTTCompatibility-class),
13
- softwareManufacturers_assnreflist
(zzz-DTTCompatibility-class),
13
- softwareManufacturers_assnreflist,Software-method
(zzz-DTTCompatibility-class),
13
- SoftwareManufacturers_assnreflist-class
(zzz-DTTCompatibility-class),
13
- softwares_assnreflist
(zzz-DTTCompatibility-class),
13
- softwares_assnreflist,Hardware-method
(zzz-DTTCompatibility-class),
13

- softwares_assnreflist,Protocol-method (zzz-DTTCompatibility-class), 13
- softwares_assnreflist,Software-method (zzz-DTTCompatibility-class), 13
- Softwares_assnreflist-class (zzz-DTTCompatibility-class), 13
- sourceBioAssays_assnreflist (zzz-DTTCompatibility-class), 13
- sourceBioAssays_assnreflist,BioAssayMap-method (zzz-DTTCompatibility-class), 13
- SourceBioAssays_assnreflist-class (zzz-DTTCompatibility-class), 13
- sourceBioMaterialMeasurements_assnlist (zzz-DTTCompatibility-class), 13
- sourceBioMaterialMeasurements_assnlist,BioAssayCreation-method (zzz-DTTCompatibility-class), 13
- sourceBioMaterialMeasurements_assnlist,Hybridization-method (zzz-DTTCompatibility-class), 13
- sourceBioMaterialMeasurements_assnlist,Treatment-method (zzz-DTTCompatibility-class), 13
- SourceBioMaterialMeasurements_assnlist-class (zzz-DTTCompatibility-class), 13
- sourceContact_assnreflist (zzz-DTTCompatibility-class), 13
- sourceContact_assnreflist,BioSource-method (zzz-DTTCompatibility-class), 13
- SourceContact_assnreflist-class (zzz-DTTCompatibility-class), 13
- sourcesQuantitationType_assnreflist (zzz-DTTCompatibility-class), 13
- sourcesQuantitationType_assnreflist,QuantitationTypeMap-method (zzz-DTTCompatibility-class), 13
- SourcesQuantitationType_assnreflist-class (zzz-DTTCompatibility-class), 13
- spacingBetweenCols (zzz-DTTCompatibility-class), 13
- spacingBetweenCols,ZoneLayout-method (zzz-DTTCompatibility-class), 13
- spacingBetweenRows (zzz-DTTCompatibility-class), 13
- spacingBetweenRows,ZoneLayout-method (zzz-DTTCompatibility-class), 13
- spacingsBetweenZonesX (zzz-DTTCompatibility-class), 13
- spacingsBetweenZonesX,ZoneGroup-method (zzz-DTTCompatibility-class), 13
- spacingsBetweenZonesY (zzz-DTTCompatibility-class), 13
- spacingsBetweenZonesY,ZoneGroup-method (zzz-DTTCompatibility-class), 13
- specializedQuantitationType (zzz-DTTCompatibility-class), 13
- specializedQuantitationType,QuantitationType_assnreflist-method (zzz-DTTCompatibility-class), 13
- SpecializedQuantitationType-class (zzz-DTTCompatibility-class), 13
- specializedQuantitationType_ref (zzz-DTTCompatibility-class), 13
- specializedQuantitationType_ref,QuantitationType_assnreflist-method (zzz-DTTCompatibility-class), 13
- specializedQuantitationType_ref,QuantitationTypes_assnreflist-method (zzz-DTTCompatibility-class), 13
- specializedQuantitationType_ref,SourcesQuantitationType_assnreflist-method (zzz-DTTCompatibility-class), 13
- specializedQuantitationType_ref,TargetQuantitationType_assnreflist-method (zzz-DTTCompatibility-class), 13
- SpecializedQuantitationType_ref-class (zzz-DTTCompatibility-class), 13
- species_assn (zzz-DTTCompatibility-class),

