

Searching for Pattern

You can search for a pattern inside an alignment.

Enter a query string in the edit box under the *Sequence area*.

The image shows a sequence alignment viewer. It displays a grid of amino acid residues (N, L, Y, Q, F, G, E, M, I, S, K, K, T, G, T, -, F, G, L, F) arranged in rows and columns. Each residue is highlighted with a different color. Below the grid is a search bar with the label "Find:" and a text input field. To the left of the input field is a left-pointing arrow, and to the right is a right-pointing arrow. The search bar is currently empty.

1	N	L	Y	Q	F	G	E	M	I	S	K	K	T	G	T	-	F	G	L	F
1	S	L	L	E	F	G	M	M	I	L	G	E	T	G	K	-	N	P	L	T
1	H	L	L	Q	F	R	K	M	I	K	K	M	T	G	K	-	E	P	V	I
1	H	L	L	Q	F	R	K	M	I	K	K	M	T	G	K	-	E	P	I	V
1																				

Find:

Press the right arrow to search in the direction "From left to right, from top to bottom". Press the left arrow to search in the direction "From right to left, from bottom to top". If the pattern is found, the result will be focused and highlighted in the *Sequence area*. You can continue the search in any direction from this position.