

AffyCompatible

March 24, 2012

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.

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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 "NetAffx-Compatibility", 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, and PSI files to data frames, and reads FASTA files to a list of description / sequence pairs (see `Biostrings::readFASTA`).

The ... argument is passed to corresponding read functions. CSV files are parsed using `read.csv(conn, ...)`, Tabular files with `read.delim(conn, ...)`, PSI files with `read.delim(conn, header=FALSE, skip=1, sep="\t", ...)`, and FASTA with `readFASTA(conn, ...)`.

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

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

Author(s)

Martin Morgan <mtmorgan@fhcrc.org>

See Also

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

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

Examples

```
## password must be obtained by the user
rsrc <- NetAffxResource(user="mtmorgan@fhcrc.org", password=password)
head(names(rsrc))
affxDescription(rsrc[["Bovine"]])
## paradigm to display all elements of 'annotation'
sapply(affxAnnotation(rsrc[["Bovine"]]), force)
## obtain and read a single annotation
anno <- rsrc[["Porcine", "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(R) Command Console. Objects are represented as R classes. Navigation and extraction of data are described in the vignette 'Retrieving MAGE and ARR sample attributes'.

Usage

```
readArr(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	<i>Update or read a NetAffx data base</i>
-------------	---

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>
---------	--

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(R) 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 "NetAffx-Compatibility", 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, recursive = TRUE, encoding)
```

Arguments

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

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-method
 (*Internal-class*), [11](#)
 [, .TypedAssociation, numeric, missing, ANY-method
 (*Internal-class*), [11](#)
 [, .TypedAssociation, numeric, missing-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](#)

Accessions_assnlist-class

 (*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](#)

Action_assn-class

 (*zzz-DTTCompatibility-class*),
[12](#)

Measurement_assn

 (*zzz-DTTCompatibility-class*),
[12](#)

- actionMeasurement_assn, Treatment-method (zzz-NetAffxCompatibility-class),
(zzz-DTTCompatibility-class), 12
- ActionMeasurement_assn-class (zzz-DTTCompatibility-class), 12
- activityDate (zzz-DTTCompatibility-class), 12
- activityDate, ProtocolApplication-method (zzz-DTTCompatibility-class), 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
- Adjustments_assnlist-class (zzz-DTTCompatibility-class), 12
- affiliation_assnref (zzz-DTTCompatibility-class), 12
- affiliation_assnref, Person-method (zzz-DTTCompatibility-class), 12
- Affiliation_assnref-class (zzz-DTTCompatibility-class), 12
- affxAnnotation (zzz-NetAffxCompatibility-class), 13
- affxAnnotation, AffxArray-method (zzz-NetAffxCompatibility-class), 13
- AffxAnnotation-class (zzz-NetAffxCompatibility-class), 13
- affxArray (zzz-NetAffxCompatibility-class), 13
- affxArray, AffxNetAffxAnnotFileList-method (zzz-NetAffxCompatibility-class), 13
- affxArray, Array_assnlist-method (zzz-DTTCompatibility-class), 12
- AffxArray-class (zzz-NetAffxCompatibility-class), 13
- 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
- AffxComment-class (zzz-NetAffxCompatibility-class), 13
- 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

13 (zzz-DTTCompatibility-class),
 affxDescription, AffxAnnotation-method 12
 (zzz-NetAffxCompatibility-class), affxOrder, BioDataCube-method
 13 (zzz-DTTCompatibility-class),
 affxDescription, AffxArray-method 12
 (zzz-NetAffxCompatibility-class), affxOrder, Treatment-method
 13 (zzz-DTTCompatibility-class),
 affxEnd 12
 (zzz-DTTCompatibility-class), affxPerson
 12 (zzz-DTTCompatibility-class),
 affxEnd, CompositePosition-method 12
 (zzz-DTTCompatibility-class), affxPerson, Contact_assnlist-method
 12 (zzz-DTTCompatibility-class),
 affxEnd, ReporterPosition-method 12
 (zzz-DTTCompatibility-class), affxPerson, Person_assnlist-method
 12 (zzz-DTTCompatibility-class),
 affxEnd, SequencePosition-method 12
 (zzz-DTTCompatibility-class), affxRow
 12 (zzz-DTTCompatibility-class),
 affxFile 12
 (zzz-NetAffxCompatibility-class), affxRow, FeatureLocation-method
 13 (zzz-DTTCompatibility-class),
 affxFile, AffxAnnotation-method 12
 (zzz-NetAffxCompatibility-class), affxRow, Zone-method
 13 (zzz-DTTCompatibility-class),
 AffxFile-class 12
 (zzz-NetAffxCompatibility-class), affxSequence
 13 (zzz-DTTCompatibility-class),
 affxImage 12
 (zzz-DTTCompatibility-class), affxSequence, BioSequence-method
 12 (zzz-DTTCompatibility-class),
 affxImage, PhysicalBioAssayData_assnlist-method 12
 (zzz-DTTCompatibility-class), affxSize
 12 (zzz-NetAffxCompatibility-class),
 affxLicence 13
 (zzz-NetAffxCompatibility-class), affxSize, AffxFile-method
 13 (zzz-NetAffxCompatibility-class),
 affxLicence, NetAffxResource-method 13
 (NetAffxResource-class), 2 affxSize, AffxUrl-method
 affxName (zzz-NetAffxCompatibility-class),
 (zzz-NetAffxCompatibility-class), 13
 13 affxStart
 affxName, AffxArray-method (zzz-DTTCompatibility-class),
 (zzz-NetAffxCompatibility-class), 12
 13 affxStart, CompositePosition-method
 affxName, AffxFile-method (zzz-DTTCompatibility-class),
 (zzz-NetAffxCompatibility-class), 12
 13 affxStart, ReporterPosition-method
 AffxNetAffxAnnotFileList-class (zzz-DTTCompatibility-class),
 (zzz-NetAffxCompatibility-class), 12
 13 affxStart, SequencePosition-method
 affxOrder (zzz-DTTCompatibility-class),

- 12
 affxStatus
 (*zzz-NetAffxCompatibility-class*),
 13
 affxStatus, *AffxNetAffxAnnotFileList*-method (*AffyCompatible-package*), 2
 (*zzz-NetAffxCompatibility-class*),
 13
AffxStatus-class
 (*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
 affxUrl
 (*zzz-NetAffxCompatibility-class*),
 13
 affxUrl, *AffxFile*-method
 (*zzz-NetAffxCompatibility-class*),
 13
 affxUrl, *NetAffxResource*-method
 (*NetAffxResource-class*), 2
AffxUrl-class
 (*zzz-NetAffxCompatibility-class*),
 13
 affyBarcode
 (*zzz-ARRCompatibility-class*),
 11
 affyBarcode, *PhysicalArray*-method
 (*zzz-ARRCompatibility-class*),
 11
 (*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
AnalysisResults_assnreflist-class
 (*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
Annotations_assnlist-class
 (*zzz-DTTCompatibility-class*),
 12
annotationsFile (accessors), 6
annotationsFile, *NetAffxResource*-method
 (*NetAffxResource-class*), 2
Array-class
 (*zzz-DTTCompatibility-class*),
 12
array_assnlist
 (*zzz-DTTCompatibility-class*),
 12
array_assnlist, *Array-package*-method
 (*zzz-DTTCompatibility-class*),
 12
Array_assnlist-class
 (*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

- Array_assnref-class (zzz-DTTCompatibility-class), 12
- array_package (zzz-DTTCompatibility-class), 12
- array_package, MAGE-ML-method (zzz-DTTCompatibility-class), 12
- Array_package-class (zzz-DTTCompatibility-class), 12
- array_ref (zzz-DTTCompatibility-class), 12
- array_ref, Array_assnref-method (zzz-DTTCompatibility-class), 12
- array_ref, Arrays_assnreflist-method (zzz-DTTCompatibility-class), 12
- Array_ref-class (zzz-DTTCompatibility-class), 12
- arrayAttribute (zzz-ARRCompatibility-class), 11
- arrayAttribute, PhysicalArray-method (zzz-ARRCompatibility-class), 11
- ArrayAttribute-class (zzz-ARRCompatibility-class), 11
- arrayDesign (zzz-DTTCompatibility-class), 12
- arrayDesign, ArrayDesign_assnlist-method (zzz-DTTCompatibility-class), 12
- ArrayDesign-class (zzz-DTTCompatibility-class), 12
- arrayDesign_assnlist (zzz-DTTCompatibility-class), 12
- arrayDesign_assnlist, ArrayDesign_package-method (zzz-DTTCompatibility-class), 12
- ArrayDesign_assnlist-class (zzz-DTTCompatibility-class), 12
- arrayDesign_assnref (zzz-DTTCompatibility-class), 12
- arrayDesign_assnref, Array-method (zzz-DTTCompatibility-class), 12
- ArrayDesign_assnref-class (zzz-DTTCompatibility-class), 12
- arrayDesign_assnref, ArrayDesign_assnref-method (zzz-DTTCompatibility-class), 12
- ArrayDesign_ref-class (zzz-DTTCompatibility-class), 12
- arrayDesign_ref (zzz-DTTCompatibility-class), 12
- arrayDesign_ref, ArrayDesign_assnref-method (zzz-DTTCompatibility-class), 12
- ArrayDesign_ref-class (zzz-DTTCompatibility-class), 12
- arrayGroup (zzz-DTTCompatibility-class), 12
- arrayGroup, ArrayGroup_assnlist-method (zzz-DTTCompatibility-class), 12
- ArrayGroup-class (zzz-DTTCompatibility-class), 12
- arrayGroup_assnlist (zzz-DTTCompatibility-class), 12
- arrayGroup_assnlist, Array_package-method (zzz-DTTCompatibility-class), 12
- ArrayGroup_assnlist-class (zzz-DTTCompatibility-class), 12
- arrayGroup_assnref (zzz-DTTCompatibility-class), 12
- arrayGroup_assnref, Array-method (zzz-DTTCompatibility-class), 12
- ArrayGroup_assnref-class (zzz-DTTCompatibility-class), 12

- 12
- arrayGroup_ref
 - (zzz-DTTCompatibility-class), 12
- arrayGroup_ref, ArrayGroup_assnref-method
 - (zzz-DTTCompatibility-class), 12
- ArrayGroup_ref-class
 - (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-class
 - (zzz-DTTCompatibility-class), 12
- arrayManufacture_assnlist
 - (zzz-DTTCompatibility-class), 12
- arrayManufacture_assnlist, Array_package-method
 - (zzz-DTTCompatibility-class), 12
- ArrayManufacture_assnlist-class
 - (zzz-DTTCompatibility-class), 12
- arrayManufacture_ref
 - (zzz-DTTCompatibility-class), 12
- arrayManufacture_ref, Information_assnref-method
 - (zzz-DTTCompatibility-class), 12
- ArrayManufacture_ref-class
 - (zzz-DTTCompatibility-class), 12
- arrayManufactureDeviation
 - (zzz-DTTCompatibility-class), 12
- arrayManufactureDeviation, ArrayManufactureDeviation_assnlist-method
 - (zzz-DTTCompatibility-class), 12
- ArrayManufactureDeviation-class
 - (zzz-DTTCompatibility-class), 12
- arrayManufactureDeviations_assnlist
 - (zzz-DTTCompatibility-class), 12
- arrayManufactureDeviations_assnlist, Array-method
 - (zzz-DTTCompatibility-class), 12
- ArrayManufactureDeviations_assnlist-class
 - (zzz-DTTCompatibility-class), 12
- arrayManufacturers_assnreflist
 - (zzz-DTTCompatibility-class), 12
- arrayManufacturers_assnreflist, ArrayManufacturers-assnreflist-method
 - (zzz-DTTCompatibility-class), 12
- ArrayManufacturers_assnreflist-class
 - (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
- Arrays_assnreflist-class
 - (zzz-DTTCompatibility-class), 12
- ArraySetFile-class
 - (zzz-ARRCompatibility-class), 11
- 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
- Associations_assnlist-class
 - (*zzz-DTTCompatibility-class*),
12
- audit
 - (*zzz-DTTCompatibility-class*),
12
- audit, AuditTrail_assnlist-method
 - (*zzz-DTTCompatibility-class*),
12
- Audit-class
 - (*zzz-DTTCompatibility-class*),
12
- auditAndSecurity_package
 - (*zzz-DTTCompatibility-class*),
12
- auditAndSecurity_package, MAGE-ML-method
 - (*zzz-DTTCompatibility-class*),
12
- AuditAndSecurity_package-class
 - (*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
- 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
- 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	(zzz-DTTCompatibility-class), 12
auditTrail_assnlist, DerivedBioAssay-method (zzz-DTTCompatibility-class), 12	auditTrail_assnlist, Hardware-method (zzz-DTTCompatibility-class), 12
auditTrail_assnlist, DerivedBioAssayData-method (zzz-DTTCompatibility-class), 12	auditTrail_assnlist, HardwareApplication-method (zzz-DTTCompatibility-class), 12
auditTrail_assnlist, DerivedSignal-method (zzz-DTTCompatibility-class), 12	auditTrail_assnlist, Hybridization-method (zzz-DTTCompatibility-class), 12
auditTrail_assnlist, Description-method (zzz-DTTCompatibility-class), 12	auditTrail_assnlist, Image-method (zzz-DTTCompatibility-class), 12
auditTrail_assnlist, Error-method (zzz-DTTCompatibility-class), 12	auditTrail_assnlist, ImageAcquisition-method (zzz-DTTCompatibility-class), 12
auditTrail_assnlist, ExpectedValue-method (zzz-DTTCompatibility-class), 12	auditTrail_assnlist, LabeledExtract-method (zzz-DTTCompatibility-class), 12
auditTrail_assnlist, Experiment-method (zzz-DTTCompatibility-class), 12	auditTrail_assnlist, MAGE-ML-method (zzz-DTTCompatibility-class), 12
auditTrail_assnlist, ExperimentalFactor-method (zzz-DTTCompatibility-class), 12	auditTrail_assnlist, ManufactureLIMS-method (zzz-DTTCompatibility-class), 12
auditTrail_assnlist, ExperimentDesign-method (zzz-DTTCompatibility-class), 12	auditTrail_assnlist, ManufactureLIMSBiomaterial-method (zzz-DTTCompatibility-class), 12
auditTrail_assnlist, FactorValue-method (zzz-DTTCompatibility-class), 12	auditTrail_assnlist, MeasuredBioAssay-method (zzz-DTTCompatibility-class), 12
auditTrail_assnlist, Failed-method (zzz-DTTCompatibility-class), 12	auditTrail_assnlist, MeasuredBioAssayData-method (zzz-DTTCompatibility-class), 12
auditTrail_assnlist, Feature-method (zzz-DTTCompatibility-class), 12	auditTrail_assnlist, MeasuredSignal-method (zzz-DTTCompatibility-class), 12
auditTrail_assnlist, FeatureDimension-method (zzz-DTTCompatibility-class), 12	auditTrail_assnlist, Node-method (zzz-DTTCompatibility-class), 12
auditTrail_assnlist, FeatureExtraction-method (zzz-DTTCompatibility-class), 12	auditTrail_assnlist, NodeContents-method (zzz-DTTCompatibility-class), 12
auditTrail_assnlist, FeatureGroup-method (zzz-DTTCompatibility-class), 12	auditTrail_assnlist, Organization-method (zzz-DTTCompatibility-class), 12
auditTrail_assnlist, FeatureReporterMap-method (zzz-DTTCompatibility-class), 12	auditTrail_assnlist, Parameter-method (zzz-DTTCompatibility-class), 12
auditTrail_assnlist, Fiducial-method	auditTrail_assnlist, Person-method (zzz-DTTCompatibility-class),

[12](#) [auditTrail_assnlist, SoftwareApplication-method](#)
[auditTrail_assnlist, PhysicalArrayDesign-method](#) ([zzz-DTTCompatibility-class](#)),
[\(zzz-DTTCompatibility-class\)](#), [12](#)
[12](#) [auditTrail_assnlist, SpecializedQuantitationType](#)
[auditTrail_assnlist, PhysicalBioAssay-method](#) ([zzz-DTTCompatibility-class](#)),
[\(zzz-DTTCompatibility-class\)](#), [12](#)
[12](#) [auditTrail_assnlist, Transformation-method](#)
[auditTrail_assnlist, PresentAbsent-method](#) ([zzz-DTTCompatibility-class](#)),
[\(zzz-DTTCompatibility-class\)](#), [12](#)
[12](#) [auditTrail_assnlist, Treatment-method](#)