- 13
- species_assn, BioSequence-method
(zzz-DTTCompatibility-class),
13
- species_assn, CompositeGroup-method
(zzz-DTTCompatibility-class),
13
- species_assn, FeatureGroup-method
(zzz-DTTCompatibility-class),
13
- species_assn, ReporterGroup-method
(zzz-DTTCompatibility-class),
13
- Species_assn-class
(zzz-DTTCompatibility-class),
13
- StandardQuantitationType_ref-class
(zzz-DTTCompatibility-class),
13
- startCoord
(zzz-DTTCompatibility-class),
13
- startCoord, MismatchInformation-method
(zzz-DTTCompatibility-class),
13
- strandType
(zzz-DTTCompatibility-class),
13
- strandType, SeqFeatureLocation-method
(zzz-DTTCompatibility-class),
13
- subregions_assnlist
(zzz-DTTCompatibility-class),
13
- subregions_assnlist, SeqFeatureLocation-method
(zzz-DTTCompatibility-class),
13
- Subregions_assnlist-class
(zzz-DTTCompatibility-class),
13
- substrateType_assn
(zzz-DTTCompatibility-class),
13
- substrateType_assn, ArrayGroup-method
(zzz-DTTCompatibility-class),
13
- SubstrateType_assn-class
(zzz-DTTCompatibility-class),
13
- summaryStatistics_assnlist
(zzz-DTTCompatibility-class),
13
- summaryStatistics_assnlist, DerivedBioAssayData-method
(zzz-DTTCompatibility-class),
13
- summaryStatistics_assnlist, MeasuredBioAssayData-method
(zzz-DTTCompatibility-class),
13
- SummaryStatistics_assnlist-class
(zzz-DTTCompatibility-class),
13
- surfaceType_assn
(zzz-DTTCompatibility-class),
13
- surfaceType_assn, PhysicalArrayDesign-method
(zzz-DTTCompatibility-class),
13
- SurfaceType_assn-class
(zzz-DTTCompatibility-class),
13
- target_assnref
(zzz-DTTCompatibility-class),
13
- target_assnref, BioAssayTreatment-method
(zzz-DTTCompatibility-class),
13
- target_assnref, ImageAcquisition-method
(zzz-DTTCompatibility-class),
13
- Target_assnref-class
(zzz-DTTCompatibility-class),
13
- targetQuantitationType_assnref
(zzz-DTTCompatibility-class),
13
- targetQuantitationType_assnref, Error-method
(zzz-DTTCompatibility-class),
13
- targetQuantitationType_assnref, ExpectedValue-method
(zzz-DTTCompatibility-class),
13
- targetQuantitationType_assnref, PValue-method
(zzz-DTTCompatibility-class),
13
- targetQuantitationType_assnref, QuantitationTypeMap-method
(zzz-DTTCompatibility-class),
13
- TargetQuantitationType_assnref-class
(zzz-DTTCompatibility-class),
13
- technologyType_assn
(zzz-DTTCompatibility-class),
13
- technologyType_assn, FeatureGroup-method

