

# zebrafishcdf

March 12, 2025

---

`i2xy`

*Convert (x,y)-coordinates to single-number indices and back.*

---

## 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	<i>zebrafishcdf</i>
--------------	---------------------

---

**Description**

environment describing the CDF file

---

zebrafishdim	<i>zebrafishdim</i>
--------------	---------------------

---

**Description**

environment describing the CDF dimensions

# Index

## \* datasets

i2xy, 1

zebrafishcdf, 2

zebrafishdim, 2

i2xy, 1

xy2i (i2xy), 1

zebrafishcdf, 1, 2

zebrafishdim, 2