[auditTrail_assnlist, Protocol-method](#) ([zzz-DTTCompatibility-class](#)),
[\(zzz-DTTCompatibility-class\)](#), [12](#)
[12](#) [auditTrail_assnlist, Zone-method](#)
[auditTrail_assnlist, ProtocolApplication-method](#) ([zzz-DTTCompatibility-class](#)),
[\(zzz-DTTCompatibility-class\)](#), [12](#)
[12](#) [AuditTrail_assnlist-class](#)
[auditTrail_assnlist, PValue-method](#) ([zzz-DTTCompatibility-class](#)),
[\(zzz-DTTCompatibility-class\)](#), [12](#)
[12](#) [authors](#)
[auditTrail_assnlist, QuantitationTypeDimension-method](#) ([zzz-DTTCompatibility-class](#)),
[\(zzz-DTTCompatibility-class\)](#), [12](#)
[12](#) [authors, BibliographicReference-method](#)
[auditTrail_assnlist, QuantitationTypeMap-method](#) ([zzz-DTTCompatibility-class](#)),
[\(zzz-DTTCompatibility-class\)](#), [12](#)
[12](#)
[auditTrail_assnlist, Ratio-method](#) [barcode](#)
[\(zzz-DTTCompatibility-class\)](#), [12](#)
[12](#) [barcode, ArrayGroup-method](#)
[auditTrail_assnlist, Reporter-method](#) ([zzz-DTTCompatibility-class](#)),
[\(zzz-DTTCompatibility-class\)](#), [12](#)
[12](#)
[auditTrail_assnlist, ReporterCompositeMap-method](#) [basis](#)
[\(zzz-DTTCompatibility-class\)](#), [12](#)
[12](#) [basis, SeqFeature-method](#)
[auditTrail_assnlist, ReporterDimension-method](#) ([zzz-DTTCompatibility-class](#)),
[\(zzz-DTTCompatibility-class\)](#), [12](#)
[12](#)
[auditTrail_assnlist, ReporterGroup-method](#) [bibliographicReference](#)
[\(zzz-DTTCompatibility-class\)](#), [12](#)
[12](#) [bibliographicReference, BibliographicReferences](#)
[auditTrail_assnlist, Security-method](#) ([zzz-DTTCompatibility-class](#)),
[\(zzz-DTTCompatibility-class\)](#), [12](#)
[12](#) [BibliographicReference-class](#)
[auditTrail_assnlist, SecurityGroup-method](#) ([zzz-DTTCompatibility-class](#)),
[\(zzz-DTTCompatibility-class\)](#), [12](#)
[12](#) [bibliographicReferences_assnlist](#)
[auditTrail_assnlist, SeqFeature-method](#) ([zzz-DTTCompatibility-class](#)),
[\(zzz-DTTCompatibility-class\)](#), [12](#)
[12](#) [bibliographicReferences_assnlist, Description-r](#)
[auditTrail_assnlist, Software-method](#) ([zzz-DTTCompatibility-class](#)),
[\(zzz-DTTCompatibility-class\)](#), [12](#)
[12](#) [BibliographicReferences_assnlist-class](#)

- (zzz-DTTCompatibility-class),
12
- bioAssay_assnlist
(zzz-DTTCompatibility-class),
12
- bioAssay_assnlist, BioAssay_package-method
(zzz-DTTCompatibility-class),
12
- BioAssay_assnlist-class
(zzz-DTTCompatibility-class),
12
- bioAssay_assnref
(zzz-DTTCompatibility-class),
12
- bioAssay_assnref, BioAssayTuple-method
(zzz-DTTCompatibility-class),
12
- BioAssay_assnref-class
(zzz-DTTCompatibility-class),
12
- bioAssay_package
(zzz-DTTCompatibility-class),
12
- bioAssay_package, MAGE-ML-method
(zzz-DTTCompatibility-class),
12
- BioAssay_package-class
(zzz-DTTCompatibility-class),
12
- BioAssay_ref-class
(zzz-DTTCompatibility-class),
12
- bioAssayCreation
(zzz-DTTCompatibility-class),
12
- bioAssayCreation, BioAssayCreation_assn-method
(zzz-DTTCompatibility-class),
12
- BioAssayCreation-class
(zzz-DTTCompatibility-class),
12
- bioAssayCreation_assn
(zzz-DTTCompatibility-class),
12
- bioAssayCreation_assn, PhysicalBioAssay-method
(zzz-DTTCompatibility-class),
12
- BioAssayCreation_assn-class
(zzz-DTTCompatibility-class),
12
- bioAssayData_assnlist
(zzz-DTTCompatibility-class),
12
- bioAssayData_assnlist, BioAssayData_package-method
(zzz-DTTCompatibility-class),
12
- BioAssayData_assnlist-class
(zzz-DTTCompatibility-class),
12
- bioAssayData_assnreflist
(zzz-DTTCompatibility-class),
12
- bioAssayData_assnreflist, Experiment-method
(zzz-DTTCompatibility-class),
12
- BioAssayData_assnreflist-class
(zzz-DTTCompatibility-class),
12
- bioAssayData_package
(zzz-DTTCompatibility-class),
12
- bioAssayData_package, MAGE-ML-method
(zzz-DTTCompatibility-class),
12
- BioAssayData_package-class
(zzz-DTTCompatibility-class),
12
- BioAssayData_ref-class
(zzz-DTTCompatibility-class),
12
- bioAssayDataCluster
(zzz-DTTCompatibility-class),
12
- bioAssayDataCluster, BioAssayDataCluster_assnlist
(zzz-DTTCompatibility-class),
12
- BioAssayDataCluster-class
(zzz-DTTCompatibility-class),
12
- bioAssayDataCluster_assnlist
(zzz-DTTCompatibility-class),
12
- bioAssayDataCluster_assnlist, HigherLevelAnalysis
(zzz-DTTCompatibility-class),
12
- BioAssayDataCluster_assnlist-class
(zzz-DTTCompatibility-class),
12
- bioAssayDataCluster_ref
(zzz-DTTCompatibility-class),
12
- bioAssayDataCluster_ref, AnalysisResults_assnref
(zzz-DTTCompatibility-class),
12

BioAssayDataCluster_ref-class	(zzz-DTTCompatibility-class), 12	(zzz-DTTCompatibility-class), 12
bioAssayDataSources_assnreflist	(zzz-DTTCompatibility-class), 12	bioAssayFactorValues_assnreflist (zzz-DTTCompatibility-class), 12
bioAssayDataSources_assnreflist,Transformation-method	(zzz-DTTCompatibility-class), 12	bioAssayFactorValues_assnreflist,DerivedBioAssayDataCluster_ref-class (zzz-DTTCompatibility-class), 12
BioAssayDataSources_assnreflist-class	(zzz-DTTCompatibility-class), 12	bioAssayFactorValues_assnreflist,MeasuredBioAssayDataCluster_ref-class (zzz-DTTCompatibility-class), 12
bioAssayDimension	(zzz-DTTCompatibility-class), 12	bioAssayFactorValues_assnreflist,PhysicalBioAssayDataCluster_ref-class (zzz-DTTCompatibility-class), 12
bioAssayDimension,BioAssayDimension_assnlist-method	(zzz-DTTCompatibility-class), 12	BioAssayFactorValues_assnreflist-class (zzz-DTTCompatibility-class), 12
BioAssayDimension-class	(zzz-DTTCompatibility-class), 12	bioAssayMap (zzz-DTTCompatibility-class), 12
bioAssayDimension_assnlist	(zzz-DTTCompatibility-class), 12	bioAssayMap,BioAssayMap_assnlist-method (zzz-DTTCompatibility-class), 12
bioAssayDimension_assnlist,BioAssayData_package-method	(zzz-DTTCompatibility-class), 12	BioAssayMap-class (zzz-DTTCompatibility-class), 12
BioAssayDimension_assnlist-class	(zzz-DTTCompatibility-class), 12	bioAssayMap_assnlist (zzz-DTTCompatibility-class), 12
bioAssayDimension_assnref	(zzz-DTTCompatibility-class), 12	bioAssayMap_assnlist,BioAssayData_package-method (zzz-DTTCompatibility-class), 12
bioAssayDimension_assnref,DerivedBioAssayDataCluster_ref-class	(zzz-DTTCompatibility-class), 12	BioAssayMap_assnlist-class (zzz-DTTCompatibility-class), 12
bioAssayDimension_assnref,MeasuredBioAssayDataCluster_ref-class	(zzz-DTTCompatibility-class), 12	bioAssayMap_ref (zzz-DTTCompatibility-class), 12
bioAssayDimension_assnref,NodeContents-method	(zzz-DTTCompatibility-class), 12	bioAssayMap_ref,BioAssayMaps_assnreflist-method (zzz-DTTCompatibility-class), 12
BioAssayDimension_assnref-class	(zzz-DTTCompatibility-class), 12	bioAssayMap_ref,DerivedBioAssayMap_assnreflist (zzz-DTTCompatibility-class), 12
bioAssayDimension_ref	(zzz-DTTCompatibility-class), 12	BioAssayMap_ref-class (zzz-DTTCompatibility-class), 12
bioAssayDimension_ref,BioAssayDimension_assnlist-method	(zzz-DTTCompatibility-class), 12	bioAssayMapping (zzz-DTTCompatibility-class), 12
BioAssayDimension_ref-class	(zzz-DTTCompatibility-class), 12	bioAssayMapping,BioAssayMapping_assn-method (zzz-DTTCompatibility-class), 12

- 12
 BioAssayMapping-class
 (*zzz-DTTCompatibility-class*),
 12
 bioAssayMapping_assn
 (*zzz-DTTCompatibility-class*),
 12
 bioAssayMapping_assn, Transformation-method (*zzz-DTTCompatibility-class*),
 12
 BioAssayMapping_assn-class
 (*zzz-DTTCompatibility-class*),
 12
 bioAssayMaps_assnreflist
 (*zzz-DTTCompatibility-class*),
 12
 bioAssayMaps_assnreflist, BioAssayMapping-method (*zzz-DTTCompatibility-class*),
 12
 BioAssayMaps_assnreflist-class
 (*zzz-DTTCompatibility-class*),
 12
 bioAssayMapTarget_assnref
 (*zzz-DTTCompatibility-class*),
 12
 bioAssayMapTarget_assnref, BioAssayMap-method (*zzz-DTTCompatibility-class*),
 12
 BioAssayMapTarget_assnref-class
 (*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
 BioAssays_assnreflist-class
 (*zzz-DTTCompatibility-class*),
 12
 bioAssayTreatment
 (*zzz-DTTCompatibility-class*),
 12
 bioAssayTreatment, BioAssayTreatments_assnlist-method (*zzz-DTTCompatibility-class*),
 12
 BioAssayTreatment-class
 (*zzz-DTTCompatibility-class*),
 12
 bioAssayTreatments_assnlist
 (*zzz-DTTCompatibility-class*),
 12
 bioAssayTreatments_assnlist, PhysicalBioAssay-method (*zzz-DTTCompatibility-class*),
 12
 BioAssayTreatments_assnlist-class
 (*zzz-DTTCompatibility-class*),
 12
 bioAssayTuple
 (*zzz-DTTCompatibility-class*),
 12
 bioAssayTuple, BioAssayTuples_assnlist-method (*zzz-DTTCompatibility-class*),
 12
 BioAssayTuple-class
 (*zzz-DTTCompatibility-class*),
 12
 bioAssayTuples_assnlist
 (*zzz-DTTCompatibility-class*),
 12
 bioAssayTuples_assnlist, BioDataTuples-method (*zzz-DTTCompatibility-class*),
 12
 BioAssayTuples_assnlist-class
 (*zzz-DTTCompatibility-class*),
 12
 bioDataCube
 (*zzz-DTTCompatibility-class*),
 12
 bioDataCube, BioDataValues_assn-method (*zzz-DTTCompatibility-class*),
 12
 BioDataCube-class
 (*zzz-DTTCompatibility-class*),
 12
 bioDataTuples
 (*zzz-DTTCompatibility-class*),
 12
 bioDataTuples, BioDataValues_assn-method (*zzz-DTTCompatibility-class*),
 12
 BioDataTuples-class
 (*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
- BioDataValues_assn-class
(zzz-*DTTCompatibility-class*),
12
- bioEvent_package
(zzz-*DTTCompatibility-class*),
12
- bioEvent_package, MAGE-ML-method
(zzz-*DTTCompatibility-class*),
12
- BioEvent_package-class
(zzz-*DTTCompatibility-class*),
12
- biologicalCharacteristics_assnreflist
(zzz-*DTTCompatibility-class*),
12
- biologicalCharacteristics_assnreflist, ComposedOfBioMaterialMethod
(zzz-*DTTCompatibility-class*),
12
- BiologicalCharacteristics_assnreflist-class
(zzz-*DTTCompatibility-class*),
12
- bioMaterial_assnlist
(zzz-*DTTCompatibility-class*),
12
- bioMaterial_assnlist, BioMaterial_package-method
(zzz-*DTTCompatibility-class*),
12
- BioMaterial_assnlist-class
(zzz-*DTTCompatibility-class*),
12
- bioMaterial_assnref
(zzz-*DTTCompatibility-class*),
12
- bioMaterial_assnref, BioMaterialMeasurement-method
(zzz-*DTTCompatibility-class*),
12
- bioMaterial_assnref, ManufactureLIMS-method
(zzz-*DTTCompatibility-class*),
12
- bioMaterial_assnref, ManufactureLIMSBiomaterial-method
(zzz-*DTTCompatibility-class*),
12
- BioMaterial_assnref-class
(zzz-*DTTCompatibility-class*),
12
- bioMaterial_package
(zzz-*DTTCompatibility-class*),
12
- bioMaterial_package, MAGE-ML-method
(zzz-*DTTCompatibility-class*),
12
- BioMaterial_package-class
(zzz-*DTTCompatibility-class*),
12
- BioMaterial_ref-class
(zzz-*DTTCompatibility-class*),
12
- bioMaterialMeasurement
(zzz-*DTTCompatibility-class*),
12
- bioMaterialMeasurement, SourceBioMaterialMeasurement
(zzz-*DTTCompatibility-class*),
12
- BioMaterialMeasurement-class
(zzz-*DTTCompatibility-class*),
12
- bioMaterialPlateCol
(zzz-*DTTCompatibility-class*),
12
- bioMaterialPlateCol, ManufactureLIMSBiomaterial
(zzz-*DTTCompatibility-class*),
12
- bioMaterialPlateIdentifier
(zzz-*DTTCompatibility-class*),
12
- bioMaterialPlateIdentifier, ManufactureLIMSBiomaterial
(zzz-*DTTCompatibility-class*),
12
- bioMaterialPlateRow
(zzz-*DTTCompatibility-class*),
12
- bioMaterialPlateRow, ManufactureLIMSBiomaterial
(zzz-*DTTCompatibility-class*),
12
- bioSample
(zzz-*DTTCompatibility-class*),
12
- bioSample, BioMaterial_assnlist-method
(zzz-*DTTCompatibility-class*),
12
- BioSample-class
(zzz-*DTTCompatibility-class*),
12
- bioSample_ref
(zzz-*DTTCompatibility-class*),
12
- bioSample_ref, BioMaterial_assnref-method
(zzz-*DTTCompatibility-class*),
12
- BioSample_ref-class
(zzz-*DTTCompatibility-class*),
12

- bioSequence (zzz-DTTCompatibility-class), 12
- bioSequence, BioSequence_assnlist-method (zzz-DTTCompatibility-class), 12
- BioSequence-class (zzz-DTTCompatibility-class), 12
- bioSequence_assnlist (zzz-DTTCompatibility-class), 12
- bioSequence_assnlist, BioSequence_package-method (zzz-DTTCompatibility-class), 12
- BioSequence_assnlist-class (zzz-DTTCompatibility-class), 12
- bioSequence_package (zzz-DTTCompatibility-class), 12
- bioSequence_package, MAGE-ML-method (zzz-DTTCompatibility-class), 12
- BioSequence_package-class (zzz-DTTCompatibility-class), 12
- bioSequence_ref (zzz-DTTCompatibility-class), 12
- bioSequence_ref, BiologicalCharacteristics_assnreflist-method (zzz-DTTCompatibility-class), 12
- bioSequence_ref, ImmobilizedCharacteristics_assnreflist-method (zzz-DTTCompatibility-class), 12
- BioSequence_ref-class (zzz-DTTCompatibility-class), 12
- bioSource (zzz-DTTCompatibility-class), 12
- bioSource, BioMaterial_assnlist-method (zzz-DTTCompatibility-class), 12
- BioSource-class (zzz-DTTCompatibility-class), 12
- bioSource_ref (zzz-DTTCompatibility-class), 12
- bioSource_ref, BioMaterial_assnref-method (zzz-DTTCompatibility-class), 12
- (zzz-DTTCompatibility-class), 12
- BioSource_ref-class (zzz-DTTCompatibility-class), 12
- bqs_package (zzz-DTTCompatibility-class), 12
- bqs_package, MAGE-ML-method (zzz-DTTCompatibility-class), 12
- BQS_package-class (zzz-DTTCompatibility-class), 12
- category_assn (zzz-DTTCompatibility-class), 12
- category_assn, ExperimentalFactor-method (zzz-DTTCompatibility-class), 12
- Category_assn-class (zzz-DTTCompatibility-class), 12
- channel (zzz-DTTCompatibility-class), 12
- channel, Channel_assnlist-method (zzz-DTTCompatibility-class), 12
- Channel-class (zzz-DTTCompatibility-class), 12
- channel_assnlist (zzz-DTTCompatibility-class), 12
- channel_assnlist, BioAssay_package-method (zzz-DTTCompatibility-class), 12
- Channel_assnlist-class (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

- 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_assnref-class
(*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
- Channel_ref-class
(*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
- Channels_assnreflist-class
(*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
- Characteristics_assnlist-class
(*zzz-DTTCompatibility-class*),
12
- clusterBioAssayData_assnref
(*zzz-DTTCompatibility-class*),
12
- clusterBioAssayData_assnref, BioAssayDataCluster
(*zzz-DTTCompatibility-class*),
12
- ClusterBioAssayData_assnref-class
(*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
- ComponentCompounds_assnlist-class
(*zzz-DTTCompatibility-class*),
12
- composite_assnref
(*zzz-DTTCompatibility-class*),
12
- composite_assnref, CompositePosition-method
(*zzz-DTTCompatibility-class*),
12
- Composite_assnref-class
(*zzz-DTTCompatibility-class*),
12
- compositeCompositeMap

(zzz-DTTCompatibility-class),	12
12	compositeGroup_assnlist, ArrayDesign_package-me
compositeCompositeMap, CompositeCompositeMap_assnlist-method	(zzz-DTTCompatibility-class),
