

caArray 036 - Working with Array Design Files in caArray

Working with Array Design Files in caArray

Topic: caArray Usage

Release: Up to caArray 2.X

Date entered: 12/18/2009

Introduction

One of the most frequently asked questions from the new caArray users is: "Where can I obtain the array design file for my experiment?". Although the array providers' websites are still the best places to obtain the most updated array design files (for more details, see [caArray 010 - Where are the Array Design Files for caArray? Where are the Array Design Files for caArray?](#)), many array design files are now available to download at the NCI public [caArray server](#), a new feature made available in caArray **version 2.3**.

In this KB entry, we will explain how to download, edit, delete and replace an array design file in caArray.

Download an existing array design file

After login to caArray, the option to manage an array design will be available in the **Curation** section in the left menu. To select a file to download, follow the steps below:

- Click "Manage Array Design" in the left menu section (Arrow 1).
- All the design files that have been imported into caArray will be displayed in the main data area (Arrow 2).
- Locate the array design file of interest, select "Download" (Arrow 3) to launch the download.

Download of an existing array design file

Build CAARRAY_R2_3_0_GA(17213) Thu Nov 05 11:14:47 EST 2009 | Node: NCICB | Welcome, inf | Logout

WELCOME TO CAARRAY

HOME

BROWSE

EXPERIMENT MANAGEMENT

MY EXPERIMENT WORKSPACE

CREATE/PROPOSE EXPERIMENT

USER MANAGEMENT

MANAGE COLLABORATION GROUPS

CURATION

MANAGE ARRAY DESIGNS

MANAGE PROTOCOLS

MANAGE VOCABULARY

ABOUT CAARRAY

WHAT IS CAARRAY?

RELEASE NOTES

TECHNICAL DOCUMENTATION

TRAINING

Manage Array Designs

Array Designs Import a New Array Design

84 items found, displaying 21 to 40. [First/Prev] 1, 2, 3, 4, 5 [Next/Last]

Array Design Name	Provider	Assay Type	Version Number	Feature Type	Organism	Edit	Edit File	Delete	Status	Download
HG-U133A_2	Affymetrix	Gene Expression	2	in_situ_oligo_features	Homo sapiens				Imported	
HG-U133B	Affymetrix	Gene Expression	1	in_situ_oligo_features	Homo sapiens				Imported	
hjay	Affymetrix	Exon	1	in_situ_oligo_features	Homo sapiens				Imported	
hjay_binary	Affymetrix	Exon	1	in_situ_oligo_features	Homo sapiens				Imported	
H-miRNA_8x15K	Agilent	microRNA	2	in_situ_oligo_features	Homo sapiens				Imported (Not Parsed)	
Hs-ATC7_6k-v5px_16Bx22Cx22R	GenePix	Gene Expression	1	spotted_ds_DNA_features	Homo sapiens				Imported	
HT_HG-U133A	Affymetrix	Gene Expression	1	in_situ_oligo_features	Homo sapiens				Imported	

Upload a new array design file

Any registered caArray user can upload, validate, and import the array designs from a supported array provider. If a desired array design file is not found in the caArray, the user can initiate the file upload into caArray:

- Click **Import a New Array Design** (Arrow 4 in the preceding illustration).
- This opens the **New Array Design** dialog box.
- Enter the appropriate information in the fields provided on **Array Design Details** form.

New Array Design Dialog with Form Completed

New Array Design (Step 1)

Array Design Details

Description: Array Design test file

Assay Type:

- + aCGH
- + Exon
- + Gene Expression
- + Methylation
- + microRNA

Selected Assay Type:

- Gene Expression

Provider: Affymetrix

Version Number: 1

Feature Type: in_situ_oligo_features (MO)

