

**New and updated features in the 2015a release:**

*MATLAB Graphics Engine*

- TrueAllele VUIer has been updated to use the latest MATLAB graphics engine

*Interface Improvements*

- Tables automatically adjust column width based on current font size
- Dialogs, such as Lane, Request, and Genotype Info, can be resized to display longer text strings

*General*

- Support for NGM™ and NGM SElect™ kits
- Improved support for match comparisons in Report across different kits such as comparing Identifiler® evidence and GlobalFiler® references

*Cloud*

- Support for TrueAllele Cloud servers

*Report*

- Improved Specificity feature including additional graphics and statistics
- Improved archive files now support the .tas file extension

*Tools*

- New feature allowing deletion of requests via the Interpretation Request menu

*Populations*

- Amended FBI population files available upon request
- New NIST 1036 population files available upon request

## Before you install the software

### *Supported Operating Systems*

The latest MATLAB requires the following operating systems.

PC: Windows 7 or 8

MAC: OS X 10.9.5 (Mavericks) or newer

Please confirm your OS before installing this software. If you have questions about updating your Operating System, please contact Cybergenetics for additional information.

### *Preserve your customized files and templates*

Cybergenetics recommends that you perform a clean installation of the Analyze and VUIer software. Before removing your current software, please preserve the following folders by copying these folders to another location. In doing so, you will preserve files that have been customized to your site.

	Located in:	
	Mac	Windows
Preferences	/Library/TrueAllele	/Program Files/VUIer
trueallele	/Documents	/My Documents (for every user)

**Table 1 - Folder location**

### *Uninstall the current versions*

After preserving these files, it is safe to uninstall the software. Be sure to remove all files and folders when uninstalling the software as indicated by Table 1.

### *Install the MATLAB 8.5 engine*

To install the MATLAB 8.5 engine, please follow the Mac- or Windows-specific instructions found in the **Getting Started** manual. Please contact Cybergenetics if you have any questions about installing the software.

*Install Analyze and VUler 2015a*

To install the 2015a VUler and Analyze versions, please follow the Mac- or Windows-specific instructions found in the **Getting Started** manual. Please contact Cybergeneics if you have any questions about installing the software.

*Replace customized Analyze files and templates, and update the Init Preferences*

Once Analyze is installed, copy the *operators.txt* and *db\_prefs.txt* files from your preserved files back into the /trueallele/user directory. If your site uses a customized DataDisk template, move this template to the /trueallele/template directory.

Start Analyze and in the Command Window select Edit > Preferences > Init and reset your initial preferences.

*Replace customized VUler files*

Once VUler is installed, copy the *connect.txt* file from your preserved files into the VUler Preferences folder. If you have customized other Preferences files, replace them as well.