(zzz-DTTCompatibility-class),	12
12	CompositeGroup_assnlist-class
compositeCompositeMap, DesignElementMap_assnlist-method	(zzz-DTTCompatibility-class),
(zzz-DTTCompatibility-class),	12
12	compositeGroup_ref
CompositeCompositeMap-class	(zzz-DTTCompatibility-class),
(zzz-DTTCompatibility-class),	12
12	compositeGroup_ref, CompositeGroups_assnreflist
compositeCompositeMap_assnlist	(zzz-DTTCompatibility-class),
(zzz-DTTCompatibility-class),	12
12	CompositeGroup_ref-class
compositeCompositeMap_assnlist, DesignElement_package-method	(zzz-DTTCompatibility-class),
(zzz-DTTCompatibility-class),	12
12	compositeGroups_assnreflist
CompositeCompositeMap_assnlist-class	(zzz-DTTCompatibility-class),
(zzz-DTTCompatibility-class),	12
12	compositeGroups_assnreflist, ArrayDesign-method
compositeCompositeMap_ref	(zzz-DTTCompatibility-class),
(zzz-DTTCompatibility-class),	12
12	compositeGroups_assnreflist, PhysicalArrayDesign
compositeCompositeMap_ref, CompositeCompositeMaps_assnreflist-method	(zzz-DTTCompatibility-class),
(zzz-DTTCompatibility-class),	12
12	CompositeGroups_assnreflist-class
compositeCompositeMap_ref, DesignElementMaps_assnreflist-method	(zzz-DTTCompatibility-class),
(zzz-DTTCompatibility-class),	12
12	compositePosition
CompositeCompositeMap_ref-class	(zzz-DTTCompatibility-class),
(zzz-DTTCompatibility-class),	12
12	compositePosition, CompositePositionSources_assn
compositeCompositeMaps_assnreflist	(zzz-DTTCompatibility-class),
(zzz-DTTCompatibility-class),	12
12	compositePosition, Coordinate_assn-method
compositeCompositeMaps_assnreflist, CompositeCompositeMaps_assnreflist-method	(zzz-DTTCompatibility-class),
(zzz-DTTCompatibility-class),	12
12	CompositePosition-class
CompositeCompositeMaps_assnreflist-class	(zzz-DTTCompatibility-class),
(zzz-DTTCompatibility-class),	12
12	compositePositionSources_assnlist
compositeGroup	(zzz-DTTCompatibility-class),
(zzz-DTTCompatibility-class),	12
12	compositePositionSources_assnlist, CompositeCor
compositeGroup, CompositeGroup_assnlist-method	(zzz-DTTCompatibility-class),
(zzz-DTTCompatibility-class),	12
12	CompositePositionSources_assnlist-class
CompositeGroup-class	(zzz-DTTCompatibility-class),
(zzz-DTTCompatibility-class),	12
12	compositeSequence
compositeGroup_assnlist	(zzz-DTTCompatibility-class),
(zzz-DTTCompatibility-class),	12

compositeSequence, CompositeSequence_assnlist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 CompositeSequenceDimension-class
 compositeSequence, DesignElement_assnlist-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 compositeSequenceDimension_ref
 CompositeSequence-class (zzz-DTTCompatibility-class), 12
 compositeSequenceDimension_ref, DesignElementDimension-class (zzz-DTTCompatibility-class), 12
 compositeSequence_assnlist (zzz-DTTCompatibility-class), 12
 CompositeSequenceDimension_ref-class
 compositeSequence_assnlist, DesignElement_package-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 compositeSequences_assnreflist
 CompositeSequence_assnlist-class (zzz-DTTCompatibility-class), 12
 compositeSequences_assnreflist, CompositeGroup-class (zzz-DTTCompatibility-class), 12
 compositeSequence_assnref (zzz-DTTCompatibility-class), 12
 compositeSequences_assnreflist, CompositeSequence-class (zzz-DTTCompatibility-class), 12
 compositeSequence_assnref, CompositeCompositeMap-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 CompositeSequences_assnreflist-class
 compositeSequence_assnref, ReporterCompositeMap-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 compound
 CompositeSequence_assnref-class (zzz-DTTCompatibility-class), 12
 compound, Compound_assnlist-method (zzz-DTTCompatibility-class), 12
 compositeSequence_ref (zzz-DTTCompatibility-class), 12
 Compound-class
 compositeSequence_ref, Composite_assnref-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 compound_assnlist
 compositeSequence_ref, CompositeSequence_assnref-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 compound_assnlist, BioMaterial_package-method (zzz-DTTCompatibility-class), 12
 compositeSequence_ref, CompositeSequences_assnref-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 Compound_assnlist-class
 compositeSequence_ref, DesignElement_assnref-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 compound_assnref
 CompositeSequence_ref-class (zzz-DTTCompatibility-class), 12
 compound_assnref, CompoundMeasurement-method (zzz-DTTCompatibility-class), 12
 compositeSequenceDimension (zzz-DTTCompatibility-class), 12
 Compound_assnref-class
 compositeSequenceDimension, DesignElementDimension_method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12

- 12
- compound_ref
(*zzz-DTTCompatibility-class*),
12
- compound_ref,Compound_assnref-method
(*zzz-DTTCompatibility-class*),
12
- compound_ref,Labels_assnreflist-method
(*zzz-DTTCompatibility-class*),
12
- Compound_ref-class
(*zzz-DTTCompatibility-class*),
12
- compoundIndices_assnlist
(*zzz-DTTCompatibility-class*),
12
- compoundIndices_assnlist,Compound-method
(*zzz-DTTCompatibility-class*),
12
- CompoundIndices_assnlist-class
(*zzz-DTTCompatibility-class*),
12
- compoundMeasurement
(*zzz-DTTCompatibility-class*),
12
- compoundMeasurement,ComponentCompounds_assnlist-method
(*zzz-DTTCompatibility-class*),
12
- compoundMeasurement,CompoundMeasurements_assnlist-method
(*zzz-DTTCompatibility-class*),
12
- CompoundMeasurement-class
(*zzz-DTTCompatibility-class*),
12
- compoundMeasurements_assnlist
(*zzz-DTTCompatibility-class*),
12
- compoundMeasurements_assnlist,Treatment-method
(*zzz-DTTCompatibility-class*),
12
- CompoundMeasurements_assnlist-class
(*zzz-DTTCompatibility-class*),
12
- concentrationUnit
(*zzz-DTTCompatibility-class*),
12
- concentrationUnit,Unit_assn-method
(*zzz-DTTCompatibility-class*),
12
- ConcentrationUnit-class
(*zzz-DTTCompatibility-class*),
12
- ConfidenceIndicator_assnlist-class
(*zzz-DTTCompatibility-class*),
12
- ConfidenceIndicator_ref-class
(*zzz-DTTCompatibility-class*),
12
- confidenceIndicators_assnreflist
(*zzz-DTTCompatibility-class*),
12
- confidenceIndicators_assnreflist,DerivedSignal
(*zzz-DTTCompatibility-class*),
12
- confidenceIndicators_assnreflist,Error-method
(*zzz-DTTCompatibility-class*),
12
- confidenceIndicators_assnreflist,ExpectedValue
(*zzz-DTTCompatibility-class*),
12
- confidenceIndicators_assnreflist,Failed-method
(*zzz-DTTCompatibility-class*),
12
- confidenceIndicators_assnreflist,MeasuredSignal
(*zzz-DTTCompatibility-class*),
12
- confidenceIndicators_assnreflist,PresentAbsent
(*zzz-DTTCompatibility-class*),
12
- confidenceIndicators_assnreflist,PValue-method
(*zzz-DTTCompatibility-class*),
12
- confidenceIndicators_assnreflist,Ratio-method
(*zzz-DTTCompatibility-class*),
12
- confidenceIndicators_assnreflist,SpecializedQu
(*zzz-DTTCompatibility-class*),
12
- ConfidenceIndicators_assnreflist-class
(*zzz-DTTCompatibility-class*),
12
- contact_assnlist
(*zzz-DTTCompatibility-class*),
12
- contact_assnlist,AuditAndSecurity_package-meth
(*zzz-DTTCompatibility-class*),
12
- Contact_assnlist-class
(*zzz-DTTCompatibility-class*),
12
- Contact_ref-class
(*zzz-DTTCompatibility-class*),
12
- contacts_assnreflist

- (zzz-*DTTCompatibility-class*), 12
- contacts_assnreflist, Database-method (zzz-*DTTCompatibility-class*), 12
- Contacts_assnreflist-class (zzz-*DTTCompatibility-class*), 12
- containedFeatures_assnreflist (zzz-*DTTCompatibility-class*), 12
- containedFeatures_assnreflist, FeatureDimension-method (zzz-*DTTCompatibility-class*), 12
- ContainedFeatures_assnreflist-class (zzz-*DTTCompatibility-class*), 12
- control (zzz-*ARRCompatibility-class*), 11
- control, UserAttribute-method (zzz-*ARRCompatibility-class*), 11
- Control-class (zzz-*ARRCompatibility-class*), 11
- controlFeatures_assnreflist (zzz-*DTTCompatibility-class*), 12
- controlFeatures_assnreflist, Feature-method (zzz-*DTTCompatibility-class*), 12
- ControlFeatures_assnreflist-class (zzz-*DTTCompatibility-class*), 12
- controlledFeatures_assnreflist (zzz-*DTTCompatibility-class*), 12
- controlledFeatures_assnreflist, Feature-method (zzz-*DTTCompatibility-class*), 12
- ControlledFeatures_assnreflist-class (zzz-*DTTCompatibility-class*), 12
- controlType_assn (zzz-*DTTCompatibility-class*), 12
- controlType_assn, CompositeSequence-method (zzz-*DTTCompatibility-class*), 12
- controlType_assn, Feature-method (zzz-*DTTCompatibility-class*), 12
- controlType_assn, Reporter-method (zzz-*DTTCompatibility-class*), 12
- ControlType_assn-class (zzz-*DTTCompatibility-class*), 12
- coordinate_assn (zzz-*DTTCompatibility-class*), 12
- coordinate_assn, SeqFeatureLocation-method (zzz-*DTTCompatibility-class*), 12
- Coordinate_assn-class (zzz-*DTTCompatibility-class*), 12
- createdBy (zzz-*ARRCompatibility-class*), 11
- createdBy, ArraySetFile-method (zzz-*ARRCompatibility-class*), 11
- createdBy, PhysicalArray-method (zzz-*ARRCompatibility-class*), 11
- createdBy, TemplateFile-method (zzz-*ARRCompatibility-class*), 11
- createdDateTime (zzz-*ARRCompatibility-class*), 11
- createdDateTime, ArraySetFile-method (zzz-*ARRCompatibility-class*), 11
- createdDateTime, PhysicalArray-method (zzz-*ARRCompatibility-class*), 11
- createdDateTime, TemplateFile-method (zzz-*ARRCompatibility-class*), 11
- createdStep (zzz-*ARRCompatibility-class*), 11
- createdStep, ArraySetFile-method (zzz-*ARRCompatibility-class*), 11
- createdStep, PhysicalArray-method (zzz-*ARRCompatibility-class*), 11
- database (zzz-*DTTCompatibility-class*), 12

- database, Database_assnlist-method (zzz-DTTCompatibility-class), 12
 Database-class (zzz-DTTCompatibility-class), 12
 database_assnlist (zzz-DTTCompatibility-class), 12
 database_assnlist, Description_package-method(zzz-DTTCompatibility-class), 12
 Database_assnlist-class (zzz-DTTCompatibility-class), 12
 database_assnref (zzz-DTTCompatibility-class), 12
 database_assnref, DatabaseEntry-method (zzz-DTTCompatibility-class), 12
 Database_assnref-class (zzz-DTTCompatibility-class), 12
 database_ref (zzz-DTTCompatibility-class), 12
 database_ref, Database_assnref-method (zzz-DTTCompatibility-class), 12
 Database_ref-class (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
 DatabaseEntry-class (zzz-DTTCompatibility-class), 12
 databaseReferences_assnlist (zzz-DTTCompatibility-class), 12
 databaseReferences_assnlist, Description-method(zzz-DTTCompatibility-class), 12
 DatabaseReferences_assnlist-class (zzz-DTTCompatibility-class), 12
 dataExternal (zzz-DTTCompatibility-class), 12
 dataExternal, DataExternal_assn-method (zzz-DTTCompatibility-class), 12
 DataExternal-class (zzz-DTTCompatibility-class), 12
 dataExternal_assn (zzz-DTTCompatibility-class), 12
 dataExternal_assn, BioDataCube-method (zzz-DTTCompatibility-class), 12
 DataExternal_assn-class (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-class (zzz-DTTCompatibility-class), 12

- 12
- dataInternal_assn
(*zzz-DTTCompatibility-class*),
12
- dataInternal_assn, BioDataCube-method
(*zzz-DTTCompatibility-class*),
12
- DataInternal_assn-class
(*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
- dataType_assn, PValue-method
(*zzz-DTTCompatibility-class*),
12
- dataType_assn, Ratio-method
(*zzz-DTTCompatibility-class*),
12
- dataType_assn, SpecializedQuantitationType-method
(*zzz-DTTCompatibility-class*),
12
- DataType_assn-class
(*zzz-DTTCompatibility-class*),
12
- datum
(*zzz-DTTCompatibility-class*),
12
- datum, Datum_assn-method
(*zzz-DTTCompatibility-class*),
12
- Datum-class
(*zzz-DTTCompatibility-class*),
12
- datum_assn
(*zzz-DTTCompatibility-class*),
12
- datum_assn, QuantitationTypeTuple-method
(*zzz-DTTCompatibility-class*),
12
- Datum_assn-class
(*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
- DefaultValue_assn-class
(*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
- DefectType_assn-class
(*zzz-DTTCompatibility-class*),
12
- deltaX
(*zzz-DTTCompatibility-class*),
12
- deltaX, PositionDelta-method
(*zzz-DTTCompatibility-class*),
12
- deltaY
(*zzz-DTTCompatibility-class*),
12
- deltaY, PositionDelta-method

	(<i>zzz-DTTCompatibility-class</i>),	12	
derivedBioAssay	(<i>zzz-DTTCompatibility-class</i>),	12	derivedBioAssayData_assnreflist
	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
derivedBioAssay, BioAssay_assnlist-method	(<i>zzz-DTTCompatibility-class</i>),	12	derivedBioAssayData_assnreflist, DerivedBioAssay
	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
derivedBioAssay, DerivedBioAssay_assnlist-method	(<i>zzz-DTTCompatibility-class</i>),	12	DerivedBioAssayData_assnreflist-class
	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
DerivedBioAssay-class	(<i>zzz-DTTCompatibility-class</i>),	12	derivedBioAssayData_ref
	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
DerivedBioAssay_assnlist-class	(<i>zzz-DTTCompatibility-class</i>),	12	derivedBioAssayData_ref, BioAssayData_assnreflist
	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
derivedBioAssay_ref	(<i>zzz-DTTCompatibility-class</i>),	12	derivedBioAssayData_ref, BioAssayDataSources_a
	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
derivedBioAssay_ref, BioAssay_assnref-method	(<i>zzz-DTTCompatibility-class</i>),	12	derivedBioAssayData_ref, ClusterBioAssayData_a
	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
derivedBioAssay_ref, BioAssayMapTarget_assnref-method	(<i>zzz-DTTCompatibility-class</i>),	12	derivedBioAssayData_ref, DerivedBioAssayData_a
	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
derivedBioAssay_ref, BioAssays_assnreflist-method	(<i>zzz-DTTCompatibility-class</i>),	12	derivedBioAssayData_ref, DerivedBioAssayDataTa
	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
derivedBioAssay_ref, SourceBioAssays_assnreflist-method	(<i>zzz-DTTCompatibility-class</i>),	12	DerivedBioAssayData_ref-class
	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
derivedBioAssay_ref, TopLevelBioAssays_assnreflist-method	(<i>zzz-DTTCompatibility-class</i>),	12	DerivedBioAssayDataTarget_assnref-class
	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
DerivedBioAssay_ref-class	(<i>zzz-DTTCompatibility-class</i>),	12	derivedBioAssayMap_assnreflist
	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
derivedBioAssayData	(<i>zzz-DTTCompatibility-class</i>),	12	derivedBioAssayMap_assnreflist, DerivedBioAssay
	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
derivedBioAssayData, BioAssayData_assnlist-method	(<i>zzz-DTTCompatibility-class</i>),	12	DerivedBioAssayMap_assnreflist-class
	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
