

AffyCompatible

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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")))
```

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>

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 i 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, PSI to data frames, and reads FASTA files to a list of description / sequence pairs (see Biostrings::readFASTA).

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", "BLASTP annotations, CSV Format"]]
df <- readAnnotation(rsrc, annotation=anno)
```

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

<code>user</code>	Length 1 character vector containing the Affymetrix user name authorized to access the NetAffx resource.
<code>password</code>	Length 1 character vector containing the Affymetrix password of the user.
<code>affxLicence</code>	Length 1 character vector containing the Affymetrix licence (typically application specific, i.e., the specified licence covers AffyCompatible).
<code>directory</code>	(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.
<code>...</code>	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’.

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](#)

readArr

Read ARR sample attribute files

Description

This functions reads the content of Affymetrix ARR sample attribute files created by GeneChip® 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(fl, ...)
```

Arguments

`fl` 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(fls)
readArr(fls[[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 <code>logical(1)</code> 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>
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Description

This functions reads the content of Affymetrix DTT (MAGE) or ARR sample files, or AffxNetAffxAnnotFile 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(fl, ...)
```

Arguments

<code>fl</code>	A length-1 character vector giving the full path to the file to be parsed.
<code>...</code>	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

<code>doc</code>	An <code>XMLInternalDocument</code> retrieved using <code>readXml</code> .
<code>xpathq</code>	A length-1 character string representing an xpath query into the document.
<code>prefix</code>	A prefix used to map, when necessary, between the XML class name and the corresponding R class. For <code>NetAffx</code> -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)
```

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

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® Command Console) application. Specifically, the classes are R representations of the Affymetrix classes defined in ArraySetAndTemplate-File.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 "[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"))
```

zzz-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)
```

`xmlValue.XMLAttributeValue`

Additional functions defined in AffyCompatible

Description

This page documents additional, miscellaneous, functions defined by `AffyCompatible`

Usage

```
xmlValue.XMLAttributeValue(x, ignoreComments = FALSE)
```

Arguments

<code>x</code>	An object of class <code>XMLAttributeValue</code> .
<code>ignoreComments</code>	Logical; ignored by the <code>xmlValue.XMLAttributeValue</code> method

Value

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

Index

*Topic classes

AffyCompatible-class, 1
 Internal-class, 11
 NetAffxResource-class, 2
 readArr, 6
 readMage, 7
 zzz-ARRCompatibility-class, 11
 zzz-DTTCompatibility-class, 12
 zzz-NetAffxCompatibility-class, 13

*Topic database

NetAffxResource, 5
 readNetAffx, 8

*Topic manip

readXml, 9
 xclass, 10
 xmlValue.XMLAttributeValue, 15

*Topic methods

accessors, 6

*Topic package

AffyCompatible-package, 2
 .Rprofile, 4
 .TypedAssociation, 12-14
 .TypedAssociation-class
 (*Internal-class*), 11
 [, .TypedAssociation, ANY, ANY, ANY-method
 (*Internal-class*), 11
 [, .TypedAssociation, numeric, missing, ANY-method
 (*Internal-class*), 11
 [, NetAffxResource, character, character-method
 (*NetAffxResource-class*), 2
 [, NetAffxResource, character, missing-method
 (*NetAffxResource-class*), 2
 [, NetAffxResource, character, numeric-method
 (*NetAffxResource-class*), 2
 [, NetAffxResource, numeric, character-method
 (*NetAffxResource-class*), 2
 [, NetAffxResource, numeric, missing-method
 (*NetAffxResource-class*), 2
 [, NetAffxResource, numeric, numeric-method

(*NetAffxResource-class*), 2
 accession
 (*zzz-DTTCompatibility-class*), 12
 accession, DatabaseEntry-method
 (*zzz-DTTCompatibility-class*), 12
 accessions_assnlist
 (*zzz-DTTCompatibility-class*), 12
 accessions_assnlist, BibliographicReference-method
 (*zzz-DTTCompatibility-class*), 12
 accessionVersion
 (*zzz-DTTCompatibility-class*), 12
 accessionVersion, DatabaseEntry-method
 (*zzz-DTTCompatibility-class*), 12
 accessors, 6
 action
 (*zzz-DTTCompatibility-class*), 12
 action, Audit-method
 (*zzz-DTTCompatibility-class*), 12
 action_assn
 (*zzz-DTTCompatibility-class*), 12
 action_assn, Treatment-method
 (*zzz-DTTCompatibility-class*), 12
 actionMeasurement_assn
 (*zzz-DTTCompatibility-class*), 12
 actionMeasurement_assn, Treatment-method
 (*zzz-DTTCompatibility-class*), 12
 activityDate
 (*zzz-DTTCompatibility-class*), 12
 activityDate, ProtocolApplication-method
 (*zzz-DTTCompatibility-class*), 12

- 12
- address
(*zzz-DTTCompatibility-class*),
12
- address, Organization-method
(*zzz-DTTCompatibility-class*),
12
- address, Person-method
(*zzz-DTTCompatibility-class*),
12
- adjustments_assnlist
(*zzz-DTTCompatibility-class*),
12
- adjustments_assnlist, ArrayManufactureDeviation-method
(*zzz-DTTCompatibility-class*),
12
- affiliation_assnref
(*zzz-DTTCompatibility-class*),
12
- affiliation_assnref, Person-method
(*zzz-DTTCompatibility-class*),
12
- affxAnnotation
(*zzz-NetAffxCompatibility-class*),
13
- affxAnnotation, AffxArray-method
(*zzz-NetAffxCompatibility-class*),
13
- affxArray
(*zzz-NetAffxCompatibility-class*),
13
- affxArray, AffxNetAffxAnnotFileList-method
(*zzz-NetAffxCompatibility-class*),
13
- affxArray, Array_assnlist-method
(*zzz-DTTCompatibility-class*),
12
- affxCategory
(*zzz-DTTCompatibility-class*),
12
- affxCategory, OntologyEntry-method
(*zzz-DTTCompatibility-class*),
12
- affxComment
(*zzz-NetAffxCompatibility-class*),
13
- affxComment, AffxAnnotation-method
(*zzz-NetAffxCompatibility-class*),
13
- affxComment, PhysicalArray-method
(*zzz-ARRCompatibility-class*),
11
- affxCompression
(*zzz-NetAffxCompatibility-class*),
13
- affxCompression, AffxUrl-method
(*zzz-NetAffxCompatibility-class*),
13
- affxCrc32
(*zzz-NetAffxCompatibility-class*),
13
- affxCrc32, AffxFile-method
(*zzz-NetAffxCompatibility-class*),
13
- affxCrc32, AffxUrl-method
(*zzz-NetAffxCompatibility-class*),
13
- affxDate
(*zzz-NetAffxCompatibility-class*),
13
- affxDate, AffxFile-method
(*zzz-NetAffxCompatibility-class*),
13
- affxDate, Audit-method
(*zzz-DTTCompatibility-class*),
12
- affxDescription
(*zzz-NetAffxCompatibility-class*),
13
- affxDescription, AffxAnnotation-method
(*zzz-NetAffxCompatibility-class*),
13
- affxDescription, AffxArray-method
(*zzz-NetAffxCompatibility-class*),
13
- affxEnd
(*zzz-DTTCompatibility-class*),
12
- affxEnd, CompositePosition-method
(*zzz-DTTCompatibility-class*),
12
- affxEnd, ReporterPosition-method
(*zzz-DTTCompatibility-class*),
12
- affxEnd, SequencePosition-method
(*zzz-DTTCompatibility-class*),
12
- affxFile
(*zzz-NetAffxCompatibility-class*),
13
- affxFile, AffxAnnotation-method
(*zzz-NetAffxCompatibility-class*),
13
- affxImage

- (*zzz-DTTCompatibility-class*), [12](#)
- affxImage*, *PhysicalBioAssayData_assnlist*-method
(*zzz-DTTCompatibility-class*), [12](#)
- affxLicence*
(*zzz-NetAffxCompatibility-class*), [13](#)
- affxLicence*, *NetAffxResource*-method
(*NetAffxResource-class*), [2](#)
- affxName*
(*zzz-NetAffxCompatibility-class*), [13](#)
- affxName*, *AffxArray*-method
(*zzz-NetAffxCompatibility-class*), [13](#)
- affxName*, *AffxFile*-method
(*zzz-NetAffxCompatibility-class*), [13](#)
- affxOrder*
(*zzz-DTTCompatibility-class*), [12](#)
- affxOrder*, *BioDataCube*-method
(*zzz-DTTCompatibility-class*), [12](#)
- affxOrder*, *Treatment*-method
(*zzz-DTTCompatibility-class*), [12](#)
- affxPerson*
(*zzz-DTTCompatibility-class*), [12](#)
- affxPerson*, *Contact_assnlist*-method
(*zzz-DTTCompatibility-class*), [12](#)
- affxPerson*, *Person_assnlist*-method
(*zzz-DTTCompatibility-class*), [12](#)
- affxRow*
(*zzz-DTTCompatibility-class*), [12](#)
- affxRow*, *FeatureLocation*-method
(*zzz-DTTCompatibility-class*), [12](#)
- affxRow*, *Zone*-method
(*zzz-DTTCompatibility-class*), [12](#)
- affxSequence*
(*zzz-DTTCompatibility-class*), [12](#)
- affxSequence*, *BioSequence*-method
(*zzz-DTTCompatibility-class*), [12](#)
- affxSize*
(*zzz-NetAffxCompatibility-class*), [13](#)
- affxSize*, *AffxFile*-method
(*zzz-NetAffxCompatibility-class*), [13](#)
- affxSize*, *AffxUrl*-method
(*zzz-NetAffxCompatibility-class*), [13](#)
- affxStart*
(*zzz-DTTCompatibility-class*), [12](#)
- affxStart*, *CompositePosition*-method
(*zzz-DTTCompatibility-class*), [12](#)
- affxStart*, *ReporterPosition*-method
(*zzz-DTTCompatibility-class*), [12](#)
- affxStart*, *SequencePosition*-method
(*zzz-DTTCompatibility-class*), [12](#)
- affxStatus*
(*zzz-NetAffxCompatibility-class*), [13](#)
- affxStatus*, *AffxNetAffxAnnotFileList*-method
(*zzz-NetAffxCompatibility-class*), [13](#)
- affxText*
(*zzz-DTTCompatibility-class*), [12](#)
- affxText*, *Description*-method
(*zzz-DTTCompatibility-class*), [12](#)
- affxText*, *Protocol*-method
(*zzz-DTTCompatibility-class*), [12](#)
- affxTitle*
(*zzz-DTTCompatibility-class*), [12](#)
- affxTitle*, *BibliographicReference*-method
(*zzz-DTTCompatibility-class*), [12](#)
- affxTitle*, *Protocol*-method
(*zzz-DTTCompatibility-class*), [12](#)
- affxType*
(*zzz-NetAffxCompatibility-class*), [13](#)
- affxType*, *AffxAnnotation*-method
(*zzz-NetAffxCompatibility-class*), [13](#)
- affxType*, *AffxFile*-method

- (*zzz-NetAffxCompatibility-class*), 13
- (*zzz-DTTCompatibility-class*), 12
- affxUrl*
 - (*zzz-NetAffxCompatibility-class*), 13
 - array_package*, MAGE-ML-method (*zzz-DTTCompatibility-class*), 12
 - array_ref*, *NetAffxResource*-method (*NetAffxResource-class*), 2
 - array_ref*, *ArrayDesign*-method (*zzz-DTTCompatibility-class*), 12
 - array_ref*, *ArrayDesign_assnlist*-method (*zzz-DTTCompatibility-class*), 12
 - array_ref*, *Arrays_assnreflist*-method (*zzz-DTTCompatibility-class*), 12
 - arrayAttribute* (*zzz-ARRCompatibility-class*), 11
 - arrayAttribute*, *PhysicalArray*-method (*zzz-ARRCompatibility-class*), 11
 - arrayDesign* (*zzz-DTTCompatibility-class*), 12
 - arrayDesign*, *ArrayDesign_assnlist*-method (*zzz-DTTCompatibility-class*), 12
 - arrayDesign_assnlist* (*zzz-DTTCompatibility-class*), 12
 - arrayDesign_assnlist*, *ArrayDesign_package*-method (*zzz-DTTCompatibility-class*), 12
 - arrayDesign_assnref* (*zzz-DTTCompatibility-class*), 12
 - arrayDesign_assnref*, *Array*-method (*zzz-DTTCompatibility-class*), 12
 - arrayDesign_package* (*zzz-DTTCompatibility-class*), 12
 - arrayDesign_package*, MAGE-ML-method (*zzz-DTTCompatibility-class*), 12
 - arrayDesign_ref* (*zzz-DTTCompatibility-class*), 12
 - arrayDesign_ref*, *ArrayDesign_assnref*-method (*zzz-DTTCompatibility-class*), 12
- affxUrl*, *AffxFile*-method (*zzz-NetAffxCompatibility-class*), 13
- affxUrl*, *NetAffxResource*-method (*NetAffxResource-class*), 2
- affyBarcode* (*zzz-ARRCompatibility-class*), 11
- affyBarcode*, *PhysicalArray*-method (*zzz-ARRCompatibility-class*), 11
- AffyCompatible*, 3, 12-14
- AffyCompatible* (*AffyCompatible-package*), 2
- AffyCompatible-class*, 1
- AffyCompatible-package*, 2
- analysisResults_assnreflist* (*zzz-DTTCompatibility-class*), 12
- analysisResults_assnreflist*, *Experiment*-method (*zzz-DTTCompatibility-class*), 12
- annotations_assnlist* (*zzz-DTTCompatibility-class*), 12
- annotations_assnlist*, *Description*-method (*zzz-DTTCompatibility-class*), 12
- annotations_assnlist*, *ExperimentalFactor*-method (*zzz-DTTCompatibility-class*), 12
- annotationsFile (accessors)*, 6
- annotationsFile*, *NetAffxResource*-method (*NetAffxResource-class*), 2
- array_assnlist* (*zzz-DTTCompatibility-class*), 12
- array_assnlist*, *Array_package*-method (*zzz-DTTCompatibility-class*), 12
- array_assnref* (*zzz-DTTCompatibility-class*), 12
- array_assnref*, *BioAssayCreation*-method (*zzz-DTTCompatibility-class*), 12
- array_assnref*, *Hybridization*-method (*zzz-DTTCompatibility-class*), 12

- 12
- arrayGroup
(*zzz-DTTCompatibility-class*),
12
- arrayGroup, ArrayGroup_assnlist-method
(*zzz-DTTCompatibility-class*),
12
- arrayGroup_assnlist
(*zzz-DTTCompatibility-class*),
12
- arrayGroup_assnlist, Array_package-method
(*zzz-DTTCompatibility-class*),
12
- arrayGroup_assnref
(*zzz-DTTCompatibility-class*),
12
- arrayGroup_assnref, Array-method
(*zzz-DTTCompatibility-class*),
12
- arrayGroup_ref
(*zzz-DTTCompatibility-class*),
12
- arrayGroup_ref, ArrayGroup_assnref-method
(*zzz-DTTCompatibility-class*),
12
- arrayIdentifier
(*zzz-DTTCompatibility-class*),
12
- arrayIdentifier, Array-method
(*zzz-DTTCompatibility-class*),
12
- arrayManufacture
(*zzz-DTTCompatibility-class*),
12
- arrayManufacture, ArrayManufacture_assnlist-method
(*zzz-DTTCompatibility-class*),
12
- arrayManufacture_assnlist
(*zzz-DTTCompatibility-class*),
12
- arrayManufacture_assnlist, Array_package-method
(*zzz-DTTCompatibility-class*),
12
- arrayManufacture_ref
(*zzz-DTTCompatibility-class*),
12
- arrayManufacture_ref, Information_assnref-method
(*zzz-DTTCompatibility-class*),
12
- arrayManufactureDeviation
(*zzz-DTTCompatibility-class*),
12
- arrayManufactureDeviation, ArrayManufactureDeviation
(*zzz-DTTCompatibility-class*),
12
- arrayManufactureDeviations_assnlist
(*zzz-DTTCompatibility-class*),
12
- arrayManufactureDeviations_assnlist, Array-method
(*zzz-DTTCompatibility-class*),
12
- arrayManufacturers_assnreflist
(*zzz-DTTCompatibility-class*),
12
- arrayManufacturers_assnreflist, ArrayManufacturers
(*zzz-DTTCompatibility-class*),
12
- arrayName
(*zzz-ARRCompatibility-class*),
11
- arrayName, PhysicalArray-method
(*zzz-ARRCompatibility-class*),
11
- arrays_assnreflist
(*zzz-DTTCompatibility-class*),
12
- arrays_assnreflist, ArrayGroup-method
(*zzz-DTTCompatibility-class*),
12
- arrays_assnreflist, ArrayManufacture-method
(*zzz-DTTCompatibility-class*),
12
- arraySpacingX
(*zzz-DTTCompatibility-class*),
12
- arraySpacingX, ArrayGroup-method
(*zzz-DTTCompatibility-class*),
12
- arraySpacingY
(*zzz-DTTCompatibility-class*),
12
- arraySpacingY, ArrayGroup-method
(*zzz-DTTCompatibility-class*),
12
- arrayXOrigin
(*zzz-DTTCompatibility-class*),
12
- arrayXOrigin, Array-method
(*zzz-DTTCompatibility-class*),
12
- arrayYOrigin
(*zzz-DTTCompatibility-class*),
12
- arrayYOrigin, Array-method

- (zzz-*DTTCompatibility-class*),
12
- ARRCompatibility, 12
- ARRCompatibility-class
(*AffyCompatible-class*), 1
- associations_assnlist
(zzz-*DTTCompatibility-class*),
12
- associations_assnlist, *OntologyEntry*-method
(zzz-*DTTCompatibility-class*),
12
- audit
(zzz-*DTTCompatibility-class*),
12
- audit, *AuditTrail_assnlist*-method
(zzz-*DTTCompatibility-class*),
12
- auditAndSecurity_package
(zzz-*DTTCompatibility-class*),
12
- auditAndSecurity_package, *MAGE-ML*-method
(zzz-*DTTCompatibility-class*),
12
- auditTrail_assnlist
(zzz-*DTTCompatibility-class*),
12
- auditTrail_assnlist, *Array*-method
(zzz-*DTTCompatibility-class*),
12
- auditTrail_assnlist, *ArrayDesign*-method
(zzz-*DTTCompatibility-class*),
12
- auditTrail_assnlist, *ArrayGroup*-method
(zzz-*DTTCompatibility-class*),
12
- auditTrail_assnlist, *ArrayManufacture*-method
(zzz-*DTTCompatibility-class*),
12
- auditTrail_assnlist, *Audit*-method
(zzz-*DTTCompatibility-class*),
12
- auditTrail_assnlist, *BibliographicReference*-method
(zzz-*DTTCompatibility-class*),
12
- auditTrail_assnlist, *BioAssayCreation*-method
(zzz-*DTTCompatibility-class*),
12
- auditTrail_assnlist, *BioAssayDataCluster*-method
(zzz-*DTTCompatibility-class*),
12
- auditTrail_assnlist, *BioAssayDimension*-method
(zzz-*DTTCompatibility-class*),
12
- 12
- auditTrail_assnlist, *BioAssayMap*-method
(zzz-*DTTCompatibility-class*),
12
- auditTrail_assnlist, *BioAssayTreatment*-method
(zzz-*DTTCompatibility-class*),
12
- auditTrail_assnlist, *BioSample*-method
(zzz-*DTTCompatibility-class*),
12
- auditTrail_assnlist, *BioSequence*-method
(zzz-*DTTCompatibility-class*),
12
- auditTrail_assnlist, *BioSource*-method
(zzz-*DTTCompatibility-class*),
12
- auditTrail_assnlist, *Channel*-method
(zzz-*DTTCompatibility-class*),
12
- auditTrail_assnlist, *CompositeCompositeMap*-method
(zzz-*DTTCompatibility-class*),
12
- auditTrail_assnlist, *CompositeGroup*-method
(zzz-*DTTCompatibility-class*),
12
- auditTrail_assnlist, *CompositeSequence*-method
(zzz-*DTTCompatibility-class*),
12
- auditTrail_assnlist, *CompositeSequenceDimension*-method
(zzz-*DTTCompatibility-class*),
12
- auditTrail_assnlist, *Compound*-method
(zzz-*DTTCompatibility-class*),
12
- auditTrail_assnlist, *Database*-method
(zzz-*DTTCompatibility-class*),
12
- auditTrail_assnlist, *DerivedBioAssay*-method
(zzz-*DTTCompatibility-class*),
12
- auditTrail_assnlist, *DerivedBioAssayData*-method
(zzz-*DTTCompatibility-class*),
12
- auditTrail_assnlist, *DerivedSignal*-method
(zzz-*DTTCompatibility-class*),
12
- auditTrail_assnlist, *Description*-method
(zzz-*DTTCompatibility-class*),
12
- auditTrail_assnlist, *Error*-method
(zzz-*DTTCompatibility-class*),
12

