

# CytoGate Gating Methods

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## Overview of CytoGate Features

### Introduction

The `Cytogate` package provides an interactive interface for drawing gates to analyse flow cytometry data through the `DrawGate` function. The package also contains a plugin function `gate_draw` to incorporate these interactive features into the `openCyto` gating pipeline. Here we aim to demonstrate the main gating features of `CytoGate` and how these features can be used with `openCyto`.

#### 1. Adding CytoGate to openCyto as a Plugin.

The features of `CytoGate` can be easily incorporated into `openCyto` as a plugin. To do this we will need to supply the name of the gating function `gate_draw` and give the gating method a name for use within `openCyto`, here we will use `DrawGate` as the gating method name. To check whether the function has been successfully added we use `listgtMethods()` to return a list of supported gating methods

```
library(openCyto)
library(CytoGate)

registerPlugins(fun = gate_draw, methodName = "DrawGate")
```

```
## [1] TRUE
```

```
listgtMethods()
```

```
## Gating Functions:
## === DrawGate
## === quantileGate
## === rangeGate
## === flowClust.2d
## === mindensity
## === mindensity2
## === cytokine
## === flowClust.1d
## === boundary
## === singletGate
## === tailgate
## === quadGate.tmix
## === quadGate.seq
## Preprocessing Functions:
## === prior_flowClust
## === warpSet
## === standardize_flowset
```

## 2. Loading in Data for Analysis

To demonstrate the main features of **CytoGate** we will use a data set supplied by CytoGate called **Activation**. This data set contains eight **.fcs** files used to determine the activation state of OT-I and OT-II transgenic T cells. We will add these samples to a **GatingSet** called **gs** for subsequent gating using the **openCyto** framework.

```
fs <- Activation
gs <- GatingSet(fs)
```

## 3. Preparing Data for Analysis

Once the data has been added to the **GatingSet** we perform necessary compensation using the attached spillover matrix and perform logicle transformation on all fluorescent channels.

```
spill <- fs[[1]]@description$SPILL
gs <- compensate(gs, spill)

channels <- colnames(fs)
trans.channels <- channels[!channels %in% c("FSC-A", "FSC-H", "FSC-W", "SSC-A", "SSC-H", "SSC-W", "Time")]

trans <- estimateLogicle(gs[[1]], trans.channels)
gs <- transform(gs, trans)
```

## 4. Supported Gate Types & Expected Inputs

The **DrawGate** gating method will automatically open a new interactive plotting window where the user can specify the coordinates of gates manually. The type of gate to constructed is supplied to the **GatingTemplate** as the **gate\_type** argument - this will be demonstrated later. Currently the supported gate types and expected inputs are as follows:

