

zebrafishcdf

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R topics documented:

<code>i2xy</code>	1
<code>zebrafishcdf</code>	2
<code>zebrafishdim</code>	2

Index	3
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<code>i2xy</code>	<i>Convert (x,y)-coordinates to single-number indices and back.</i>
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Description

Convert (x,y)-coordinates on the chip (and in the CEL file) to the single-number indices used in AffyBatch and CDF environment, and back.

Usage

```
i2xy(i)  
xy2i(x, y)
```

Arguments

<code>x</code>	numeric. x-coordinate (from 1 to 712)
<code>y</code>	numeric. y-coordinate (from 1 to 712)
<code>i</code>	numeric. single-number index (from 1 to 506944)

Details

Type `i2xy` and `xy2i` at the R prompt to view the function definitions.

See Also

[zebrafishcdf](#)

Examples

```
xy2i(5,5)
i      = 1:(712*712)
coord = i2xy(i)
j      = xy2i(coord[, "x"], coord[, "y"])
stopifnot(all(i==j))
range(coord[, "x"])
range(coord[, "y"])
```

zebrafishcdf *zebrafishcdf*

Description

environment describing the CDF file

zebrafishdim *zebrafishdim*

Description

environment describing the CDF dimensions

Index

*Topic **datasets**

`i2xy`, 1

`zebrafishcdf`, 2

`zebrafishdim`, 2

`i2xy`, 1

`xy2i` (`i2xy`), 1

`zebrafishcdf`, 1, 2

`zebrafishdim`, 2