- auditTrail_assnlist, ExpectedValue-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, Experiment-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, ExperimentalFactor-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, ExperimentDesign-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, FactorValue-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, Failed-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, Feature-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, FeatureDimension-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, FeatureExtraction-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, FeatureGroup-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, FeatureReporterMap-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, Fiducial-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, Hardware-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, HardwareApplication-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, Hybridization-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, Image-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, ImageAcquisition-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, LabeledExtract-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, MAGE-ML-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, ManufactureLIMS-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, ManufactureLIMSBiomaterial-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, MeasuredBioAssay-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, MeasuredBioAssayData-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, MeasuredSignal-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, Node-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, NodeContents-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, Organization-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, Parameter-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, Person-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, PhysicalArrayDesign-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, PhysicalBioAssay-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, PresentAbsent-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, Protocol-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, ProtocolApplication-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- auditTrail_assnlist, PValue-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12

[12](#) authors, BibliographicReference-method
 auditTrail_assnlist, QuantitationTypeDimensionalMethod-method
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 auditTrail_assnlist, QuantitationTypeMap-method
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 auditTrail_assnlist, Ratio-method
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 auditTrail_assnlist, Reporter-method
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 auditTrail_assnlist, ReporterCompositeMap-method
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 auditTrail_assnlist, ReporterDimensionalMethod-method
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 auditTrail_assnlist, ReporterGroup-method
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 auditTrail_assnlist, Security-method
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 auditTrail_assnlist, SecurityGroup-method
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 auditTrail_assnlist, SeqFeature-method
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 auditTrail_assnlist, Software-method
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 auditTrail_assnlist, SoftwareApplication-method
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 auditTrail_assnlist, SpecializedQuantitationType-method
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 auditTrail_assnlist, Transformation-method
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 auditTrail_assnlist, Treatment-method
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 auditTrail_assnlist, Zone-method
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 authors
 (zzz-DTTCompatibility-class), [12](#)
[12](#)

authors, BibliographicReference-method
 barcode
 barcode, ArrayGroup-method
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 basis
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 basis, SeqFeature-method
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 bibliographicReference
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 bibliographicReference, BibliographicReferences
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 bibliographicReferences_assnlist
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 bibliographicReferences_assnlist, Description-r
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 bioAssay_assnlist
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 bioAssay_assnlist, BioAssay_package-method
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 bioAssay_assnref
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 bioAssay_assnref, BioAssayTuple-method
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 bioAssay_package
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 bioAssay_package, MAGE-ML-method
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 bioAssayCreation
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 bioAssayCreation, BioAssayCreation_assn-method
 (zzz-DTTCompatibility-class), [12](#)
[12](#)
 bioAssayCreation_assn

	(<i>zzz-DTTCompatibility-class</i>),	12	
	12		bioAssayDimension_assnlist
bioAssayCreation_assn,PhysicalBioAssay-method	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
	12		bioAssayDimension_assnlist,BioAssayData_package
bioAssayData_assnlist	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
	12		bioAssayDimension_assnref
bioAssayData_assnlist,BioAssayData_package-method	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
	12		bioAssayDimension_assnref,DerivedBioAssayData
bioAssayData_assnreflist	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
	12		bioAssayDimension_assnref,MeasuredBioAssayData
bioAssayData_assnreflist,Experiment-method	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
	12		bioAssayDimension_assnref,NodeContents-method
bioAssayData_package	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
	12		bioAssayDimension_ref
bioAssayData_package,MAGE-ML-method	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
	12		bioAssayDimension_ref,BioAssayDimension_assnref
bioAssayDataCluster	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
	12		bioAssayFactorValues_assnreflist
bioAssayDataCluster,BioAssayDataCluster_assnlist-method	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
	12		bioAssayFactorValues_assnreflist,DerivedBioAss
bioAssayDataCluster_assnlist	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
	12		bioAssayFactorValues_assnreflist,MeasuredBioAss
bioAssayDataCluster_assnlist,HigherLevelAnalysis-method	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
	12		bioAssayFactorValues_assnreflist,PhysicalBioAss
bioAssayDataCluster_ref	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
	12		bioAssayMap
bioAssayDataCluster_ref,AnalysisResults_assnreflist-method	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
	12		bioAssayMap,BioAssayMap_assnlist-method
bioAssayDataSources_assnreflist	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
	12		bioAssayMap_assnlist
bioAssayDataSources_assnreflist,Transformation-method	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
	12		bioAssayMap_assnlist,BioAssayData_package-method
bioAssayDimension	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
	12		bioAssayMap_ref
bioAssayDimension,BioAssayDimension_assnlist-method	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
	12		12

bioAssayMap_ref, BioAssayMaps_assnreflist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 bioAssayMap_ref, DerivedBioAssayMap_assnreflist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 bioAssayMapping (zzz-DTTCompatibility-class), 12
 bioAssayMapping, BioAssayMapping_assn-method (zzz-DTTCompatibility-class), 12
 bioAssayMapping_assn (zzz-DTTCompatibility-class), 12
 bioAssayMapping_assn, Transformation-method (zzz-DTTCompatibility-class), 12
 bioAssayMaps_assnreflist (zzz-DTTCompatibility-class), 12
 bioAssayMaps_assnreflist, BioAssayMapping-method (zzz-DTTCompatibility-class), 12
 bioAssayMapTarget_assnref (zzz-DTTCompatibility-class), 12
 bioAssayMapTarget_assnref, BioAssayMap-method (zzz-DTTCompatibility-class), 12
 bioAssays_assnreflist (zzz-DTTCompatibility-class), 12
 bioAssays_assnreflist, BioAssayDimension-method (zzz-DTTCompatibility-class), 12
 bioAssays_assnreflist, Experiment-method (zzz-DTTCompatibility-class), 12
 bioAssayTreatment (zzz-DTTCompatibility-class), 12
 bioAssayTreatment, BioAssayTreatments_assnlist-method (zzz-DTTCompatibility-class), 12
 bioAssayTreatments_assnlist (zzz-DTTCompatibility-class), 12
 bioAssayTreatments_assnlist, PhysicalBioAssay-method (zzz-DTTCompatibility-class), 12
 bioAssayTuple (zzz-DTTCompatibility-class),

bioAssayTuple, BioAssayTuples_assnlist-method (zzz-DTTCompatibility-class),
 bioAssayTuples_assnlist (zzz-DTTCompatibility-class),
 bioAssayTuples_assnlist, BioDataTuples-method (zzz-DTTCompatibility-class),
 bioDataCube (zzz-DTTCompatibility-class), 12
 bioDataCube, BioDataValues_assn-method (zzz-DTTCompatibility-class), 12
 bioDataTuples (zzz-DTTCompatibility-class), 12
 bioDataTuples, BioDataValues_assn-method (zzz-DTTCompatibility-class), 12
 bioDataValues_assn (zzz-DTTCompatibility-class), 12
 bioDataValues_assn, DerivedBioAssayData-method (zzz-DTTCompatibility-class), 12
 bioDataValues_assn, MeasuredBioAssayData-method (zzz-DTTCompatibility-class), 12
 bioEvent_package (zzz-DTTCompatibility-class), 12
 bioEvent_package, MAGE-ML-method (zzz-DTTCompatibility-class), 12
 biologicalCharacteristics_assnreflist (zzz-DTTCompatibility-class), 12
 biologicalCharacteristics_assnreflist, Composit (zzz-DTTCompatibility-class), 12
 bioMaterial_assnlist (zzz-DTTCompatibility-class), 12
 bioMaterial_assnlist, BioMaterial_package-method (zzz-DTTCompatibility-class), 12
 bioMaterial_assnref (zzz-DTTCompatibility-class),

12
 bioMaterial_assnref, BioMaterialMeasurement-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 bioMaterial_assnref, BioSequence, BioSequence_assnlist-method
 (zzz-DTTCompatibility-class), 12
 bioMaterial_assnref, ManufactureLIMS-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 bioMaterial_assnref, ManufactureLIMSBiomaterial-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 bioMaterial_package
 (zzz-DTTCompatibility-class), 12
 12
 bioMaterial_package, MAGE-ML-method
 (zzz-DTTCompatibility-class), 12
 12
 bioMaterialMeasurement
 (zzz-DTTCompatibility-class), 12
 12
 bioMaterialMeasurement, SourceBioMaterialMeasurements_assnlist-method
 (zzz-DTTCompatibility-class), 12
 12
 bioMaterialPlateCol
 (zzz-DTTCompatibility-class), 12
 12
 bioMaterialPlateCol, ManufactureLIMSBiomaterial-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 bioMaterialPlateIdentifier
 (zzz-DTTCompatibility-class), 12
 12
 bioMaterialPlateIdentifier, ManufactureLIMSBiomaterial-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 bioMaterialPlateRow
 (zzz-DTTCompatibility-class), 12
 12
 bioMaterialPlateRow, ManufactureLIMSBiomaterial-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 bioSample
 (zzz-DTTCompatibility-class), 12
 12
 bioSample, BioMaterial_assnlist-method
 (zzz-DTTCompatibility-class), 12
 12
 bioSample_ref
 (zzz-DTTCompatibility-class), 12
 12
 bioSample_ref, BioMaterial_assnref-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 bioSequence
 (zzz-DTTCompatibility-class), 12
 bioSequence, BioSequence_assnlist-method
 (zzz-DTTCompatibility-class), 12
 bioSequence_assnlist
 (zzz-DTTCompatibility-class), 12
 bioSequence_assnlist, BioSequence_package-method
 (zzz-DTTCompatibility-class), 12
 bioSequence_package
 (zzz-DTTCompatibility-class), 12
 bioSequence_package, MAGE-ML-method
 (zzz-DTTCompatibility-class), 12
 bioSequence_ref
 (zzz-DTTCompatibility-class), 12
 bioSequence_ref, BiologicalCharacteristics_assnlist-method
 (zzz-DTTCompatibility-class), 12
 bioSequence_ref, ImmobilizedCharacteristics_assnlist-method
 (zzz-DTTCompatibility-class), 12
 bioSource
 (zzz-DTTCompatibility-class), 12
 bioSource, BioMaterial_assnlist-method
 (zzz-DTTCompatibility-class), 12
 bioSource_ref
 (zzz-DTTCompatibility-class), 12
 bioSource_ref, BioMaterial_assnref-method
 (zzz-DTTCompatibility-class), 12
 bqs_package
 (zzz-DTTCompatibility-class), 12
 bqs_package, MAGE-ML-method
 (zzz-DTTCompatibility-class), 12
 category_assn
 (zzz-DTTCompatibility-class), 12
 category_assn, ExperimentalFactor-method
 (zzz-DTTCompatibility-class), 12
 channel

- (zzz-*DTTCompatibility-class*),
12
- channel, Channel_assnlist-method
(zzz-*DTTCompatibility-class*),
12
- channel_assnlist
(zzz-*DTTCompatibility-class*),
12
- channel_assnlist, BioAssay_package-method
(zzz-*DTTCompatibility-class*),
12
- channel_assnref
(zzz-*DTTCompatibility-class*),
12
- channel_assnref, DerivedSignal-method
(zzz-*DTTCompatibility-class*),
12
- channel_assnref, Error-method
(zzz-*DTTCompatibility-class*),
12
- channel_assnref, ExpectedValue-method
(zzz-*DTTCompatibility-class*),
12
- channel_assnref, Failed-method
(zzz-*DTTCompatibility-class*),
12
- channel_assnref, MeasuredSignal-method
(zzz-*DTTCompatibility-class*),
12
- channel_assnref, PresentAbsent-method
(zzz-*DTTCompatibility-class*),
12
- channel_assnref, PValue-method
(zzz-*DTTCompatibility-class*),
12
- channel_assnref, Ratio-method
(zzz-*DTTCompatibility-class*),
12
- channel_assnref, SpecializedQuantitationType-method
(zzz-*DTTCompatibility-class*),
12
- channel_ref
(zzz-*DTTCompatibility-class*),
12
- channel_ref, Channel_assnref-method
(zzz-*DTTCompatibility-class*),
12
- channel_ref, Channels_assnreflist-method
(zzz-*DTTCompatibility-class*),
12
- channels_assnreflist
(zzz-*DTTCompatibility-class*),
12
- channels_assnreflist, DerivedBioAssay-method
(zzz-*DTTCompatibility-class*),
12
- channels_assnreflist, Image-method
(zzz-*DTTCompatibility-class*),
12
- channels_assnreflist, MeasuredBioAssay-method
(zzz-*DTTCompatibility-class*),
12
- channels_assnreflist, PhysicalBioAssay-method
(zzz-*DTTCompatibility-class*),
12
- characteristics_assnlist
(zzz-*DTTCompatibility-class*),
12
- characteristics_assnlist, BioSample-method
(zzz-*DTTCompatibility-class*),
12
- characteristics_assnlist, BioSource-method
(zzz-*DTTCompatibility-class*),
12
- characteristics_assnlist, LabeledExtract-method
(zzz-*DTTCompatibility-class*),
12
- clusterBioAssayData_assnref
(zzz-*DTTCompatibility-class*),
12
- clusterBioAssayData_assnref, BioAssayDataCluster-method
(zzz-*DTTCompatibility-class*),
12
- column
(zzz-*DTTCompatibility-class*),
12
- column, FeatureLocation-method
(zzz-*DTTCompatibility-class*),
12
- column, Zone-method
(zzz-*DTTCompatibility-class*),
12
- componentCompounds_assnlist
(zzz-*DTTCompatibility-class*),
12
- componentCompounds_assnlist, Compound-method
(zzz-*DTTCompatibility-class*),
12
- composite_assnref
(zzz-*DTTCompatibility-class*),
12
- composite_assnref, CompositePosition-method
(zzz-*DTTCompatibility-class*),
12

- 12
 compositeSequence_ref, CompositeSequences_assnref-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- 12
 compositeSequence_ref, DesignElement_assnref-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- 12
 compositeSequenceDimension (zzz-DTTCompatibility-class), 12
- 12
 compositeSequenceDimension, DesignElementDimension_assnref-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- 12
 compositeSequenceDimension_ref (zzz-DTTCompatibility-class), 12
- 12
 compositeSequenceDimension_ref, DesignElementDimension_assnref-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- 12
 compositeSequences_assnreflist (zzz-DTTCompatibility-class), 12
- 12
 compositeSequences_assnreflist, CompositeGroup-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- 12
 compositeSequences_assnreflist, CompositeSequenceGroup-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- 12
 compound (zzz-DTTCompatibility-class), 12
- 12
 compound, Compound_assnlist-method (zzz-DTTCompatibility-class), 12
- 12
 compound_assnlist (zzz-DTTCompatibility-class), 12
- 12
 compound_assnlist, BioMaterial_package-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- 12
 compound_assnref (zzz-DTTCompatibility-class), 12
- 12
 compound_assnref, CompoundMeasurement-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- 12
 compound_ref (zzz-DTTCompatibility-class), 12
- 12
 compound_ref, Compound_assnref-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- 12
 compound_ref, Labels_assnreflist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- 12
 compoundIndices_assnlist (zzz-DTTCompatibility-class), 12
- 12
 compoundIndices_assnlist, Compound-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- 12
 compoundMeasurement (zzz-DTTCompatibility-class), 12
- 12
 compoundMeasurement, ComponentCompounds_assnlist (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- 12
 compoundMeasurement, CompoundMeasurements_assnlist (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- 12
 compoundMeasurements_assnlist (zzz-DTTCompatibility-class), 12
- 12
 compoundMeasurements_assnlist, Treatment-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- 12
 concentrationUnit (zzz-DTTCompatibility-class), 12
- 12
 concentrationUnit, Unit_assn-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- 12
 confidenceIndicators_assnreflist (zzz-DTTCompatibility-class), 12
- 12
 confidenceIndicators_assnreflist, DerivedSignal-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- 12
 confidenceIndicators_assnreflist, Error-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- 12
 confidenceIndicators_assnreflist, ExpectedValue-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- 12
 confidenceIndicators_assnreflist, Failed-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- 12
 confidenceIndicators_assnreflist, MeasuredSignal-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- 12
 confidenceIndicators_assnreflist, PresentAbsent-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
- 12
 confidenceIndicators_assnreflist, PValue-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12

- (*zzz-DTTCompatibility-class*), 12
 12
 confidenceIndicators_assnreflist, Ratio-method (*zzz-DTTCompatibility-class*), 12
 (*zzz-DTTCompatibility-class*), 12
 confidenceIndicators_assnreflist, SpecializedQuantitativeControlType-method (*zzz-DTTCompatibility-class*), 12
 (*zzz-DTTCompatibility-class*), 12
 contact_assnlist
 (*zzz-DTTCompatibility-class*), 12
 12
 contact_assnlist, AuditAndSecurity_package-method (*zzz-ARRCompatibility-class*), 11
 (*zzz-DTTCompatibility-class*), 12
 12
 contacts_assnreflist
 (*zzz-DTTCompatibility-class*), 12
 12
 contacts_assnreflist, Database-method
 (*zzz-DTTCompatibility-class*), 12
 12
 containedFeatures_assnreflist
 (*zzz-DTTCompatibility-class*), 12
 12
 containedFeatures_assnreflist, FeatureDimension-method
 (*zzz-DTTCompatibility-class*), 12
 12
 control
 (*zzz-ARRCompatibility-class*), 11
 11
 control, UserAttribute-method
 (*zzz-ARRCompatibility-class*), 11
 11
 controlFeatures_assnreflist
 (*zzz-DTTCompatibility-class*), 12
 12
 controlFeatures_assnreflist, Feature-method
 (*zzz-DTTCompatibility-class*), 12
 12
 controlledFeatures_assnreflist
 (*zzz-DTTCompatibility-class*), 12
 12
 controlledFeatures_assnreflist, Feature-method
 (*zzz-DTTCompatibility-class*), 12
 12
 controlType_assn