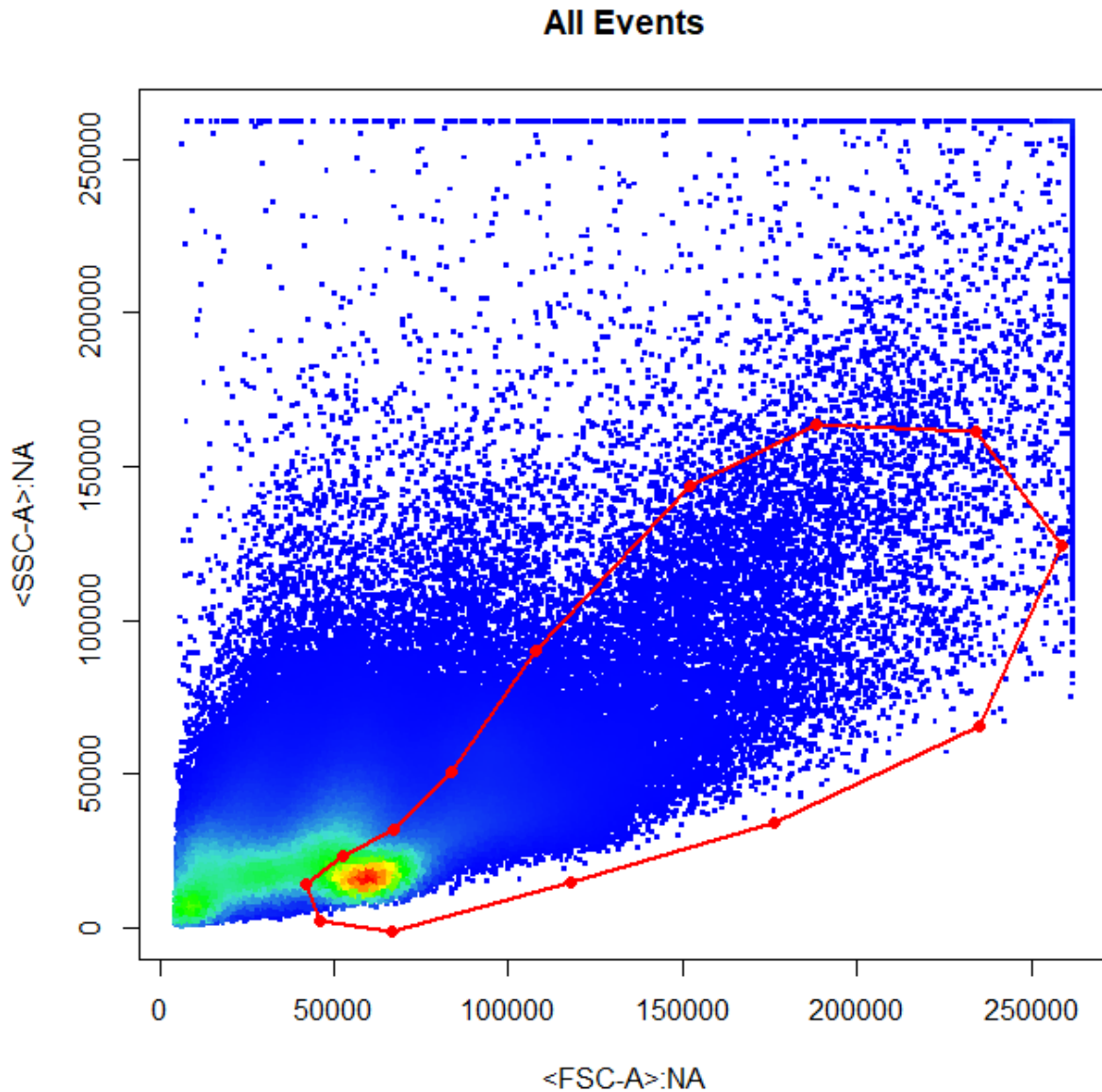
- **"polygon"** constructs a polygon gate from **n** selected points where **n** must be greater than 3. To seal the polygon the user must **right click** and select **"stop"**. This is the default **gate\_type** used by **DrawGate**. **n>3**
- **"rectangle"** constructs a rectangle gate parallel to the x and y axes from 2 selected points along the diagonal of the rectangle. **n=2**
- **"interval"** constructs a gate based on user defined lower and upper bounds. 1D and 2D interval gates are supported, For 1D gates a density distribution is plotted and 2 points selected indicating the lower and upper bounds of the gate. 2D interval gates require an additional argument **axis** used to specify whether the **"x"** or **"y"** axis should be gated. **n=2**
- **"threshold"** constructs a gate that selects all events above a selected lower bound. 1D and 2D threshold gates are supported. For 2D threshold gates a rectangle gate is constructed which selects events above the selected x and y co-ordinates. **n=1**
- **"ellipse"** constructs an ellipse gate from 4 selected points indicating the limits of the gate in 2 dimensions. **n=4**
- **"quadrant"** constructs 4 rectangle gates based on selection of a single point indicating the center of the crosshairs. **n=1**

Multi-gates are supported for all gate types except **"threshold"** and **"quadrant"** through an additional argument **N** defining the number of gates to be constructed. This multi-gate feature will be demonstrated later.

