

# Wizard2011

Dependency

Information

Close

This page enables you to define Dependencies between different parameters in your tool. For a valid dependency description you have to provide a name, a short description and the dependency itself. To define definitions, write them into the "Definition"-text field in the center of the screen. You also can use the buttons for all existing parameters to conveniently include them in the dependency definition. The language to describe dependencies is Jan Kruegers DependencyLanguage. The BNF-definition of the language can be found in the BiBiServ-wiki at the TechFak Bielefeld. You can also test your dependency if it is technically valid for a given function. The result of the test will be shown in the "Log" window at the bottom of the screen. On the right side of the screen you have an overview of existing dependencies. You can edit and delete existing dependencies and add new ones. Please note that if you use the "edit" or the "+" button your current dependency is not saved!

Name:\*

Short Description: \*

Description:

Open Text-Editor

Definition: \*

```

1  and(
2    def(<nucleic acid sequence handling>),
3    def(<threshold>)
4  )
5
6
7
8
9

```

nucleic acid  
sequence  
handling

threshold

Test Function:

DNA alignment

▼

Test Function

No Errors.

Return

Save and return

Save

Dependencies +

No dependency defined yet!