 (*zzz-DTTCompatibility-class*), 12
 12
 controlType_assn, CompositeSequence-method (*zzz-DTTCompatibility-class*), 12
 (*zzz-DTTCompatibility-class*), 12
 12
 controlType_assn, Feature-method
 (*zzz-DTTCompatibility-class*), 12
 12
- controlType_assn, Reporter-method
 (*zzz-DTTCompatibility-class*), 12
 coordinate_assn
 (*zzz-DTTCompatibility-class*), 12
 coordinate_assn, SeqFeatureLocation-method
 (*zzz-DTTCompatibility-class*), 12
 12
 createdBy
 (*zzz-ARRCompatibility-class*), 11
 11
 createdBy, ArraySetFile-method
 (*zzz-ARRCompatibility-class*), 11
 11
 createdBy, PhysicalArray-method
 (*zzz-ARRCompatibility-class*), 11
 11
 createdBy, TemplateFile-method
 (*zzz-ARRCompatibility-class*), 11
 11
 createdBy
 (*zzz-ARRCompatibility-class*), 11
 11
 createdBy, ArraySetFile-method
 (*zzz-ARRCompatibility-class*), 11
 11
 createdBy, PhysicalArray-method
 (*zzz-ARRCompatibility-class*), 11
 11
 createdBy, TemplateFile-method
 (*zzz-ARRCompatibility-class*), 11
 11
 createdDateTime
 (*zzz-ARRCompatibility-class*), 11
 11
 createdDateTime, ArraySetFile-method
 (*zzz-ARRCompatibility-class*), 11
 11
 createdDateTime, PhysicalArray-method
 (*zzz-ARRCompatibility-class*), 11
 11
 createdDateTime, TemplateFile-method
 (*zzz-ARRCompatibility-class*), 11
 11
 createdStep
 (*zzz-ARRCompatibility-class*), 11
 11
 createdStep, ArraySetFile-method
 (*zzz-ARRCompatibility-class*), 11
 11
 createdStep, PhysicalArray-method
 (*zzz-ARRCompatibility-class*), 11
 11
 createdStep
 (*zzz-ARRCompatibility-class*), 11
 11
- database
 (*zzz-DTTCompatibility-class*), 12
 12
 database, Database_assnlist-method
 (*zzz-DTTCompatibility-class*), 12
 12
 database_assnlist
 (*zzz-DTTCompatibility-class*), 12
 12

database_assnlist, Description_package-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 database_assnref
 (zzz-DTTCompatibility-class), 12
 database_assnref, DatabaseEntry-method
 (zzz-DTTCompatibility-class), 12
 database_ref
 (zzz-DTTCompatibility-class), 12
 database_ref, Database_assnref-method
 (zzz-DTTCompatibility-class), 12
 databaseEntry
 (zzz-DTTCompatibility-class), 12
 databaseEntry, Accessions_assnlist-method
 (zzz-DTTCompatibility-class), 12
 databaseEntry, DatabaseReferences_assnlist-method
 (zzz-DTTCompatibility-class), 12
 databaseEntry, ExternalLIMS_assn-method
 (zzz-DTTCompatibility-class), 12
 databaseEntry, IdentifierLIMS_assn-method
 (zzz-DTTCompatibility-class), 12
 databaseEntry, OntologyReference_assn-method
 (zzz-DTTCompatibility-class), 12
 databaseEntry, SequenceDatabases_assnlist-method
 (zzz-DTTCompatibility-class), 12
 databaseReferences_assnlist
 (zzz-DTTCompatibility-class), 12
 databaseReferences_assnlist, Description-method
 (zzz-DTTCompatibility-class), 12
 dataExternal
 (zzz-DTTCompatibility-class), 12
 dataExternal, DataExternal_assn-method
 (zzz-DTTCompatibility-class), 12
 dataExternal_assn
 (zzz-DTTCompatibility-class), 12
 dataExternal_assn, BioDataCube-method
 (zzz-DTTCompatibility-class), 12
 dataFormat
 (zzz-DTTCompatibility-class), 12
 dataFormat, DataExternal-method
 (zzz-DTTCompatibility-class), 12
 dataFormatInfoURI
 (zzz-DTTCompatibility-class), 12
 dataFormatInfoURI, DataExternal-method
 (zzz-DTTCompatibility-class), 12
 dataInternal
 (zzz-DTTCompatibility-class), 12
 dataInternal, DataInternal_assn-method
 (zzz-DTTCompatibility-class), 12
 dataInternal_assn
 (zzz-DTTCompatibility-class), 12
 dataInternal_assn, BioDataCube-method
 (zzz-DTTCompatibility-class), 12
 dataType_assn
 (zzz-DTTCompatibility-class), 12
 dataType_assn, DerivedSignal-method
 (zzz-DTTCompatibility-class), 12
 dataType_assn, Error-method
 (zzz-DTTCompatibility-class), 12
 dataType_assn, ExpectedValue-method
 (zzz-DTTCompatibility-class), 12
 dataType_assn, Failed-method
 (zzz-DTTCompatibility-class), 12
 dataType_assn, MeasuredSignal-method
 (zzz-DTTCompatibility-class), 12
 dataType_assn, NodeValue-method
 (zzz-DTTCompatibility-class), 12
 dataType_assn, Parameter-method
 (zzz-DTTCompatibility-class), 12
 dataType_assn, PresentAbsent-method
 (zzz-DTTCompatibility-class), 12

- 12
- dataType_assn, PValue-method
(zzz-DTTCompatibility-class),
12
- dataType_assn, Ratio-method
(zzz-DTTCompatibility-class),
12
- dataType_assn, SpecializedQuantitationType-method
(zzz-DTTCompatibility-class),
12
- datum
(zzz-DTTCompatibility-class),
12
- datum, Datum_assn-method
(zzz-DTTCompatibility-class),
12
- datum_assn
(zzz-DTTCompatibility-class),
12
- datum_assn, QuantitationTypeTuple-method
(zzz-DTTCompatibility-class),
12
- defaultValue
(zzz-ARRCompatibility-class),
11
- defaultValue, UserAttribute-method
(zzz-ARRCompatibility-class),
11
- defaultValue_assn
(zzz-DTTCompatibility-class),
12
- defaultValue_assn, Parameter-method
(zzz-DTTCompatibility-class),
12
- defectType_assn
(zzz-DTTCompatibility-class),
12
- defectType_assn, FeatureDefect-method
(zzz-DTTCompatibility-class),
12
- defectType_assn, ZoneDefect-method
(zzz-DTTCompatibility-class),
12
- deltaX
(zzz-DTTCompatibility-class),
12
- deltaX, PositionDelta-method
(zzz-DTTCompatibility-class),
12
- deltaY
(zzz-DTTCompatibility-class),
12
- deltaY, PositionDelta-method
(zzz-DTTCompatibility-class),
12
- derivedBioAssay
(zzz-DTTCompatibility-class),
12
- derivedBioAssay, BioAssay_assnlist-method
(zzz-DTTCompatibility-class),
12
- derivedBioAssay, DerivedBioAssay_assnlist-method
(zzz-DTTCompatibility-class),
12
- derivedBioAssay_ref
(zzz-DTTCompatibility-class),
12
- derivedBioAssay_ref, BioAssay_assnref-method
(zzz-DTTCompatibility-class),
12
- derivedBioAssay_ref, BioAssayMapTarget_assnref-method
(zzz-DTTCompatibility-class),
12
- derivedBioAssay_ref, BioAssays_assnreflist-method
(zzz-DTTCompatibility-class),
12
- derivedBioAssay_ref, SourceBioAssays_assnreflist-method
(zzz-DTTCompatibility-class),
12
- derivedBioAssay_ref, TopLevelBioAssays_assnreflist-method
(zzz-DTTCompatibility-class),
12
- derivedBioAssayData
(zzz-DTTCompatibility-class),
12
- derivedBioAssayData, BioAssayData_assnlist-method
(zzz-DTTCompatibility-class),
12
- derivedBioAssayData, DerivedBioAssayData_assnlist-method
(zzz-DTTCompatibility-class),
12
- derivedBioAssayData_assnreflist
(zzz-DTTCompatibility-class),
12
- derivedBioAssayData_assnreflist, DerivedBioAssayData_assnreflist-method
(zzz-DTTCompatibility-class),
12
- derivedBioAssayData_ref
(zzz-DTTCompatibility-class),
12
- derivedBioAssayData_ref, BioAssayData_assnreflist-method
(zzz-DTTCompatibility-class),
12
- derivedBioAssayData_ref, BioAssayDataSources_assnreflist-method
(zzz-DTTCompatibility-class),
12

(zzz-DTTCompatibility-class), 12
 12 description, ReplicateDescription_assn-method
 derivedBioAssayData_ref, ClusterBioAssayData_assnref-method
 (zzz-DTTCompatibility-class), 12
 12 description_package
 derivedBioAssayData_ref, DerivedBioAssayData_assnref-method
 (zzz-DTTCompatibility-class), 12
 12 description_package, MAGE-ML-method
 derivedBioAssayData_ref, DerivedBioAssayData_assnref-method
 (zzz-DTTCompatibility-class), 12
 12 descriptions_assnlist
 derivedBioAssayMap_assnreflist (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12 descriptions_assnlist, Array-method
 derivedBioAssayMap_assnreflist, DerivedBioAssayMap_assnreflist
 (zzz-DTTCompatibility-class), 12
 12 descriptions_assnlist, ArrayDesign-method
 derivedSignal (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12 descriptions_assnlist, ArrayGroup-method
 derivedSignal, QuantitationType_assnlist-method
 (zzz-DTTCompatibility-class), 12
 12 descriptions_assnlist, ArrayManufacture-method
 derivedSignal_ref (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12 descriptions_assnlist, Audit-method
 derivedSignal_ref, QuantitationType_assnref-method
 (zzz-DTTCompatibility-class), 12
 12 descriptions_assnlist, BibliographicReference-method
 derivedSignal_ref, QuantitationTypes_assnreflist
 (zzz-DTTCompatibility-class), 12
 12 descriptions_assnlist, BioAssayCreation-method
 derivedSignal_ref, SourcesQuantitationType_assnreflist
 (zzz-DTTCompatibility-class), 12
 12 descriptions_assnlist, BioAssayDataCluster-method
 derivedSignal_ref, TargetQuantitationType_assnreflist
 (zzz-DTTCompatibility-class), 12
 12 descriptions_assnlist, BioAssayDimension-method
 description (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12 descriptions_assnlist, BioAssayMap-method
 description, Descriptions_assnlist-method
 (zzz-DTTCompatibility-class), 12
 12 descriptions_assnlist, BioAssayTreatment-method
 description, NormalizationDescription_assn-method
 (zzz-DTTCompatibility-class), 12
 12 descriptions_assnlist, BioSample-method
 description, OntologyEntry-method
 (zzz-DTTCompatibility-class), 12
 12 descriptions_assnlist, BioSequence-method
 description, QualityControlDescription_assn-method
 (zzz-DTTCompatibility-class), 12
 12

descriptions_assnlist, BioSource-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, Channel-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, CompositeCompositeMap-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, CompositeGroup-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, CompositeSequence-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, CompositeSequenceDimension-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, Compound-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, Database-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, DerivedBioAssay-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, DerivedBioAssayData-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, DerivedSignal-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, Description-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, Error-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, ExpectedValue-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, Experiment-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, ExperimentalFactor-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, ExperimentDesign-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, FactorValue-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, Failed-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, Feature-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, FeatureDimension-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, FeatureExtraction-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, FeatureGroup-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, FeatureReporterMap-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, Fiducial-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, Hardware-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, HardwareApplication-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, Hybridization-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, Image-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, ImageAcquisition-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, LabeledExtract-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, MAGE-ML-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, ManufactureLIMS-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, ManufactureLIMSBiomaterial-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 descriptions_assnlist, MeasuredBioAssay-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12

[12](#) descriptions_assnlist, ReporterCompositeMap-method
 descriptions_assnlist, MeasuredBioAssayData-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), [12](#)
[12](#) descriptions_assnlist, ReporterDimension-method
 descriptions_assnlist, MeasuredSignal-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), [12](#)
[12](#) descriptions_assnlist, ReporterGroup-method
 descriptions_assnlist, Node-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), [12](#)
[12](#) descriptions_assnlist, Security-method
 descriptions_assnlist, NodeContents-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), [12](#)
[12](#) descriptions_assnlist, SecurityGroup-method
 descriptions_assnlist, Organization-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), [12](#)
[12](#) descriptions_assnlist, SeqFeature-method
 descriptions_assnlist, Parameter-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), [12](#)
[12](#) descriptions_assnlist, Software-method
 descriptions_assnlist, Person-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), [12](#)
[12](#) descriptions_assnlist, SoftwareApplication-method
 descriptions_assnlist, PhysicalArrayDesign-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), [12](#)
[12](#) descriptions_assnlist, SpecializedQuantitationMethod
 descriptions_assnlist, PhysicalBioAssay-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), [12](#)
[12](#) descriptions_assnlist, Transformation-method
 descriptions_assnlist, PresentAbsent-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), [12](#)
[12](#) descriptions_assnlist, Treatment-method
 descriptions_assnlist, Protocol-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), [12](#)
[12](#) descriptions_assnlist, Zone-method
 descriptions_assnlist, ProtocolApplication-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), [12](#)
[12](#) designElement_assnref
 descriptions_assnlist, PValue-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), [12](#)
[12](#) designElement_assnref, DesignElementTuple-method
 descriptions_assnlist, QuantitationTypeDimensionMethod (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), [12](#)
[12](#) designElement_package
 descriptions_assnlist, QuantitationTypeMap-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), [12](#)
[12](#) designElement_package, MAGE-ML-method
 descriptions_assnlist, Ratio-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), [12](#)
[12](#) designElementDimension_assnlist
 descriptions_assnlist, Reporter-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), [12](#)
[12](#) designElementDimension_assnlist, BioAssayData_p

- (zzz-DTTCompatibility-class), 12
- designElementDimension_assnref (zzz-DTTCompatibility-class), 12
- designElementDimension_assnref, DerivedBioAssayData-method (zzz-DTTCompatibility-class), 12
- designElementDimension_assnref, MeasuredBioAssayData-method (zzz-DTTCompatibility-class), 12
- designElementDimension_assnref, NodeContents-method (zzz-DTTCompatibility-class), 12
- designElementMapping (zzz-DTTCompatibility-class), 12
- designElementMapping, DesignElementMapping_assn-method (zzz-DTTCompatibility-class), 12
- designElementMapping_assn (zzz-DTTCompatibility-class), 12
- designElementMapping_assn, Transformation-method (zzz-DTTCompatibility-class), 12
- designElementMaps_assnreflist (zzz-DTTCompatibility-class), 12
- designElementMaps_assnreflist, DesignElementMapping-method (zzz-DTTCompatibility-class), 12
- designElementTuple (zzz-DTTCompatibility-class), 12
- designElementTuple, DesignElementTuples_assnlist-method (zzz-DTTCompatibility-class), 12
- designElementTuples_assnlist (zzz-DTTCompatibility-class), 12
- designElementTuples_assnlist, BioAssayTuple-method (zzz-DTTCompatibility-class), 12
- designProviders_assnreflist (zzz-DTTCompatibility-class), 12
- designProviders_assnreflist, ArrayDesign-method (zzz-DTTCompatibility-class), 12
- designProviders_assnreflist, PhysicalArrayDesign-method (zzz-DTTCompatibility-class), 12
- DTTCompatibility, 13
- DTTCompatibility-class (AffyCompatible-class), 1
- editor (zzz-DTTCompatibility-class), 12
- editor, BibliographicReference-method (zzz-DTTCompatibility-class), 12
- email (zzz-DTTCompatibility-class), 12
- directory (accessors), 6
- directory, NetAffxResource-method (NetAffxResource-class), 2
- distanceUnit (zzz-DTTCompatibility-class), 12
- distanceUnit, DistanceUnit_assn-method (zzz-DTTCompatibility-class), 12
- distanceUnit, Unit_assn-method (zzz-DTTCompatibility-class), 12
- distanceUnit_assn (zzz-DTTCompatibility-class), 12
- distanceUnit_assn, ArrayGroup-method (zzz-DTTCompatibility-class), 12
- distanceUnit_assn, FeatureGroup-method (zzz-DTTCompatibility-class), 12
- distanceUnit_assn, Fiducial-method (zzz-DTTCompatibility-class), 12
- distanceUnit_assn, Position-method (zzz-DTTCompatibility-class), 12
- distanceUnit_assn, PositionDelta-method (zzz-DTTCompatibility-class), 12
- distanceUnit_assn, Zone-method (zzz-DTTCompatibility-class), 12
- distanceUnit_assn, ZoneGroup-method (zzz-DTTCompatibility-class), 12
- distanceUnit_assn, ZoneLayout-method (zzz-DTTCompatibility-class), 12

email, Organization-method (zzz-DTTCompatibility-class), 12
 email, Person-method (zzz-DTTCompatibility-class), 12
 error (zzz-DTTCompatibility-class), 12
 error, ConfidenceIndicator_assnlist-method (zzz-DTTCompatibility-class), 12
 error, QuantitationType_assnlist-method (zzz-DTTCompatibility-class), 12
 error_ref (zzz-DTTCompatibility-class), 12
 error_ref, ConfidenceIndicators_assnreflist-method (zzz-DTTCompatibility-class), 12
 error_ref, QuantitationType_assnref-method (zzz-DTTCompatibility-class), 12
 error_ref, QuantitationTypes_assnreflist-method (zzz-DTTCompatibility-class), 12
 error_ref, SourcesQuantitationType_assnreflist-method (zzz-DTTCompatibility-class), 12
 error_ref, TargetQuantitationType_assnref-method (zzz-DTTCompatibility-class), 12
 expectedValue (zzz-DTTCompatibility-class), 12
 expectedValue, ConfidenceIndicator_assnlist-method (zzz-DTTCompatibility-class), 12
 expectedValue, QuantitationType_assnlist-method (zzz-DTTCompatibility-class), 12
 expectedValue_ref (zzz-DTTCompatibility-class), 12
 expectedValue_ref, ConfidenceIndicators_assnreflist-method (zzz-DTTCompatibility-class), 12
 expectedValue_ref, QuantitationType_assnref-method (zzz-DTTCompatibility-class), 12
 expectedValue_ref, QuantitationTypes_assnreflist-method (zzz-DTTCompatibility-class), 12
 (zzz-DTTCompatibility-class), 12
 expectedValue_ref, SourcesQuantitationType_assnreflist-method (zzz-DTTCompatibility-class), 12
 expectedValue_ref, TargetQuantitationType_assnreflist-method (zzz-DTTCompatibility-class), 12
 experiment
 experiment (zzz-DTTCompatibility-class), 12
 experiment, Experiment_assnlist-method (zzz-DTTCompatibility-class), 12
 experiment_assnlist (zzz-DTTCompatibility-class), 12
 experiment_assnlist, Experiment_package-method (zzz-DTTCompatibility-class), 12
 experiment_package
 experiment_package, MAGE-ML-method (zzz-DTTCompatibility-class), 12
 experimentalFactor
 experimentalFactor, ExperimentalFactors_assnlist-method (zzz-DTTCompatibility-class), 12
 experimentalFactor_ref (zzz-DTTCompatibility-class), 12
 experimentalFactor_ref, ExperimentalFactor_assnlist-method (zzz-DTTCompatibility-class), 12
 experimentalFactors_assnlist (zzz-DTTCompatibility-class), 12
