## CellTrajectory

Load libraries

```
library("ggplot2") # general plotting
library("igraph") # working with graphs
library("ggpubr") # combining plots
```

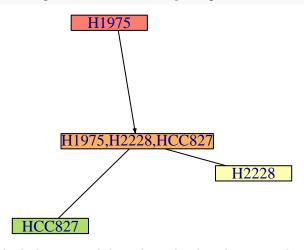
Load functions

```
source("R/kNNgraph.R")
source("R/get_edges2D.R")
```

Load the data

```
cells_info <- readRDS("Data/cellbench-SC1_luyitian.rds")</pre>
```

View the ground truth model



View the kNN graph of the high-dimensional data through a low-dimensional embedding

```
fit <- data.frame(prcomp(cells_info$expression, rank.=2)$x)
fit[["group"]] <- factor(cells_info$grouping)
if(mean(fit[fit[["group"]] == "HCC827", 1]) > 0){
  fit[,1] <- -fit[,1]
}
if(mean(fit[fit[["group"]] == "H2228", 2]) < 0){
  fit[,2] <- -fit[,2]</pre>
```

View the kNN graph of the low-dimensional embedding

Combine the plots

ggarrange(P1, P2, ncol=2)

