

# zebrafishcdf

January 14, 2026

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|------|---------------------------------------------------------------------|
| i2xy | <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

|   |                                                 |
|---|-------------------------------------------------|
| x | numeric. x-coordinate (from 1 to 712)           |
| y | numeric. y-coordinate (from 1 to 712)           |
| i | 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"])
```

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|--------------|---------------------|
| zebrafishcdf | <i>zebrafishcdf</i> |
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**Description**

environment describing the CDF file

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| zebrafishdim | <i>zebrafishdim</i> |
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**Description**

environment describing the CDF dimensions

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