 experimentalFactors_assnlist, ExperimentDesign (zzz-DTTCompatibility-class), 12
 experimentDesign
 experimentDesign, ExperimentDesigns_assnlist-method (zzz-DTTCompatibility-class), 12
 experimentDesigns_assnlist (zzz-DTTCompatibility-class), 12

12
 experimentDesigns_assnlist, Experiment-method(*zzz-DTTCompatibility-class*),
 (*zzz-DTTCompatibility-class*), 12
 12
 exportedFromDB
 (*zzz-DTTCompatibility-class*), 12
 12
 exportedFromDB, ExternalReference-method
 (*zzz-DTTCompatibility-class*), 12
 12
 exportedFromServer
 (*zzz-DTTCompatibility-class*), 12
 12
 exportedFromServer, ExternalReference-method
 (*zzz-DTTCompatibility-class*), 12
 12
 exportID
 (*zzz-DTTCompatibility-class*), 12
 12
 exportID, ExternalReference-method
 (*zzz-DTTCompatibility-class*), 12
 12
 exportName
 (*zzz-DTTCompatibility-class*), 12
 12
 exportName, ExternalReference-method
 (*zzz-DTTCompatibility-class*), 12
 12
 externalLIMS_assn
 (*zzz-DTTCompatibility-class*), 12
 12
 externalLIMS_assn, Compound-method
 (*zzz-DTTCompatibility-class*), 12
 12
 externalReference
 (*zzz-DTTCompatibility-class*), 12
 12
 externalReference, ExternalReference_assn-method
 (*zzz-DTTCompatibility-class*), 12
 12
 externalReference_assn
 (*zzz-DTTCompatibility-class*), 12
 12
 externalReference_assn, Description-method
 (*zzz-DTTCompatibility-class*), 12
 12
 factorValue
 (*zzz-DTTCompatibility-class*), 12
 12
 factorValue, FactorValues_assnlist-method
 (*zzz-DTTCompatibility-class*), 12
 12
 factorValue_ref
 (*zzz-DTTCompatibility-class*), 12
 12
 factorValue_ref, BioAssayFactorValues_assnrefl
 (*zzz-DTTCompatibility-class*), 12
 12
 factorValues_assnlist
 (*zzz-DTTCompatibility-class*), 12
 12
 factorValues_assnlist, ExperimentalFactor-metho
 (*zzz-DTTCompatibility-class*), 12
 12
 failed
 (*zzz-DTTCompatibility-class*), 12
 12
 failed, QuantitationType_assnlist-method
 (*zzz-DTTCompatibility-class*), 12
 12
 failed_ref
 (*zzz-DTTCompatibility-class*), 12
 12
 failed_ref, QuantitationType_assnref-method
 (*zzz-DTTCompatibility-class*), 12
 12
 failed_ref, QuantitationTypes_assnrefl
 (*zzz-DTTCompatibility-class*), 12
 12
 failed_ref, SourcesQuantitationType_assnrefl
 (*zzz-DTTCompatibility-class*), 12
 12
 failed_ref, TargetQuantitationType_assnref-meth
 (*zzz-DTTCompatibility-class*), 12
 12
 failTypes_assnlist
 (*zzz-DTTCompatibility-class*), 12
 12
 failTypes_assnlist, Reporter-method
 (*zzz-DTTCompatibility-class*), 12
 12
 fax(*zzz-DTTCompatibility-class*), 12
 12
 fax, Organization-method
 (*zzz-DTTCompatibility-class*), 12
 12
 fax, Person-method
 (*zzz-DTTCompatibility-class*), 12
 12
 feature
 (*zzz-DTTCompatibility-class*), 12
 12
 feature, DesignElement_assnlist-method
 (*zzz-DTTCompatibility-class*), 12

- 12
- feature, Features_assnlist-method
(zzz-DTTCompatibility-class),
12
- feature_assnref
(zzz-DTTCompatibility-class),
12
- feature_assnref, FeatureDefect-method
(zzz-DTTCompatibility-class),
12
- feature_assnref, FeatureInformation-method
(zzz-DTTCompatibility-class),
12
- feature_assnref, ManufactureLIMS-method
(zzz-DTTCompatibility-class),
12
- feature_assnref, ManufactureLIMSBiomaterial-method
(zzz-DTTCompatibility-class),
12
- feature_ref
(zzz-DTTCompatibility-class),
12
- feature_ref, ContainedFeatures_assnreflist-method
(zzz-DTTCompatibility-class),
12
- feature_ref, ControlFeatures_assnreflist-method
(zzz-DTTCompatibility-class),
12
- feature_ref, ControlledFeatures_assnreflist-method
(zzz-DTTCompatibility-class),
12
- feature_ref, DesignElement_assnref-method
(zzz-DTTCompatibility-class),
12
- feature_ref, Feature_assnref-method
(zzz-DTTCompatibility-class),
12
- featureDefect
(zzz-DTTCompatibility-class),
12
- featureDefect, FeatureDefects_assnlist-method
(zzz-DTTCompatibility-class),
12
- featureDefects_assnlist
(zzz-DTTCompatibility-class),
12
- featureDefects_assnlist, ArrayManufactureDevice_assnlist-method
(zzz-DTTCompatibility-class),
12
- featureDimension
(zzz-DTTCompatibility-class),
12
- featureDimension, DesignElementDimension_assnlist-method
(zzz-DTTCompatibility-class),
12
- featureDimension_ref
(zzz-DTTCompatibility-class),
12
- featureDimension_ref, DesignElementDimension_assnlist-method
(zzz-DTTCompatibility-class),
12
- featureExtraction
(zzz-DTTCompatibility-class),
12
- featureExtraction, FeatureExtraction_assnlist-method
(zzz-DTTCompatibility-class),
12
- featureExtraction_assnlist
(zzz-DTTCompatibility-class),
12
- featureExtraction_assnlist, MeasuredBioAssay-method
(zzz-DTTCompatibility-class),
12
- featureGroup
(zzz-DTTCompatibility-class),
12
- featureGroup, FeatureGroups_assnlist-method
(zzz-DTTCompatibility-class),
12
- featureGroup_ref
(zzz-DTTCompatibility-class),
12
- featureGroup_ref, FeatureGroup_assnref-method
(zzz-DTTCompatibility-class),
12
- featureGroups_assnlist
(zzz-DTTCompatibility-class),
12
- featureGroups_assnlist, ArrayDesign-method
(zzz-DTTCompatibility-class),
12
- featureGroups_assnlist, PhysicalArrayDesign-method
(zzz-DTTCompatibility-class),
12
- featureHeight
(zzz-DTTCompatibility-class),
12
- featureHeight, FeatureGroup-method
(zzz-DTTCompatibility-class),
12
- featureInformation
(zzz-DTTCompatibility-class),
12
- featureInformation, FeatureInformationSources_assnlist-method
(zzz-DTTCompatibility-class),
12

- (zzz-DTTCompatibility-class), 12
- featureInformationSources_assnlist (zzz-DTTCompatibility-class), 12
- featureInformationSources_assnlist, FeatureReporterMap-method (zzz-DTTCompatibility-class), 12
- featureLength (zzz-DTTCompatibility-class), 12
- featureLength, FeatureGroup-method (zzz-DTTCompatibility-class), 12
- featureLIMSS_assnlist (zzz-DTTCompatibility-class), 12
- featureLIMSS_assnlist, ArrayManufacture-method (zzz-DTTCompatibility-class), 12
- featureLocation (zzz-DTTCompatibility-class), 12
- featureLocation, FeatureLocation_assn-method (zzz-DTTCompatibility-class), 12
- featureLocation_assn (zzz-DTTCompatibility-class), 12
- featureLocation_assn, Feature-method (zzz-DTTCompatibility-class), 12
- featureReporterMap (zzz-DTTCompatibility-class), 12
- featureReporterMap, DesignElementMap_assnlist (zzz-DTTCompatibility-class), 12
- featureReporterMap, FeatureReporterMap_assnlist (zzz-DTTCompatibility-class), 12
- featureReporterMap_assnlist (zzz-DTTCompatibility-class), 12
- featureReporterMap_assnlist, DesignElement_map-method (zzz-DTTCompatibility-class), 12
- featureReporterMap_ref (zzz-DTTCompatibility-class), 12
- featureReporterMap_ref, DesignElementMaps_assnlist (zzz-DTTCompatibility-class), 12
- featureReporterMap_ref, FeatureReporterMaps_assnlist (zzz-DTTCompatibility-class), 12
- featureReporterMaps_assnreflist (zzz-DTTCompatibility-class), 12
- featureReporterMaps_assnreflist, Reporter-method (zzz-DTTCompatibility-class), 12
- features_assnlist (zzz-DTTCompatibility-class), 12
- features_assnlist, FeatureGroup-method (zzz-DTTCompatibility-class), 12
- featureShape_assn (zzz-DTTCompatibility-class), 12
- featureShape_assn, FeatureGroup-method (zzz-DTTCompatibility-class), 12
- featureWidth (zzz-DTTCompatibility-class), 12
- featureWidth, FeatureGroup-method (zzz-DTTCompatibility-class), 12
- fiducial (zzz-DTTCompatibility-class), 12
- fiducial, Fiducials_assnlist-method (zzz-DTTCompatibility-class), 12
- fiducials_assnlist (zzz-DTTCompatibility-class), 12
- fiducials_assnlist, ArrayGroup-method (zzz-DTTCompatibility-class), 12
- fiducialType_assn (zzz-DTTCompatibility-class), 12
- fiducialType_assn, Fiducial-method (zzz-DTTCompatibility-class), 12
- filenameURI (zzz-DTTCompatibility-class), 12
- filenameURI, DataExternal-method (zzz-DTTCompatibility-class), 12

- firstName
 (*zzz-DTTCompatibility-class*),
 12
- firstName, Person-method
 (*zzz-DTTCompatibility-class*),
 12
- format_assn
 (*zzz-DTTCompatibility-class*),
 12
- format_assn, Image-method
 (*zzz-DTTCompatibility-class*),
 12
- guid
 (*zzz-ARRCompatibility-class*),
 11
- guid, ArraySetFile-method
 (*zzz-ARRCompatibility-class*),
 11
- guid, PhysicalArray-method
 (*zzz-ARRCompatibility-class*),
 11
- guid, TemplateFile-method
 (*zzz-ARRCompatibility-class*),
 11
- hardware
 (*zzz-DTTCompatibility-class*),
 12
- hardware, Hardware_assnlist-method
 (*zzz-DTTCompatibility-class*),
 12
- hardware_assnlist
 (*zzz-DTTCompatibility-class*),
 12
- hardware_assnlist, Protocol_package-method
 (*zzz-DTTCompatibility-class*),
 12
- hardware_assnref
 (*zzz-DTTCompatibility-class*),
 12
- hardware_assnref, HardwareApplication-method
 (*zzz-DTTCompatibility-class*),
 12
- hardware_assnref, Software-method
 (*zzz-DTTCompatibility-class*),
 12
- hardware_ref
 (*zzz-DTTCompatibility-class*),
 12
- hardware_ref, Hardware_assnref-method
 (*zzz-DTTCompatibility-class*),
 12
- hardware_ref, Hardware_assnreflist-method
 (*zzz-DTTCompatibility-class*),
 12
- hardwareApplication
 (*zzz-DTTCompatibility-class*),
 12
- hardwareApplication, HardwareApplications_assnlist
 (*zzz-DTTCompatibility-class*),
 12
- hardwareApplications_assnlist
 (*zzz-DTTCompatibility-class*),
 12
- hardwareApplications_assnlist, ProtocolApplication
 (*zzz-DTTCompatibility-class*),
 12
- hardwareManufacturers_assnreflist
 (*zzz-DTTCompatibility-class*),
 12
- hardwareManufacturers_assnreflist, Hardware-man
 (*zzz-DTTCompatibility-class*),
 12
- hardwarewares_assnreflist
 (*zzz-DTTCompatibility-class*),
 12
- hardwarewares_assnreflist, Protocol-method
 (*zzz-DTTCompatibility-class*),
 12
- higherLevelAnalysis_package
 (*zzz-DTTCompatibility-class*),
 12
- higherLevelAnalysis_package, MAGE-ML-method
 (*zzz-DTTCompatibility-class*),
 12
- hybridization
 (*zzz-DTTCompatibility-class*),
 12
- hybridization, BioAssayCreation_assn-method
 (*zzz-DTTCompatibility-class*),
 12
- identifier
 (*zzz-DTTCompatibility-class*),
 12
- identifier, Array-method
 (*zzz-DTTCompatibility-class*),
 12
- identifier, Array_ref-method
 (*zzz-DTTCompatibility-class*),
 12
- identifier, ArrayDesign-method
 (*zzz-DTTCompatibility-class*),
 12
- identifier, ArrayDesign_ref-method

- (*zzz-DTTCompatibility-class*),
12
- identifier, ArrayGroup-method
(*zzz-DTTCompatibility-class*),
12
- identifier, ArrayGroup_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier, ArrayManufacture-method
(*zzz-DTTCompatibility-class*),
12
- identifier, ArrayManufacture_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier, BioAssay_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier, BioAssayCreation-method
(*zzz-DTTCompatibility-class*),
12
- identifier, BioAssayData_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier, BioAssayDataCluster-method
(*zzz-DTTCompatibility-class*),
12
- identifier, BioAssayDataCluster_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier, BioAssayDimension-method
(*zzz-DTTCompatibility-class*),
12
- identifier, BioAssayDimension_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier, BioAssayMap-method
(*zzz-DTTCompatibility-class*),
12
- identifier, BioAssayMap_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier, BioAssayTreatment-method
(*zzz-DTTCompatibility-class*),
12
- identifier, BioMaterial_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier, BioSample-method
(*zzz-DTTCompatibility-class*),
12
- identifier, BioSample_ref-method
(*zzz-DTTCompatibility-class*),
12
- 12
- identifier, BioSequence-method
(*zzz-DTTCompatibility-class*),
12
- identifier, BioSequence_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier, BioSource-method
(*zzz-DTTCompatibility-class*),
12
- identifier, BioSource_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier, Channel-method
(*zzz-DTTCompatibility-class*),
12
- identifier, Channel_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier, CompositeCompositeMap-method
(*zzz-DTTCompatibility-class*),
12
- identifier, CompositeCompositeMap_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier, CompositeGroup-method
(*zzz-DTTCompatibility-class*),
12
- identifier, CompositeGroup_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier, CompositeSequence-method
(*zzz-DTTCompatibility-class*),
12
- identifier, CompositeSequence_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier, CompositeSequenceDimension-method
(*zzz-DTTCompatibility-class*),
12
- identifier, CompositeSequenceDimension_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier, Compound-method
(*zzz-DTTCompatibility-class*),
12
- identifier, Compound_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier, ConfidenceIndicator_ref-method
(*zzz-DTTCompatibility-class*),
12

- identifier, Contact_ref-method
(zzz-DTTCompatibility-class),
12
- identifier, Database-method
(zzz-DTTCompatibility-class),
12
- identifier, Database_ref-method
(zzz-DTTCompatibility-class),
12
- identifier, DerivedBioAssay-method
(zzz-DTTCompatibility-class),
12
- identifier, DerivedBioAssay_ref-method
(zzz-DTTCompatibility-class),
12
- identifier, DerivedBioAssayData-method
(zzz-DTTCompatibility-class),
12
- identifier, DerivedBioAssayData_ref-method
(zzz-DTTCompatibility-class),
12
- identifier, DerivedSignal-method
(zzz-DTTCompatibility-class),
12
- identifier, DerivedSignal_ref-method
(zzz-DTTCompatibility-class),
12
- identifier, DesignElement_ref-method
(zzz-DTTCompatibility-class),
12
- identifier, DesignElementDimension_ref-method
(zzz-DTTCompatibility-class),
12
- identifier, DesignElementMap_ref-method
(zzz-DTTCompatibility-class),
12
- identifier, Error-method
(zzz-DTTCompatibility-class),
12
- identifier, Error_ref-method
(zzz-DTTCompatibility-class),
12
- identifier, ExpectedValue-method
(zzz-DTTCompatibility-class),
12
- identifier, ExpectedValue_ref-method
(zzz-DTTCompatibility-class),
12
- identifier, Experiment-method
(zzz-DTTCompatibility-class),
12
- identifier, ExperimentalFactor-method
(zzz-DTTCompatibility-class),
12
- identifier, ExperimentalFactor_ref-method
(zzz-DTTCompatibility-class),
12
- identifier, FactorValue-method
(zzz-DTTCompatibility-class),
12
- identifier, FactorValue_ref-method
(zzz-DTTCompatibility-class),
12
- identifier, Failed-method
(zzz-DTTCompatibility-class),
12
- identifier, Failed_ref-method
(zzz-DTTCompatibility-class),
12
- identifier, Feature-method
(zzz-DTTCompatibility-class),
12
- identifier, Feature_ref-method
(zzz-DTTCompatibility-class),
12
- identifier, FeatureDimension-method
(zzz-DTTCompatibility-class),
12
- identifier, FeatureDimension_ref-method
(zzz-DTTCompatibility-class),
12
- identifier, FeatureExtraction-method
(zzz-DTTCompatibility-class),
12
- identifier, FeatureGroup-method
(zzz-DTTCompatibility-class),
12
- identifier, FeatureGroup_ref-method
(zzz-DTTCompatibility-class),
12
- identifier, FeatureReporterMap-method
(zzz-DTTCompatibility-class),
12
- identifier, FeatureReporterMap_ref-method
(zzz-DTTCompatibility-class),
12
- identifier, Hardware-method
(zzz-DTTCompatibility-class),
12
- identifier, Hardware_ref-method
(zzz-DTTCompatibility-class),
12
- identifier, Hybridization-method
(zzz-DTTCompatibility-class),
12

- 12
- identifier, Image-method
(*zzz-DTTCompatibility-class*),
12
- identifier, Image_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier, ImageAcquisition-method
(*zzz-DTTCompatibility-class*),
12
- identifier, LabeledExtract-method
(*zzz-DTTCompatibility-class*),
12
- identifier, LabeledExtract_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier, MAGE-ML-method
(*zzz-DTTCompatibility-class*),
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- identifier, MeasuredBioAssay-method
(*zzz-DTTCompatibility-class*),
12
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(*zzz-DTTCompatibility-class*),
12
- identifier, MeasuredBioAssayData-method
(*zzz-DTTCompatibility-class*),
12
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(*zzz-DTTCompatibility-class*),
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- identifier, MeasuredSignal-method
(*zzz-DTTCompatibility-class*),
12
- identifier, MeasuredSignal_ref-method
(*zzz-DTTCompatibility-class*),
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- identifier, Organization-method
(*zzz-DTTCompatibility-class*),
12
- identifier, Organization_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier, Parameter-method
(*zzz-DTTCompatibility-class*),
12
- identifier, Parameter_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier, Person-method
(*zzz-DTTCompatibility-class*),
12
- identifier, Person_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier, PhysicalArrayDesign-method
(*zzz-DTTCompatibility-class*),
12
- identifier, PhysicalArrayDesign_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier, PhysicalBioAssay-method
(*zzz-DTTCompatibility-class*),
12
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(*zzz-DTTCompatibility-class*),
12
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(*zzz-DTTCompatibility-class*),
12
- identifier, PresentAbsent_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier, Protocol-method
(*zzz-DTTCompatibility-class*),
12
- identifier, Protocol_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier, PValue-method
(*zzz-DTTCompatibility-class*),
12
- identifier, PValue_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier, QuantitationType_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier, QuantitationTypeDimension-method
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- identifier, QuantitationTypeDimension_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier, QuantitationTypeMap-method
(*zzz-DTTCompatibility-class*),
12
- identifier, QuantitationTypeMap_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier, Ratio-method
(*zzz-DTTCompatibility-class*),
12
- identifier, Ratio_ref-method

- (zzz-DTTCompatibility-class),
12
- identifier, Reporter-method
(zzz-DTTCompatibility-class),
12
- identifier, Reporter_ref-method
(zzz-DTTCompatibility-class),
12
- identifier, ReporterCompositeMap-method
(zzz-DTTCompatibility-class),
12
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(zzz-DTTCompatibility-class),
12
- identifier, ReporterDimension-method
(zzz-DTTCompatibility-class),
12
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(zzz-DTTCompatibility-class),
12
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(zzz-DTTCompatibility-class),
12
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(zzz-DTTCompatibility-class),
12
- identifier, Security-method
(zzz-DTTCompatibility-class),
12
- identifier, Security_ref-method
(zzz-DTTCompatibility-class),
12
- identifier, SecurityGroup-method
(zzz-DTTCompatibility-class),
12
- identifier, SecurityGroup_ref-method
(zzz-DTTCompatibility-class),
12
- identifier, Software-method
(zzz-DTTCompatibility-class),
12
- identifier, Software_ref-method
(zzz-DTTCompatibility-class),
