

# Package ‘biOpsGUI’

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**Type** Package

**Title** GUI for Basic image operations

**Version** 0.1.2

**Depends** R (>= 2.4.0), RGtk2, biOps

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**Encoding** UTF-8

**Description** GUI (using GTK+) for the basic image operations package. At the moment, just provides a function that displays imagedata from biOps package in a new window, keeping original size and with information on the coords and values of the pixels in the image. It requires RGtk2.

**License** GPL

**Repository** CRAN

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<code>imgDisplay</code>	<i>Display an imagedata object</i>
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**Description**

This function shows a window that displays the imagedata.

**Usage**

```
imgDisplay(imgdata)
```

**Arguments**

<code>imgdata</code>	The image
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**Examples**

```
x <- readJpeg(system.file("samples", "violet.jpg", package="biOps"))  
imgDisplay(x)
```

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