## 5. Polygon Gates (n)

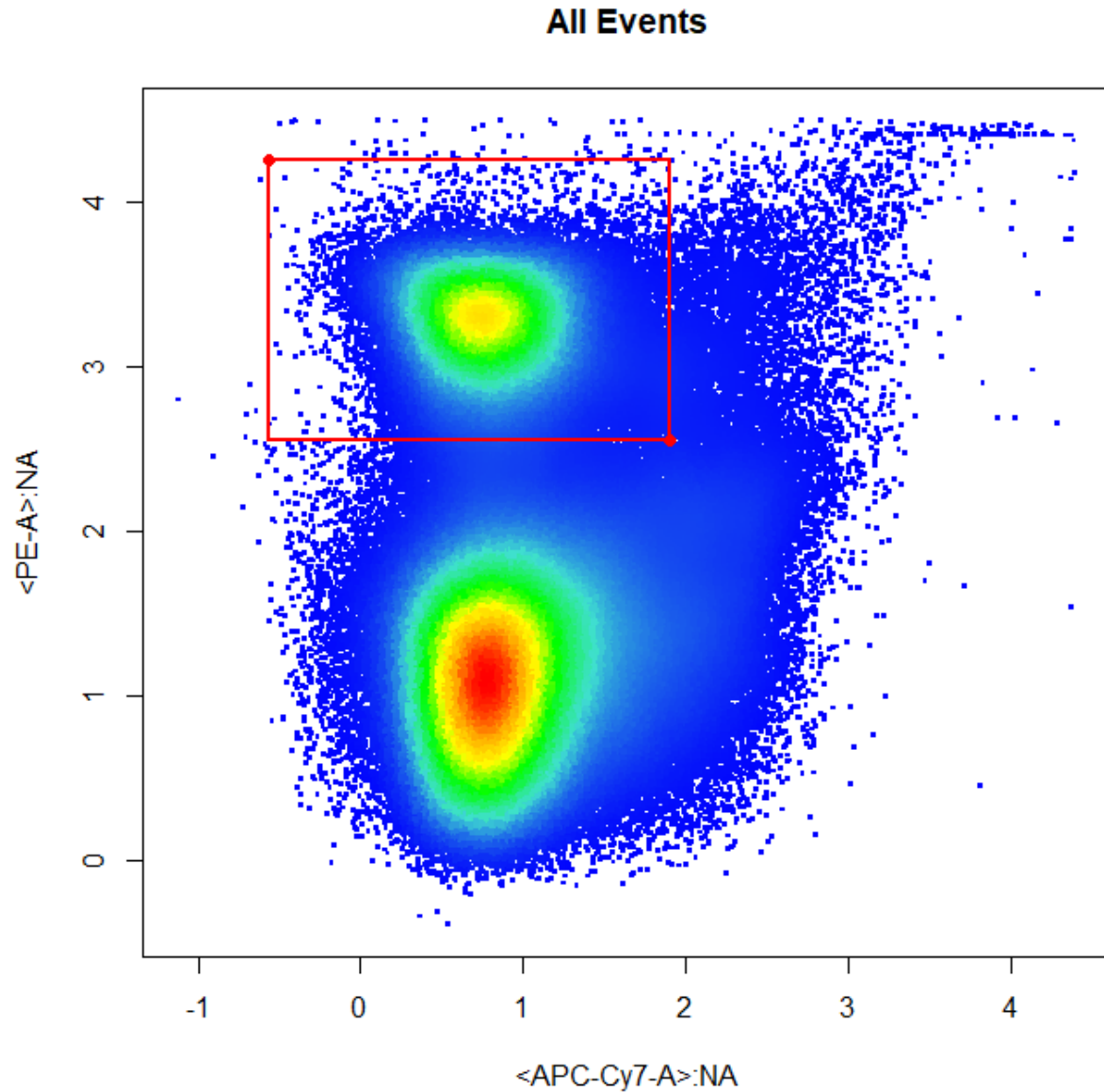
Below is an example of the entry in the `GatingTemplate`. When using `DrawGate` as an `openCyto` plugin it is recommended that pooled using the `collapseDataForGating` and `groupBy` arguments to ensure that the selected gates encompass the population in all samples. Additionally, to speed up the gating process user should also supply the `subSample` argument to restrict the data to a subset of the pooled data for plotting. Notice how the `gate_type` is supplied as a `gating_args` using `'gatetype'`.

```
template <- add_pop(  
  gs, alias = "Cells", parent = "root", pop = "+", dims = "FSC-A,SSC-A", gating_method = "DrawGate",  
  gating_args = "subSample=25000,gate_type='polygon'", collapseDataForGating = TRUE, groupBy = 8  
)
```



## 6. Rectangle Gates (n=2)

```
template <- add_pop(  
  gs, alias = "PE+", parent = "root", pop = "+", dims = "FSC-A,SSC-A", gating_method = "DrawGate",  
  gating_args = "subSample=25000,gate_type='rectangle'", collapseDataForGating = TRUE, groupBy = 2  
)
```