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- identifier, SpecializedQuantitationType-method
(zzz-DTTCompatibility-class),
12
- identifier, SpecializedQuantitationType_ref-method
(zzz-DTTCompatibility-class),
12
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(zzz-DTTCompatibility-class),
12
- identifier, Transformation-method
(zzz-DTTCompatibility-class),
12
- identifier, Treatment-method
(zzz-DTTCompatibility-class),
12
- identifier, Zone-method
(zzz-DTTCompatibility-class),
12
- identifier, Zone_ref-method
(zzz-DTTCompatibility-class),
12
- identifierLIMS_assn
(zzz-DTTCompatibility-class),
12
- identifierLIMS_assn, ManufactureLIMS-method
(zzz-DTTCompatibility-class),
12
- identifierLIMS_assn, ManufactureLIMSBiomaterial-method
(zzz-DTTCompatibility-class),
12
- image_ref
(zzz-DTTCompatibility-class),
12
- image_ref, Images_assnreflist-method
(zzz-DTTCompatibility-class),
12
- imageAcquisition
(zzz-DTTCompatibility-class),
12
- imageAcquisition, BioAssayTreatments_assnreflist-method
(zzz-DTTCompatibility-class),
12
- images_assnreflist
(zzz-DTTCompatibility-class),
12
- images_assnreflist, ImageAcquisition-method
(zzz-DTTCompatibility-class),
12
- immobilizedCharacteristics_assnreflist
(zzz-DTTCompatibility-class),
12
- immobilizedCharacteristics_assnreflist, Reporter-method
(zzz-DTTCompatibility-class),
12
- information_assnref
(zzz-DTTCompatibility-class),
12
- information_assnref, Array-method
(zzz-DTTCompatibility-class),
12

- Internal-class, 11
- isApproximateLength
(*zzz-DTTCompatibility-class*),
12
- isApproximateLength, BioSequence-method
(*zzz-DTTCompatibility-class*),
12
- isBackground
(*zzz-DTTCompatibility-class*),
12
- isBackground, DerivedSignal-method
(*zzz-DTTCompatibility-class*),
12
- isBackground, Error-method
(*zzz-DTTCompatibility-class*),
12
- isBackground, ExpectedValue-method
(*zzz-DTTCompatibility-class*),
12
- isBackground, Failed-method
(*zzz-DTTCompatibility-class*),
12
- isBackground, MeasuredSignal-method
(*zzz-DTTCompatibility-class*),
12
- isBackground, PresentAbsent-method
(*zzz-DTTCompatibility-class*),
12
- isBackground, PValue-method
(*zzz-DTTCompatibility-class*),
12
- isBackground, Ratio-method
(*zzz-DTTCompatibility-class*),
12
- isBackground, SpecializedQuantitationType-method
(*zzz-DTTCompatibility-class*),
12
- isCircular
(*zzz-DTTCompatibility-class*),
12
- isCircular, BioSequence-method
(*zzz-DTTCompatibility-class*),
12
- isSolvent
(*zzz-DTTCompatibility-class*),
12
- isSolvent, Compound-method
(*zzz-DTTCompatibility-class*),
12
- issue
(*zzz-DTTCompatibility-class*),
12
- issue, BibliographicReference-method
(*zzz-DTTCompatibility-class*),
12
- kindCV
(*zzz-DTTCompatibility-class*),
12
- kindCV, Measurement-method
(*zzz-DTTCompatibility-class*),
12
- labeledExtract
(*zzz-DTTCompatibility-class*),
12
- labeledExtract, BioMaterial_assnlist-method
(*zzz-DTTCompatibility-class*),
12
- labeledExtract_ref
(*zzz-DTTCompatibility-class*),
12
- labeledExtract_ref, BioMaterial_assnref-method
(*zzz-DTTCompatibility-class*),
12
- labels_assnreflist
(*zzz-DTTCompatibility-class*),
12
- labels_assnreflist, Channel-method
(*zzz-DTTCompatibility-class*),
12
- labels_assnreflist, LabeledExtract-method
(*zzz-DTTCompatibility-class*),
12
- lastName
(*zzz-DTTCompatibility-class*),
12
- lastName, Person-method
(*zzz-DTTCompatibility-class*),
12
- length, ArrayGroup-method
(*zzz-DTTCompatibility-class*),
12
- length, BioSequence-method
(*zzz-DTTCompatibility-class*),
12
- libraryPackageName
(*zzz-ARRCompatibility-class*),
11
- libraryPackageName, PhysicalArray-method
(*zzz-ARRCompatibility-class*),
11
- list, 11
- lowerRightX
(*zzz-DTTCompatibility-class*),

- 12
 lowerRightX, Zone-method
 (*zzz-DTTCompatibility-class*),
 12
 lowerRightY
 (*zzz-DTTCompatibility-class*),
 12
 lowerRightY, Zone-method
 (*zzz-DTTCompatibility-class*),
 12
 make
 (*zzz-DTTCompatibility-class*),
 12
 make, Hardware-method
 (*zzz-DTTCompatibility-class*),
 12
 manufactureLIMS
 (*zzz-DTTCompatibility-class*),
 12
 manufactureLIMS, FeatureLIMSS_assnlist-method
 (*zzz-DTTCompatibility-class*),
 12
 manufactureLIMSBiomaterial
 (*zzz-DTTCompatibility-class*),
 12
 manufactureLIMSBiomaterial, FeatureLIMSS_assnlist-method
 (*zzz-DTTCompatibility-class*),
 12
 manufacturingDate
 (*zzz-DTTCompatibility-class*),
 12
 manufacturingDate, ArrayManufacture-method
 (*zzz-DTTCompatibility-class*),
 12
 massUnit
 (*zzz-DTTCompatibility-class*),
 12
 massUnit, Unit_assn-method
 (*zzz-DTTCompatibility-class*),
 12
 masterFileGUID
 (*zzz-ARRCompatibility-class*),
 11
 masterFileGUID, PhysicalArray-method
 (*zzz-ARRCompatibility-class*),
 11
 masterFileName
 (*zzz-ARRCompatibility-class*),
 11
 masterFileName, PhysicalArray-method
 (*zzz-ARRCompatibility-class*),
 11
 materialType_assn
 (*zzz-DTTCompatibility-class*),
 12
 materialType_assn, BioSample-method
 (*zzz-DTTCompatibility-class*),
 12
 materialType_assn, BioSource-method
 (*zzz-DTTCompatibility-class*),
 12
 materialType_assn, LabeledExtract-method
 (*zzz-DTTCompatibility-class*),
 12
 measuredBioAssay
 (*zzz-DTTCompatibility-class*),
 12
 measuredBioAssay, BioAssay_assnlist-method
 (*zzz-DTTCompatibility-class*),
 12
 measuredBioAssay, MeasuredBioAssay_assnlist-method
 (*zzz-DTTCompatibility-class*),
 12
 measuredBioAssay_ref
 (*zzz-DTTCompatibility-class*),
 12
 measuredBioAssay_ref, BioAssay_assnref-method
 (*zzz-DTTCompatibility-class*),
 12
 measuredBioAssay_ref, BioAssays_assnreflist-method
 (*zzz-DTTCompatibility-class*),
 12
 measuredBioAssay_ref, MeasuredBioAssayTarget_assnreflist-method
 (*zzz-DTTCompatibility-class*),
 12
 measuredBioAssay_ref, SourceBioAssays_assnreflist-method
 (*zzz-DTTCompatibility-class*),
 12
 measuredBioAssay_ref, TopLevelBioAssays_assnreflist-method
 (*zzz-DTTCompatibility-class*),
 12
 measuredBioAssayData
 (*zzz-DTTCompatibility-class*),
 12
 measuredBioAssayData, BioAssayData_assnlist-method
 (*zzz-DTTCompatibility-class*),
 12
 measuredBioAssayData, MeasuredBioAssayData_assnlist-method
 (*zzz-DTTCompatibility-class*),
 12
 measuredBioAssayData_assnreflist
 (*zzz-DTTCompatibility-class*),
 12
 measuredBioAssayData_assnreflist, MeasuredBioAssayData_assnreflist-method
 (*zzz-DTTCompatibility-class*),
 12

	(zzz-DTTCompatibility-class),	12	
measuredBioAssayData_ref	(zzz-DTTCompatibility-class),	12	measurement_assn, BioMaterialMeasurement-method
measuredBioAssayData_ref, BioAssayData_assnref	(zzz-DTTCompatibility-class),	12	measurement_assn, CompoundMeasurement-method
measuredBioAssayData_ref, BioAssayDataSources	(zzz-DTTCompatibility-class),	12	measurement_assn, FactorValue-method
measuredBioAssayData_ref, ClusterBioAssayData	(zzz-DTTCompatibility-class),	12	measurement_package
measuredBioAssayData_ref, MeasuredBioAssayData	(zzz-DTTCompatibility-class),	12	measurement_package, MAGE-ML-method
measuredSignal	(zzz-DTTCompatibility-class),	12	measurement_package, MAGE-ML-method
measuredSignal, QuantitationType_assnref	(zzz-DTTCompatibility-class),	12	mediaCol
measuredSignal_ref	(zzz-DTTCompatibility-class),	12	(zzz-ARRCompatibility-class),
measuredSignal_ref, QuantitationType_assnref	(zzz-DTTCompatibility-class),	12	mediaCol, PhysicalArray-method
measuredSignal_ref, QuantitationTypes_assnref	(zzz-DTTCompatibility-class),	12	(zzz-ARRCompatibility-class),
measuredSignal_ref, SourcesQuantitationType_assnref	(zzz-DTTCompatibility-class),	12	mediaFileGUID
measuredSignal_ref, TargetQuantitationType_assnref	(zzz-DTTCompatibility-class),	12	(zzz-ARRCompatibility-class),
measurement	(zzz-DTTCompatibility-class),	12	mediaFileGUID, PhysicalArray-method
measurement, ActionMeasurement_assn-method	(zzz-DTTCompatibility-class),	12	(zzz-ARRCompatibility-class),
measurement, DefaultValue_assn-method	(zzz-DTTCompatibility-class),	12	mediaFileName
measurement, Measurement_assn-method	(zzz-DTTCompatibility-class),	12	(zzz-ARRCompatibility-class),
measurement_assn	(zzz-DTTCompatibility-class),	12	mediaFileName, PhysicalArray-method
			mediaRow
			(zzz-ARRCompatibility-class),
			mediaRow, PhysicalArray-method
			mediaType
			(zzz-ARRCompatibility-class),
			mediaType, PhysicalArray-method
			members_assnreflist
			(zzz-DTTCompatibility-class),
			12
			members_assnreflist, SecurityGroup-method
			(zzz-DTTCompatibility-class),
			12

midInitials	(zzz-DTTCompatibility-class), 12	(zzz-DTTCompatibility-class), 12
midInitials, Person-method	(zzz-DTTCompatibility-class), 12	name, ArrayManufacture-method (zzz-DTTCompatibility-class), 12
mismatchInformation	(zzz-DTTCompatibility-class), 12	name, ArrayManufacture_ref-method (zzz-DTTCompatibility-class), 12
mismatchInformation, MismatchInformation_assnlist-method	(zzz-DTTCompatibility-class), 12	name, BioAssay_ref-method (zzz-DTTCompatibility-class), 12
mismatchInformation_assnlist	(zzz-DTTCompatibility-class), 12	name, BioAssayCreation-method (zzz-DTTCompatibility-class), 12
mismatchInformation_assnlist, CompositePosition-method	(zzz-DTTCompatibility-class), 12	name, BioAssayData_ref-method (zzz-DTTCompatibility-class), 12
mismatchInformation_assnlist, FeatureInformation-method	(zzz-DTTCompatibility-class), 12	name, BioAssayDataCluster-method (zzz-DTTCompatibility-class), 12
mismatchInformation_assnlist, ReporterPosition-method	(zzz-DTTCompatibility-class), 12	name, BioAssayDataCluster_ref-method (zzz-DTTCompatibility-class), 12
model	(zzz-DTTCompatibility-class), 12	name, BioAssayDimension-method (zzz-DTTCompatibility-class), 12
model, Hardware-method	(zzz-DTTCompatibility-class), 12	name, BioAssayDimension_ref-method (zzz-DTTCompatibility-class), 12
name	(zzz-DTTCompatibility-class), 12	name, BioAssayMap-method (zzz-DTTCompatibility-class), 12
name, Array-method	(zzz-DTTCompatibility-class), 12	name, BioAssayMap_ref-method (zzz-DTTCompatibility-class), 12
name, Array_ref-method	(zzz-DTTCompatibility-class), 12	name, BioAssayTreatment-method (zzz-DTTCompatibility-class), 12
name, ArrayAttribute-method	(zzz-ARRCompatibility-class), 11	name, BioMaterial_ref-method (zzz-DTTCompatibility-class), 12
name, ArrayDesign-method	(zzz-DTTCompatibility-class), 12	name, BioSample-method (zzz-DTTCompatibility-class), 12
name, ArrayDesign_ref-method	(zzz-DTTCompatibility-class), 12	name, BioSample_ref-method (zzz-DTTCompatibility-class), 12
name, ArrayGroup-method	(zzz-DTTCompatibility-class), 12	name, BioSequence-method (zzz-DTTCompatibility-class), 12
name, ArrayGroup_ref-method		name, BioSequence_ref-method (zzz-DTTCompatibility-class),

- 12
- name, BioSource-method
(*zzz-DTTCompatibility-class*),
12
- name, BioSource_ref-method
(*zzz-DTTCompatibility-class*),
12
- name, Channel-method
(*zzz-DTTCompatibility-class*),
12
- name, Channel_ref-method
(*zzz-DTTCompatibility-class*),
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- name, CompositeCompositeMap-method
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12
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(*zzz-DTTCompatibility-class*),
12
- name, CompositeGroup-method
(*zzz-DTTCompatibility-class*),
12
- name, CompositeGroup_ref-method
(*zzz-DTTCompatibility-class*),
12
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(*zzz-DTTCompatibility-class*),
12
- name, CompositeSequenceDimension-method
(*zzz-DTTCompatibility-class*),
12
- name, CompositeSequenceDimension_ref-method
(*zzz-DTTCompatibility-class*),
12
- name, Compound-method
(*zzz-DTTCompatibility-class*),
12
- name, Compound_ref-method
(*zzz-DTTCompatibility-class*),
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- name, ConfidenceIndicator_ref-method
(*zzz-DTTCompatibility-class*),
12
- name, Contact_ref-method
(*zzz-DTTCompatibility-class*),
12
- name, Database-method
(*zzz-DTTCompatibility-class*),
12
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(*zzz-DTTCompatibility-class*),
12
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(*zzz-DTTCompatibility-class*),
12
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(*zzz-DTTCompatibility-class*),
12
- name, DerivedBioAssayData-method
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12
- name, DerivedBioAssayData_ref-method
(*zzz-DTTCompatibility-class*),
12
- name, DerivedSignal-method
(*zzz-DTTCompatibility-class*),
12
- name, DerivedSignal_ref-method
(*zzz-DTTCompatibility-class*),
12
- name, DesignElement_ref-method
(*zzz-DTTCompatibility-class*),
12
- name, DesignElementDimension_ref-method
(*zzz-DTTCompatibility-class*),
12
- name, DesignElementMap_ref-method
(*zzz-DTTCompatibility-class*),
12
- name, Error-method
(*zzz-DTTCompatibility-class*),
12
- name, Error_ref-method
(*zzz-DTTCompatibility-class*),
12
- name, ExpectedValue-method
(*zzz-DTTCompatibility-class*),
12
- name, ExpectedValue_ref-method
(*zzz-DTTCompatibility-class*),
12
- name, Experiment-method
(*zzz-DTTCompatibility-class*),
12
- name, ExperimentalFactor-method
(*zzz-DTTCompatibility-class*),
12
- name, ExperimentalFactor_ref-method
(*zzz-DTTCompatibility-class*),
12
- name, FactorValue-method

- (*zzz-DTTCompatibility-class*),
12
- name, FactorValue_ref-method
(*zzz-DTTCompatibility-class*),
12
- name, Failed-method
(*zzz-DTTCompatibility-class*),
12
- name, Failed_ref-method
(*zzz-DTTCompatibility-class*),
12
- name, Feature-method
(*zzz-DTTCompatibility-class*),
12
- name, Feature_ref-method
(*zzz-DTTCompatibility-class*),
12
- name, FeatureDimension-method
(*zzz-DTTCompatibility-class*),
12
- name, FeatureDimension_ref-method
(*zzz-DTTCompatibility-class*),
12
- name, FeatureExtraction-method
(*zzz-DTTCompatibility-class*),
12
- name, FeatureGroup-method
(*zzz-DTTCompatibility-class*),
12
- name, FeatureGroup_ref-method
(*zzz-DTTCompatibility-class*),
12
- name, FeatureReporterMap-method
(*zzz-DTTCompatibility-class*),
12
- name, FeatureReporterMap_ref-method
(*zzz-DTTCompatibility-class*),
12
- name, Hardware-method
(*zzz-DTTCompatibility-class*),
12
- name, Hardware_ref-method
(*zzz-DTTCompatibility-class*),
12
- name, Hybridization-method
(*zzz-DTTCompatibility-class*),
12
- name, Image-method
(*zzz-DTTCompatibility-class*),
12
- name, Image_ref-method
(*zzz-DTTCompatibility-class*),
12
- name, ImageAcquisition-method
(*zzz-DTTCompatibility-class*),
12
- name, LabeledExtract-method
(*zzz-DTTCompatibility-class*),
12
- name, LabeledExtract_ref-method
(*zzz-DTTCompatibility-class*),
12
- name, MAGE-ML-method
(*zzz-DTTCompatibility-class*),
12
- name, MeasuredBioAssay-method
(*zzz-DTTCompatibility-class*),
12
- name, MeasuredBioAssay_ref-method
(*zzz-DTTCompatibility-class*),
12
- name, MeasuredBioAssayData-method
(*zzz-DTTCompatibility-class*),
12
- name, MeasuredBioAssayData_ref-method
(*zzz-DTTCompatibility-class*),
12
- name, MeasuredSignal-method
(*zzz-DTTCompatibility-class*),
12
- name, MeasuredSignal_ref-method
(*zzz-DTTCompatibility-class*),
12
- name, NameValueType-method
(*zzz-DTTCompatibility-class*),
12
- name, NodeValue-method
(*zzz-DTTCompatibility-class*),
12
- name, Organization-method
(*zzz-DTTCompatibility-class*),
12
- name, Organization_ref-method
(*zzz-DTTCompatibility-class*),
12
- name, Parameter-method
(*zzz-DTTCompatibility-class*),
12
- name, Parameter_ref-method
(*zzz-DTTCompatibility-class*),
12
- name, Person-method
(*zzz-DTTCompatibility-class*),
12

name, Person_ref-method (zzz-DTTCompatibility-class),
 12
 name, PhysicalArrayDesign-method (zzz-DTTCompatibility-class),
 12
 name, PhysicalArrayDesign_ref-method (zzz-DTTCompatibility-class),
 12
 name, PhysicalBioAssay-method (zzz-DTTCompatibility-class),
 12
 name, PhysicalBioAssay_ref-method (zzz-DTTCompatibility-class),
 12
 name, PresentAbsent-method (zzz-DTTCompatibility-class),
 12
 name, PresentAbsent_ref-method (zzz-DTTCompatibility-class),
 12
 name, Protocol-method (zzz-DTTCompatibility-class),
 12
 name, Protocol_ref-method (zzz-DTTCompatibility-class),
 12
 name, PValue-method (zzz-DTTCompatibility-class),
 12
 name, PValue_ref-method (zzz-DTTCompatibility-class),
 12
 name, QuantitationType_ref-method (zzz-DTTCompatibility-class),
 12
 name, QuantitationTypeDimension-method (zzz-DTTCompatibility-class),
 12
 name, QuantitationTypeDimension_ref-method (zzz-DTTCompatibility-class),
 12
 name, QuantitationTypeMap-method (zzz-DTTCompatibility-class),
 12
 name, QuantitationTypeMap_ref-method (zzz-DTTCompatibility-class),
 12
 name, Ratio-method (zzz-DTTCompatibility-class),
 12
 name, Ratio_ref-method (zzz-DTTCompatibility-class),
 12
 name, Reporter-method (zzz-DTTCompatibility-class),
 12
 name, Reporter_ref-method (zzz-DTTCompatibility-class),
 12
 name, ReporterCompositeMap-method (zzz-DTTCompatibility-class),
 12
 name, ReporterCompositeMap_ref-method (zzz-DTTCompatibility-class),
 12
 name, ReporterDimension-method (zzz-DTTCompatibility-class),
 12
 name, ReporterDimension_ref-method (zzz-DTTCompatibility-class),
 12
 name, ReporterGroup-method (zzz-DTTCompatibility-class),
 12
 name, ReporterGroup_ref-method (zzz-DTTCompatibility-class),
 12
 name, Security-method (zzz-DTTCompatibility-class),
 12
 name, Security_ref-method (zzz-DTTCompatibility-class),
 12
 name, SecurityGroup-method (zzz-DTTCompatibility-class),
 12
 name, SecurityGroup_ref-method (zzz-DTTCompatibility-class),
 12
 name, Software-method (zzz-DTTCompatibility-class),
 12
 name, Software_ref-method (zzz-DTTCompatibility-class),
 12
 name, SpecializedQuantitationType-method (zzz-DTTCompatibility-class),
 12
 name, SpecializedQuantitationType_ref-method (zzz-DTTCompatibility-class),
 12
 name, StandardQuantitationType_ref-method (zzz-DTTCompatibility-class),
 12

- 12
- name, Transformation-method
(*zzz-DTTCompatibility-class*),
12
- name, Treatment-method
(*zzz-DTTCompatibility-class*),
12
- name, UserAttribute-method
(*zzz-ARRCompatibility-class*),
11
- name, Zone-method
(*zzz-DTTCompatibility-class*),
12
- name, Zone_ref-method
(*zzz-DTTCompatibility-class*),
12
- names, AffxArray-method
(*zzz-NetAffxCompatibility-class*),
13
- names, NetAffxResource-method
(*NetAffxResource-class*), 2
- namespace
(*zzz-ARRCompatibility-class*),
11
- namespace, UserAttribute-method
(*zzz-ARRCompatibility-class*),
11
- nameValueType
(*zzz-DTTCompatibility-class*),
12
- nameValueType, PropertySets_assnlist-method
(*zzz-DTTCompatibility-class*),
12
- nameValueType, QualityControlStatistics_assnlist-method
(*zzz-DTTCompatibility-class*),
12
- nameValueType, SummaryStatistics_assnlist-method
(*zzz-DTTCompatibility-class*),
12
- NetAffxCompatibility, 3, 14
- NetAffxCompatibility-class
(*AffyCompatible-class*), 1
- NetAffxResource, 5, 5, 6, 8, 14
- NetAffxResource-accessors
(*accessors*), 6
- NetAffxResource-class, 2
- newSequence
(*zzz-DTTCompatibility-class*),
12
- newSequence, MismatchInformation-method