derivedBioAssayData, DerivedBioAssayData_assnlist-method	(<i>zzz-DTTCompatibility-class</i>),	12	derivedSignal
	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
DerivedBioAssayData-class	(<i>zzz-DTTCompatibility-class</i>),	12	derivedSignal, QuantitationType_assnlist-method
	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),
DerivedBioAssayData_assnlist-class	(<i>zzz-DTTCompatibility-class</i>),	12	DerivedSignal-class
	(<i>zzz-DTTCompatibility-class</i>),	12	(<i>zzz-DTTCompatibility-class</i>),

derivedSignal_ref	(zzz-DTTCompatibility-class),	(zzz-DTTCompatibility-class),
12	12	12
derivedSignal_ref, QuantitationType_assnref-method	(zzz-DTTCompatibility-class),	descriptions_assnlist, ArrayDesign-method
12	12	12
derivedSignal_ref, QuantitationTypes_assnref-method	(zzz-DTTCompatibility-class),	descriptions_assnlist, ArrayGroup-method
12	12	12
derivedSignal_ref, SourcesQuantitationType_assnref-method	(zzz-DTTCompatibility-class),	descriptions_assnlist, ArrayManufacture-method
12	12	12
derivedSignal_ref, TargetQuantitationType_assnref-method	(zzz-DTTCompatibility-class),	descriptions_assnlist, Audit-method
12	12	12
DerivedSignal_ref-class	(zzz-DTTCompatibility-class),	descriptions_assnlist, BibliographicReference-method
12	12	12
description	(zzz-DTTCompatibility-class),	descriptions_assnlist, BioAssayCreation-method
12	12	12
description, Descriptions_assnlist-method	(zzz-DTTCompatibility-class),	descriptions_assnlist, BioAssayDataCluster-method
12	12	12
description, NormalizationDescription_assn-method	(zzz-DTTCompatibility-class),	descriptions_assnlist, BioAssayDimension-method
12	12	12
description, OntologyEntry-method	(zzz-DTTCompatibility-class),	descriptions_assnlist, BioAssayMap-method
12	12	12
description, QualityControlDescription_assn-method	(zzz-DTTCompatibility-class),	descriptions_assnlist, BioAssayTreatment-method
12	12	12
description, ReplicateDescription_assn-method	(zzz-DTTCompatibility-class),	descriptions_assnlist, BioSample-method
12	12	12
Description-class	(zzz-DTTCompatibility-class),	descriptions_assnlist, BioSequence-method
12	12	12
description_package	(zzz-DTTCompatibility-class),	descriptions_assnlist, BioSource-method
12	12	12
description_package, MAGE-ML-method	(zzz-DTTCompatibility-class),	descriptions_assnlist, Channel-method
12	12	12
Description_package-class	(zzz-DTTCompatibility-class),	descriptions_assnlist, CompositeCompositeMap-method
12	12	12
descriptions_assnlist	(zzz-DTTCompatibility-class),	descriptions_assnlist, CompositeGroup-method
12	12	12
descriptions_assnlist, Array-method	(zzz-DTTCompatibility-class),	descriptions_assnlist, CompositeSequence-method
		12

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

(zzz-*DTTCompatibility-class*), 12
 12
 descriptions_assnlist, Parameter-method (zzz-*DTTCompatibility-class*), 12
 (zzz-*DTTCompatibility-class*), 12
 descriptions_assnlist, Person-method (zzz-*DTTCompatibility-class*), 12
 (zzz-*DTTCompatibility-class*), 12
 descriptions_assnlist, PhysicalArrayDesign-method (zzz-*DTTCompatibility-class*), 12
 (zzz-*DTTCompatibility-class*), 12
 descriptions_assnlist, PhysicalBioAssay-method (zzz-*DTTCompatibility-class*), 12
 (zzz-*DTTCompatibility-class*), 12
 descriptions_assnlist, PresentAbsent-method (zzz-*DTTCompatibility-class*), 12
 (zzz-*DTTCompatibility-class*), 12
 descriptions_assnlist, Protocol-method (zzz-*DTTCompatibility-class*), 12
 (zzz-*DTTCompatibility-class*), 12
 descriptions_assnlist, ProtocolApplication-method (zzz-*DTTCompatibility-class*), 12
 (zzz-*DTTCompatibility-class*), 12
 descriptions_assnlist, PValue-method (zzz-*DTTCompatibility-class*), 12
 (zzz-*DTTCompatibility-class*), 12
 descriptions_assnlist, QuantitationTypeDimensions-method (zzz-*DTTCompatibility-class*), 12
 (zzz-*DTTCompatibility-class*), 12
 descriptions_assnlist, QuantitationTypeMap-method (zzz-*DTTCompatibility-class*), 12
 (zzz-*DTTCompatibility-class*), 12
 descriptions_assnlist, Ratio-method (zzz-*DTTCompatibility-class*), 12
 (zzz-*DTTCompatibility-class*), 12
 descriptions_assnlist, Reporter-method (zzz-*DTTCompatibility-class*), 12
 (zzz-*DTTCompatibility-class*), 12
 descriptions_assnlist, ReporterCompositeMap-method (zzz-*DTTCompatibility-class*), 12
 (zzz-*DTTCompatibility-class*), 12
 descriptions_assnlist, ReporterDimension-method (zzz-*DTTCompatibility-class*), 12
 (zzz-*DTTCompatibility-class*), 12
 descriptions_assnlist, ReporterGroup-method (zzz-*DTTCompatibility-class*), 12
 (zzz-*DTTCompatibility-class*), 12
 descriptions_assnlist, Security-method (zzz-*DTTCompatibility-class*), 12
 (zzz-*DTTCompatibility-class*), 12
 descriptions_assnlist, SecurityGroup-method (zzz-*DTTCompatibility-class*), 12
 (zzz-*DTTCompatibility-class*), 12
 descriptions_assnlist, SeqFeature-method (zzz-*DTTCompatibility-class*), 12
 (zzz-*DTTCompatibility-class*), 12
 descriptions_assnlist, Software-method (zzz-*DTTCompatibility-class*), 12
 (zzz-*DTTCompatibility-class*), 12
 descriptions_assnlist, SoftwareApplication-method (zzz-*DTTCompatibility-class*), 12
 (zzz-*DTTCompatibility-class*), 12
 descriptions_assnlist, SpecializedQuantitation-method (zzz-*DTTCompatibility-class*), 12
 (zzz-*DTTCompatibility-class*), 12
 descriptions_assnlist, Transformation-method (zzz-*DTTCompatibility-class*), 12
 (zzz-*DTTCompatibility-class*), 12
 descriptions_assnlist, Treatment-method (zzz-*DTTCompatibility-class*), 12
 (zzz-*DTTCompatibility-class*), 12
 descriptions_assnlist, Zone-method (zzz-*DTTCompatibility-class*), 12
 (zzz-*DTTCompatibility-class*), 12
 Descriptions_assnlist-class (zzz-*DTTCompatibility-class*), 12
 DesignElement_assnlist-class (zzz-*DTTCompatibility-class*), 12
 designElement_assnref (zzz-*DTTCompatibility-class*), 12
 designElement_assnref, DesignElementTuple-method (zzz-*DTTCompatibility-class*), 12
 (zzz-*DTTCompatibility-class*), 12
 DesignElement_assnref-class (zzz-*DTTCompatibility-class*), 12
 designElement_package (zzz-*DTTCompatibility-class*), 12
 designElement_package, MAGE-ML-method (zzz-*DTTCompatibility-class*), 12
 (zzz-*DTTCompatibility-class*), 12
 DesignElement_package-class (zzz-*DTTCompatibility-class*), 12
 DesignElement_ref-class (zzz-*DTTCompatibility-class*), 12
 designElementDimension_assnlist (zzz-*DTTCompatibility-class*), 12
 (zzz-*DTTCompatibility-class*), 12

designElementDimension_assnlist, BioAssayData (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 DesignElementDimension_assnlist-class (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 designElementDimension_assnref (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 designElementDimension_assnref, DerivedBioAssayData (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 designElementDimension_assnref, MeasuredBioAssayData (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 designElementDimension_assnref, NodeContents (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 DesignElementDimension_assnref-class (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 DesignElementDimension_ref-class (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 DesignElementMap_assnlist-class (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 DesignElementMap_ref-class (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 designElementMapping (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 designElementMapping, DesignElementMapping_assn (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 DesignElementMapping-class (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 designElementMapping_assn (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 designElementMapping_assn, Transformation-method (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 DesignElementMapping_assn-class (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 designElementMaps_assnreflist (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 designElementMaps_assnreflist, DesignElementMapping (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 DesignElementMaps_assnreflist-class (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 designElementTuple (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 designElementTuple, DesignElementTuples_assnlist (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 DesignElementTuple-class (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 designElementTuples_assnlist (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 designElementTuples_assnlist, BioAssayTuple-method (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 DesignElementTuples_assnlist-class (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 designProviders_assnreflist (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 designProviders_assnreflist, ArrayDesign-method (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 designProviders_assnreflist, PhysicalArrayDesign-method (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 DesignProviders_assnreflist-class (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 directory (accessors), 6
 directory, NetAffxResource-method (NetAffxResource-class), 2
 distanceUnit (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 distanceUnit, DistanceUnit_assn-method (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 distanceUnit, Unit_assn-method (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 DistanceUnit-class (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12
 distanceUnit_assn (z3z-DTTCompatibility-class),
 (z3z-DTTCompatibility-class), 12

- 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
- DistanceUnit_assn-class
(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
- 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-class
(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
- Error_ref-class
(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-class
(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
- expectedValue_ref, SourcesQuantitationType_assnreflist-method
(zzz-DTTCompatibility-class),
12

- (zzz-DTTCompatibility-class), 12
- expectedValue_ref, TargetQuantitationType_assnref-method (zzz-DTTCompatibility-class), 12
- ExpectedValue_ref-class (zzz-DTTCompatibility-class), 12
- experiment (zzz-DTTCompatibility-class), 12
- experiment, Experiment_assnlist-method (zzz-DTTCompatibility-class), 12
- Experiment-class (zzz-DTTCompatibility-class), 12
- experiment_assnlist (zzz-DTTCompatibility-class), 12
- experiment_assnlist, Experiment_package-method (zzz-DTTCompatibility-class), 12
- Experiment_assnlist-class (zzz-DTTCompatibility-class), 12
- experiment_package (zzz-DTTCompatibility-class), 12
- experiment_package, MAGE-ML-method (zzz-DTTCompatibility-class), 12
- Experiment_package-class (zzz-DTTCompatibility-class), 12
- experimentalFactor (zzz-DTTCompatibility-class), 12
- experimentalFactor, ExperimentalFactors_assnlist-method (zzz-DTTCompatibility-class), 12
- ExperimentalFactor-class (zzz-DTTCompatibility-class), 12
- ExperimentalFactor_assnref-class (zzz-DTTCompatibility-class), 12
- experimentalFactor_ref (zzz-DTTCompatibility-class), 12
- experimentalFactor_ref, ExperimentalFactor_assnref-method (zzz-DTTCompatibility-class), 12
- ExperimentalFactor_ref-class (zzz-DTTCompatibility-class), 12
- experimentalFactors_assnlist (zzz-DTTCompatibility-class), 12
- experimentalFactors_assnlist, ExperimentDesign (zzz-DTTCompatibility-class), 12
- ExperimentalFactors_assnlist-class (zzz-DTTCompatibility-class), 12
- experimentDesign (zzz-DTTCompatibility-class), 12
- experimentDesign, ExperimentDesigns_assnlist-method (zzz-DTTCompatibility-class), 12
- ExperimentDesign-class (zzz-DTTCompatibility-class), 12
- experimentDesigns_assnlist (zzz-DTTCompatibility-class), 12
- experimentDesigns_assnlist, Experiment-method (zzz-DTTCompatibility-class), 12
- ExperimentDesigns_assnlist-class (zzz-DTTCompatibility-class), 12
- exportedFromDB (zzz-DTTCompatibility-class), 12
- exportedFromDB, ExternalReference-method (zzz-DTTCompatibility-class), 12
- exportedFromServer (zzz-DTTCompatibility-class), 12
- exportedFromServer, ExternalReference-method (zzz-DTTCompatibility-class), 12
- exportID (zzz-DTTCompatibility-class), 12
- exportID, ExternalReference-method (zzz-DTTCompatibility-class), 12
- exportName (zzz-DTTCompatibility-class), 12

exportName, ExternalReference-method	(zzz-DTTCompatibility-class), 12	failed	(zzz-DTTCompatibility-class), 12
externalLIMS_assn	(zzz-DTTCompatibility-class), 12	failed, QuantitationType_assnlist-method	(zzz-DTTCompatibility-class), 12
externalLIMS_assn, Compound-method	(zzz-DTTCompatibility-class), 12	Failed-class	(zzz-DTTCompatibility-class), 12
ExternalLIMS_assn-class	(zzz-DTTCompatibility-class), 12	failed_ref	(zzz-DTTCompatibility-class), 12
externalReference	(zzz-DTTCompatibility-class), 12	failed_ref, QuantitationType_assnref-method	(zzz-DTTCompatibility-class), 12
externalReference, ExternalReference_assn-method	(zzz-DTTCompatibility-class), 12	failed_ref, QuantitationTypes_assnreflist-method	(zzz-DTTCompatibility-class), 12
externalReference_assn	(zzz-DTTCompatibility-class), 12	failed_ref, SourcesQuantitationType_assnreflist-method	(zzz-DTTCompatibility-class), 12
externalReference_assn, Description-method	(zzz-DTTCompatibility-class), 12	Failed_ref-class	(zzz-DTTCompatibility-class), 12
ExternalReference_assn-class	(zzz-DTTCompatibility-class), 12	factorValue	(zzz-DTTCompatibility-class), 12
factorValue	(zzz-DTTCompatibility-class), 12	factorValue, FactorValues_assnlist-method	(zzz-DTTCompatibility-class), 12
factorValue, FactorValues_assnlist-method	(zzz-DTTCompatibility-class), 12	FactorValue-class	(zzz-DTTCompatibility-class), 12
FactorValue-class	(zzz-DTTCompatibility-class), 12	factorValue_ref	(zzz-DTTCompatibility-class), 12
factorValue_ref	(zzz-DTTCompatibility-class), 12	factorValue_ref, BioAssayFactorValues_assnreflist-method	(zzz-DTTCompatibility-class), 12
factorValue_ref, BioAssayFactorValues_assnreflist-method	(zzz-DTTCompatibility-class), 12	FactorValue_ref-class	(zzz-DTTCompatibility-class), 12
FactorValue_ref-class	(zzz-DTTCompatibility-class), 12	factorValues_assnlist	(zzz-DTTCompatibility-class), 12
factorValues_assnlist	(zzz-DTTCompatibility-class), 12	factorValues_assnlist, ExperimentalFactor-method	(zzz-DTTCompatibility-class), 12
factorValues_assnlist, ExperimentalFactor-method	(zzz-DTTCompatibility-class), 12	FactorValues_assnlist-class	(zzz-DTTCompatibility-class), 12
FactorValues_assnlist-class	(zzz-DTTCompatibility-class), 12	feature	(zzz-DTTCompatibility-class), 12
