

# DNA Statistics Element

Evaluates statistic for DNA sequences.

**Element type:** dna-stats

## Parameters

Parameter	Description	Default value	Parameter in Workflow File	Type
GC-content	Evaluate GC-content.	True	gc-content	boolean
GC1-content	Evaluate GC1-content.	True	gc1-content	boolean
GC2-content	Evaluate GC2-content.	True	gc2-content	boolean
GC3-content	Evaluate GC3-content.	True	gc3-content	boolean

## Input/Output Ports

The element has 1 *input port*:

**Name in GUI:** *Input sequence*

**Name in Workflow File:** in-sequence

**Slots:**

Slot In GUI	Slot in Workflow File	Type
Sequence	sequence	sequence

The element has 1 *output port*:

**Name in GUI:** *Result annotation*

**Name in Workflow File:** out-annotations

**Slots:**

Slot In GUI	Slot in Workflow File	Type
Set of annotations	annotations	annotation-table-list