(*zzz-DTTCompatibility-class*),
12
- node
(*zzz-DTTCompatibility-class*),
12
- node, Nodes_assnlist-method
(*zzz-DTTCompatibility-class*),
12
- nodeContents
(*zzz-DTTCompatibility-class*),
12
- nodeContents, NodeContents_assnlist-method
(*zzz-DTTCompatibility-class*),
12
- nodeContents_assnlist
(*zzz-DTTCompatibility-class*),
12
- nodeContents_assnlist, Node-method
(*zzz-DTTCompatibility-class*),
12
- nodes_assnlist
(*zzz-DTTCompatibility-class*),
12
- nodes_assnlist, BioAssayDataCluster-method
(*zzz-DTTCompatibility-class*),
12
- nodes_assnlist, Node-method
(*zzz-DTTCompatibility-class*),
12
- nodeValue
(*zzz-DTTCompatibility-class*),
12
- nodeValue, NodeValue_assnlist-method
(*zzz-DTTCompatibility-class*),
12
- nodes_value_assnlist
(*zzz-DTTCompatibility-class*),
12
- nodeValue_assnlist, Node-method
(*zzz-DTTCompatibility-class*),
12
- normalizationDescription_assn
(*zzz-DTTCompatibility-class*),
12
- normalizationDescription_assn, ExperimentDesign
(*zzz-DTTCompatibility-class*),
12
- numArrays
(*zzz-DTTCompatibility-class*),
12
- numArrays, ArrayGroup-method
(*zzz-DTTCompatibility-class*),
12
- numberOfFeatures

- (*zzz-DTTCompatibility-class*),
12
- numberOfFeatures, ArrayDesign-method
(*zzz-DTTCompatibility-class*),
12
- numberOfFeatures, PhysicalArrayDesign-method
(*zzz-DTTCompatibility-class*),
12
- numFeaturesPerCol
(*zzz-DTTCompatibility-class*),
12
- numFeaturesPerCol, ZoneLayout-method
(*zzz-DTTCompatibility-class*),
12
- numFeaturesPerRow
(*zzz-DTTCompatibility-class*),
12
- numFeaturesPerRow, ZoneLayout-method
(*zzz-DTTCompatibility-class*),
12
- ontologyEntries_assnlist
(*zzz-DTTCompatibility-class*),
12
- ontologyEntries_assnlist, BioSequence-method
(*zzz-DTTCompatibility-class*),
12
- ontologyEntry
(*zzz-DTTCompatibility-class*),
12
- ontologyEntry, Action_assn-method
(*zzz-DTTCompatibility-class*),
12
- ontologyEntry, Annotations_assnlist-method
(*zzz-DTTCompatibility-class*),
12
- ontologyEntry, Associations_assnlist-method
(*zzz-DTTCompatibility-class*),
12
- ontologyEntry, Category_assn-method
(*zzz-DTTCompatibility-class*),
12
- ontologyEntry, Characteristics_assnlist-method
(*zzz-DTTCompatibility-class*),
12
- ontologyEntry, CompoundIndices_assnlist-method
(*zzz-DTTCompatibility-class*),
12
- ontologyEntry, ControlType_assn-method
(*zzz-DTTCompatibility-class*),
12
- ontologyEntry, DataType_assn-method
(*zzz-DTTCompatibility-class*),
12
- ontologyEntry, DefectType_assn-method
(*zzz-DTTCompatibility-class*),
12
- ontologyEntry, FailTypes_assnlist-method
(*zzz-DTTCompatibility-class*),
12
- ontologyEntry, FeatureShape_assn-method
(*zzz-DTTCompatibility-class*),
12
- ontologyEntry, FiducialType_assn-method
(*zzz-DTTCompatibility-class*),
12
- ontologyEntry, Format_assn-method
(*zzz-DTTCompatibility-class*),
12
- ontologyEntry, MaterialType_assn-method
(*zzz-DTTCompatibility-class*),
12
- ontologyEntry, OntologyEntries_assnlist-method
(*zzz-DTTCompatibility-class*),
12
- ontologyEntry, Parameters_assnlist-method
(*zzz-DTTCompatibility-class*),
12
- ontologyEntry, PolymerType_assn-method
(*zzz-DTTCompatibility-class*),
12
- ontologyEntry, Roles_assnlist-method
(*zzz-DTTCompatibility-class*),
12
- ontologyEntry, Scale_assn-method
(*zzz-DTTCompatibility-class*),
12
- ontologyEntry, Species_assn-method
(*zzz-DTTCompatibility-class*),
12
- ontologyEntry, SubstrateType_assn-method
(*zzz-DTTCompatibility-class*),
12
- ontologyEntry, SurfaceType_assn-method
(*zzz-DTTCompatibility-class*),
12
- ontologyEntry, TechnologyType_assn-method
(*zzz-DTTCompatibility-class*),
12
- ontologyEntry, Type_assn-method
(*zzz-DTTCompatibility-class*),
12
- ontologyEntry, Types_assnlist-method
(*zzz-DTTCompatibility-class*),
12

- ontologyEntry, Value_assn-method (zzz-DTTCompatibility-class), 12
- ontologyEntry, WarningType_assn-method (zzz-DTTCompatibility-class), 12
- ontologyReference_assn (zzz-DTTCompatibility-class), 12
- ontologyReference_assn, OntologyEntry-method (zzz-DTTCompatibility-class), 12
- organization (zzz-DTTCompatibility-class), 12
- organization, Contact_assnlist-method (zzz-DTTCompatibility-class), 12
- organization, Organization_assnlist-method (zzz-DTTCompatibility-class), 12
- organization_ref (zzz-DTTCompatibility-class), 12
- organization_ref, Affiliation_assnref-method (zzz-DTTCompatibility-class), 12
- organization_ref, ArrayManufacturers_assnreflist-method (zzz-DTTCompatibility-class), 12
- organization_ref, Contacts_assnreflist-method (zzz-DTTCompatibility-class), 12
- organization_ref, DesignProviders_assnreflist-method (zzz-DTTCompatibility-class), 12
- organization_ref, HardwareManufacturers_assnreflist-method (zzz-DTTCompatibility-class), 12
- organization_ref, Members_assnreflist-method (zzz-DTTCompatibility-class), 12
- organization_ref, Owner_assnreflist-method (zzz-DTTCompatibility-class), 12
- organization_ref, Parent_assnref-method (zzz-DTTCompatibility-class), 12
- organization_ref, Performer_assnref-method (zzz-DTTCompatibility-class), 12
- organization_ref, Providers_assnreflist-method (zzz-DTTCompatibility-class), 12
- organization_ref, SoftwareManufacturers_assnreflist-method (zzz-DTTCompatibility-class), 12
- organization_ref, SourceContact_assnreflist-method (zzz-DTTCompatibility-class), 12
- orientationMark
- orientationMark, ArrayGroup-method (zzz-DTTCompatibility-class), 12
- orientationMarkPosition
- orientationMarkPosition, ArrayGroup-method (zzz-DTTCompatibility-class), 12
- originalProjectName (zzz-ARRCompatibility-class), 11
- originalProjectName, ArraySetFile-method (zzz-ARRCompatibility-class), 11
- originRelativeTo (zzz-DTTCompatibility-class), 12
- originRelativeTo, Array-method (zzz-DTTCompatibility-class), 12
- otherKind (zzz-DTTCompatibility-class), 12
- otherKind, Measurement-method (zzz-DTTCompatibility-class), 12
- owner_assnreflist (zzz-DTTCompatibility-class), 12
- owner_assnreflist, Security-method (zzz-DTTCompatibility-class), 12
- pages
- pages, BibliographicReference-method (zzz-DTTCompatibility-class), 12
- parameter

- 12
 parameter, ParameterTypes_assnlist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 parameter_ref (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 parameter_ref, ParameterType_assnref-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 parameters_assnlist (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 parameters_assnlist, BibliographicReference-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 parameterType_assnref (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 parameterType_assnref, ParameterValue-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 parameterTypes_assnlist (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 parameterTypes_assnlist, Hardware-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 parameterTypes_assnlist, Protocol-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 parameterTypes_assnlist, Software-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 parameterValue (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 parameterValue, ParameterValues_assnlist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 parameterValues_assnlist (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 parameterValues_assnlist, HardwareApplication-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 parameterValues_assnlist, ProtocolApplication-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 parameterValues_assnlist, SoftwareApplication-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 12
 parent_assnref (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 parent_assnref, Organization-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 password(accessors), 6
 password, NetAffxResource-method (NetAffxResource-class), 2
 patassignmentMethod (zzz-ARRCompatibility-class),
 (zzz-ARRCompatibility-class), 11
 patassignmentMethod, PhysicalArray-method (zzz-ARRCompatibility-class),
 (zzz-ARRCompatibility-class), 11
 performer_assnref (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 performer_assnref, Audit-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 performers_assnreflist (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 performers_assnreflist, ProtocolApplication-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 person_ref (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 person_ref, ArrayManufacturers_assnreflist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 person_ref, Contacts_assnreflist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 person_ref, DesignProviders_assnreflist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 person_ref, HardwareManufacturers_assnreflist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 person_ref, Members_assnreflist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 person_ref, Owner_assnreflist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 person_ref, Performer_assnref-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 person_ref, Performers_assnreflist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12

- (zzz-DTTCompatibility-class), 12
- 12
- person_ref, Providers_assnreflist-method (zzz-DTTCompatibility-class), 12
- (zzz-DTTCompatibility-class), 12
- person_ref, SoftwareManufacturers_assnreflist-method (zzz-DTTCompatibility-class), 12
- (zzz-DTTCompatibility-class), 12
- person_ref, SourceContact_assnreflist-method (zzz-DTTCompatibility-class), 12
- (zzz-DTTCompatibility-class), 12
- phone (zzz-DTTCompatibility-class), 12
- 12
- phone, Organization-method (zzz-DTTCompatibility-class), 12
- (zzz-DTTCompatibility-class), 12
- phone, Person-method (zzz-DTTCompatibility-class), 12
- (zzz-DTTCompatibility-class), 12
- physicalArray (zzz-ARRCompatibility-class), 11
- 11
- physicalArray, PhysicalArrays-method (zzz-ARRCompatibility-class), 11
- (zzz-ARRCompatibility-class), 11
- physicalArrayDesign (zzz-DTTCompatibility-class), 12
- 12
- physicalArrayDesign, ArrayDesign_assnlist-method (zzz-DTTCompatibility-class), 12
- (zzz-DTTCompatibility-class), 12
- physicalArrayDesign_ref (zzz-DTTCompatibility-class), 12
- (zzz-DTTCompatibility-class), 12
- physicalArrayDesign_ref, ArrayDesign_assnref-method (zzz-DTTCompatibility-class), 12
- (zzz-DTTCompatibility-class), 12
- physicalArrays (zzz-ARRCompatibility-class), 11
- 11
- physicalArrays, ArraySetFile-method (zzz-ARRCompatibility-class), 11
- (zzz-ARRCompatibility-class), 11
- physicalBioAssay (zzz-DTTCompatibility-class), 12
- 12
- physicalBioAssay, BioAssay_assnlist-method (zzz-DTTCompatibility-class), 12
- (zzz-DTTCompatibility-class), 12
- physicalBioAssay, PhysicalBioAssay_assnlist-method (zzz-DTTCompatibility-class), 12
- (zzz-DTTCompatibility-class), 12
- physicalBioAssay_ref (zzz-DTTCompatibility-class), 12
- 12
- physicalBioAssay_ref, BioAssay_assnref-method (zzz-DTTCompatibility-class), 12
- 12
- physicalBioAssay_ref, BioAssays_assnreflist-method (zzz-DTTCompatibility-class), 12
- 12
- physicalBioAssay_ref, PhysicalBioAssay_assnref-method (zzz-DTTCompatibility-class), 12
- 12
- physicalBioAssay_ref, PhysicalBioAssaySource_assnref-method (zzz-DTTCompatibility-class), 12
- 12
- physicalBioAssay_ref, PhysicalBioAssayTarget_assnref-method (zzz-DTTCompatibility-class), 12
- 12
- physicalBioAssay_ref, SourceBioAssays_assnreflist-method (zzz-DTTCompatibility-class), 12
- 12
- physicalBioAssay_ref, Target_assnref-method (zzz-DTTCompatibility-class), 12
- 12
- physicalBioAssay_ref, TopLevelBioAssays_assnreflist-method (zzz-DTTCompatibility-class), 12
- 12
- physicalBioAssayData_assnlist (zzz-DTTCompatibility-class), 12
- 12
- physicalBioAssayData_assnlist, PhysicalBioAssayAssayData_assnlist-method (zzz-DTTCompatibility-class), 12
- 12
- physicalBioAssaySource_assnref (zzz-DTTCompatibility-class), 12
- 12
- physicalBioAssaySource_assnref, FeatureExtractAssaySource_assnref-method (zzz-DTTCompatibility-class), 12
- 12
- polymerType_assn (zzz-DTTCompatibility-class), 12
- 12
- polymerType_assn, BioSequence-method (zzz-DTTCompatibility-class), 12
- 12
- position (zzz-DTTCompatibility-class), 12
- 12
- position, Position_assn-method (zzz-DTTCompatibility-class), 12
- 12

- position_assn (zzz-DTTCompatibility-class), 12
 position_assn, Feature-method (zzz-DTTCompatibility-class), 12
 position_assn, Fiducial-method (zzz-DTTCompatibility-class), 12
 positionDelta (zzz-DTTCompatibility-class), 12
 positionDelta, PositionDelta_assn-method (zzz-DTTCompatibility-class), 12
 positionDelta_assn (zzz-DTTCompatibility-class), 12
 positionDelta_assn, FeatureDefect-method (zzz-DTTCompatibility-class), 12
 positionDelta_assn, ZoneDefect-method (zzz-DTTCompatibility-class), 12
 presentAbsent (zzz-DTTCompatibility-class), 12
 presentAbsent, QuantitationType_assnlist-method (zzz-DTTCompatibility-class), 12
 presentAbsent_ref (zzz-DTTCompatibility-class), 12
 presentAbsent_ref, QuantitationType_assnref-method (zzz-DTTCompatibility-class), 12
 presentAbsent_ref, QuantitationTypes_assnreflist-method (zzz-DTTCompatibility-class), 12
 presentAbsent_ref, SourcesQuantitationType_assnreflist-method (zzz-DTTCompatibility-class), 12
 presentAbsent_ref, TargetQuantitationType_assnreflist-method (zzz-DTTCompatibility-class), 12
 producerTransformation_assn (zzz-DTTCompatibility-class), 12
 producerTransformation_assn, DerivedBioAssayData-assnreflist-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, Array-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, ArrayDesign-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, ArrayGroup-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, ArrayManufacture-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, ArrayManufactureDeviation-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, Audit-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, BibliographicReference-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, BioAssayCreation-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, BioAssayDataCluster-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, BioAssayDimension-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, BioAssayMap-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, BioAssayMapping-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, BioAssayTreatment-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, BioDataCube-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, BioDataTuples-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, BioMaterialMeasurement-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, BioSample-method (zzz-DTTCompatibility-class), 12

12
 propertySets_assnlist, BioSequence-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, BioSource-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, Channel-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, CompositeCompositeMap-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, CompositeGroup-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, CompositePosition-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, CompositeSequence-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, CompositeSequenceDimension-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, Compound-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, CompoundMeasurement-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, ConcentrationUnit-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, Database-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, DatabaseEntry-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, DerivedBioAssay-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, DerivedBioAssayData-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, DerivedSignal-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, Description-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, DesignElementMapping-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, DistanceUnit-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, Error-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, ExpectedValue-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, Experiment-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, ExperimentalFactor-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, ExperimentDesign-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, ExternalReference-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, FactorValue-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, Failed-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, Feature-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, FeatureDefect-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, FeatureDimension-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, FeatureExtraction-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, FeatureGroup-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, FeatureInformation-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, FeatureLocation-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, FeatureReporterMap-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12

(zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, Fiducial-method (zzz-DTTCompatibility-class), 12
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, Hardware-method (zzz-DTTCompatibility-class), 12
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, HardwareApplication-method (zzz-DTTCompatibility-class), 12
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, Hybridization-method (zzz-DTTCompatibility-class), 12
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, Image-method (zzz-DTTCompatibility-class), 12
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, ImageAcquisition-method (zzz-DTTCompatibility-class), 12
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, LabeledExtract-method (zzz-DTTCompatibility-class), 12
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, MAGE-ML-method (zzz-DTTCompatibility-class), 12
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, ManufactureLIMS-method (zzz-DTTCompatibility-class), 12
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, ManufactureLIMSBiomaterial-method (zzz-DTTCompatibility-class), 12
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, MassUnit-method (zzz-DTTCompatibility-class), 12
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, MeasuredBioAssay-method (zzz-DTTCompatibility-class), 12
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, MeasuredBioAssayData-method (zzz-DTTCompatibility-class), 12
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, MeasuredSignal-method (zzz-DTTCompatibility-class), 12