		feature, DesignElement_assnlist-method	(zzz-DTTCompatibility-class), 12

- feature, Features_assnlist-method (zzz-DTTCompatibility-class), 12
- Feature-class (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_assnref-class (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
- Feature_ref-class (zzz-DTTCompatibility-class), 12
- featureDefect (zzz-DTTCompatibility-class), 12
- featureDefect, FeatureDefects_assnlist-method (zzz-DTTCompatibility-class), 12
- FeatureDefect-class (zzz-DTTCompatibility-class), 12
- featureDefects_assnlist (zzz-DTTCompatibility-class), 12
- featureDefects_assnlist, ArrayManufactureDeviation-method (zzz-DTTCompatibility-class), 12
- FeatureDefects_assnlist-class (zzz-DTTCompatibility-class), 12
- featureDimension (zzz-DTTCompatibility-class), 12
- featureDimension, DesignElementDimension_assnlist-method (zzz-DTTCompatibility-class), 12
- FeatureDimension-class (zzz-DTTCompatibility-class), 12
- featureDimension_ref (zzz-DTTCompatibility-class), 12
- featureDimension_ref, DesignElementDimension_assnlist-method (zzz-DTTCompatibility-class), 12
- FeatureDimension_ref-class (zzz-DTTCompatibility-class), 12
- featureExtraction (zzz-DTTCompatibility-class), 12
- featureExtraction, FeatureExtraction_assnlist-method (zzz-DTTCompatibility-class), 12
- FeatureExtraction-class (zzz-DTTCompatibility-class), 12
- featureExtraction_assn (zzz-DTTCompatibility-class), 12
- featureExtraction_assn, MeasuredBioAssay-method (zzz-DTTCompatibility-class), 12
- FeatureExtraction_assn-class (zzz-DTTCompatibility-class), 12
- featureGroup (zzz-DTTCompatibility-class), 12
- featureGroup, FeatureGroups_assnlist-method (zzz-DTTCompatibility-class), 12

- 12
- FeatureGroup-class
(zzz-DTTCompatibility-class),
12
- FeatureGroup_assnref-class
(zzz-DTTCompatibility-class),
12
- featureGroup_ref
(zzz-DTTCompatibility-class),
12
- featureGroup_ref, FeatureGroup_assnref-method(zzz-DTTCompatibility-class),
12
- FeatureGroup_ref-class
(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
- FeatureGroups_assnlist-class
(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
- FeatureInformation-class
(zzz-DTTCompatibility-class),
12
- featureInformationSources_assnlist
(zzz-DTTCompatibility-class),
12
- featureInformationSources_assnlist, FeatureReporterMap-method(zzz-DTTCompatibility-class),
12
- FeatureInformationSources_assnlist-class
(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
- FeatureLIMSs_assnlist-class
(zzz-DTTCompatibility-class),
12
- featureLocation
(zzz-DTTCompatibility-class),
12
- featureLocation, FeatureLocation_assn-method(zzz-DTTCompatibility-class),
12
- FeatureLocation-class
(zzz-DTTCompatibility-class),
12
- featureLocation_assn
(zzz-DTTCompatibility-class),
12
- featureLocation_assn, Feature-method
(zzz-DTTCompatibility-class),
12
- FeatureLocation_assn-class
(zzz-DTTCompatibility-class),
12
- featureReporterMap
(zzz-DTTCompatibility-class),
12
- featureReporterMap, DesignElementMap_assnlist-method(zzz-DTTCompatibility-class),
12
- featureReporterMap, FeatureReporterMap_assnlist-class(zzz-DTTCompatibility-class),
12
- FeatureReporterMap-class
(zzz-DTTCompatibility-class),
12
- featureReporterMap_assnlist
(zzz-DTTCompatibility-class),
12
- featureReporterMap_assnlist, DesignElement_package-method(zzz-DTTCompatibility-class),
12
- FeatureReporterMap_assnlist-class
(zzz-DTTCompatibility-class),
12

- (zzz-*DTTCompatibility-class*), 12
- featureReporterMap_ref
 - (zzz-*DTTCompatibility-class*), 12
- featureReporterMap_ref, DesignElementMaps_assnreflist-method
 - (zzz-*DTTCompatibility-class*), 12
- featureReporterMap_ref, FeatureReporterMaps_assnreflist-method
 - (zzz-*DTTCompatibility-class*), 12
- FeatureReporterMap_ref-class
 - (zzz-*DTTCompatibility-class*), 12
- featureReporterMaps_assnreflist
 - (zzz-*DTTCompatibility-class*), 12
- featureReporterMaps_assnreflist, Reporter-method
 - (zzz-*DTTCompatibility-class*), 12
- FeatureReporterMaps_assnreflist-class
 - (zzz-*DTTCompatibility-class*), 12
- features_assnlist
 - (zzz-*DTTCompatibility-class*), 12
- features_assnlist, FeatureGroup-method
 - (zzz-*DTTCompatibility-class*), 12
- Features_assnlist-class
 - (zzz-*DTTCompatibility-class*), 12
- featureShape_assn
 - (zzz-*DTTCompatibility-class*), 12
- featureShape_assn, FeatureGroup-method
 - (zzz-*DTTCompatibility-class*), 12
- FeatureShape_assn-class
 - (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
- Fiducial-class
 - (zzz-*DTTCompatibility-class*), 12
- fiducials_assnlist
 - (zzz-*DTTCompatibility-class*), 12
- fiducials_assnlist, ArrayGroup-method
 - (zzz-*DTTCompatibility-class*), 12
- Fiducials_assnlist-class
 - (zzz-*DTTCompatibility-class*), 12
- fiducialType_assn
 - (zzz-*DTTCompatibility-class*), 12
- fiducialType_assn, Fiducial-method
 - (zzz-*DTTCompatibility-class*), 12
- FiducialType_assn-class
 - (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
- Format_assn-class
 - (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-class
 (*zzz-DTTCompatibility-class*),
 12
- hardware_assnlist
 (*zzz-DTTCompatibility-class*),
 12
- hardware_assnlist, Protocol_package-method
 (*zzz-DTTCompatibility-class*),
 12
- Hardware_assnlist-class
 (*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_assnref-class
 (*zzz-DTTCompatibility-class*),
 12
- hardware_ref
 (*zzz-DTTCompatibility-class*),
 12
- hardware_ref, Hardware_assnref-method
 (*zzz-DTTCompatibility-class*),
 12
- hardware_ref, Hardwares_assnreflist-method
 (*zzz-DTTCompatibility-class*),
 12
- Hardware_ref-class
 (*zzz-DTTCompatibility-class*),
 12
- hardwareApplication
 (*zzz-DTTCompatibility-class*),
 12
- hardwareApplication, HardwareApplications_assnreflist-method
 (*zzz-DTTCompatibility-class*),
 12
- HardwareApplication-class
 (*zzz-DTTCompatibility-class*),
 12
- hardwareApplications_assnlist
 (*zzz-DTTCompatibility-class*),
 12
- hardwareApplications_assnlist, ProtocolApplication-class
 (*zzz-DTTCompatibility-class*),
 12
- HardwareApplications_assnlist-class
 (*zzz-DTTCompatibility-class*),
 12
- hardwareManufacturers_assnreflist
 (*zzz-DTTCompatibility-class*),
 12
- hardwareManufacturers_assnreflist, Hardware-manufacturer-class
 (*zzz-DTTCompatibility-class*),
 12
- HardwareManufacturers_assnreflist-class
 (*zzz-DTTCompatibility-class*),
 12
- hardwares_assnreflist
 (*zzz-DTTCompatibility-class*),
 12
- hardwares_assnreflist, Protocol-method
 (*zzz-DTTCompatibility-class*),
 12
- Hardwares_assnreflist-class
 (*zzz-DTTCompatibility-class*),
 12
- higherLevelAnalysis_package
 (*zzz-DTTCompatibility-class*),
 12
- higherLevelAnalysis_package, MAGE-ML-method
 (*zzz-DTTCompatibility-class*),
 12
- HigherLevelAnalysis_package-class
 (*zzz-DTTCompatibility-class*),
 12
- hybridization
 (*zzz-DTTCompatibility-class*),
 12
- hybridization, BioAssayCreation_assnreflist-method
 (*zzz-DTTCompatibility-class*),
 12
- Hybridization-class
 (*zzz-DTTCompatibility-class*),
 12
- identifier
 (*zzz-DTTCompatibility-class*),
 12
- identifier, Array-method
 (*zzz-DTTCompatibility-class*),
 12

- 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
- 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
- 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*),
12
- identifier,MeasuredBioAssay-method
(*zzz-DTTCompatibility-class*),
12
- identifier,MeasuredBioAssay_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier,MeasuredBioAssayData-method
(*zzz-DTTCompatibility-class*),
12
- identifier,MeasuredBioAssayData_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier,MeasuredSignal-method
(*zzz-DTTCompatibility-class*),
12
- identifier,MeasuredSignal_ref-method
(*zzz-DTTCompatibility-class*),
12
- 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
- identifier,PhysicalBioAssay_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier,PresentAbsent-method
(*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
(*zzz-DTTCompatibility-class*),
12
- identifier,QuantitationTypeDimension_ref-method
(*zzz-DTTCompatibility-class*),
12
- identifier,QuantitationTypeMap-method
(*zzz-DTTCompatibility-class*),
12

- 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
- identifier, ReporterCompositeMap_ref-method (zzz-DTTCompatibility-class), 12
- identifier, ReporterDimension-method (zzz-DTTCompatibility-class), 12
- identifier, ReporterDimension_ref-method (zzz-DTTCompatibility-class), 12
- identifier, ReporterGroup-method (zzz-DTTCompatibility-class), 12
- identifier, ReporterGroup_ref-method (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), 12
- identifier, SpecializedQuantitationType-method (zzz-DTTCompatibility-class), 12
- identifier, SpecializedQuantitationType_ref-method (zzz-DTTCompatibility-class), 12
- identifier, StandardQuantitationType_ref-method (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 (zzz-DTTCompatibility-class), 12
- IdentifierLIMS_assn-class (zzz-DTTCompatibility-class), 12
- Image-class (zzz-DTTCompatibility-class), 12
- image_ref (zzz-DTTCompatibility-class), 12
- image_ref, Images_assnreflist-method (zzz-DTTCompatibility-class), 12
- Image_ref-class (zzz-DTTCompatibility-class), 12
- imageAcquisition (zzz-DTTCompatibility-class), 12
- imageAcquisition, BioAssayTreatments_assnlist-method (zzz-DTTCompatibility-class), 12
- ImageAcquisition-class

- (zzz-DTTCompatibility-class),
12
- images_assnreflist
(zzz-DTTCompatibility-class),
12
- images_assnreflist, ImageAcquisition-method
(zzz-DTTCompatibility-class),
12
- Images_assnreflist-class
(zzz-DTTCompatibility-class),
12
- immobilizedCharacteristics_assnreflist
(zzz-DTTCompatibility-class),
12
- immobilizedCharacteristics_assnreflist, Reporter-method
(zzz-DTTCompatibility-class),
12
- ImmobilizedCharacteristics_assnreflist-class
(zzz-DTTCompatibility-class),
12
- information_assnref
(zzz-DTTCompatibility-class),
12
- information_assnref, Array-method
(zzz-DTTCompatibility-class),
12
- Information_assnref-class
(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
- (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-class
(zzz-DTTCompatibility-class),
12
- labeledExtract_ref
(zzz-DTTCompatibility-class),
12

- labeledExtract_ref, BioMaterial_assnrefl-class
 (*zzz-DTTCompatibility-class*),
 12
- LabeledExtract_ref-class
 (*zzz-DTTCompatibility-class*),
 12
- labels_assnrefl-class
 (*zzz-DTTCompatibility-class*),
 12
- labels_assnrefl-class, Channel-method
 (*zzz-DTTCompatibility-class*),
 12
- labels_assnrefl-class, LabeledExtract-method
 (*zzz-DTTCompatibility-class*),
 12
- Labels_assnrefl-class
 (*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, //
- lowerRightX
 (*zzz-DTTCompatibility-class*),
 12
- lowerRightX, Zone-method
 (*zzz-DTTCompatibility-class*),
 12
- lowerRightY
 (*zzz-DTTCompatibility-class*),
 12
- lowerRightY, Zone-method
 (*zzz-DTTCompatibility-class*),
 12
- MAGE-ML-class
 (*zzz-DTTCompatibility-class*),
 12
- make, Hardware-method
 (*zzz-DTTCompatibility-class*),
 12
- manufactureLIMS
 (*zzz-DTTCompatibility-class*),
 12
- manufactureLIMS, FeatureLIMSSs_assnrefl-method
 (*zzz-DTTCompatibility-class*),
 12
- ManufactureLIMS-class
 (*zzz-DTTCompatibility-class*),
 12
- manufactureLIMSBiomaterial
 (*zzz-DTTCompatibility-class*),
 12
- manufactureLIMSBiomaterial, FeatureLIMSSs_assnrefl-method
 (*zzz-DTTCompatibility-class*),
 12
- ManufactureLIMSBiomaterial-class
 (*zzz-DTTCompatibility-class*),
 12
- manufacturingDate
 (*zzz-DTTCompatibility-class*),
 12
- manufacturingDate, ArrayManufacture-method
 (*zzz-DTTCompatibility-class*),
 12
- massUnit
 (*zzz-DTTCompatibility-class*),
 12
- massUnit, Unit_assnrefl-method
 (*zzz-DTTCompatibility-class*),
 12
- MassUnit-class
 (*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_assnrefl

- (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
- MaterialType_assn-class (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-class (zzz-DTTCompatibility-class), 12
- MeasuredBioAssay_assnlist-class (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_assnref-method (zzz-DTTCompatibility-class), 12
- measuredBioAssay_ref, SourceBioAssays_assnref-method (zzz-DTTCompatibility-class), 12
- measuredBioAssay_ref, TopLevelBioAssays_assnref-method (zzz-DTTCompatibility-class), 12
- MeasuredBioAssay_ref-class (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-class (zzz-DTTCompatibility-class), 12
- MeasuredBioAssayData_assnlist-class (zzz-DTTCompatibility-class), 12
- measuredBioAssayData_assnreflist (zzz-DTTCompatibility-class), 12
- measuredBioAssayData_assnreflist, MeasuredBioAssayData_assnreflist-method (zzz-DTTCompatibility-class), 12
- MeasuredBioAssayData_assnreflist-class (zzz-DTTCompatibility-class), 12
- measuredBioAssayData_ref (zzz-DTTCompatibility-class), 12
- measuredBioAssayData_ref, BioAssayData_assnreflist-method (zzz-DTTCompatibility-class), 12
- measuredBioAssayData_ref, BioAssayDataSources_assnreflist-method (zzz-DTTCompatibility-class), 12
- measuredBioAssayData_ref, ClusterBioAssayData_assnreflist-method (zzz-DTTCompatibility-class), 12
- measuredBioAssayData_ref, MeasuredBioAssayData_assnreflist-method (zzz-DTTCompatibility-class), 12
- MeasuredBioAssayData_ref-class (zzz-DTTCompatibility-class), 12
- MeasuredBioAssayTarget_assnref-class (zzz-DTTCompatibility-class), 12
- measuredSignal (zzz-DTTCompatibility-class), 12
- measuredSignal, QuantitationType_assnlist-method (zzz-DTTCompatibility-class), 12
- MeasuredSignal-class (zzz-DTTCompatibility-class), 12

measuredSignal_ref (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 Measurement_package-class
 measuredSignal_ref, QuantitationType_assnref-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 mediaCol
 measuredSignal_ref, QuantitationTypes_assnref-method (zzz-ARRCompatibility-class),
 (zzz-DTTCompatibility-class), 11
 mediaCol, PhysicalArray-method
 measuredSignal_ref, SourcesQuantitationType_assnref-method (zzz-ARRCompatibility-class),
 (zzz-DTTCompatibility-class), 11
 mediaFileGUID
 measuredSignal_ref, TargetQuantitationType_assnref-method (zzz-ARRCompatibility-class),
 (zzz-DTTCompatibility-class), 11
 mediaFileGUID, PhysicalArray-method
 MeasuredSignal_ref-class (zzz-DTTCompatibility-class), 11
 mediaFileName
 measurement (zzz-ARRCompatibility-class), 11
 mediaFileName, PhysicalArray-method
 measurement, ActionMeasurement_assn-method (zzz-ARRCompatibility-class),
 (zzz-DTTCompatibility-class), 11
 mediaRow
 measurement, DefaultValue_assn-method (zzz-ARRCompatibility-class), 11
 mediaRow, PhysicalArray-method
 measurement, Measurement_assn-method (zzz-ARRCompatibility-class), 11