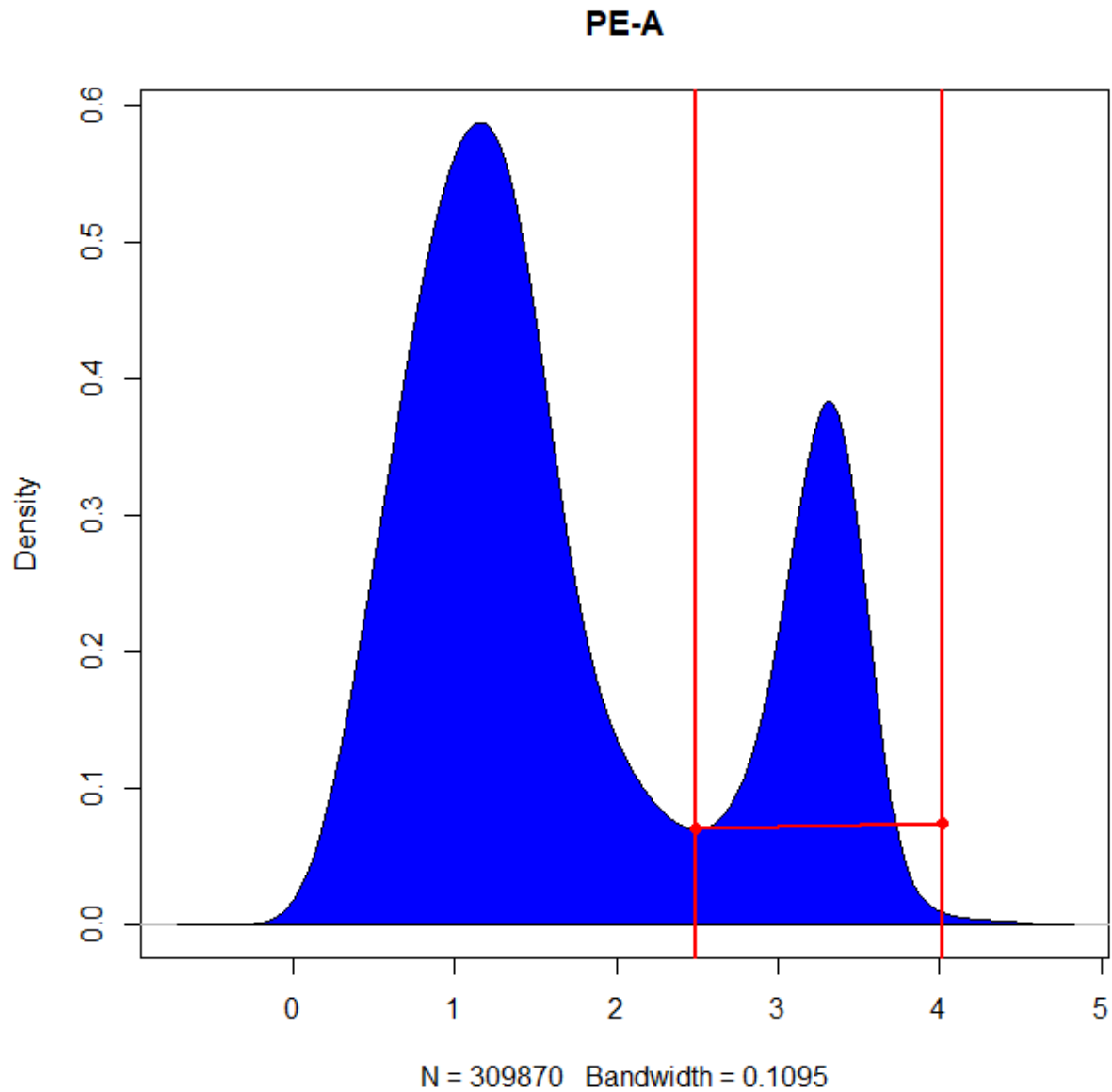


## 6. Interval Gates

1-Dimensional Interval Gates (n=2)

```
template <- add_pop(  
  gs, alias = "PE+", parent = "root", pop = "+", dims = "PE-A", gating_method = "DrawGate",
```

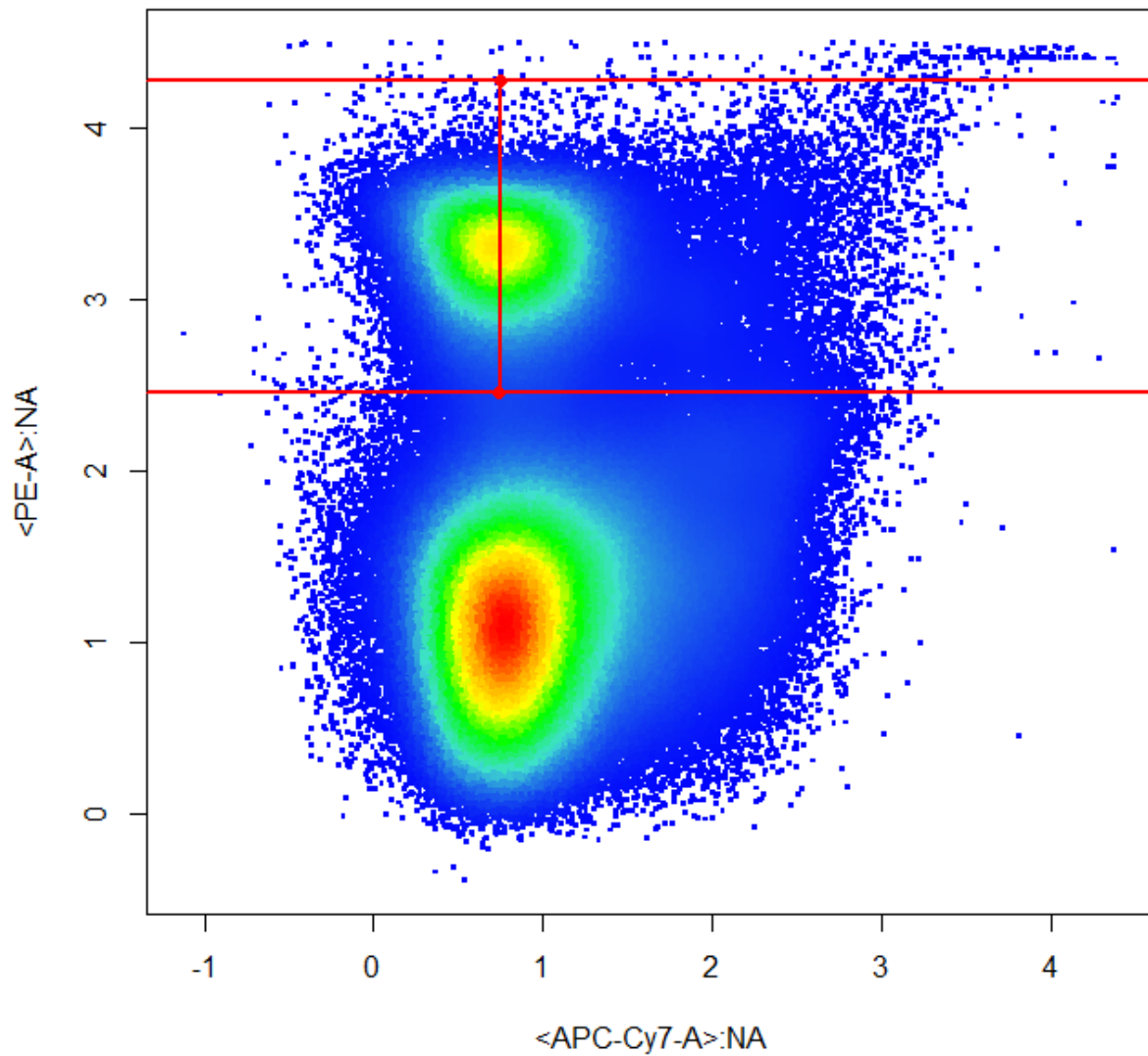
```
gating_args = "subSample=25000,gate_type='interval'", collapseDataForGating = TRUE, groupBy = 2
)
```



