

`bubblegrid` helps you see your data when a plain scatterplot would be overplotted. To do that, it aggregates points within a hexagonal grid, then shows, via size, the point count. It also shows, via color, the intensity (e.g. gene expression) averaged within bins.

```
devtools::install_github("maehrlab/bubblegrid")
library("bubblegrid")
data(test_data)
bubblegrid(x = test_data[[1]], y = test_data[[2]], color = test_data[[3]])
```