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, Measurement-method (zzz-DTTCompatibility-class), 12
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, MismatchInformation-method (zzz-DTTCompatibility-class), 12
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, NameValueType-method (zzz-DTTCompatibility-class), 12
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, Node-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, NodeContents-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, NodeValue-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, OntologyEntry-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, Organization-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, Parameter-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, ParameterValue-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, PhysicalArrayDesign-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, PhysicalBioAssay-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, Position-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, PositionDelta-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, PresentAbsent-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, Protocol-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, ProtocolApplication-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, PValue-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, QuantitationTypeDimensionality-method (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, QuantitationTypeDimensionality-method (zzz-DTTCompatibility-class), 12

propertySets_assnlist, QuantitationTypeMap-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, QuantitationTypeMapping-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, QuantityUnit-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, Ratio-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, Reporter-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, ReporterCompositeMap-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, ReporterDimension-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, ReporterGroup-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, ReporterPosition-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, Security-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, SecurityGroup-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, SeqFeature-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, SeqFeatureLocation-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, SequencePosition-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, Software-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, SoftwareApplication-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, SpecializedQuantitationType-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, TemperatureUnit-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, TimeUnit-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, Transformation-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, Treatment-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, VolumeUnit-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, Zone-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, ZoneDefect-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, ZoneGroup-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, ZoneLayout-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 protocol
 protocol, Protocol_assnlist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 protocol_assnlist
 protocol_assnlist, Protocol_package-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 protocol_assnref
 protocol_assnref, ProtocolApplication-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 protocol_package
 protocol_package, MAGE-ML-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 protocol_ref

- 12
 protocol_ref, Protocol_assnref-method
 (zzz-DTTCompatibility-class),
 12
- protocolApplication
 (zzz-DTTCompatibility-class),
 12
- protocolApplication, ProtocolApplications_assnlist-method
 (zzz-DTTCompatibility-class),
 12
- protocolApplications_assnlist
 (zzz-DTTCompatibility-class),
 12
- protocolApplications_assnlist, ArrayDesign-method
 (zzz-DTTCompatibility-class),
 12
- protocolApplications_assnlist, ArrayManufacture-method
 (zzz-DTTCompatibility-class),
 12
- protocolApplications_assnlist, BioAssayCreation-method
 (zzz-DTTCompatibility-class),
 12
- protocolApplications_assnlist, BioAssayMap-method
 (zzz-DTTCompatibility-class),
 12
- protocolApplications_assnlist, BioAssayTreatment-method
 (zzz-DTTCompatibility-class),
 12
- protocolApplications_assnlist, CompositeCompositeMap-method
 (zzz-DTTCompatibility-class),
 12
- protocolApplications_assnlist, FeatureExtraction-method
 (zzz-DTTCompatibility-class),
 12
- protocolApplications_assnlist, FeatureReporterMap-method
 (zzz-DTTCompatibility-class),
 12
- protocolApplications_assnlist, Hybridization-method
 (zzz-DTTCompatibility-class),
 12
- protocolApplications_assnlist, ImageAcquisition-method
 (zzz-DTTCompatibility-class),
 12
- protocolApplications_assnlist, PhysicalArrayDesign-method
 (zzz-DTTCompatibility-class),
 12
- protocolApplications_assnlist, QuantitationTypeMap-method
 (zzz-DTTCompatibility-class),
 12
- protocolApplications_assnlist, ReporterCompositeMap-method
 (zzz-DTTCompatibility-class),
 12
- 12
 protocolApplications_assnlist, Transformation-method
 (zzz-DTTCompatibility-class),
 12
- protocolApplications_assnlist, Treatment-method
 (zzz-DTTCompatibility-class),
 12
- providers_assnreflist
 (zzz-DTTCompatibility-class),
 12
- providers_assnreflist, Experiment-method
 (zzz-DTTCompatibility-class),
 12
- publication
 (zzz-DTTCompatibility-class),
 12
- publication, BibliographicReference-method
 (zzz-DTTCompatibility-class),
 12
- publisher
 (zzz-DTTCompatibility-class),
 12
- publisher, BibliographicReference-method
 (zzz-DTTCompatibility-class),
 12
- pvalue
 (zzz-DTTCompatibility-class),
 12
- pvalue, ConfidenceIndicator_assnlist-method
 (zzz-DTTCompatibility-class),
 12
- pvalue, QuantitationType_assnlist-method
 (zzz-DTTCompatibility-class),
 12
- pvalue_ref
 (zzz-DTTCompatibility-class),
 12
- pvalue_ref, ConfidenceIndicators_assnreflist-method
 (zzz-DTTCompatibility-class),
 12
- pvalue_ref, QuantitationType_assnref-method
 (zzz-DTTCompatibility-class),
 12
- pvalue_ref, QuantitationTypes_assnreflist-method
 (zzz-DTTCompatibility-class),
 12
- pvalue_ref, SourcesQuantitationType_assnreflist
 (zzz-DTTCompatibility-class),
 12
- pvalue_ref, TargetQuantitationType_assnref-method
 (zzz-DTTCompatibility-class),
 12
- quality

(zzz-DTTCompatibility-class), 12	12
quality, ManufactureLIMS-method (zzz-DTTCompatibility-class), 12	quantitationTypeDimension (zzz-DTTCompatibility-class), 12
quality, ManufactureLIMSBiomaterial-method (zzz-DTTCompatibility-class), 12	quantitationTypeDimension, QuantitationTypeDimension (zzz-DTTCompatibility-class), 12
qualityControlDescription_assn (zzz-DTTCompatibility-class), 12	quantitationTypeDimension_assnlist (zzz-DTTCompatibility-class), 12
qualityControlDescription_assn, ExperimentDescription (zzz-DTTCompatibility-class), 12	quantitationTypeDimension_assnlist, BioAssayData (zzz-DTTCompatibility-class), 12
qualityControlStatistics_assnlist (zzz-DTTCompatibility-class), 12	quantitationTypeDimension_assnref (zzz-DTTCompatibility-class), 12
qualityControlStatistics_assnlist, ArrayManufacturing (zzz-DTTCompatibility-class), 12	quantitationTypeDimension_assnref, DerivedBioAssay (zzz-DTTCompatibility-class), 12
qualityControlStatistics_assnlist, BioSample-method (zzz-DTTCompatibility-class), 12	quantitationTypeDimension_assnref, MeasuredBioAssay (zzz-DTTCompatibility-class), 12
qualityControlStatistics_assnlist, BioSource-method (zzz-DTTCompatibility-class), 12	quantitationTypeDimension_ref (zzz-DTTCompatibility-class), 12
qualityControlStatistics_assnlist, LabeledExtension (zzz-DTTCompatibility-class), 12	quantitationTypeDimension_ref, QuantitationDimension (zzz-DTTCompatibility-class), 12
quantitationDimension_assnref (zzz-DTTCompatibility-class), 12	quantitationTypeDimension_ref, QuantitationTypeDimension (zzz-DTTCompatibility-class), 12
quantitationDimension_assnref, NodeContents-method (zzz-DTTCompatibility-class), 12	quantitationTypeMap (zzz-DTTCompatibility-class), 12
quantitationType_assnlist (zzz-DTTCompatibility-class), 12	quantitationTypeMap, QuantitationTypeMap_assnlist (zzz-DTTCompatibility-class), 12
quantitationType_assnlist, QuantitationType_package (zzz-DTTCompatibility-class), 12	quantitationTypeMap_assnlist (zzz-DTTCompatibility-class), 12
quantitationType_assnref (zzz-DTTCompatibility-class), 12	quantitationTypeMap_assnlist, BioAssayData_package (zzz-DTTCompatibility-class), 12
quantitationType_assnref, QuantitationTypeTuple (zzz-DTTCompatibility-class), 12	quantitationTypeMap_ref (zzz-DTTCompatibility-class), 12
quantitationType_package (zzz-DTTCompatibility-class), 12	quantitationTypeMap_ref, QuantitationTypeMaps (zzz-DTTCompatibility-class), 12
quantitationType_package, MAGE-ML-method (zzz-DTTCompatibility-class), 12	quantitationTypeMapping (zzz-DTTCompatibility-class), 12

quantitationTypeMapping, QuantitationTypeMapping (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 quantitationTypeMapping_assnlist
 (zzz-DTTCompatibility-class), 12
 quantitationTypeMapping_assn, Transformation (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 quantitationTypeMaps_assnreflist
 (zzz-DTTCompatibility-class), 12
 quantitationTypeMaps_assnreflist, DerivedSignal (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 quantitationTypeMaps_assnreflist, Error (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 quantitationTypeMaps_assnreflist, ExpectedValue (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 quantitationTypeMaps_assnreflist, Failed (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 quantitationTypeMaps_assnreflist, MeasuredSignal, quantitationType_assnref-method
 (zzz-DTTCompatibility-class), (zzz-DTTCompatibility-class), 12
 quantitationTypeMaps_assnreflist, PresentAbsence, quantitationTypes_assnreflist-method
 (zzz-DTTCompatibility-class), (zzz-DTTCompatibility-class), 12
 quantitationTypeMaps_assnreflist, PValue, quantitationType_assnreflist-method
 (zzz-DTTCompatibility-class), (zzz-DTTCompatibility-class), 12
 quantitationTypeMaps_assnreflist, QuantitationTypeMapping, quantitationType_assnref-method
 (zzz-DTTCompatibility-class), (zzz-DTTCompatibility-class), 12
 quantitationTypeMaps_assnreflist, Ratio, readAnnotation
 (zzz-DTTCompatibility-class), (NetAffxResource-class), 2
 quantitationTypeMaps_assnreflist, SpecializedQuantitationType (NetAffxResource-class), 2
 (zzz-DTTCompatibility-class), readAnnotation, NetAffxResource, character, character, character
 (NetAffxResource-class), 2
 quantitationTypes_assnreflist
 (zzz-DTTCompatibility-class), readAnnotation, NetAffxResource, missing, AffxAnnotation
 (NetAffxResource-class), 2
 quantitationTypes_assnreflist, QuantitationType (NetAffxResource-class), 2
 (zzz-DTTCompatibility-class), readAnnotation, NetAffxResource, numeric, character, character
 (NetAffxResource-class), 2
 quantitationTypeTuple
 (zzz-DTTCompatibility-class), readArr, 6
 12 readMage, 7
 readNetAffx, 8
 quantitationTypeTuple, QuantitationTypeTuples, readAnnotation, 7, 9, 10

regions_assnlist (zzz-DTTCompatibility-class), 12
regions_assnlist, SeqFeature-method (zzz-DTTCompatibility-class), 12
releaseDate (zzz-DTTCompatibility-class), 12
releaseDate, SoftwareApplication-method (zzz-DTTCompatibility-class), 12
replacedLength (zzz-DTTCompatibility-class), 12
replacedLength, MismatchInformation-method (zzz-DTTCompatibility-class), 12
replicateDescription_assn (zzz-DTTCompatibility-class), 12
replicateDescription_assn, ExperimentDesign-method (zzz-DTTCompatibility-class), 12
reporter (zzz-DTTCompatibility-class), 12
reporter, DesignElement_assnlist-method (zzz-DTTCompatibility-class), 12
reporter, Reporter_assnlist-method (zzz-DTTCompatibility-class), 12
reporter_assnlist (zzz-DTTCompatibility-class), 12
reporter_assnlist, DesignElement_package-method (zzz-DTTCompatibility-class), 12
reporter_assnref (zzz-DTTCompatibility-class), 12
reporter_assnref, FeatureReporterMap-method (zzz-DTTCompatibility-class), 12
reporter_assnref, ReporterPosition-method (zzz-DTTCompatibility-class), 12
reporter_ref (zzz-DTTCompatibility-class), 12
reporter_ref, DesignElement_assnref-method (zzz-DTTCompatibility-class), 12
reporter_ref, Reporter_assnref-method (zzz-DTTCompatibility-class), 12
reporter_ref, Reporters_assnreflist-method (zzz-DTTCompatibility-class), 12
reporterCompositeMap (zzz-DTTCompatibility-class), 12
reporterCompositeMap, DesignElementMap_assnlist (zzz-DTTCompatibility-class), 12
reporterCompositeMap, ReporterCompositeMap_assnlist (zzz-DTTCompatibility-class), 12
reporterCompositeMap_assnlist (zzz-DTTCompatibility-class), 12
reporterCompositeMap_assnlist, DesignElement_package-method (zzz-DTTCompatibility-class), 12
reporterCompositeMap_ref (zzz-DTTCompatibility-class), 12
reporterCompositeMap_ref, DesignElementMaps_assnlist (zzz-DTTCompatibility-class), 12
reporterCompositeMap_ref, ReporterCompositeMaps_assnreflist (zzz-DTTCompatibility-class), 12
reporterCompositeMaps_assnreflist (zzz-DTTCompatibility-class), 12
reporterCompositeMaps_assnreflist, CompositeSeq-method (zzz-DTTCompatibility-class), 12
reporterDimension (zzz-DTTCompatibility-class), 12
reporterDimension, DesignElementDimension_assnlist (zzz-DTTCompatibility-class), 12
reporterDimension_ref (zzz-DTTCompatibility-class), 12
reporterDimension_ref, DesignElementDimension_assnlist (zzz-DTTCompatibility-class), 12
reporterGroup (zzz-DTTCompatibility-class), 12

12 required, UserAttribute-method
 reporterGroup, ReporterGroup_assnlist-method (zzz-ARRCompatibility-class),
 (zzz-DTTCompatibility-class), 11
 12 roles_assnlist
 reporterGroup_assnlist (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12 roles_assnlist, Organization-method
 reporterGroup_assnlist, ArrayDesign_package-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12 roles_assnlist, Person-method
 reporterGroup_ref (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12 scale_assn
 reporterGroup_ref, ReporterGroups_assnreflist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12 scale_assn, DerivedSignal-method
 reporterGroups_assnreflist (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12 scale_assn, Error-method
 reporterGroups_assnreflist, ArrayDesign-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12 scale_assn, ExpectedValue-method
 reporterGroups_assnreflist, PhysicalArrayDesign-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12 scale_assn, Failed-method
 reporterPosition (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12 scale_assn, MeasuredSignal-method
 reporterPosition, Coordinate_assn-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12 scale_assn, NodeValue-method
 reporterPosition, ReporterPositionSources_assnlist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12 scale_assn, PresentAbsent-method
 reporterPositionSources_assnlist (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12 scale_assn, PValue-method
 reporterPositionSources_assnlist, ReporterCompositeMap-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12 scale_assn, Ratio-method
 reporters_assnreflist (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12 scale_assn, SpecializedQuantitationType-method
 reporters_assnreflist, ReporterDimension-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12 security
 reporters_assnreflist, ReporterGroup-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12 security, Security_assnlist-method
 required (zzz-ARRCompatibility-class),
 (zzz-ARRCompatibility-class), 12
 11 security_assnlist

- (zzz-*DTTCompatibility-class*), 12
- security_assnlist, *AuditAndSecurity_package-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref (zzz-*DTTCompatibility-class*), 12
- security_assnref, *Array-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *ArrayDesign-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *ArrayGroup-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *ArrayManufacture-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *Audit-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *BibliographicReference-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *BioAssayCreation-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *BioAssayDataCluster-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *BioAssayDimension-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *BioAssayMap-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *BioAssayTreatment-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *BioSample-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *BioSequence-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *BioSource-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *Channel-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *CompositeCompositeMap-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *CompositeGroup-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *CompositeSequence-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *CompositeSequenceDimension-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *Compound-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *Database-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *DerivedBioAssay-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *DerivedBioAssayData-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *DerivedSignal-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *Description-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *Error-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *ExpectedValue-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *Experiment-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *ExperimentalFactor-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *ExperimentDesign-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *FactorValue-method* (zzz-*DTTCompatibility-class*), 12
- security_assnref, *Failed-method* (zzz-*DTTCompatibility-class*), 12

- security_assnref, Feature-method (zzz-DTTCompatibility-class), 12
 security_assnref, FeatureDimension-method (zzz-DTTCompatibility-class), 12
 security_assnref, FeatureExtraction-method (zzz-DTTCompatibility-class), 12
 security_assnref, FeatureGroup-method (zzz-DTTCompatibility-class), 12
 security_assnref, FeatureReporterMap-method (zzz-DTTCompatibility-class), 12
 security_assnref, Fiducial-method (zzz-DTTCompatibility-class), 12