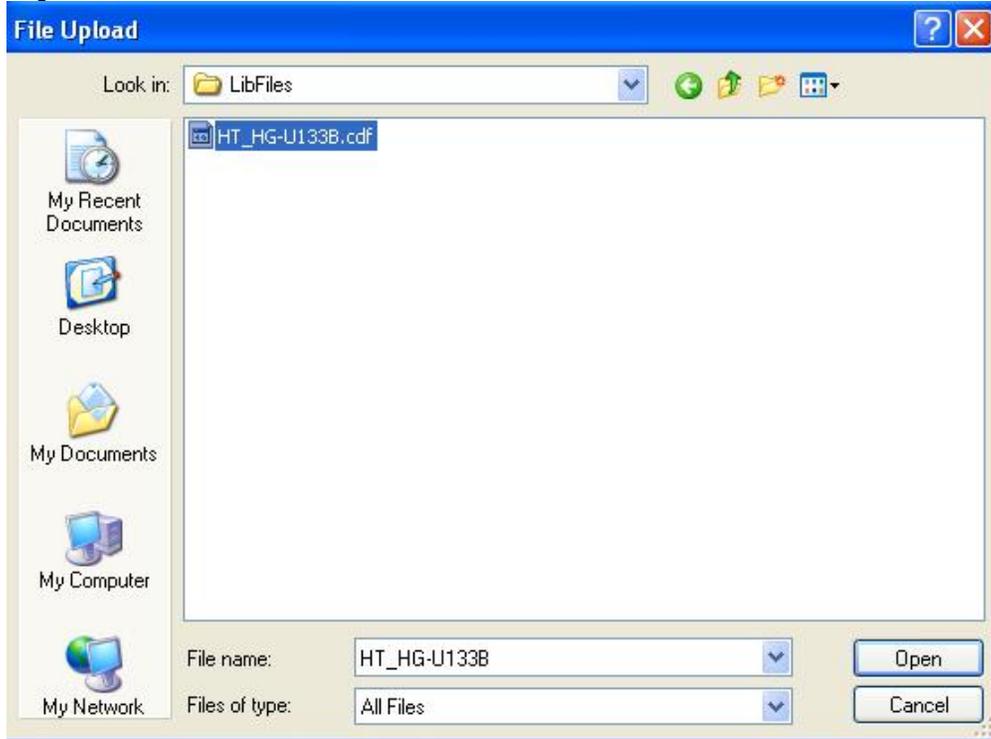
Organism: Homo sapiens (ncbitax)

Cancel Next

The required fields are the Description (Array Design test file in the example), Assay Type (select, Gene Expression in the example), Provider (Affymetrix in the example), Version Number (1 in the example), Feature Type (in_situ_oligo_features (MO) in the example) and Organism (Homo sapiens (ncbitax)).

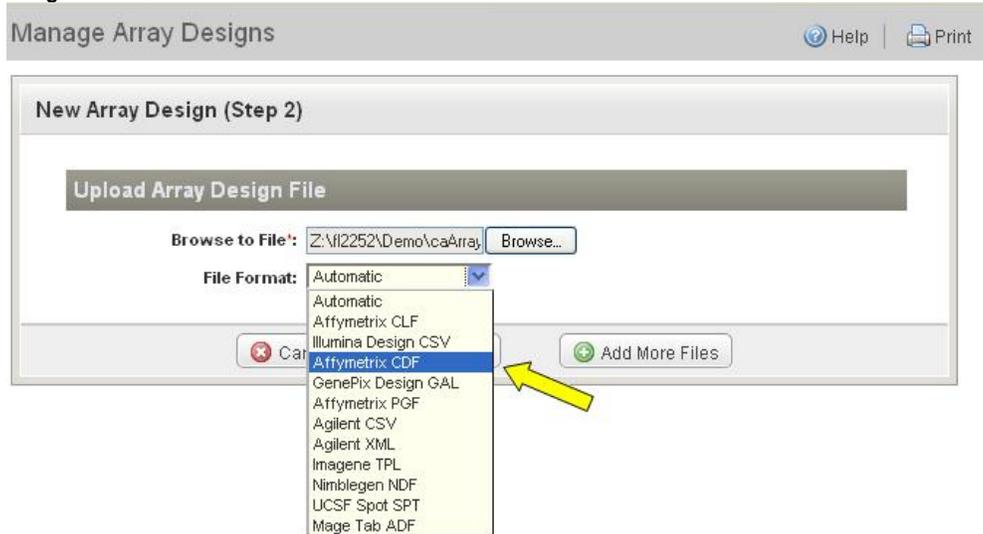
- Click "Next" to select an array design file of interest.