- (zzz-DTTCompatibility-class),
13
- TechnologyType_assn-class
(zzz-DTTCompatibility-class),
13
- temperatureUnit
(zzz-DTTCompatibility-class),
13
- temperatureUnit,Unit_assn-method
(zzz-DTTCompatibility-class),
13
- TemperatureUnit-class
(zzz-DTTCompatibility-class),
13
- TemplateFile-class
(zzz-ARRCompatibility-class),
12
- timeUnit (zzz-DTTCompatibility-class),
13
- timeUnit,Unit_assn-method
(zzz-DTTCompatibility-class),
13
- TimeUnit-class
(zzz-DTTCompatibility-class),
13
- tolerance (zzz-DTTCompatibility-class),
13
- tolerance,ArrayManufacture-method
(zzz-DTTCompatibility-class),
13
- tollFreePhone
(zzz-DTTCompatibility-class),
13
- tollFreePhone,Organization-method
(zzz-DTTCompatibility-class),
13
- tollFreePhone,Person-method
(zzz-DTTCompatibility-class),
13
- topLevelBioAssays_assnreflist
(zzz-DTTCompatibility-class),
13
- topLevelBioAssays_assnreflist,ExperimentDesign-method(zzz-ARRCompatibility-class),
(zzz-DTTCompatibility-class),
13
- TopLevelBioAssays_assnreflist-class
(zzz-DTTCompatibility-class),
13
- transformation
(zzz-DTTCompatibility-class),
13
- transformation,ProducerTransformation_assn-method
(zzz-DTTCompatibility-class),
13
- Transformation-class
(zzz-DTTCompatibility-class),
13
- treatment (zzz-DTTCompatibility-class),
13
- treatment,Treatments_assnlist-method
(zzz-DTTCompatibility-class),
13
- Treatment-class
(zzz-DTTCompatibility-class),
13
- treatments_assnlist
(zzz-DTTCompatibility-class),
13
- treatments_assnlist,BioSample-method
(zzz-DTTCompatibility-class),
13
- treatments_assnlist,BioSource-method
(zzz-DTTCompatibility-class),
13
- treatments_assnlist,LabeledExtract-method
(zzz-DTTCompatibility-class),
13
- Treatments_assnlist-class
(zzz-DTTCompatibility-class),
13
- type (zzz-DTTCompatibility-class), 13
- type,ArraySetFile-method
(zzz-ARRCompatibility-class),
12
- type,Measurement-method
(zzz-DTTCompatibility-class),
13
- type,NameValueType-method
(zzz-DTTCompatibility-class),
13
- type,PhysicalArray-method
(zzz-ARRCompatibility-class),
12
- type,TemplateFile-method
(zzz-ARRCompatibility-class),
12
- type,UserAttribute-method
(zzz-ARRCompatibility-class),
12
- type_assn (zzz-DTTCompatibility-class),
13
- type_assn,BioSample-method
(zzz-DTTCompatibility-class),
13

- type_assn, BioSequence-method
(zzz-DTTCompatibility-class),
[13](#)
- type_assn, DatabaseEntry-method
(zzz-DTTCompatibility-class),
[13](#)
- type_assn, DerivedBioAssay-method
(zzz-DTTCompatibility-class),
[13](#)
- type_assn, Hardware-method
(zzz-DTTCompatibility-class),
[13](#)
- type_assn, NodeValue-method
(zzz-DTTCompatibility-class),
[13](#)
- type_assn, Protocol-method
(zzz-DTTCompatibility-class),
[13](#)
- type_assn, Software-method
(zzz-DTTCompatibility-class),
[13](#)
- Type_assn-class
(zzz-DTTCompatibility-class),
[13](#)
- types_assnlist
(zzz-DTTCompatibility-class),
[13](#)
- types_assnlist, CompositeGroup-method
(zzz-DTTCompatibility-class),
[13](#)
- types_assnlist, ExperimentDesign-method
(zzz-DTTCompatibility-class),
[13](#)
- types_assnlist, FeatureGroup-method
(zzz-DTTCompatibility-class),
[13](#)
- types_assnlist, ReporterGroup-method
(zzz-DTTCompatibility-class),
[13](#)
- Types_assnlist-class
(zzz-DTTCompatibility-class),
[13](#)
- unit_assn (zzz-DTTCompatibility-class),
[13](#)
- unit_assn, Measurement-method
(zzz-DTTCompatibility-class),
[13](#)
- Unit_assn-class
(zzz-DTTCompatibility-class),
[13](#)
- unitName (zzz-DTTCompatibility-class),
[13](#)
- unitName, ConcentrationUnit-method
(zzz-DTTCompatibility-class),
[13](#)
- unitName, DistanceUnit-method
(zzz-DTTCompatibility-class),
[13](#)
- unitName, MassUnit-method
(zzz-DTTCompatibility-class),
[13](#)
- unitName, QuantityUnit-method
(zzz-DTTCompatibility-class),
[13](#)
- unitName, TemperatureUnit-method
(zzz-DTTCompatibility-class),
[13](#)
- unitName, TimeUnit-method
(zzz-DTTCompatibility-class),
[13](#)
- unitName, VolumeUnit-method
(zzz-DTTCompatibility-class),
[13](#)
- unitNameCV
(zzz-DTTCompatibility-class),
[13](#)
- unitNameCV, ConcentrationUnit-method
(zzz-DTTCompatibility-class),
[13](#)
- unitNameCV, DistanceUnit-method
(zzz-DTTCompatibility-class),
[13](#)
- unitNameCV, MassUnit-method
(zzz-DTTCompatibility-class),
[13](#)
- unitNameCV, QuantityUnit-method
(zzz-DTTCompatibility-class),
[13](#)
- unitNameCV, TemperatureUnit-method
(zzz-DTTCompatibility-class),
[13](#)
- unitNameCV, TimeUnit-method
(zzz-DTTCompatibility-class),
[13](#)
- unitNameCV, VolumeUnit-method
(zzz-DTTCompatibility-class),
[13](#)
- upperLeftX
(zzz-DTTCompatibility-class),
[13](#)
- upperLeftX, Zone-method
(zzz-DTTCompatibility-class),
[13](#)
- upperLeftY