 security_assnref, Hardware-method (zzz-DTTCompatibility-class), 12
 security_assnref, HardwareApplication-method (zzz-DTTCompatibility-class), 12
 security_assnref, Hybridization-method (zzz-DTTCompatibility-class), 12
 security_assnref, Image-method (zzz-DTTCompatibility-class), 12
 security_assnref, ImageAcquisition-method (zzz-DTTCompatibility-class), 12
 security_assnref, LabeledExtract-method (zzz-DTTCompatibility-class), 12
 security_assnref, MAGE-ML-method (zzz-DTTCompatibility-class), 12
 security_assnref, ManufactureLIMS-method (zzz-DTTCompatibility-class), 12
 security_assnref, ManufactureLIMSBiomaterial-method (zzz-DTTCompatibility-class), 12
 security_assnref, MeasuredBioAssay-method (zzz-DTTCompatibility-class), 12
 security_assnref, MeasuredBioAssayData-method (zzz-DTTCompatibility-class), 12
 security_assnref, MeasuredSignal-method (zzz-DTTCompatibility-class), 12
 security_assnref, Node-method (zzz-DTTCompatibility-class), 12
 security_assnref, NodeContents-method (zzz-DTTCompatibility-class), 12
 security_assnref, Organization-method (zzz-DTTCompatibility-class), 12
 security_assnref, Parameter-method (zzz-DTTCompatibility-class), 12
 security_assnref, Person-method (zzz-DTTCompatibility-class), 12
 security_assnref, PhysicalArrayDesign-method (zzz-DTTCompatibility-class), 12
 security_assnref, PhysicalBioAssay-method (zzz-DTTCompatibility-class), 12
 security_assnref, PresentAbsent-method (zzz-DTTCompatibility-class), 12
 security_assnref, Protocol-method (zzz-DTTCompatibility-class), 12
 security_assnref, ProtocolApplication-method (zzz-DTTCompatibility-class), 12
 security_assnref, PValue-method (zzz-DTTCompatibility-class), 12
 security_assnref, QuantitationTypeDimension-method (zzz-DTTCompatibility-class), 12
 security_assnref, QuantitationTypeMap-method (zzz-DTTCompatibility-class), 12
 security_assnref, Ratio-method (zzz-DTTCompatibility-class), 12
 security_assnref, Reporter-method (zzz-DTTCompatibility-class), 12
 security_assnref, ReporterCompositeMap-method (zzz-DTTCompatibility-class), 12
 security_assnref, ReporterDimension-method (zzz-DTTCompatibility-class), 12

- 12
- security_assnref, ReporterGroup-method
(zzz-DTTCompatibility-class),
12
- security_assnref, Security-method
(zzz-DTTCompatibility-class),
12
- security_assnref, SecurityGroup-method
(zzz-DTTCompatibility-class),
12
- security_assnref, SeqFeature-method
(zzz-DTTCompatibility-class),
12
- security_assnref, Software-method
(zzz-DTTCompatibility-class),
12
- security_assnref, SoftwareApplication-method
(zzz-DTTCompatibility-class),
12
- security_assnref, SpecializedQuantitationType-method
(zzz-DTTCompatibility-class),
12
- security_assnref, Transformation-method
(zzz-DTTCompatibility-class),
12
- security_assnref, Treatment-method
(zzz-DTTCompatibility-class),
12
- security_assnref, Zone-method
(zzz-DTTCompatibility-class),
12
- security_ref
(zzz-DTTCompatibility-class),
12
- security_ref, Security_assnref-method
(zzz-DTTCompatibility-class),
12
- securityGroup
(zzz-DTTCompatibility-class),
12
- securityGroup, SecurityGroup_assnlist-method
(zzz-DTTCompatibility-class),
12
- securityGroup_assnlist
(zzz-DTTCompatibility-class),
12
- securityGroup_assnlist, AuditAndSecurity_package-method
(zzz-DTTCompatibility-class),
12
- securityGroup_ref
(zzz-DTTCompatibility-class),
12
- securityGroup_ref, SecurityGroups_assnreflist-method
(zzz-DTTCompatibility-class),
12
- securityGroups_assnreflist
(zzz-DTTCompatibility-class),
12
- securityGroups_assnreflist, Security-method
(zzz-DTTCompatibility-class),
12
- seqFeature
(zzz-DTTCompatibility-class),
12
- seqFeature, SeqFeatures_assnlist-method
(zzz-DTTCompatibility-class),
12
- seqFeatureLocation
(zzz-DTTCompatibility-class),
12
- seqFeatureLocation, Regions_assnlist-method
(zzz-DTTCompatibility-class),
12
- seqFeatureLocation, Subregions_assnlist-method
(zzz-DTTCompatibility-class),
12
- seqFeatures_assnlist
(zzz-DTTCompatibility-class),
12
- seqFeatures_assnlist, BioSequence-method
(zzz-DTTCompatibility-class),
12
- sequenceDatabases_assnlist
(zzz-DTTCompatibility-class),
12
- sequenceDatabases_assnlist, BioSequence-method
(zzz-DTTCompatibility-class),
12
- sequencePosition
(zzz-DTTCompatibility-class),
12
- sequencePosition, Coordinate_assn-method
(zzz-DTTCompatibility-class),
12
- serialNumber
(zzz-DTTCompatibility-class),
12
- serialNumber, HardwareApplication-method
(zzz-DTTCompatibility-class),
12
- show, .TypedAssociation-method
(Internal-class), 11
- show, AffyCompatible-method
(AffyCompatible-class), 1

- show, NetAffxResource-method
(NetAffxResource-class), 2
- software
(zzz-DTTCompatibility-class),
12
- software, Software_assnlist-method
(zzz-DTTCompatibility-class),
12
- software_assnlist
(zzz-DTTCompatibility-class),
12
- software_assnlist, Protocol_package-method
(zzz-DTTCompatibility-class),
12
- software_assnref
(zzz-DTTCompatibility-class),
12
- software_assnref, SoftwareApplication-method
(zzz-DTTCompatibility-class),
12
- software_ref
(zzz-DTTCompatibility-class),
12
- software_ref, Software_assnref-method
(zzz-DTTCompatibility-class),
12
- software_ref, Softwares_assnreflist-method
(zzz-DTTCompatibility-class),
12
- softwareApplication
(zzz-DTTCompatibility-class),
12
- softwareApplication, SoftwareApplications_assnreflist-method
(zzz-DTTCompatibility-class),
12
- softwareApplications_assnlist
(zzz-DTTCompatibility-class),
12
- softwareApplications_assnlist, ProtocolApplication-assnreflist-method
(zzz-DTTCompatibility-class),
12
- softwareManufacturers_assnreflist
(zzz-DTTCompatibility-class),
12
- softwareManufacturers_assnreflist, Software-manufacturer-method
(zzz-DTTCompatibility-class),
12
- softwares_assnreflist
(zzz-DTTCompatibility-class),
12
- softwares_assnreflist, Hardware-method
(zzz-DTTCompatibility-class),
12
- 12
- softwares_assnreflist, Protocol-method
(zzz-DTTCompatibility-class),
12
- softwares_assnreflist, Software-method
(zzz-DTTCompatibility-class),
12
- sourceBioAssays_assnreflist
(zzz-DTTCompatibility-class),
12
- sourceBioAssays_assnreflist, BioAssayMap-method
(zzz-DTTCompatibility-class),
12
- sourceBioMaterialMeasurements_assnlist
(zzz-DTTCompatibility-class),
12
- sourceBioMaterialMeasurements_assnlist, BioAssayMap-method
(zzz-DTTCompatibility-class),
12
- sourceBioMaterialMeasurements_assnlist, Hybridization-method
(zzz-DTTCompatibility-class),
12
- sourceBioMaterialMeasurements_assnlist, Treatment-method
(zzz-DTTCompatibility-class),
12
- sourceContact_assnreflist
(zzz-DTTCompatibility-class),
12
- sourceContact_assnreflist, BioSource-method
(zzz-DTTCompatibility-class),
12
- sourcesQuantitationType_assnreflist
(zzz-DTTCompatibility-class),
12
- sourcesQuantitationType_assnreflist, Quantitation-method
(zzz-DTTCompatibility-class),
12
- spacingBetweenCols
(zzz-DTTCompatibility-class),
12
- spacingBetweenCols, ZoneLayout-method
(zzz-DTTCompatibility-class),
12
- spacingBetweenRows
(zzz-DTTCompatibility-class),
12
- spacingBetweenRows, ZoneLayout-method
(zzz-DTTCompatibility-class),
12
- spacingsBetweenZonesX
(zzz-DTTCompatibility-class),
12

- spacingsBetweenZonesX, ZoneGroup-method (zzz-DTTCompatibility-class),
 12
 spacingsBetweenZonesY (zzz-DTTCompatibility-class),
 12
 spacingsBetweenZonesY, ZoneGroup-method (zzz-DTTCompatibility-class),
 12
 specializedQuantitationType (zzz-DTTCompatibility-class),
 12
 specializedQuantitationType, QuantitationType_assnref-method (zzz-DTTCompatibility-class),
 12
 specializedQuantitationType_ref (zzz-DTTCompatibility-class),
 12
 specializedQuantitationType_ref, QuantitationType_assnref-method (zzz-DTTCompatibility-class),
 12
 specializedQuantitationType_ref, QuantitationTypes_assnrefref-list-method (zzz-DTTCompatibility-class),
 12
 specializedQuantitationType_ref, SourcesQuantitationType_assnrefref-list-method (zzz-DTTCompatibility-class),
 12
 specializedQuantitationType_ref, TargetQuantitationType_assnref-method (zzz-DTTCompatibility-class),
 12
 species_assn (zzz-DTTCompatibility-class),
 12
 species_assn, BioSequence-method (zzz-DTTCompatibility-class),
 12
 species_assn, CompositeGroup-method (zzz-DTTCompatibility-class),
 12
 species_assn, FeatureGroup-method (zzz-DTTCompatibility-class),
 12
 species_assn, ReporterGroup-method (zzz-DTTCompatibility-class),
 12
 startCoord (zzz-DTTCompatibility-class),
 12
 startCoord, MismatchInformation-method (zzz-DTTCompatibility-class),
 12
 strandType (zzz-DTTCompatibility-class),
 12
 strandType, SeqFeatureLocation-method (zzz-DTTCompatibility-class),
 12
 subregions_assnlist (zzz-DTTCompatibility-class),
 12
 subregions_assnlist, SeqFeatureLocation-method (zzz-DTTCompatibility-class),
 12
 substrateType_assn (zzz-DTTCompatibility-class),
 12
 substrateType_assn, ArrayGroup-method (zzz-DTTCompatibility-class),
 12
 summaryStatistics_assnlist (zzz-DTTCompatibility-class),
 12
 summaryStatistics_assnlist, DerivedBioAssayData (zzz-DTTCompatibility-class),
 12
 summaryStatistics_assnlist, MeasuredBioAssayData (zzz-DTTCompatibility-class),
 12
 surfaceType_assn (zzz-DTTCompatibility-class),
 12
 surfaceType_assn, PhysicalArrayDesign-method (zzz-DTTCompatibility-class),
 12
 target_assnref (zzz-DTTCompatibility-class),
 12
 target_assnref, BioAssayTreatment-method (zzz-DTTCompatibility-class),
 12
 target_assnref, ImageAcquisition-method (zzz-DTTCompatibility-class),
 12
 targetQuantitationType_assnref (zzz-DTTCompatibility-class),
 12
 targetQuantitationType_assnref, Error-method (zzz-DTTCompatibility-class),
 12
 targetQuantitationType_assnref, ExpectedValue-r (zzz-DTTCompatibility-class),
 12
 targetQuantitationType_assnref, PValue-method (zzz-DTTCompatibility-class),
 12

- 12
- targetQuantitationType_assnref, QuantitationTypeMap-*method* (*zzz-DTTCompatibility-class*), 12
- technologyType_assn (*zzz-DTTCompatibility-class*), 12
- technologyType_assn, FeatureGroup-*method* (*zzz-DTTCompatibility-class*), 12
- temperatureUnit (*zzz-DTTCompatibility-class*), 12
- temperatureUnit, Unit_assn-*method* (*zzz-DTTCompatibility-class*), 12
- timeUnit (*zzz-DTTCompatibility-class*), 12
- timeUnit, Unit_assn-*method* (*zzz-DTTCompatibility-class*), 12
- tolerance (*zzz-DTTCompatibility-class*), 12
- tolerance, ArrayManufacture-*method* (*zzz-DTTCompatibility-class*), 12
- tollFreePhone (*zzz-DTTCompatibility-class*), 12
- tollFreePhone, Organization-*method* (*zzz-DTTCompatibility-class*), 12
- tollFreePhone, Person-*method* (*zzz-DTTCompatibility-class*), 12
- topLevelBioAssays_assnreflist (*zzz-DTTCompatibility-class*), 12
- topLevelBioAssays_assnreflist, ExperimentDesign-*method* (*zzz-DTTCompatibility-class*), 12
- transformation (*zzz-DTTCompatibility-class*), 12
- transformation, ProducerTransformation_assn-*method* (*zzz-DTTCompatibility-class*), 12
- treatment (*zzz-DTTCompatibility-class*), 12
- treatment, Treatments_assnlist-*method* (*zzz-DTTCompatibility-class*), 12
- treatments_assnlist (*zzz-DTTCompatibility-class*), 12
- treatments_assnlist, BioSample-*method* (*zzz-DTTCompatibility-class*), 12
- treatments_assnlist, BioSource-*method* (*zzz-DTTCompatibility-class*), 12
- treatments_assnlist, LabeledExtract-*method* (*zzz-DTTCompatibility-class*), 12
- type (*zzz-DTTCompatibility-class*), 12
- type, ArraySetFile-*method* (*zzz-ARRCompatibility-class*), 11
- type, Measurement-*method* (*zzz-DTTCompatibility-class*), 12
- type, NameValueType-*method* (*zzz-DTTCompatibility-class*), 12
- type, PhysicalArray-*method* (*zzz-ARRCompatibility-class*), 11
- type, TemplateFile-*method* (*zzz-ARRCompatibility-class*), 11
- type, UserAttribute-*method* (*zzz-ARRCompatibility-class*), 11
- type_assn (*zzz-DTTCompatibility-class*), 12
- type_assn, BioSample-*method* (*zzz-DTTCompatibility-class*), 12
- type_assn, BioSequence-*method* (*zzz-DTTCompatibility-class*), 12
- type_assn, DatabaseEntry-*method* (*zzz-DTTCompatibility-class*), 12
- type_assn, DerivedBioAssay-*method* (*zzz-DTTCompatibility-class*), 12
- type_assn, Hardware-*method*

- (zzz-*DTTCompatibility-class*),
12
- type_assn, NodeValue-method
 (zzz-*DTTCompatibility-class*),
 12
- type_assn, Protocol-method
 (zzz-*DTTCompatibility-class*),
 12
- type_assn, Software-method
 (zzz-*DTTCompatibility-class*),
 12
- types_assnlist
 (zzz-*DTTCompatibility-class*),
 12
- types_assnlist, CompositeGroup-method
 (zzz-*DTTCompatibility-class*),
 12
- types_assnlist, ExperimentDesign-method
 (zzz-*DTTCompatibility-class*),
 12
- types_assnlist, FeatureGroup-method
 (zzz-*DTTCompatibility-class*),
 12
- types_assnlist, ReporterGroup-method
 (zzz-*DTTCompatibility-class*),
 12
- unit_assn
 (zzz-*DTTCompatibility-class*),
 12
- unit_assn, Measurement-method
 (zzz-*DTTCompatibility-class*),
 12
- unitName
 (zzz-*DTTCompatibility-class*),
 12
- unitName, ConcentrationUnit-method
 (zzz-*DTTCompatibility-class*),
 12
- unitName, DistanceUnit-method
 (zzz-*DTTCompatibility-class*),
 12
- unitName, MassUnit-method
 (zzz-*DTTCompatibility-class*),
 12
- unitName, QuantityUnit-method
 (zzz-*DTTCompatibility-class*),
 12
- unitName, TemperatureUnit-method
 (zzz-*DTTCompatibility-class*),
 12
- unitName, TimeUnit-method
 (zzz-*DTTCompatibility-class*),
 12
- unitName, VolumeUnit-method
 (zzz-*DTTCompatibility-class*),
 12
- unitNameCV
 (zzz-*DTTCompatibility-class*),
 12
- unitNameCV, ConcentrationUnit-method
 (zzz-*DTTCompatibility-class*),
 12
- unitNameCV, DistanceUnit-method
 (zzz-*DTTCompatibility-class*),
 12
- unitNameCV, MassUnit-method
 (zzz-*DTTCompatibility-class*),
 12
- unitNameCV, QuantityUnit-method
 (zzz-*DTTCompatibility-class*),
 12
- unitNameCV, TemperatureUnit-method
 (zzz-*DTTCompatibility-class*),
 12
- unitNameCV, TimeUnit-method
 (zzz-*DTTCompatibility-class*),
 12
- unitNameCV, VolumeUnit-method
 (zzz-*DTTCompatibility-class*),
 12
- upperLeftX
 (zzz-*DTTCompatibility-class*),
 12
- upperLeftX, Zone-method
 (zzz-*DTTCompatibility-class*),
 12
- upperLeftY
 (zzz-*DTTCompatibility-class*),
 12
- upperLeftY, Zone-method
 (zzz-*DTTCompatibility-class*),
 12
- uri (zzz-*DTTCompatibility-class*),
 12
- uri, BibliographicReference-method
 (zzz-*DTTCompatibility-class*),
 12
- uri, Database-method
 (zzz-*DTTCompatibility-class*),
 12
- uri, DatabaseEntry-method
 (zzz-*DTTCompatibility-class*),
 12
- uri, Description-method

- (*zzz-DTTCompatibility-class*),
12
- uri, Hardware-method
(*zzz-DTTCompatibility-class*),
12
- uri, Image-method
(*zzz-DTTCompatibility-class*),
12
- uri, Organization-method
(*zzz-DTTCompatibility-class*),
12
- uri, Person-method
(*zzz-DTTCompatibility-class*),
12
- uri, Protocol-method
(*zzz-DTTCompatibility-class*),
12
- uri, Software-method
(*zzz-DTTCompatibility-class*),
12
- user (*accessors*), 6
- user, NetAffxResource-method
(*NetAffxResource-class*), 2
- userAttribute
(*zzz-ARRCompatibility-class*),
11
- userAttribute, UserAttributes-method
(*zzz-ARRCompatibility-class*),
11
- userAttributes
(*zzz-ARRCompatibility-class*),
11
- userAttributes, ArraySetFile-method
(*zzz-ARRCompatibility-class*),
11
- userAttributes, TemplateFile-method
(*zzz-ARRCompatibility-class*),
11
- userAttributeValue
(*zzz-ARRCompatibility-class*),
11
- userAttributeValue, UserAttribute-method
(*zzz-ARRCompatibility-class*),
11
- value
(*zzz-ARRCompatibility-class*),
11
- value, Control-method
(*zzz-ARRCompatibility-class*),
11
- value, Datum-method
(*zzz-DTTCompatibility-class*),
12
- value, Measurement-method
(*zzz-DTTCompatibility-class*),
12
- value, NameValueType-method
(*zzz-DTTCompatibility-class*),
12
- value, NodeValue-method
(*zzz-DTTCompatibility-class*),
12
- value, OntologyEntry-method
(*zzz-DTTCompatibility-class*),
12
- value, ParameterValue-method
(*zzz-DTTCompatibility-class*),
12
- value_assn
(*zzz-DTTCompatibility-class*),
12
- value_assn, FactorValue-method
(*zzz-DTTCompatibility-class*),
12
- vector, 11
- version
(*zzz-ARRCompatibility-class*),
11
- version, ArrayDesign-method
(*zzz-DTTCompatibility-class*),
12
- version, ArraySetFile-method
(*zzz-ARRCompatibility-class*),
11
- version, Database-method
(*zzz-DTTCompatibility-class*),
12
- version, PhysicalArrayDesign-method
(*zzz-DTTCompatibility-class*),
12
- version, SoftwareApplication-method
(*zzz-DTTCompatibility-class*),
12
- version, TemplateFile-method
(*zzz-ARRCompatibility-class*),
11
- volume
(*zzz-DTTCompatibility-class*),
12
- volume, BibliographicReference-method
(*zzz-DTTCompatibility-class*),
12
- volumeUnit
(*zzz-DTTCompatibility-class*),

- 12
- volumeUnit, Unit_assn-method
(zzz-DTTCompatibility-class),
12
- warningType_assn
(zzz-DTTCompatibility-class),
12
- warningType_assn, Reporter-method
(zzz-DTTCompatibility-class),
12
- width
(zzz-DTTCompatibility-class),
12
- width, ArrayGroup-method
(zzz-DTTCompatibility-class),
12
- x(zzz-DTTCompatibility-class), 12
- x, Position-method
(zzz-DTTCompatibility-class),
12
- xclass, 7, 9, 10
- xmlValue.XMLAttributeValue, 15
- y(zzz-DTTCompatibility-class), 12
- y, Position-method
(zzz-DTTCompatibility-class),
12
- year
(zzz-DTTCompatibility-class),
12
- year, BibliographicReference-method
(zzz-DTTCompatibility-class),
12
- zone
(zzz-DTTCompatibility-class),
12
- zone, ZoneLocations_assnlist-method
(zzz-DTTCompatibility-class),
12
- zone_assnref
(zzz-DTTCompatibility-class),
12
- zone_assnref, Feature-method
(zzz-DTTCompatibility-class),
12
- zone_assnref, ZoneDefect-method
(zzz-DTTCompatibility-class),
12
- zone_ref
(zzz-DTTCompatibility-class),
12
- zone_ref, Zone_assnref-method
(zzz-DTTCompatibility-class),
12
- zoneDefect
(zzz-DTTCompatibility-class),
12
- zoneDefect, Adjustments_assnlist-method
(zzz-DTTCompatibility-class),
12
- zoneGroup
(zzz-DTTCompatibility-class),
12
- zoneGroup, ZoneGroups_assnlist-method
(zzz-DTTCompatibility-class),
12
- zoneGroups_assnlist
(zzz-DTTCompatibility-class),
12
- zoneGroups_assnlist, PhysicalArrayDesign-method
(zzz-DTTCompatibility-class),
12
- zoneLayout
(zzz-DTTCompatibility-class),
12
- zoneLayout, ZoneLayout_assn-method
(zzz-DTTCompatibility-class),
12
- zoneLayout_assn
(zzz-DTTCompatibility-class),
12
- zoneLayout_assn, ZoneGroup-method
(zzz-DTTCompatibility-class),
12
- zoneLocations_assnlist
(zzz-DTTCompatibility-class),
12
- zoneLocations_assnlist, ZoneGroup-method
(zzz-DTTCompatibility-class),
12
- zonesPerX
(zzz-DTTCompatibility-class),
12
- zonesPerX, ZoneGroup-method
(zzz-DTTCompatibility-class),
12
- zonesPerY
(zzz-DTTCompatibility-class),
12
- zonesPerY, ZoneGroup-method
(zzz-DTTCompatibility-class),
12
- zzz-.TypedAssociation-class

(Internal-class), [11](#)
zzz-AffyCompatible-class
(AffyCompatible-class), [1](#)
zzz-ARRCompatibility-class, [11](#)
zzz-DTTCompatibility-class, [12](#)
zzz-NetAffxCompatibility-class,
[13](#)