2-Dimensional Interval Gates - Axis Argument (n=2)

```
template <- add_pop(  
  gs, alias = "PE+", parent = "root", pop = "+", dims = "APC-Cy7-A,PE-A", gating_method = "DrawGate",  
  gating_args = "subSample=25000,gate_type='interval',axis='y'", collapseDataForGating = TRUE, groupBy =  
)
```

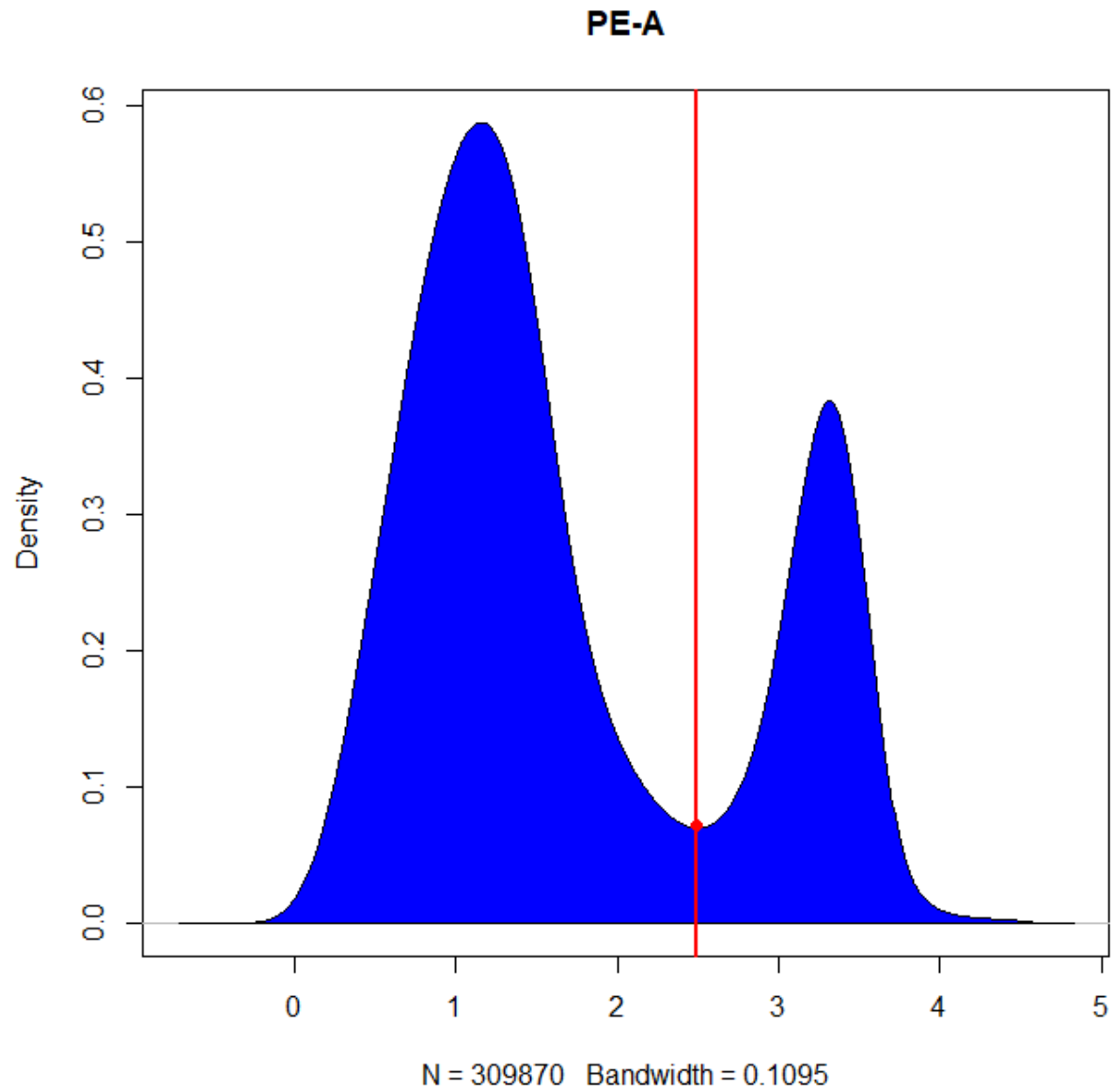