- (zzz-DTTCompatibility-class),
13
- upperLeftY, Zone-method
(zzz-DTTCompatibility-class),
13
- uri (zzz-DTTCompatibility-class), 13
- uri, BibliographicReference-method
(zzz-DTTCompatibility-class),
13
- uri, Database-method
(zzz-DTTCompatibility-class),
13
- uri, DatabaseEntry-method
(zzz-DTTCompatibility-class),
13
- uri, Description-method
(zzz-DTTCompatibility-class),
13
- uri, Hardware-method
(zzz-DTTCompatibility-class),
13
- uri, Image-method
(zzz-DTTCompatibility-class),
13
- uri, Organization-method
(zzz-DTTCompatibility-class),
13
- uri, Person-method
(zzz-DTTCompatibility-class),
13
- uri, Protocol-method
(zzz-DTTCompatibility-class),
13
- uri, Software-method
(zzz-DTTCompatibility-class),
13
- user (accessors), 2
- user, NetAffxResource-method
(NetAffxResource-class), 5
- userAttribute
(zzz-ARRCompatibility-class),
12
- userAttribute, UserAttributes-method
(zzz-ARRCompatibility-class),
12
- UserAttribute-class
(zzz-ARRCompatibility-class),
12
- userAttributes
(zzz-ARRCompatibility-class),
12
- userAttributes, ArraySetFile-method
(zzz-ARRCompatibility-class),
12
- UserAttributes-class
(zzz-ARRCompatibility-class),
12
- userAttributeValue
(zzz-ARRCompatibility-class),
12
- userAttributeValue, UserAttribute-method
(zzz-ARRCompatibility-class),
12
- UserAttributeValue-class
(zzz-ARRCompatibility-class),
12
- value (zzz-ARRCompatibility-class), 12
- value, Control-method
(zzz-ARRCompatibility-class),
12
- value, Datum-method
(zzz-DTTCompatibility-class),
13
- value, Measurement-method
(zzz-DTTCompatibility-class),
13
- value, NameValueType-method
(zzz-DTTCompatibility-class),
13
- value, NodeValue-method
(zzz-DTTCompatibility-class),
13
- value, OntologyEntry-method
(zzz-DTTCompatibility-class),
13
- value, ParameterValue-method
(zzz-DTTCompatibility-class),
13
- value_assn
(zzz-DTTCompatibility-class),
13
- value_assn, FactorValue-method
(zzz-DTTCompatibility-class),
13
- Value_assn-class
(zzz-DTTCompatibility-class),
13
- vector, 4
- version (zzz-ARRCompatibility-class), 12
- version, ArrayDesign-method
(zzz-DTTCompatibility-class),