 mediaType
 Measurement-class (zzz-ARRCompatibility-class), 11
 mediaType, PhysicalArray-method
 measurement_assn (zzz-ARRCompatibility-class), 11
 members_assnreflist
 measurement_assn, BioMaterialMeasurement-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 members_assnreflist, SecurityGroup-method
 measurement_assn, CompoundMeasurement-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 Members_assnreflist-class
 measurement_assn, FactorValue-method (zzz-DTTCompatibility-class), 12
 midInitials
 Measurement_assn-class (zzz-DTTCompatibility-class), 12
 midInitials, Person-method
 measurement_package (zzz-DTTCompatibility-class), 12
 mismatchInformation
 measurement_package, MAGE-ML-method (zzz-DTTCompatibility-class),

- 12
 mismatchInformation, MismatchInformation_assnlist-method (zzz-DTTCompatibility-class), 12
 (zzz-DTTCompatibility-class), 12
 MismatchInformation-class (zzz-DTTCompatibility-class), 12
 mismatchInformation_assnlist (zzz-DTTCompatibility-class), 12
 mismatchInformation_assnlist, CompositePosition-method (zzz-DTTCompatibility-class), 12
 mismatchInformation_assnlist, FeatureInformation-method (zzz-DTTCompatibility-class), 12
 mismatchInformation_assnlist, ReporterPosition-method (zzz-DTTCompatibility-class), 12
 MismatchInformation_assnlist-class (zzz-DTTCompatibility-class), 12
 model (zzz-DTTCompatibility-class), 12
 model, Hardware-method (zzz-DTTCompatibility-class), 12
 name (zzz-DTTCompatibility-class), 12
 name, Array-method (zzz-DTTCompatibility-class), 12
 name, Array_ref-method (zzz-DTTCompatibility-class), 12
 name, ArrayAttribute-method (zzz-ARRCompatibility-class), 11
 name, ArrayDesign-method (zzz-DTTCompatibility-class), 12
 name, ArrayDesign_ref-method (zzz-DTTCompatibility-class), 12
 name, ArrayGroup-method (zzz-DTTCompatibility-class), 12
 name, ArrayGroup_ref-method (zzz-DTTCompatibility-class), 12
 name, ArrayManufacture-method (zzz-DTTCompatibility-class), 12
 name, ArrayManufacture_ref-method (zzz-DTTCompatibility-class), 12
 name, BioAssay_ref-method (zzz-DTTCompatibility-class), 12
 name, BioAssayCreation-method (zzz-DTTCompatibility-class), 12
 name, BioAssayData_ref-method (zzz-DTTCompatibility-class), 12
 name, BioAssayDataCluster-method (zzz-DTTCompatibility-class), 12
 name, BioAssayDataCluster_ref-method (zzz-DTTCompatibility-class), 12
 name, BioAssayDimension-method (zzz-DTTCompatibility-class), 12
 name, BioAssayDimension_ref-method (zzz-DTTCompatibility-class), 12
 name, BioAssayMap-method (zzz-DTTCompatibility-class), 12
 name, BioAssayMap_ref-method (zzz-DTTCompatibility-class), 12
 name, BioAssayTreatment-method (zzz-DTTCompatibility-class), 12
 name, BioMaterial_ref-method (zzz-DTTCompatibility-class), 12
 name, BioSample-method (zzz-DTTCompatibility-class), 12
 name, BioSample_ref-method (zzz-DTTCompatibility-class), 12
 name, BioSequence-method (zzz-DTTCompatibility-class), 12
 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*),
 12
 name, CompositeCompositeMap-method
 (*zzz-DTTCompatibility-class*),
 12
 name, CompositeCompositeMap_ref-method
 (*zzz-DTTCompatibility-class*),
 12
 name, CompositeGroup-method
 (*zzz-DTTCompatibility-class*),
 12
 name, CompositeGroup_ref-method
 (*zzz-DTTCompatibility-class*),
 12
 name, CompositeSequence-method
 (*zzz-DTTCompatibility-class*),
 12
 name, CompositeSequence_ref-method
 (*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*),
 12
 name, ConfidenceIndicator_ref-method
 (*zzz-DTTCompatibility-class*),
 12
 name, Contact_ref-method
 (*zzz-DTTCompatibility-class*),
 12
 name, Database-method
 (*zzz-DTTCompatibility-class*),
 12
 name, Database_ref-method
 (*zzz-DTTCompatibility-class*),
 12
 name, DerivedBioAssay-method
 (*zzz-DTTCompatibility-class*),
 12
 name, DerivedBioAssay_ref-method
 (*zzz-DTTCompatibility-class*),
 12
 name, DerivedBioAssayData-method
 (*zzz-DTTCompatibility-class*),
 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

- 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
- 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
- NameValueType-class
(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
- Node-class
(zzz-DTTCompatibility-class),
12
- nodeContents
(zzz-DTTCompatibility-class),
12
- nodeContents, NodeContents_assnlist-method
(zzz-DTTCompatibility-class),
12
- NodeContents-class
(zzz-DTTCompatibility-class),
12
- nodeContents_assnlist
(zzz-DTTCompatibility-class),
12
- nodeContents_assnlist, Node-method
(zzz-DTTCompatibility-class),
12
- NodeContents_assnlist-class
(zzz-DTTCompatibility-class),
12
- nodes_assnlist
(zzz-DTTCompatibility-class),
12
- nodes_assnlist, NodeAssayDataCluster-method
(zzz-DTTCompatibility-class),
12
- nodes_assnlist, Node-method
(zzz-DTTCompatibility-class),
12
- Nodes_assnlist-class
(zzz-DTTCompatibility-class),
12
- nodeValue
(zzz-DTTCompatibility-class),
12
- nodeValue, NodeValue_assnlist-method
(zzz-DTTCompatibility-class),
12
- NodeValue-class
(zzz-DTTCompatibility-class),
12
- nodeValue_assnlist
(zzz-DTTCompatibility-class),
12

- nodeValue_assnlist, Node-method
(zzz-DTTCompatibility-class),
12
- nodeValue_assnlist-class
(zzz-DTTCompatibility-class),
12
- normalizationDescription_assn
(zzz-DTTCompatibility-class),
12
- normalizationDescription_assn, ExperimentDesign-method
(zzz-DTTCompatibility-class),
12
- NormalizationDescription_assn-class
(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
- OntologyEntries_assnlist-class
(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

- 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
- OntologyEntry-class
(zzz-DTTCompatibility-class),
12
- ontologyReference_assn
(zzz-DTTCompatibility-class),
12
- ontologyReference_assn, OntologyEntry-method
(zzz-DTTCompatibility-class),
12
- OntologyReference_assn-class
(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-class
(zzz-DTTCompatibility-class),
12
- Organization_assnlist-class
(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
- Organization_ref-class
(zzz-DTTCompatibility-class),
12
- orientationMark
(zzz-DTTCompatibility-class),
12
- orientationMark, ArrayGroup-method

- (*zzz-DTTCompatibility-class*),
12
- orientationMarkPosition
(*zzz-DTTCompatibility-class*),
12
- 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
- Owner_assnreflist-class
(*zzz-DTTCompatibility-class*),
12
- pages
(*zzz-DTTCompatibility-class*),
12
- pages, BibliographicReference-method
(*zzz-DTTCompatibility-class*),
12
- parameter
(*zzz-DTTCompatibility-class*),
12
- parameter, ParameterTypes_assnlist-method
(*zzz-DTTCompatibility-class*),
12
- Parameter-class
(*zzz-DTTCompatibility-class*),
12
- parameter_ref
(*zzz-DTTCompatibility-class*),
12
- parameter_ref, ParameterType_assnref-method
(*zzz-DTTCompatibility-class*),
12
- Parameter_ref-class
(*zzz-DTTCompatibility-class*),
12
- parameters_assnlist
(*zzz-DTTCompatibility-class*),
12
- parameters_assnlist, BibliographicReference-method
(*zzz-DTTCompatibility-class*),
12
- Parameters_assnlist-class
(*zzz-DTTCompatibility-class*),
12
- parameterType_assnref
(*zzz-DTTCompatibility-class*),
12
- parameterType_assnref, ParameterValue-method
(*zzz-DTTCompatibility-class*),
12
- ParameterType_assnref-class
(*zzz-DTTCompatibility-class*),
12
- parameterTypes_assnlist
(*zzz-DTTCompatibility-class*),
12
- parameterTypes_assnlist, Hardware-method
(*zzz-DTTCompatibility-class*),
12
- parameterTypes_assnlist, Protocol-method
(*zzz-DTTCompatibility-class*),
12
- parameterTypes_assnlist, Software-method
(*zzz-DTTCompatibility-class*),
12
- ParameterTypes_assnlist-class
(*zzz-DTTCompatibility-class*),
12
- parameterValue
(*zzz-DTTCompatibility-class*),
12
- parameterValue, ParameterValues_assnlist-method
(*zzz-DTTCompatibility-class*),
12
- ParameterValue-class
(*zzz-DTTCompatibility-class*),
12
- parameterValues_assnlist
(*zzz-DTTCompatibility-class*),
12

- parameterValues_assnlist, HardwareApplication-method
(zzz-DTTCompatibility-class), 12
- parameterValues_assnlist, ProtocolApplication-method
(zzz-DTTCompatibility-class), 12
- parameterValues_assnlist, SoftwareApplication-method
(zzz-DTTCompatibility-class), 12
- ParameterValues_assnlist-class
(zzz-DTTCompatibility-class), 12
- parent_assnref
(zzz-DTTCompatibility-class), 12
- parent_assnref, Organization-method
(zzz-DTTCompatibility-class), 12
- Parent_assnref-class
(zzz-DTTCompatibility-class), 12
- password(accessors), 6
- password, NetAffxResource-method
(NetAffxResource-class), 2
- patassignmentMethod
(zzz-ARRCompatibility-class), 11
- patassignmentMethod, PhysicalArray-method
(zzz-ARRCompatibility-class), 11
- performer_assnref
(zzz-DTTCompatibility-class), 12
- performer_assnref, Audit-method
(zzz-DTTCompatibility-class), 12
- Performer_assnref-class
(zzz-DTTCompatibility-class), 12
- performers_assnreflist
(zzz-DTTCompatibility-class), 12
- performers_assnreflist, ProtocolApplication-method
(zzz-DTTCompatibility-class), 12
- Performers_assnreflist-class
(zzz-DTTCompatibility-class), 12
- Person-class
(zzz-DTTCompatibility-class), 12
- Person_assnlist-class
(zzz-ARRCompatibility-class), 12
- person_ref
(zzz-DTTCompatibility-class), 12
- person_ref, ArrayManufacturers_assnreflist-method
(zzz-DTTCompatibility-class), 12
- person_ref, Contacts_assnreflist-method
(zzz-DTTCompatibility-class), 12
- person_ref, DesignProviders_assnreflist-method
(zzz-DTTCompatibility-class), 12
- person_ref, HardwareManufacturers_assnreflist-method
(zzz-DTTCompatibility-class), 12
- person_ref, Members_assnreflist-method
(zzz-DTTCompatibility-class), 12
- person_ref, Owner_assnreflist-method
(zzz-DTTCompatibility-class), 12
- person_ref, Performer_assnref-method
(zzz-DTTCompatibility-class), 12
- person_ref, Performers_assnreflist-method
(zzz-DTTCompatibility-class), 12
- person_ref, Providers_assnreflist-method
(zzz-DTTCompatibility-class), 12
- person_ref, SoftwareManufacturers_assnreflist-method
(zzz-DTTCompatibility-class), 12
- person_ref, SourceContact_assnreflist-method
(zzz-DTTCompatibility-class), 12
- Person_ref-class
(zzz-DTTCompatibility-class), 12
- phone
(zzz-DTTCompatibility-class), 12
- phone, Organization-method
(zzz-DTTCompatibility-class), 12
- phone, Person-method
(zzz-DTTCompatibility-class), 12
- physicalArray
(zzz-ARRCompatibility-class), 12

- 11
- physicalArray, PhysicalArrays-method
(zzz-ARRCompatibility-class),
11
- PhysicalArray-class
(zzz-ARRCompatibility-class),
11
- physicalArrayDesign
(zzz-DTTCompatibility-class),
12
- physicalArrayDesign, ArrayDesign_assnlist-method
(zzz-DTTCompatibility-class),
12
- PhysicalArrayDesign-class
(zzz-DTTCompatibility-class),
12
- physicalArrayDesign_ref
(zzz-DTTCompatibility-class),
12
- physicalArrayDesign_ref, ArrayDesign_assnref-method
(zzz-DTTCompatibility-class),
12
- PhysicalArrayDesign_ref-class
(zzz-DTTCompatibility-class),
12
- physicalArrays
(zzz-ARRCompatibility-class),
11
- physicalArrays, ArraySetFile-method
(zzz-ARRCompatibility-class),
11
- PhysicalArrays-class
(zzz-ARRCompatibility-class),
11
- physicalBioAssay
(zzz-DTTCompatibility-class),
12
- physicalBioAssay, BioAssay_assnlist-method
(zzz-DTTCompatibility-class),
12
- physicalBioAssay, PhysicalBioAssay_assnlist-method
(zzz-DTTCompatibility-class),
12
- PhysicalBioAssay-class
(zzz-DTTCompatibility-class),
12
- PhysicalBioAssay_assnlist-class
(zzz-DTTCompatibility-class),
12
- PhysicalBioAssay_assnref-class
(zzz-DTTCompatibility-class),
12
- physicalBioAssay_ref
(zzz-DTTCompatibility-class),
12
- physicalBioAssay_ref, BioAssay_assnref-method
(zzz-DTTCompatibility-class),
12
- physicalBioAssay_ref, BioAssays_assnreflist-method
(zzz-DTTCompatibility-class),
12
- physicalBioAssay_ref, PhysicalBioAssay_assnref-method
(zzz-DTTCompatibility-class),
12
- physicalBioAssay_ref, PhysicalBioAssaySource_assnref-method
(zzz-DTTCompatibility-class),
12
- physicalBioAssay_ref, PhysicalBioAssayTarget_assnref-method
(zzz-DTTCompatibility-class),
12
- physicalBioAssay_ref, SourceBioAssays_assnreflist-method
(zzz-DTTCompatibility-class),
12
- physicalBioAssay_ref, Target_assnref-method
(zzz-DTTCompatibility-class),
12
- physicalBioAssay_ref, TopLevelBioAssays_assnref-method
(zzz-DTTCompatibility-class),
12
- PhysicalBioAssay_ref-class
(zzz-DTTCompatibility-class),
12
- physicalBioAssayData_assnlist
(zzz-DTTCompatibility-class),
12
- physicalBioAssayData_assnlist, PhysicalBioAssayAssayData_assnlist-method
(zzz-DTTCompatibility-class),
12
- PhysicalBioAssayData_assnlist-class
(zzz-DTTCompatibility-class),
12
- physicalBioAssaySource_assnref
(zzz-DTTCompatibility-class),
12
- physicalBioAssaySource_assnref, FeatureExtractAssaySource_assnref-method
(zzz-DTTCompatibility-class),
12
- PhysicalBioAssaySource_assnref-class
(zzz-DTTCompatibility-class),
12
- PhysicalBioAssayTarget_assnref-class
(zzz-DTTCompatibility-class),
12
- polymerType_assn

- (zzz-*DTTCompatibility-class*),
12
- polymerType_assn, BioSequence-method
(zzz-*DTTCompatibility-class*),
12
- PolymerType_assn-class
(zzz-*DTTCompatibility-class*),
12
- position
(zzz-*DTTCompatibility-class*),
12
- position, Position_assn-method
(zzz-*DTTCompatibility-class*),
12
- Position-class
(zzz-*DTTCompatibility-class*),
12
- position_assn
(zzz-*DTTCompatibility-class*),
12
- position_assn, Feature-method
(zzz-*DTTCompatibility-class*),
12
- position_assn, Fiducial-method
(zzz-*DTTCompatibility-class*),
12
- Position_assn-class
(zzz-*DTTCompatibility-class*),
12
- positionDelta
(zzz-*DTTCompatibility-class*),
12
- positionDelta, PositionDelta_assn-method
(zzz-*DTTCompatibility-class*),
12
- PositionDelta-class
(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
- PositionDelta_assn-class
(zzz-*DTTCompatibility-class*),
12
- presentAbsent
(zzz-*DTTCompatibility-class*),
12
- presentAbsent, QuantitationType_assnlist-method
(zzz-*DTTCompatibility-class*),
12
- PresentAbsent-class
(zzz-*DTTCompatibility-class*),
12
- presentAbsent_ref
(zzz-*DTTCompatibility-class*),
12
- presentAbsent_ref, QuantitationType_assnref-method
(zzz-*DTTCompatibility-class*),
12
- presentAbsent_ref, QuantitationTypes_assnreflist
(zzz-*DTTCompatibility-class*),
12
- presentAbsent_ref, SourcesQuantitationType_assn
(zzz-*DTTCompatibility-class*),
12
- presentAbsent_ref, TargetQuantitationType_assn
(zzz-*DTTCompatibility-class*),
12
- PresentAbsent_ref-class
(zzz-*DTTCompatibility-class*),
12
- producerTransformation_assn
(zzz-*DTTCompatibility-class*),
12
- producerTransformation_assn, DerivedBioAssayData
(zzz-*DTTCompatibility-class*),
12
- ProducerTransformation_assn-class
