

Build DIAMOND Database Element

Build a DIAMOND formatted database from a FASTA input file.

Element type: diamond-build

Parameters

Parameter	Description	Default value	Parameter in Workflow File	Type
Database	A name of the binary DIAMOND database file that should be created.		database	<i>string</i>
Genomic library	Genomes that should be used to build the database.		genomic-library	<i>url-datasets</i>

Input/Output Ports

The element has 1 *output port*:

Name in GUI: Output DIAMOND database

Name in Workflow File: out

Slots:

SlotInGUI	Slot in Workflow File	Type
Output URL	url	<i>string</i>