- 13
- version, ArraySetFile-method
(zzz-ARRCompatibility-class),
12
- version, Database-method
(zzz-DTTCompatibility-class),
13
- version, PhysicalArrayDesign-method
(zzz-DTTCompatibility-class),
13
- version, SoftwareApplication-method
(zzz-DTTCompatibility-class),
13
- version, TemplateFile-method
(zzz-ARRCompatibility-class),
12
- volume (zzz-DTTCompatibility-class), 13
- volume, BibliographicReference-method
(zzz-DTTCompatibility-class),
13
- volumeUnit
(zzz-DTTCompatibility-class),
13
- volumeUnit, Unit_assn-method
(zzz-DTTCompatibility-class),
13
- VolumeUnit-class
(zzz-DTTCompatibility-class),
13
- warningType_assn
(zzz-DTTCompatibility-class),
13
- warningType_assn, Reporter-method
(zzz-DTTCompatibility-class),
13
- WarningType_assn-class
(zzz-DTTCompatibility-class),
13
- width (zzz-DTTCompatibility-class), 13
- width, ArrayGroup-method
(zzz-DTTCompatibility-class),
13
- x (zzz-DTTCompatibility-class), 13
- x, Position-method
(zzz-DTTCompatibility-class),
13
- xclass, 8–10, 11
- xmlValue.XMLAttributeValue, 12
- y (zzz-DTTCompatibility-class), 13
- y, Position-method
(zzz-DTTCompatibility-class),
13
- year (zzz-DTTCompatibility-class), 13
- year, BibliographicReference-method
(zzz-DTTCompatibility-class),
13
- zone (zzz-DTTCompatibility-class), 13
- zone, ZoneLocations_assnlist-method
(zzz-DTTCompatibility-class),
13
- Zone-class
(zzz-DTTCompatibility-class),
13
- zone_assnref
(zzz-DTTCompatibility-class),
13
- zone_assnref, Feature-method
(zzz-DTTCompatibility-class),
13
- zone_assnref, ZoneDefect-method
(zzz-DTTCompatibility-class),
13
- Zone_assnref-class
(zzz-DTTCompatibility-class),
13
- zone_ref (zzz-DTTCompatibility-class),
13
- zone_ref, Zone_assnref-method
(zzz-DTTCompatibility-class),
13
- Zone_ref-class
(zzz-DTTCompatibility-class),
13
- zoneDefect
(zzz-DTTCompatibility-class),
13
- zoneDefect, Adjustments_assnlist-method
(zzz-DTTCompatibility-class),
13
- ZoneDefect-class
(zzz-DTTCompatibility-class),
13
- zoneGroup (zzz-DTTCompatibility-class),
13
- zoneGroup, ZoneGroups_assnlist-method
(zzz-DTTCompatibility-class),
13
- ZoneGroup-class
(zzz-DTTCompatibility-class),
13
- zoneGroups_assnlist

(zzz-DTTCompatibility-class),
13

zoneGroups_assnlist,PhysicalArrayDesign-method
(zzz-DTTCompatibility-class),
13

ZoneGroups_assnlist-class
(zzz-DTTCompatibility-class),
13

zoneLayout
(zzz-DTTCompatibility-class),
13

zoneLayout,ZoneLayout_assn-method
(zzz-DTTCompatibility-class),
13

ZoneLayout-class
(zzz-DTTCompatibility-class),
13

zoneLayout_assn
(zzz-DTTCompatibility-class),
13

zoneLayout_assn,ZoneGroup-method
(zzz-DTTCompatibility-class),
13

ZoneLayout_assn-class
(zzz-DTTCompatibility-class),
13

zoneLocations_assnlist
(zzz-DTTCompatibility-class),
13

zoneLocations_assnlist,ZoneGroup-method
(zzz-DTTCompatibility-class),
13

ZoneLocations_assnlist-class
(zzz-DTTCompatibility-class),
13

zonesPerX (zzz-DTTCompatibility-class),
13

zonesPerX,ZoneGroup-method
(zzz-DTTCompatibility-class),
13

zonesPerY (zzz-DTTCompatibility-class),
13

zonesPerY,ZoneGroup-method
(zzz-DTTCompatibility-class),
13

zzz-.TypedAssociation-class
(Internal-class), 4

zzz-AffyCompatible-class
(AffyCompatible-class), 3

zzz-ARRCompatibility-class, 12

zzz-DTTCompatibility-class, 13

zzz-NetAffxCompatibility-class, 14