

# drosgenome1cdf

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### Description

environment describing the CDF file

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| drosgenome1dim | <i>drosgenome1dim</i> |
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### Description

environment describing the CDF dimensions

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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 640)           |
| y | numeric. y-coordinate (from 1 to 640)           |
| i | numeric. single-number index (from 1 to 409600) |

**Details**

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

**See Also**

[drosgenome1cdf](#)

**Examples**

```
xy2i(5,5)
i      = 1:(640*640)
coord = i2xy(i)
j      = xy2i(coord[, "x"], coord[, "y"])
stopifnot(all(i==j))
range(coord[, "x"])
range(coord[, "y"])
```

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