## All Events



### 7. Threshold Gates

1-Dimensional Threshold Gates (n=1)

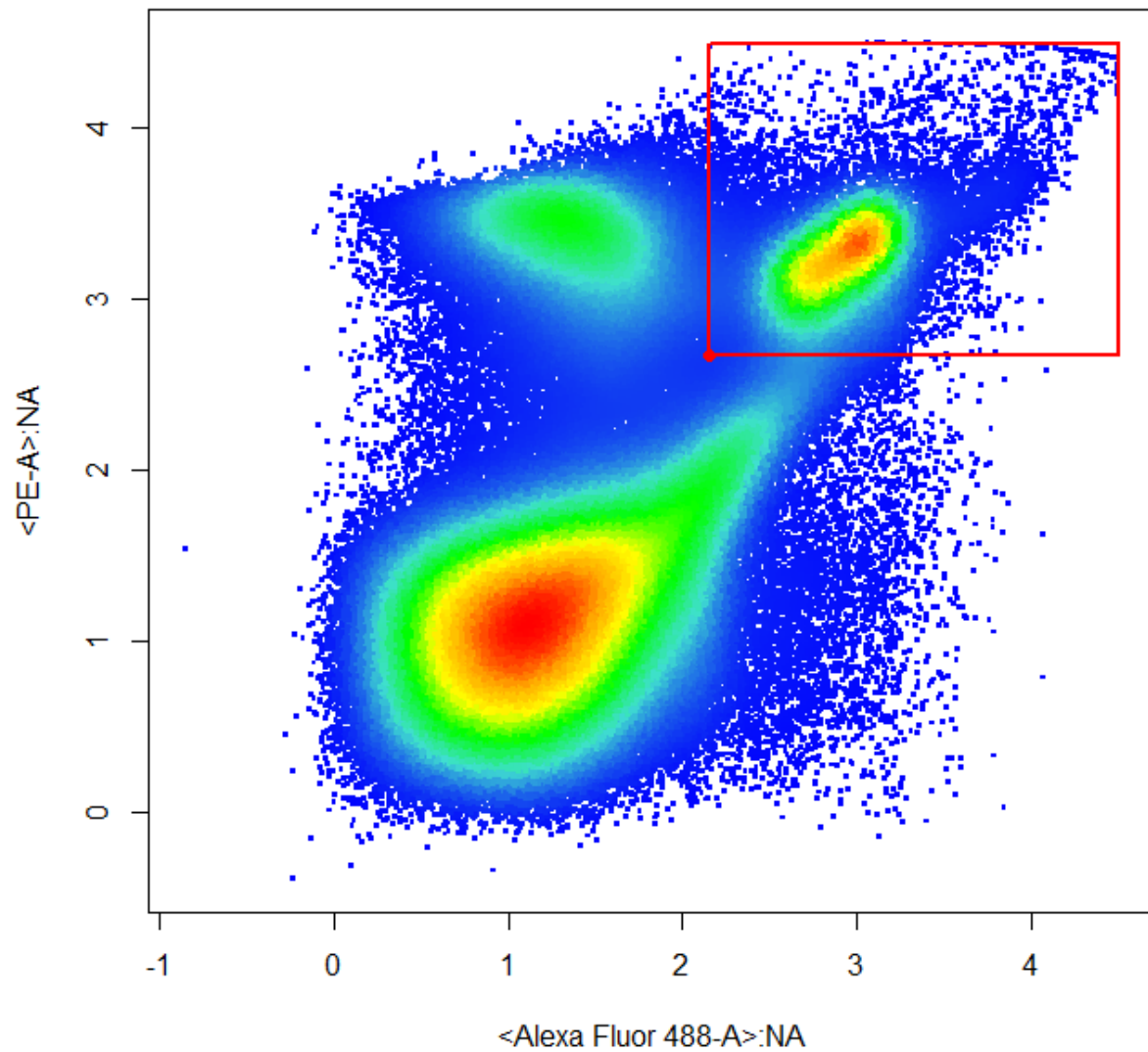
```
template <- add_pop(  
  gs, alias = "PE+", parent = "root", pop = "+", dims = "PE-A", gating_method = "DrawGate",  
  gating_args = "subSample=25000,gate_type='threshold'", collapseDataForGating = TRUE, groupBy = 2  
)
```