Selecting an array design file



- Identify the array design file format in the drop-down list.

Selecting the Array design file format



- Multiple array design Files can be submitted to upload at the same time. Click on "Add More Files" to select a new file. After all of the files are selected, submit the files to upload by clicking "Save".

Submitting files to upload

Manage Array Designs
Help | Print

New Array Design (Step 2)

Upload Array Design File

Browse to File:

File Format: Affymetrix CDF

Validate an array design file

- After the array design file is uploaded, caArray will automatically initiate the validation process. If the file is from a supported array provider, caArray will mark the file as "Imported" after the validation.

Array Design Files, Imported and Importing (Being Validated)

HG-U133A	Affymetrix	Gene Expression	1	in_situ_oligo_features	Homo sapiens			Imported
HT_HG-U133A	Affymetrix	Gene Expression	1.1	in_situ_oligo_features	Homo sapiens			Imported
HT_HG-U133B	Affymetrix	Gene Expression	1	in_situ_oligo_features	Homo sapiens			Importing
HT_Rat230_PM	Affymetrix	Gene Expression	1.0	in_situ_oligo_features	Rattus rattus			Imported
HuEx-1_0-st-ta1	Affymetrix	Exon	v2	in_situ_oligo_features	Homo sapiens			Imported

- For more information on the different states of the array design files, refer to [caArray 017 - What is the meaning of the caArray Status of Importing - Imported versus Imported Not Parsed?](#), the section "caArray Status of Importing: Imported vs Imported Not Parsed".

Array Design Files, Imported

HT_HG-U133A	Affymetrix	Gene Expression	1.1	in_situ_oligo_features	Homo sapiens			Imported
HT_HG-U133B	Affymetrix	Gene Expression	1	in_situ_oligo_features	Homo sapiens			Imported
HT_Rat230_PM	Affymetrix	Gene Expression	1.0	in_situ_oligo_features	Rattus rattus			Imported
HuEx-1_0-st-ta1	Affymetrix	Exon	v2	in_situ_oligo_features	Homo sapiens			Imported

Delete, Edit and Replace an array design file in caArray

Delete an Array Design

After data has been imported into an experiment, the array design associated with that data cannot be removed from the experiment's list of array designs. Conversely, once an array design has been imported, the file associated with it cannot be changed. Array designs that are not associated with an experiment can now be deleted. To delete an array design, follow this step:

- On the row corresponding to the array design, click the Delete icon (arrow in the image below).
- A confirmation window will pop up. Once you click the "OK" button, the file will be physically deleted from caArray.

Array Design Files with Delete Icon and Confirmation Window

The screenshot shows a table of array designs. A confirmation dialog box is displayed over the delete icon for the array design HT_HG-U133B. The dialog box contains the following text:

The page at <http://afapp1.c2b2.columbia.edu:38080> says:

You are about to delete this array design. Please note that large array designs will take longer to delete. Press OK to continue.

Buttons: OK, Cancel

Array Design Name	Provider	Assay Type	Version Number	Feature Type	Organism	Edit	Edit File	Delete	Status	Download
016436_D_G2_20070426	Agilent	Gene Expression	1	in_situ_oligo_features	Homo sapiens				Imported (Not Parsed)	
HT_HG-U133B	Affymetrix	Gene Expression	1	in_situ_oligo_features	Homo sapiens				Imported	
HT_Rat230_PM	Affymetrix	Gene Expression	1.0	in_situ_oligo_features	Rattus rattus				Imported	

Edit an Array Design

In caArray 2.2, any logged in user can edit an array design. To edit an array design, follow these steps:

- On the row corresponding to the array design, click the Edit icon.

