

ATalkARexx

COLLABORATORS

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Chapter 1

ATalkARexx

1.1 AmigaTalk© 1998 ARexx documentation:

Herein is documented the ARexx Commands that TheBrowser can issue to the AmigaTalk program.

ARexx commands recognized by AmigaTalk:

PlaceCodesInFile

PurgeClass

ReloadClass

GetClassListType

ReloadMethod

AddClass

AddMethod

Quit

GetError

GetClassFileName

ReportStatus

1.2 PlaceCodesInFile command:

SYNOPSIS

success = PlaceCodesInFile filename classname methodname

Get the byteCodes for the given method & given class & write them to the filename given.

1.3 PurgeClass command:

SYNOPSIS

success = PurgeClass classname

Remove the given class from the AmigaTalk internal Class Dictionary.

1.4 ReloadClass command:

SYNOPSIS

success = ReloadClass tempfilename

Load the contents of the given file into the system using the)i system directive of AmigaTalk.

1.5 GetClassListType command:

SYNOPSIS

typestring = GetClassListType

Get a string from a Requester that describes the first class description file set to load into TheBrowser. TheBrowser only executes this command once on startup.

1.6 ReloadMethod command:

SYNOPSIS

success = ReloadMethod filename classname methodname

Using the)i system directive, load in the given method from the given class & located in the given file.

1.7 AddClass command:

SYNOPSIS

success = AddClass filename classname

Using the)i system directive, load in the given class located in the given file.

1.8 AddMethod command:

SYNOPSIS

success = AddMethod filename classname methodname

Using the)i system directive, load in the given method from the given class & located in the given file.

1.9 Quit command:

SYNOPSIS

Quit

Tell AmigaTalk to shut down the ARexx Port, because TheBrowser has been closed.

1.10 GetError command:

SYNOPSIS

errorstring = GetError errornumber

Translate the given error number into an error description string.

1.11 GetClassFileName command:

SYNOPSIS

filename = GetClassFileName classname

Return the source filename of the given class.

1.12 ReportStatus command:

SYNOPSIS

status = ReportStatus

This command currently does nothing.