2-Dimensional Threshold Gates (n=1)

```
template <- add_pop(  
  gs, alias = "PE+FITC+", parent = "root", pop = "+", dims = "Alexa Fluor 488-A,PE-A", gating_method =  
  gating_args = "subSample=25000,gate_type='threshold'", collapseDataForGating = TRUE, groupBy = 2  
)
```

## All Events

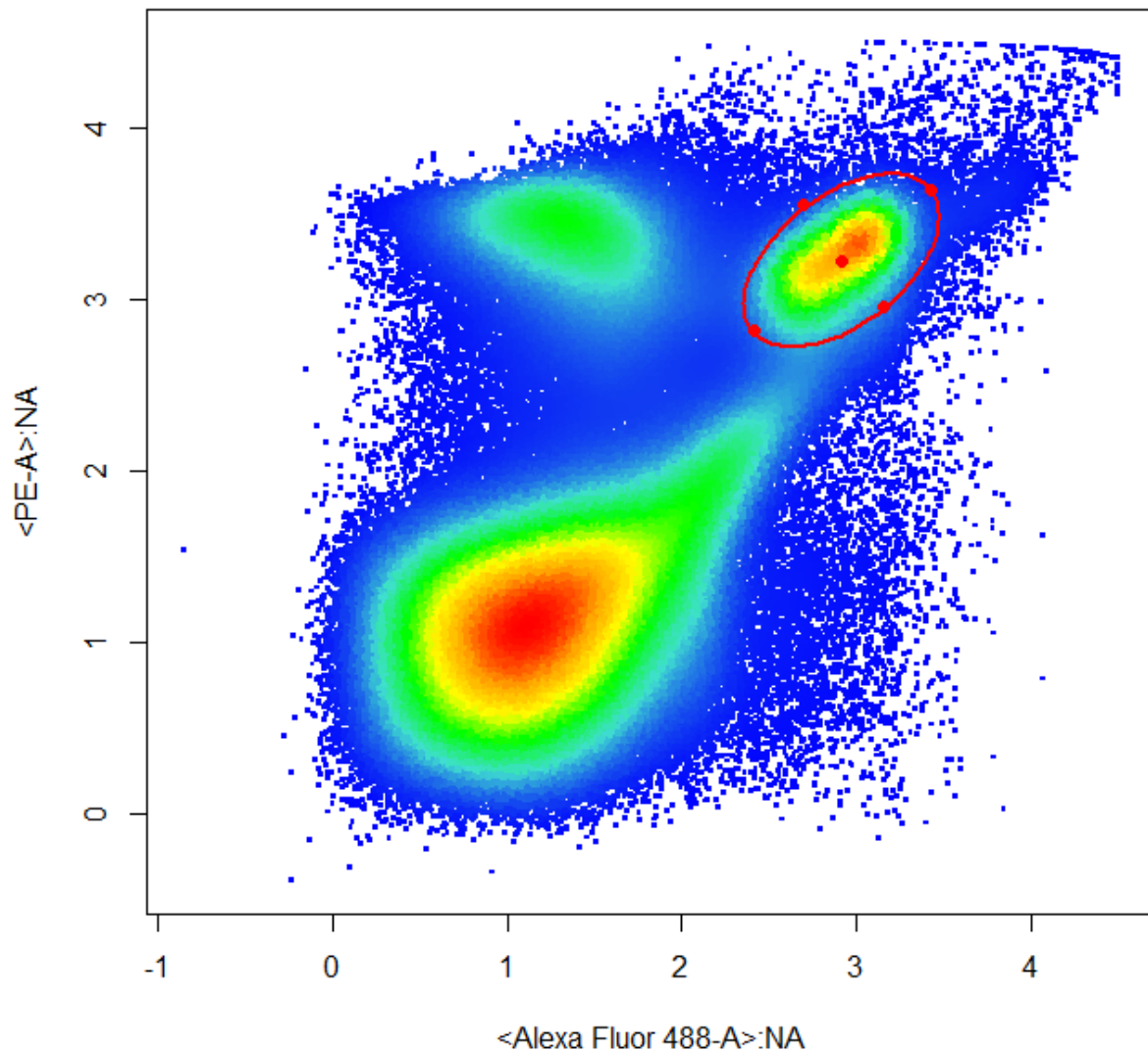


### 8. Ellipsoid Gates (n=4)

```
template <- add_pop(  
  gs, alias = "PE+FITC+", parent = "root", pop = "+", dims = "Alexa Fluor 488-A,PE-A", gating_method =  
  gating_args = "subSample=25000,gate_type='ellipse'", collapseDataForGating = TRUE, groupBy = 2  
)
```