The screenshot shows a close-up of the row for HT_HG-U133B. The Edit icon (a pencil) is highlighted with a yellow arrow.

- The required fields become editable; enter any edits.
- Please note, the editable fields are limited for the array design files uploaded prior to caArray 2.3. Starting with caArray 2.3, all required fields become editable.

Array Design Details Dialog Open for Editing in V2.3 and V2.2 and Under with Fewer Fields Open for Editing

A. caArray Version 2.3

Manage Array Designs

HT_HG-U133B

Array Design Details

Array Name: HT_HG-U133B
 Design Name: HT_HG-U133B
 Array Design LSID: URN:LSID:Affymetrix.com:PhysicalArrayDesign:HT_HG-U133B
 Description: test for array design file

Assay Type: aCGH Exon Gene Expression Methylation microRNA

Selected Assay Type: Gene Expression

Provider: Affymetrix
 Version Number: 1
 Feature Type: in_situ_oligo_features (MO)
 Organism: Homo sapiens (ncbitax)

Buttons: Cancel, Save

B. caArray Version 2.2 & Under

B. caArray Version 2.2 & Under

Manage Array Designs

HT_HG-U133A

Some details of this array design may not be modified because it is already associated with an existing experiment.

Array Design Details

Array Design Name: HT_HG-U133A
 Array Design LSID: URN:LSID:Affymetrix.com:PhysicalArrayDesign:HT_HG-U133A
 Description: [Yellowed out]

Assay Type: Gene Expression
 Provider: Affymetrix
 Version Number: 1.1
 Feature Type: in_situ_oligo_features (MO)
 Organism: Homo sapiens (ncbitax)

Buttons: Cancel, Save

Replace an Array Design

Once an array design has been imported and parsed into caArray, the array design file associated with it cannot be changed. You will have to delete the existing one and upload a new one. This can be done only if there is no experiment data associated with this array design file.

The only array design file that can be replaced is the one imported without validation and parsing. (For more information on the definition of different file types, refer to [caArray 016 - Which file can be parsed into caArray? What is the benefit of file parsing?](#) Files Recognized by caArray.)

To replace an array design file, follow these steps:

- Find the file that is replaceable (the file with a status of "Imported (Not Parsed)" (circled in red in the illustration).
- Click on "Edit File" icon (Arrow) to activate "Upload Array Design File" Window.
- Browse and find the new array design file.
- Save the new file by clicking the Save button.

Replacing an Array Design File

The screenshot shows the caArray web interface. At the top, there is a table with columns: Array Design Name, Provider, Assay Type, Version Number, Feature Type, Organism, Edit, Edit File, Delete, Status, and Download. The first row is highlighted, and its 'Status' cell, containing 'Imported (Not Parsed)', is circled in red. A yellow arrow points to the 'Edit File' icon in the same row. Below the table, a dialog box titled 'caArray - Manage Array Designs - Mozilla Firefox' is open. The dialog box has a title bar and a toolbar with 'Help' and 'Print' buttons. The main content area shows the title '016436_D_G2_20070426' and a section titled 'Upload Array Design File'. This section includes a 'Current File' field with the value '016436_D_G2_20070426.xml', a 'Browse to File:' field with a 'Browse...' button, and a 'File Format:' dropdown menu set to 'Automatic'. At the bottom of the dialog box are three buttons: 'Cancel', 'Save', and 'Add More Files'.

Array Design Name	Provider	Assay Type	Version Number	Feature Type	Organism	Edit	Edit File	Delete	Status	Download
016436_D_G2_20070426	Agilent	Gene Expression	1	in_situ_oligo_features	Homo sapiens				Imported (Not Parsed)	
					Homo sapiens				Imported	
					Homo sapiens				Imported	
					Homo sapiens				Imported	
					Homo sapiens				Imported	
					Homo sapiens				Imported	
					Homo sapiens				Imported	
					Homo sapiens				Imported	
					Homo sapiens				Imported	
					Homo sapiens				Imported	

Have a comment?

Please leave your comment in the [caArray End User Forum](#).