MacVector 12.7

for Mac OS X

System Requirements

MacVector 12.7 runs on any PowerPC or Intel Macintosh running **Mac OS X 10.5** or higher. It is a Universal Binary, meaning that it runs natively on both PowerPC and Intel based Macintosh computers. There are no specific hardware requirements for MacVector – if your machine can run OS X 10.5 or above, it can run MacVector. A complete installation of MacVector 12.7 uses approximately 180 MB of disk space.

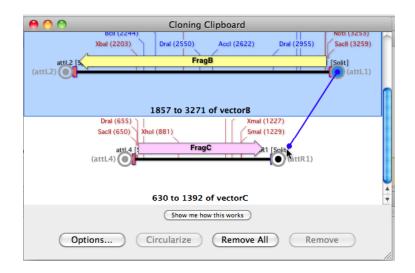
Installation and License Activation

Install MacVector 12.7 by double-clicking on the MacVector 12.7.pkg installer application. You will be prompted for a system administrator account and password during installation. Once installation is complete, you must enter a valid license owner, serial number and activation code the first time you run MacVector. This information is usually sent by e-mail and is also printed on the inside of the CD sleeve. If you previously installed an earlier version of MacVector and have a serial number with a maintenance end date of Sep 1st 2012 or later, MacVector 12.7 will automatically use your existing license and you will not be required to enter the details again.

Changes for MacVector 12.7

Cloning Clipboard

There is a new window called the **Cloning Clipboard** that can be opened and closed from the **Window** menu. Whenever a restriction fragment is generated using the **Digest** button (or **Edit** | **Digest** menu item), the fragment is placed on the cloning clipboard. You can then join fragments on the clipboard by selecting the end of one fragment and dragging it onto the end of another fragment;



When you release the mouse over an end, you are presented with a dialog that lets you fill or cut back ends if you wish. Finally, after building up a molecule from multiple pieces, you can click on the circularize button to create a new circular molecule, or just doubleclick on a fragment to open it in a new MacVector sequence window. For more information, check out the file Cloning Clipboard Tutorial.pdf in the /Applications/MacVector 12.7/Documentation/ folder.

Optimized Reverse Translations

There are new options in the **Analyze** | **Reverse Translation** dialog, allowing you to optimize codon usage when reverse translating a protein. You can now request that the generated DNA sequence uses only the most popular codons from a given codon bias file, or that the generated sequence should have an overall codon usage that most closely matches that file.

Align To Reference Performance Enhancements

A number of operations in the **Align To Reference** functionality have been optimized to be of more use for aligning large numbers of Next Generation Sequencing reads against large genomes. Importing reads is now very much faster (in many cases 100x faster) and aligning reads is from 5x to 50x faster. With the appropriate parameters, you can now align several hundred thousand reads against a typical bacterial chromosome in just a few hours. Individual edits in the **Editor** tab with several hundred thousand reads are now effectively instantaneous (previously they could take many minutes while the consensus was recalculated). In addition the text in the **SNP** tab is generated anywhere from 10x to 1,000x faster than previously. Finally, although generation of the text in the **Text** tab has also been significantly speeded up (again up to 1,000x) it also now runs as a separate thread so you can cancel the operation if it takes too long.

MacVector Free

One major change that will primarily affect users of the trial version is that, starting with MacVector 12.7, once a trial license expires, MacVector will no longer simply refuse to run. Instead, MacVector will continue to run, but all of the functions in the **Analyze** menu become disabled. This means users can still open, edit, save and print MacVector documents, or save MacVector files in other formats. They can still perform simple click cloning operations (but not the new **Cloning Clipboard**), generate new constructs and have full control over the appearance of features. **Entrez** and **BLAST** are still available

along with the **Find** functionality. This will let licensed users share MacVector files and data with more casual users without having to convert sequences into a common format, often losing information (such as graphical appearance and layout) in the process.

Miscellaneous Changes/Bug Fixes

You can now create a feature "between residues" which can be displayed as a line graphic in the **Map** tab.

Start Location:	20 At 🛟
Stop Location:	
	21 At 🛟
Between (^)	•
	Cancel OK

If you subsequently select that feature the caret becomes positioned between the residues in the **Editor** tab.

Support information

For assistance with MacVector, please contact your local MacVector, Inc office. You will need a current MacVector maintenance contract to be eligible for technical support other than for basic installation problems. New sales of MacVector include 12 months of support that also entitles you to any upgrades to MacVector released during the maintenance period.

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When contacting Customer Support with a technical problem, please be prepared to give your product serial number as well as a detailed description of your problem and any error messages you encounter. Visit the MacVector Web site for details of any available updates, and any relevant information that could not be added to these release notes in time for publication:

http://www.macvector.com

MacVector, Inc.

Software for Scientists