

Restriction Analysis

From this chapter you can learn how to search for restriction sites on a DNA sequence.

The restriction sites found are stored as automatic annotations. This means that if the automatic annotations highlighting is enabled then the restriction sites are searched and highlighted for each nucleotide sequence opened. Refer [Automatic Annotations Highlighting](#) to learn more.

Open a DNA sequence in and click the following button on the *Sequence View* toolbar:



Alternatively, select either the *Actions Analyze Find restriction sites* item in the main menu or the *Analyze Find restriction sites* item in the context menu.

The *Find restriction sites* dialog appears:

Find Restriction Sites

Filter by name:

Name	Accession	Type	Sequence	Organism / Details
▶ A (0, 264)				AaaI .. AxyI
▶ B (3, 917)				BacI .. BvuBI
▶ C (1, 193)				CaeI .. CvnI
▶ D (1, 31)				DaeI .. DspI
▶ E (1, 325)				EaeI .. EspHK30I
▶ F (0, 63)				F-CphI .. F-TevIV

Selected enzymes:

BamHI, BbsI, BglII, ClaI, DraI, EcoRI, HindIII, PstI, SalI, SmaI, XmaI

☐ Filter by number of results:

Minimum hits: Maximum hits:

☐ Exclude region:

Region: -

Total number of enzymes: 4862, selected 11

Open enzymes
Export enzymes
Select All
Select None
Select by length
Invert selection
Load selection
Save selection
REBASE Info
OK
Cancel
Help

You can see the list of restriction enzymes that can be used to search for restriction sites. The information about enzymes was obtained from the [REBASE](#) database. For each enzyme in the list a brief description is available (the accession ID in the database, the recognition sequence, etc.). If you're online you can get more detailed information about an enzyme selected by clicking the *REBASE Info* button.

- [Selecting Restriction Enzymes](#)
- [Using Custom File with Enzymes](#)
- [Filtering by Number of Hits](#)
- [Excluding Region](#)
- [Circular Molecule](#)
- [Results](#)