

Wizard2011

Input	
<div><div>Information<div>Close</div></div><p>On this site you can describe a possible input for your function. In addition to the inputs name, short description and description you also have to define the representation and the handling. The representation is the type of your input data in bioinformatic terms (e.g. a certain type of FASTA). You can simplify the search for the right representation by using filters. The handling of your input defines how the input data gets into the tool technically speaking (e.g. as a file). The "option" is a commandline prefix for this input that is used when your tool is called via commandline (e.g. -FILE=). You also can define an example for your input. On the right side of the screen you have an overview of existing inputs. You can edit and delete existing examples and add new ones. Please note that if you use the "edit" or the "+" button your current input is not saved!</p></div>	
Name: *	<input type="text"/>
Short Description: *	<div><div></div></div>
Description: *	<div><div>Open Text-Editor</div></div>
Representation: *	<div><div>Filter: All</div><div>Filter: All</div><div>Filter: All</div><div>AlignmentML_AA</div></div>
Handling: *	<div><div>File</div></div>
Option:	<input type="text"/>
Example:	<div><div>Choose</div><div></div></div>
<div><div>Return</div><div>Save and return</div><div>Save</div></div>	

Inputs +

No input defined yet!