(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
(zzz-*DTTCompatibility-class*),
12

propertySets_assnlist, Audit-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, BibliographicReference-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, BioAssayCreation-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, BioAssayDataCluster-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, BioAssayDimension-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, BioAssayMap-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, BioAssayMapping-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, BioAssayTreatment-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, BioDataCube-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, BioDataTuples-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, BioMaterialMeasurement-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, BioSample-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, BioSequence-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, BioSource-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, Channel-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, CompositeCompositeMap-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, CompositeGroup-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, CompositePosition-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, CompositeSequence-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, CompositeSequenceDimension-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, Compound-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, CompoundMeasurement-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, ConcentrationUnit-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, Database-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, DatabaseEntry-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, DerivedBioAssay-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, DerivedBioAssayData-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, DerivedSignal-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, Description-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, DesignElementMapping-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, DistanceUnit-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, Error-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, ExpectedValue-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 propertySets_assnlist, Experiment-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 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, MAGE-ML-method
 propertySets_assnlist, FactorValue-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, ManufactureLIMS-method
 propertySets_assnlist, Failed-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, ManufactureLIMSBiomateri
 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, MeasuredBioAssay-method
 propertySets_assnlist, FeatureDimension-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, MeasuredBioAssayData-met
 propertySets_assnlist, FeatureExtraction-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, MeasuredSignal-method
 propertySets_assnlist, FeatureGroup-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, Measurement-method
 propertySets_assnlist, FeatureInformation-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, MismatchInformation-meth
 propertySets_assnlist, FeatureLocation-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, NameValueType-method
 propertySets_assnlist, FeatureReporterMap-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, Node-method
 propertySets_assnlist, Fiducial-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, NodeContents-method
 propertySets_assnlist, Hardware-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, NodeValue-method
 propertySets_assnlist, HardwareApplication-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, OntologyEntry-method
 propertySets_assnlist, Hybridization-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, Organization-method
 propertySets_assnlist, Image-method (zzz-DTTCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12
 propertySets_assnlist, Parameter-method

- propertySets_assnlist, ZoneDefect-method (*zzz-DTTCompatibility-class*),
 (*zzz-DTTCompatibility-class*), 12
 propertySets_assnlist, ZoneGroup-method (*zzz-DTTCompatibility-class*),
 (*zzz-DTTCompatibility-class*), 12
 propertySets_assnlist, ZoneLayout-method (*zzz-DTTCompatibility-class*),
 (*zzz-DTTCompatibility-class*), 12
 PropertySets_assnlist-class (*zzz-DTTCompatibility-class*),
 12
 protocol (*zzz-DTTCompatibility-class*),
 12
 protocol, Protocol_assnlist-method (*zzz-DTTCompatibility-class*),
 12
 Protocol-class (*zzz-DTTCompatibility-class*),
 12
 protocol_assnlist (*zzz-DTTCompatibility-class*),
 12
 protocol_assnlist, Protocol_package-method (*zzz-DTTCompatibility-class*),
 12
 Protocol_assnlist-class (*zzz-DTTCompatibility-class*),
 12
 protocol_assnref (*zzz-DTTCompatibility-class*),
 12
 protocol_assnref, ProtocolApplication-method (*zzz-DTTCompatibility-class*),
 12
 Protocol_assnref-class (*zzz-DTTCompatibility-class*),
 12
 protocol_package (*zzz-DTTCompatibility-class*),
 12
 protocol_package, MAGE-ML-method (*zzz-DTTCompatibility-class*),
 12
 Protocol_package-class (*zzz-DTTCompatibility-class*),
 12
 protocol_ref (*zzz-DTTCompatibility-class*),
 12
 protocol_ref, Protocol_assnref-method (*zzz-DTTCompatibility-class*),
 12
 Protocol_ref-class (*zzz-DTTCompatibility-class*),
 12
 protocolApplication (*zzz-DTTCompatibility-class*),
 12
 protocolApplication, ProtocolApplications_assnlist (*zzz-DTTCompatibility-class*),
 12
 ProtocolApplication-class (*zzz-DTTCompatibility-class*),
 12
 protocolApplications_assnlist (*zzz-DTTCompatibility-class*),
 12
 protocolApplications_assnlist, ArrayDesign-method (*zzz-DTTCompatibility-class*),
 12
 protocolApplications_assnlist, ArrayManufacture (*zzz-DTTCompatibility-class*),
 12
 protocolApplications_assnlist, BioAssayCreation (*zzz-DTTCompatibility-class*),
 12
 protocolApplications_assnlist, BioAssayMap-method (*zzz-DTTCompatibility-class*),
 12
 protocolApplications_assnlist, BioAssayTreatment (*zzz-DTTCompatibility-class*),
 12
 protocolApplications_assnlist, CompositeComposition (*zzz-DTTCompatibility-class*),
 12
 protocolApplications_assnlist, FeatureExtraction (*zzz-DTTCompatibility-class*),
 12
 protocolApplications_assnlist, FeatureReporterMethod (*zzz-DTTCompatibility-class*),
 12
 protocolApplications_assnlist, HybridizationMethod (*zzz-DTTCompatibility-class*),
 12
 protocolApplications_assnlist, ImageAcquisition (*zzz-DTTCompatibility-class*),
 12
 protocolApplications_assnlist, PhysicalArrayDesign (*zzz-DTTCompatibility-class*),
 12
 protocolApplications_assnlist, QuantitationType (*zzz-DTTCompatibility-class*),
 12

- 12 pvalue_ref, QuantitationType_assnref-method
 protocolApplications_assnlist, ReporterCompositionMapCompatibility-class),
 (zzz-DTTCompatibility-class), 12
 12 pvalue_ref, QuantitationTypes_assnreflist-method
 protocolApplications_assnlist, Transformation-method Compatibility-class),
 (zzz-DTTCompatibility-class), 12
 12 pvalue_ref, SourcesQuantitationType_assnreflist
 protocolApplications_assnlist, Treatment-method Compatibility-class),
 (zzz-DTTCompatibility-class), 12
 12 pvalue_ref, TargetQuantitationType_assnref-meth
 ProtocolApplications_assnlist-class
 (zzz-DTTCompatibility-class), 12
 12 PValue_ref-class
 providers_assnreflist
 (zzz-DTTCompatibility-class), 12
 12
 providers_assnreflist, Experiment-method quality
 (zzz-DTTCompatibility-class),
 12
 12 quality, ManufactureLIMS-method
 Providers_assnreflist-class
 (zzz-DTTCompatibility-class),
 12
 12 quality, ManufactureLIMSBiomaterial-method
 publication
 (zzz-DTTCompatibility-class),
 12
 12 qualityControlDescription_assn
 publication, BibliographicReference-method
 (zzz-DTTCompatibility-class),
 12
 12 qualityControlDescription_assn, ExperimentDesign
 publisher
 (zzz-DTTCompatibility-class),
 12
 12 QualityControlDescription_assn-class
 publisher, BibliographicReference-method
 (zzz-DTTCompatibility-class),
 12
 12 qualityControlStatistics_assnlist
 pvalue
 (zzz-DTTCompatibility-class),
 12
 12 qualityControlStatistics_assnlist, ArrayManufac
 pvalue, ConfidenceIndicator_assnlist-method
 (zzz-DTTCompatibility-class),
 12
 12 qualityControlStatistics_assnlist, BioSample-me
 pvalue, QuantitationType_assnlist-method
 (zzz-DTTCompatibility-class),
 12
 12 qualityControlStatistics_assnlist, BioSource-me
 PValue-class
 (zzz-DTTCompatibility-class),
 12
 12 qualityControlStatistics_assnlist, LabeledExtra
 pvalue_ref
 (zzz-DTTCompatibility-class),
 12
 12 QualityControlStatistics_assnlist-class
 pvalue_ref, ConfidenceIndicators_assnreflist-method
 (zzz-DTTCompatibility-class),
 12
 12 quantitationDimension_assnref

- QuantitationTypeMap_ref-class
 (*zzz-DTTCompatibility-class*),
 12
- quantitationTypeMapping
 (*zzz-DTTCompatibility-class*),
 12
- quantitationTypeMapping, QuantitationTypeMapping_assn-method
 (*zzz-DTTCompatibility-class*),
 12
- QuantitationTypeMapping-class
 (*zzz-DTTCompatibility-class*),
 12
- quantitationTypeMapping_assn
 (*zzz-DTTCompatibility-class*),
 12
- quantitationTypeMapping_assn, Transformation-method
 (*zzz-DTTCompatibility-class*),
 12
- QuantitationTypeMapping_assn-class
 (*zzz-DTTCompatibility-class*),
 12
- quantitationTypeMaps_assnreflist
 (*zzz-DTTCompatibility-class*),
 12
- quantitationTypeMaps_assnreflist, DerivedSignal-method
 (*zzz-DTTCompatibility-class*),
 12
- quantitationTypeMaps_assnreflist, Error-method
 (*zzz-DTTCompatibility-class*),
 12
- quantitationTypeMaps_assnreflist, ExpectedValue-method
 (*zzz-DTTCompatibility-class*),
 12
- quantitationTypeMaps_assnreflist, Failed-method
 (*zzz-DTTCompatibility-class*),
 12
- quantitationTypeMaps_assnreflist, MeasuredSignal-method
 (*zzz-DTTCompatibility-class*),
 12
- quantitationTypeMaps_assnreflist, PresentAbsent-method
 (*zzz-DTTCompatibility-class*),
 12
- quantitationTypeMaps_assnreflist, PValue-method
 (*zzz-DTTCompatibility-class*),
 12
- quantitationTypeMaps_assnreflist, QuantitationTypeMappings-method
 (*zzz-DTTCompatibility-class*),
 12
- quantitationTypeMaps_assnreflist, Ratio-method
 (*zzz-DTTCompatibility-class*),
 12
- quantitationTypeMaps_assnreflist, SpecializedQuantitationType-method
 (*zzz-DTTCompatibility-class*),
 12
- (*zzz-DTTCompatibility-class*),
 12
- QuantitationTypeMaps_assnreflist-class
 (*zzz-DTTCompatibility-class*),
 12
- QuantitationTypeMaps_assnreflist-class
 (*zzz-DTTCompatibility-class*),
 12
- quantitationTypeTuple
 (*zzz-DTTCompatibility-class*),
 12
- quantitationTypeTuple, QuantitationTypeTuples_a
 (*zzz-DTTCompatibility-class*),
 12
- QuantitationTypeTuple-class
 (*zzz-DTTCompatibility-class*),
 12
- quantitationTypeTuples_assnlist
 (*zzz-DTTCompatibility-class*),
 12
- quantitationTypeTuples_assnlist, DesignElementT
 (*zzz-DTTCompatibility-class*),
 12
- QuantitationTypeTuples_assnlist-class
 (*zzz-DTTCompatibility-class*),
 12
- quantityUnit
 (*zzz-DTTCompatibility-class*),
 12
- quantityUnit, Unit_assn-method
 (*zzz-DTTCompatibility-class*),
 12
- QuantityUnit-class
 (*zzz-DTTCompatibility-class*),
 12
- ratio
 (*zzz-DTTCompatibility-class*),
 12
- ratio, QuantitationType_assnlist-method
 (*zzz-DTTCompatibility-class*),
 12
- Ratio-class
 (*zzz-DTTCompatibility-class*),
 12
- ratio_ref
 (*zzz-DTTCompatibility-class*),
 12

- [12](#)
 ratio_ref, QuantitationType_assnref-method
 (*zzz-DTTCompatibility-class*),
 [12](#)
- ratio_ref, QuantitationTypes_assnreflist-method
 (*zzz-DTTCompatibility-class*),
 [12](#)
- ratio_ref, SourcesQuantitationType_assnref-method
 (*zzz-DTTCompatibility-class*),
 [12](#)
- ratio_ref, TargetQuantitationType_assnref-method
 (*zzz-DTTCompatibility-class*),
 [12](#)
- Ratio_ref-class
 (*zzz-DTTCompatibility-class*),
 [12](#)
- readAnnotation
 (*NetAffxResource-class*), [2](#)
- readAnnotation, NetAffxResource, character, character-method
 (*NetAffxResource-class*), [2](#)
- readAnnotation, NetAffxResource, character, numeric-method
 (*NetAffxResource-class*), [2](#)
- readAnnotation, NetAffxResource, missing, AffxAnnotation-list-method
 (*NetAffxResource-class*), [2](#)
- readAnnotation, NetAffxResource, numeric, character-method
 (*NetAffxResource-class*), [2](#)
- readAnnotation, NetAffxResource, numeric, numeric-method
 (*NetAffxResource-class*), [2](#)
- readArr, [6](#)
- readMage, [7](#)
- readNetAffx, [8](#)
- readXml, [7](#), [9](#), [10](#)
- regions_assnlist
 (*zzz-DTTCompatibility-class*),
 [12](#)
- regions_assnlist, SeqFeature-method
 (*zzz-DTTCompatibility-class*),
 [12](#)
- Regions_assnlist-class
 (*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](#)
- [12](#)
 replicateDescription_assn
 (*zzz-DTTCompatibility-class*),
 [12](#)
- replicateDescription_assn, ExperimentDesign-method
 (*zzz-DTTCompatibility-class*),
 [12](#)
- replicateDescription_assn-class
 (*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-class
 (*zzz-DTTCompatibility-class*),
 [12](#)
- reporter_assnlist, AffxAnnotation-list-method
 (*zzz-DTTCompatibility-class*),
 [12](#)
- reporter_assnlist, DesignElement_package-method
 (*zzz-DTTCompatibility-class*),
 [12](#)
- Reporter_assnlist-class
 (*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_assnref-class
 (*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), (zzz-DTTCompatibility-class), 12	ReporterDimension-class	(zzz-DTTCompatibility-class), 12
Reporter_ref-class	(zzz-DTTCompatibility-class), 12	reporterDimension_ref	(zzz-DTTCompatibility-class), 12
reporterCompositeMap	(zzz-DTTCompatibility-class), 12	reporterDimension_ref, DesignElementDimension_a	(zzz-DTTCompatibility-class), 12
reporterCompositeMap, DesignElementMap_assnlist	(zzz-DTTCompatibility-class), 12	ReporterDimension_ref-class	(zzz-DTTCompatibility-class), 12
reporterCompositeMap, ReporterCompositeMap_as	(zzz-DTTCompatibility-class), 12	reporterGroup	(zzz-DTTCompatibility-class), 12
ReporterCompositeMap-class	(zzz-DTTCompatibility-class), 12	reporterGroup, ReporterGroup_assnlist-method	(zzz-DTTCompatibility-class), 12
reporterCompositeMap_assnlist	(zzz-DTTCompatibility-class), 12	ReporterGroup-class	(zzz-DTTCompatibility-class), 12
reporterCompositeMap_assnlist, DesignElement_pack	(zzz-DTTCompatibility-class), 12	reporterGroup_assnlist	(zzz-DTTCompatibility-class), 12
ReporterCompositeMap_assnlist-class	(zzz-DTTCompatibility-class), 12	reporterGroup_assnlist, ArrayDesign_package-met	(zzz-DTTCompatibility-class), 12
