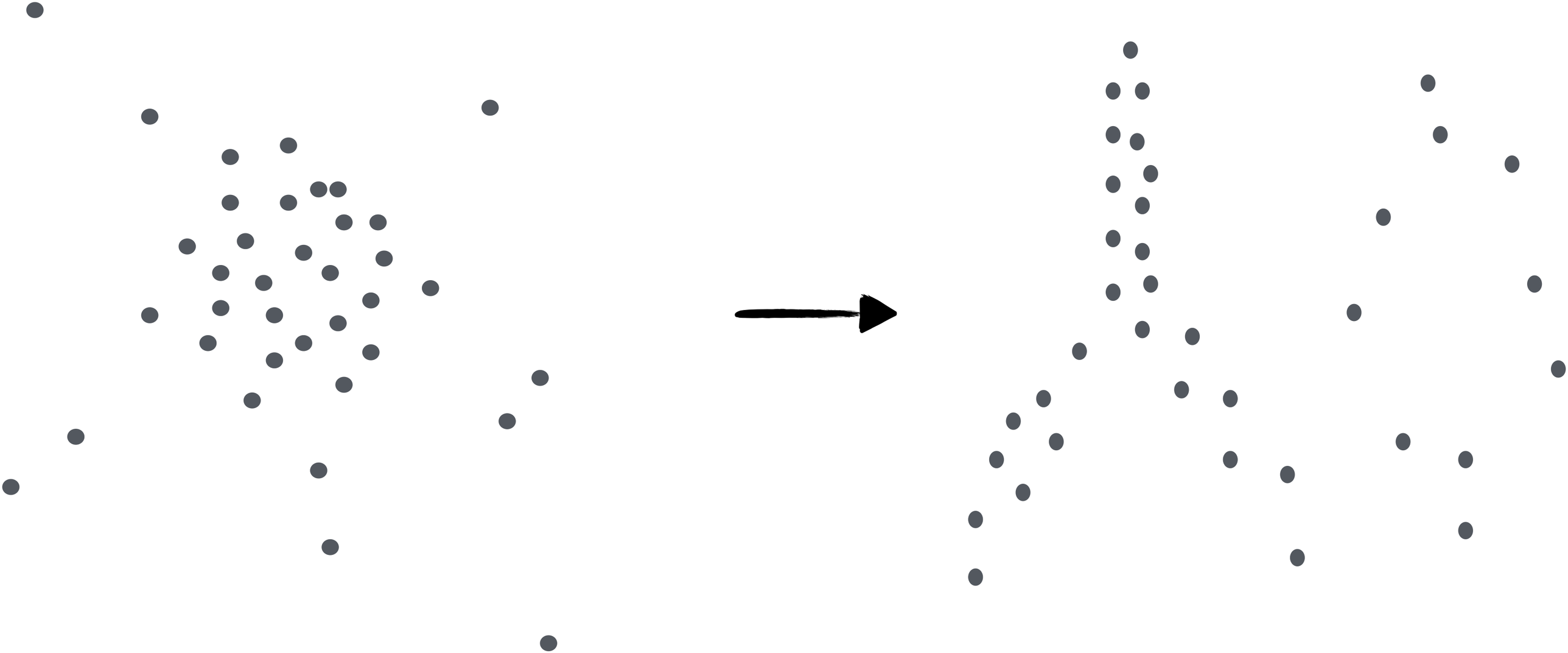
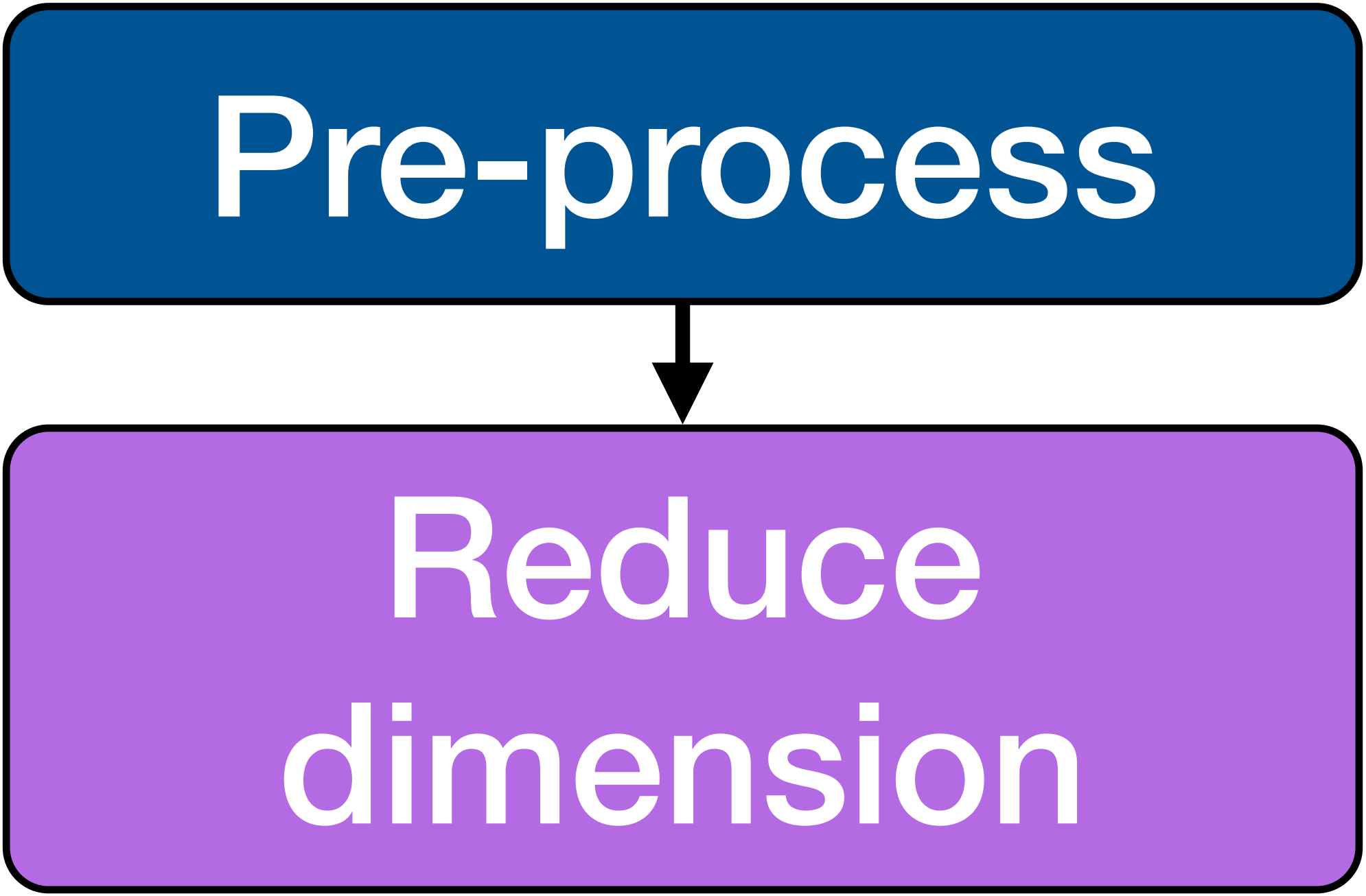