## All Events

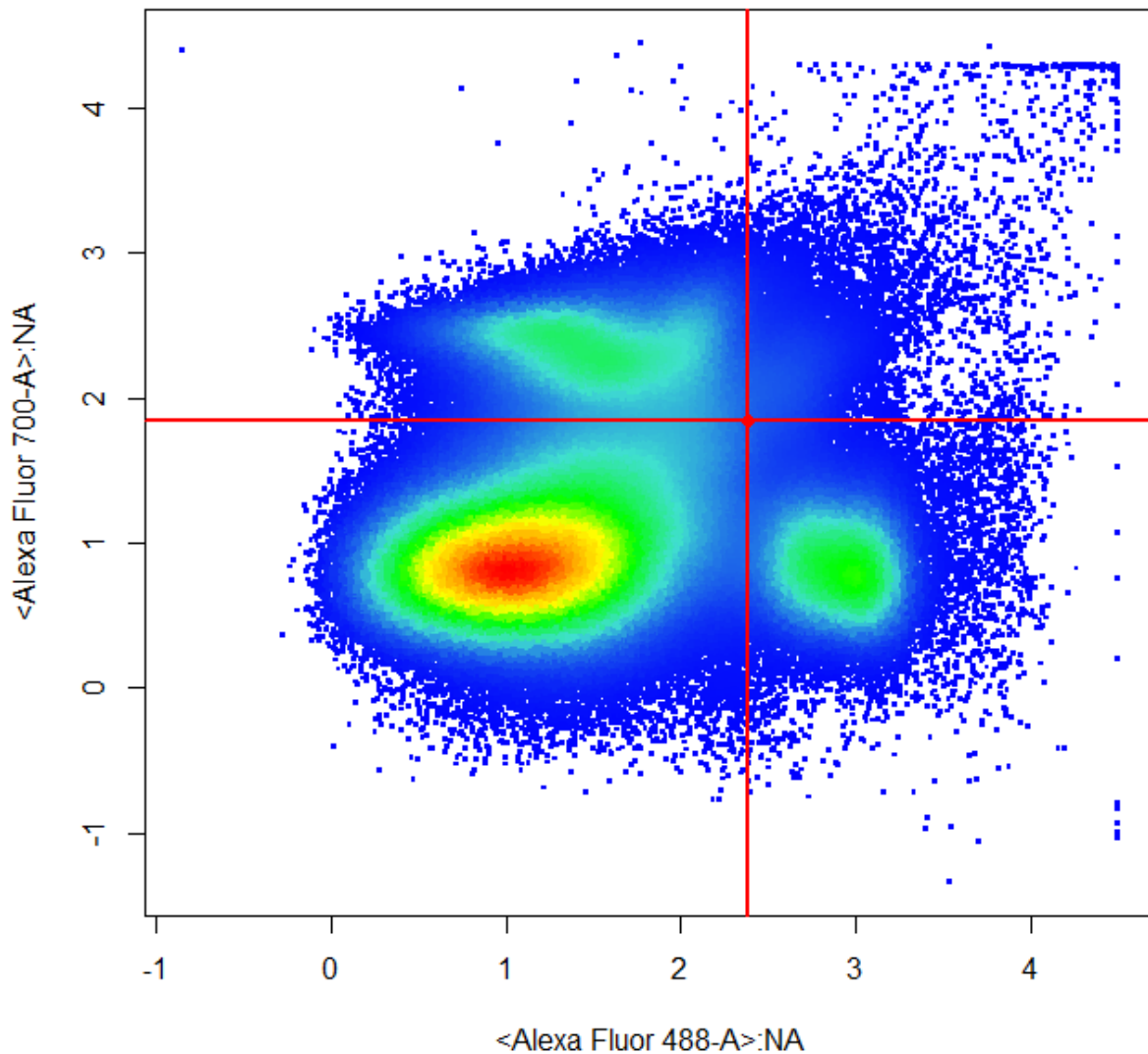


### 9. Quadrant Gates (n=1)

Quadrant gates returns 4 gates and therefore 4 populations names must be supplied as the `alias` argument and `pop` should be set to "\*" indicating that multiple gates will be returned.

```
template <- add_pop(  
  gs, alias = "DN,FITC+,AF700+FITC+,AF700+", parent = "root", pop = "*", dims = "Alexa Fluor 488-A,Alexa Fluor 488-A",  
  gating_args = "subSample=25000,gate_type='quadrant'", collapseDataForGating = TRUE, groupBy = 2  
)
```

## All Events



### 10. Multi-Gates

Here we will demonstrate the multi-gate feature of `DrawGate` for `gate_type = "ppolygon"` only although this feature is supported for all other gate types except `threshold` and `quadrant`. Simply supply the number of gates to construct as the argument `N` - for example to construct 2 polygon gates set `N=2`, supply two population names to the `alias` column and set `pop="*"` to return multiple populations.

```
template <- add_pop(  
  gs, alias = "PE+,FITC+", parent = "root", pop = "*", dims = "Alexa Fluor 488-A,PE-A", gating_method =  
  gating_args = "subSample=25000,gate_type='polygon',N=2", collapseDataForGating = TRUE, groupBy = 2  
)
```

### All Events