reporterCompositeMap_ref	(zzz-DTTCompatibility-class), 12	ReporterGroup_assnlist-class	(zzz-DTTCompatibility-class), 12
reporterCompositeMap_ref, DesignElementMaps_a	(zzz-DTTCompatibility-class), 12	reporterGroup_ref	(zzz-DTTCompatibility-class), 12
reporterCompositeMap_ref, ReporterCompositeMap	(zzz-DTTCompatibility-class), 12	reporterGroup_ref, ReporterGroups_assnreflist-r	(zzz-DTTCompatibility-class), 12
ReporterCompositeMap_ref-class	(zzz-DTTCompatibility-class), 12	ReporterGroup_ref-class	(zzz-DTTCompatibility-class), 12
reporterCompositeMaps_assnreflist	(zzz-DTTCompatibility-class), 12	reporterGroups_assnreflist	(zzz-DTTCompatibility-class), 12
reporterCompositeMaps_assnreflist, CompositeS	(zzz-DTTCompatibility-class), 12	reporterGroups_assnreflist, ArrayDesign-method	(zzz-DTTCompatibility-class), 12
ReporterCompositeMaps_assnreflist-class	(zzz-DTTCompatibility-class), 12	reporterGroups_assnreflist, PhysicalArrayDesign	(zzz-DTTCompatibility-class), 12
reporterDimension	(zzz-DTTCompatibility-class), 12	ReporterGroups_assnreflist-class	(zzz-DTTCompatibility-class), 12
reporterDimension, DesignElementDimension_ass	(zzz-DTTCompatibility-class), 12		

- 12
- reporterPosition
 - (zzz-DTTCompatibility-class), 12
- reporterPosition, Coordinate_assn-method
 - (zzz-DTTCompatibility-class), 12
- reporterPosition, ReporterPositionSources_assnlist-method
 - (zzz-DTTCompatibility-class), 12
- ReporterPosition-class
 - (zzz-DTTCompatibility-class), 12
- reporterPositionSources_assnlist
 - (zzz-DTTCompatibility-class), 12
- reporterPositionSources_assnlist, ReporterCompatibility-method
 - (zzz-DTTCompatibility-class), 12
- ReporterPositionSources_assnlist-class
 - (zzz-DTTCompatibility-class), 12
- reporters_assnreflist
 - (zzz-DTTCompatibility-class), 12
- reporters_assnreflist, ReporterDimension-method
 - (zzz-DTTCompatibility-class), 12
- reporters_assnreflist, ReporterGroup-method
 - (zzz-DTTCompatibility-class), 12
- Reporters_assnreflist-class
 - (zzz-DTTCompatibility-class), 12
- required
 - (zzz-ARRCompatibility-class), 11
- required, UserAttribute-method
 - (zzz-ARRCompatibility-class), 11
- roles_assnlist
 - (zzz-DTTCompatibility-class), 12
- roles_assnlist, Organization-method
 - (zzz-DTTCompatibility-class), 12
- roles_assnlist, Person-method
 - (zzz-DTTCompatibility-class), 12
- Roles_assnlist-class
 - (zzz-DTTCompatibility-class), 12
- scale_assn
 - (zzz-DTTCompatibility-class), 12
- scale_assn, DerivedSignal-method
 - (zzz-DTTCompatibility-class), 12
- scale_assn, Error-method
 - (zzz-DTTCompatibility-class), 12
- scale_assn, ExpectedValue-method
 - (zzz-DTTCompatibility-class), 12
- scale_assn, Failed-method
 - (zzz-DTTCompatibility-class), 12
- scale_assn, MeasuredSignal-method
 - (zzz-DTTCompatibility-class), 12
- scale_assn, NodeValue-method
 - (zzz-DTTCompatibility-class), 12
- scale_assn, PresentAbsent-method
 - (zzz-DTTCompatibility-class), 12
- scale_assn, PValue-method
 - (zzz-DTTCompatibility-class), 12
- scale_assn, Ratio-method
 - (zzz-DTTCompatibility-class), 12
- scale_assn, SpecializedQuantitationType-method
 - (zzz-DTTCompatibility-class), 12
- Scale_assn-class
 - (zzz-DTTCompatibility-class), 12
- security
 - (zzz-DTTCompatibility-class), 12
- security, Security_assnlist-method
 - (zzz-DTTCompatibility-class), 12
- Security-class
 - (zzz-DTTCompatibility-class), 12
- security_assnlist
 - (zzz-DTTCompatibility-class), 12
- security_assnlist, AuditAndSecurity_package-method
 - (zzz-DTTCompatibility-class), 12
- Security_assnlist-class

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

- 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_assnref-class
(zzz-DTTCompatibility-class),
12
- security_ref
(zzz-DTTCompatibility-class),
12
- security_ref, Security_assnref-method
(zzz-DTTCompatibility-class),
12
- Security_ref-class
(zzz-DTTCompatibility-class),
12
- securityGroup
(zzz-DTTCompatibility-class),
12
- securityGroup, SecurityGroup_assnlist-method
(zzz-DTTCompatibility-class),
12
- SecurityGroup-class
(zzz-DTTCompatibility-class),
12
- securityGroup_assnlist
(zzz-DTTCompatibility-class),
12
- securityGroup_assnlist, AuditAndSecurity_packag
(zzz-DTTCompatibility-class),
12
- SecurityGroup_assnlist-class
(zzz-DTTCompatibility-class),
12
- securityGroup_ref
(zzz-DTTCompatibility-class),
12
- securityGroup_ref, SecurityGroups_assnreflist-r
(zzz-DTTCompatibility-class),
12
- SecurityGroup_ref-class
(zzz-DTTCompatibility-class),
12
- securityGroups_assnreflist
(zzz-DTTCompatibility-class),
12
- securityGroups_assnreflist, Security-method
(zzz-DTTCompatibility-class),
12
- SecurityGroups_assnreflist-class
(zzz-DTTCompatibility-class),
12
- seqFeature
(zzz-DTTCompatibility-class),
12
- seqFeature, SeqFeatures_assnlist-method
(zzz-DTTCompatibility-class),
12
- SeqFeature-class
(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
- SeqFeatureLocation-class
(zzz-DTTCompatibility-class),
12
- seqFeatures_assnlist
(zzz-DTTCompatibility-class),
12
- seqFeatures_assnlist, BioSequence-method
(zzz-DTTCompatibility-class),
12
- SeqFeatures_assnlist-class

- (*zzz-DTTCompatibility-class*), 12
- sequenceDatabases_assnlist (*zzz-DTTCompatibility-class*), 12
- sequenceDatabases_assnlist, BioSequence-method (*zzz-DTTCompatibility-class*), 12
- SequenceDatabases_assnlist-class (*zzz-DTTCompatibility-class*), 12
- sequencePosition (*zzz-DTTCompatibility-class*), 12
- sequencePosition, Coordinate_assn-method (*zzz-DTTCompatibility-class*), 12
- SequencePosition-class (*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-class (*zzz-DTTCompatibility-class*), 12
- software_assnlist (*zzz-DTTCompatibility-class*), 12
- software_assnlist, Protocol_package-method (*zzz-DTTCompatibility-class*), 12
- Software_assnlist-class (*zzz-DTTCompatibility-class*), 12
- software_assnref (*zzz-DTTCompatibility-class*), 12
- software_assnref, SoftwareApplication-method (*zzz-DTTCompatibility-class*), 12
- Software_assnref-class (*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
- Software_ref-class (*zzz-DTTCompatibility-class*), 12
- softwareApplication (*zzz-DTTCompatibility-class*), 12
- softwareApplication, SoftwareApplications_assnlist (*zzz-DTTCompatibility-class*), 12
- SoftwareApplication-class (*zzz-DTTCompatibility-class*), 12
- softwareApplications_assnlist (*zzz-DTTCompatibility-class*), 12
- softwareApplications_assnlist, ProtocolApplication (*zzz-DTTCompatibility-class*), 12
- SoftwareApplications_assnlist-class (*zzz-DTTCompatibility-class*), 12
- softwareManufacturers_assnreflist (*zzz-DTTCompatibility-class*), 12
- softwareManufacturers_assnreflist, Software-man (*zzz-DTTCompatibility-class*), 12
- SoftwareManufacturers_assnreflist-class (*zzz-DTTCompatibility-class*), 12
- softwares_assnreflist (*zzz-DTTCompatibility-class*), 12
- softwares_assnreflist, Hardware-method (*zzz-DTTCompatibility-class*), 12

softwares_assnreflist, Protocol-method	(zzz-DTTCompatibility-class), 12	spacingBetweenCols, ZoneLayout-method	(zzz-DTTCompatibility-class), 12
softwares_assnreflist, Software-method	(zzz-DTTCompatibility-class), 12	spacingBetweenRows	(zzz-DTTCompatibility-class), 12
Softwares_assnreflist-class	(zzz-DTTCompatibility-class), 12	spacingBetweenRows, ZoneLayout-method	(zzz-DTTCompatibility-class), 12
sourceBioAssays_assnreflist	(zzz-DTTCompatibility-class), 12	spacingsBetweenZonesX	(zzz-DTTCompatibility-class), 12
sourceBioAssays_assnreflist, BioAssayMap-method	(zzz-DTTCompatibility-class), 12	spacingsBetweenZonesX, ZoneGroup-method	(zzz-DTTCompatibility-class), 12
SourceBioAssays_assnreflist-class	(zzz-DTTCompatibility-class), 12	spacingsBetweenZonesY	(zzz-DTTCompatibility-class), 12
sourceBioMaterialMeasurements_assnlist	(zzz-DTTCompatibility-class), 12	spacingsBetweenZonesY, ZoneGroup-method	(zzz-DTTCompatibility-class), 12
sourceBioMaterialMeasurements_assnlist, BioAssayMap-method	(zzz-DTTCompatibility-class), 12	specializedQuantitationType	(zzz-DTTCompatibility-class), 12
sourceBioMaterialMeasurements_assnlist, Hybridization-method	(zzz-DTTCompatibility-class), 12	specializedQuantitationType, QuantitationType_ref	(zzz-DTTCompatibility-class), 12
sourceBioMaterialMeasurements_assnlist, Treatment-method	(zzz-DTTCompatibility-class), 12	SpecializedQuantitationType-class	(zzz-DTTCompatibility-class), 12
SourceBioMaterialMeasurements_assnlist-class	(zzz-DTTCompatibility-class), 12	specializedQuantitationType_ref	(zzz-DTTCompatibility-class), 12
sourceContact_assnreflist	(zzz-DTTCompatibility-class), 12	specializedQuantitationType_ref, QuantitationType_ref	(zzz-DTTCompatibility-class), 12
sourceContact_assnreflist, BioSource-method	(zzz-DTTCompatibility-class), 12	specializedQuantitationType_ref, QuantitationType_ref, SourcesQuantitationType_ref	(zzz-DTTCompatibility-class), 12
SourceContact_assnreflist-class	(zzz-DTTCompatibility-class), 12	specializedQuantitationType_ref, TargetQuantitationType_ref	(zzz-DTTCompatibility-class), 12
sourcesQuantitationType_assnreflist	(zzz-DTTCompatibility-class), 12	SpecializedQuantitationType_ref-class	(zzz-DTTCompatibility-class), 12
sourcesQuantitationType_assnreflist, QuantitationTypeMap-method	(zzz-DTTCompatibility-class), 12	species_assn	(zzz-DTTCompatibility-class), 12
SourcesQuantitationType_assnreflist-class	(zzz-DTTCompatibility-class), 12		
spacingBetweenCols			

- 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
- Species_assn-class
(*zzz-DTTCompatibility-class*),
12
- StandardQuantitationType_ref-class
(*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
- Subregions_assnlist-class
(*zzz-DTTCompatibility-class*),
12
- substrateType_assn
(*zzz-DTTCompatibility-class*),
12
- substrateType_assn, ArrayGroup-method
(*zzz-DTTCompatibility-class*),
12
- SubstrateType_assn-class
(*zzz-DTTCompatibility-class*),
12
- summaryStatistics_assnlist
(*zzz-DTTCompatibility-class*),
12
- summaryStatistics_assnlist, DerivedBioAssayData
(*zzz-DTTCompatibility-class*),
12
- summaryStatistics_assnlist, MeasuredBioAssayData
(*zzz-DTTCompatibility-class*),
12
- SummaryStatistics_assnlist-class
(*zzz-DTTCompatibility-class*),
12
- surfaceType_assn
(*zzz-DTTCompatibility-class*),
12
- surfaceType_assn, PhysicalArrayDesign-method
(*zzz-DTTCompatibility-class*),
12
- SurfaceType_assn-class
(*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
- Target_assnref-class
(*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
- targetQuantitationType_assnref, QuantitationTyp
(*zzz-DTTCompatibility-class*),
12
- TargetQuantitationType_assnref-class
(*zzz-DTTCompatibility-class*),
12
- technologyType_assn
(*zzz-DTTCompatibility-class*),
12
- technologyType_assn, FeatureGroup-method

- (zzz-*DTTCompatibility-class*),
12
- TechnologyType_assn-class
(zzz-*DTTCompatibility-class*),
12
- temperatureUnit
(zzz-*DTTCompatibility-class*),
12
- temperatureUnit, Unit_assn-method
(zzz-*DTTCompatibility-class*),
12
- TemperatureUnit-class
(zzz-*DTTCompatibility-class*),
12
- TemplateFile-class
(zzz-*ARRCompatibility-class*),
11
- timeUnit
(zzz-*DTTCompatibility-class*),
12
- timeUnit, Unit_assn-method
(zzz-*DTTCompatibility-class*),
12
- TimeUnit-class
(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
- TopLevelBioAssays_assnreflist-class
(zzz-*DTTCompatibility-class*),
12
- transformation
(zzz-*DTTCompatibility-class*),
12
- transformation, ProducerTransformation_assn-method
(zzz-*DTTCompatibility-class*),
12
- Transformation-class
(zzz-*DTTCompatibility-class*),
12
- treatment
(zzz-*DTTCompatibility-class*),
12
- treatment, Treatments_assnlist-method
(zzz-*DTTCompatibility-class*),
12
- Treatment-class
(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
- Treatments_assnlist-class
(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
- Type_assn-class (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
- Types_assnlist-class (zzz-DTTCompatibility-class), 12
- unit_assn (zzz-DTTCompatibility-class), 12
- unit_assn, Measurement-method (zzz-DTTCompatibility-class), 12
- Unit_assn-class (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

- 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
- UserAttribute-class
 - (*zzz-ARRCompatibility-class*), 11
- userAttributes
 - (*zzz-ARRCompatibility-class*), 11
- userAttributes, ArraySetFile-method
 - (*zzz-ARRCompatibility-class*), 11
- userAttributes, TemplateFile-method
 - (*zzz-ARRCompatibility-class*), 11
- UserAttributes-class
 - (*zzz-ARRCompatibility-class*), 11
- userAttributeValue
 - (*zzz-ARRCompatibility-class*), 11
- userAttributeValue, UserAttribute-method
 - (*zzz-ARRCompatibility-class*), 11
- UserAttributeValue-class
 - (*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
- Value_assn-class
(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
- VolumeUnit-class
(zzz-DTTCompatibility-class),
12
- warningType_assn
(zzz-DTTCompatibility-class),
12
- warningType_assn, Reporter-method
(zzz-DTTCompatibility-class),
12
- WarningType_assn-class
(zzz-DTTCompatibility-class),
- 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-class
(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_assnref-class
(zzz-DTTCompatibility-class),
12
- zone_ref
(zzz-DTTCompatibility-class),
12
- zone_ref, Zone_assnref-method
(zzz-DTTCompatibility-class),
12

Zone_ref-class
 (*zzz-DTTCompatibility-class*),
 12

zoneDefect
 (*zzz-DTTCompatibility-class*),
 12

zoneDefect, Adjustments_assnlist-method
 (*zzz-DTTCompatibility-class*),
 12

ZoneDefect-class
 (*zzz-DTTCompatibility-class*),
 12

zoneGroup
 (*zzz-DTTCompatibility-class*),
 12

zoneGroup, ZoneGroups_assnlist-method
 (*zzz-DTTCompatibility-class*),
 12

ZoneGroup-class
 (*zzz-DTTCompatibility-class*),
 12

zoneGroups_assnlist
 (*zzz-DTTCompatibility-class*),
 12

zoneGroups_assnlist, PhysicalArrayDesign-method
 (*zzz-DTTCompatibility-class*),
 12

ZoneGroups_assnlist-class
 (*zzz-DTTCompatibility-class*),
 12

zoneLayout
 (*zzz-DTTCompatibility-class*),
 12

zoneLayout, ZoneLayout_assn-method
 (*zzz-DTTCompatibility-class*),
 12

ZoneLayout-class
 (*zzz-DTTCompatibility-class*),
 12

zoneLayout_assn
 (*zzz-DTTCompatibility-class*),
 12

zoneLayout_assn, ZoneGroup-method
 (*zzz-DTTCompatibility-class*),
 12

ZoneLayout_assn-class
 (*zzz-DTTCompatibility-class*),
 12

zoneLocations_assnlist
 (*zzz-DTTCompatibility-class*),
 12

zoneLocations_assnlist, ZoneGroup-method
 (*zzz-DTTCompatibility-class*),
 12

ZoneLocations_assnlist-class
 (*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