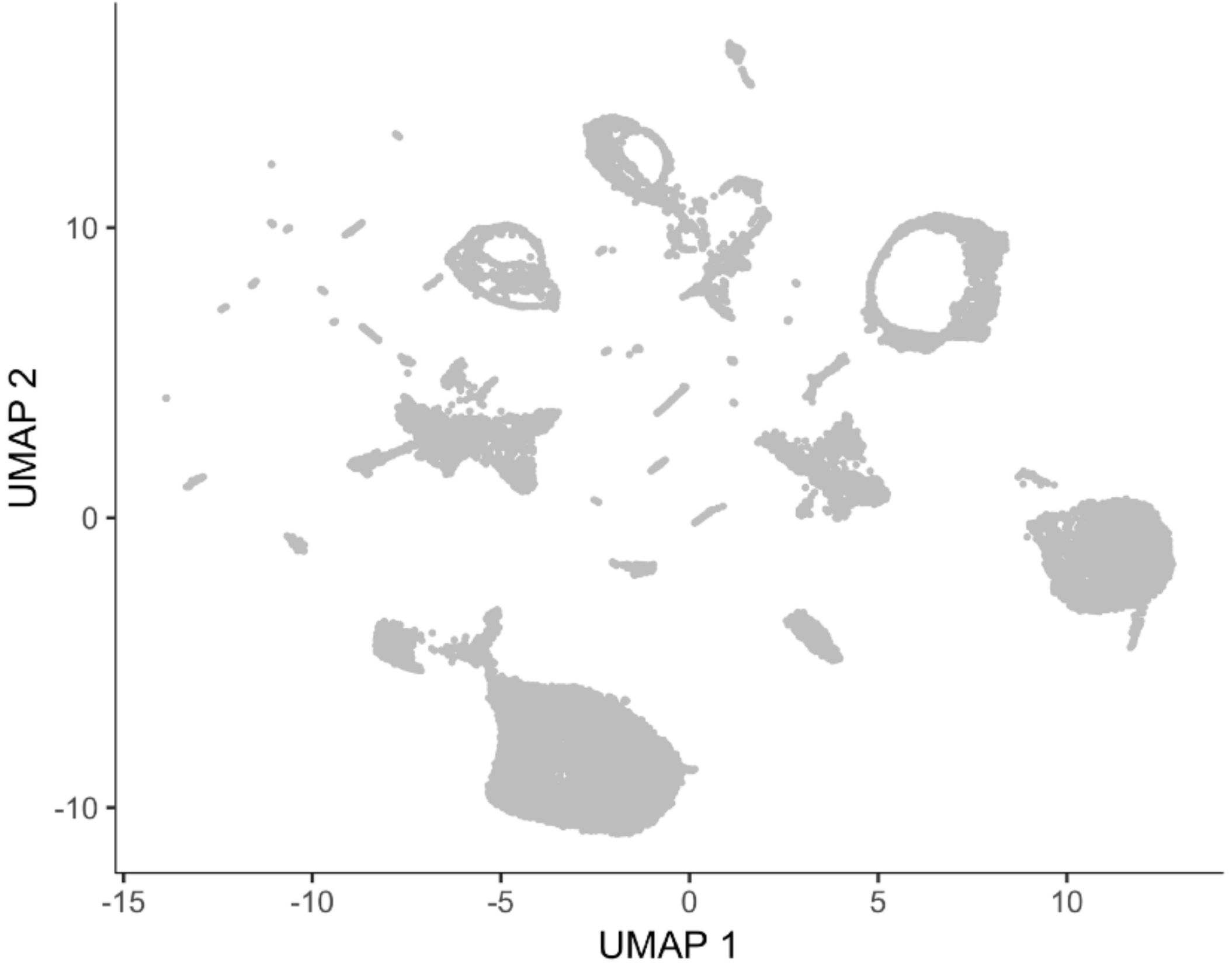
Reduce dimensions



```
cds <- reduce_dimension(cds)
plot_cells(cds)
```

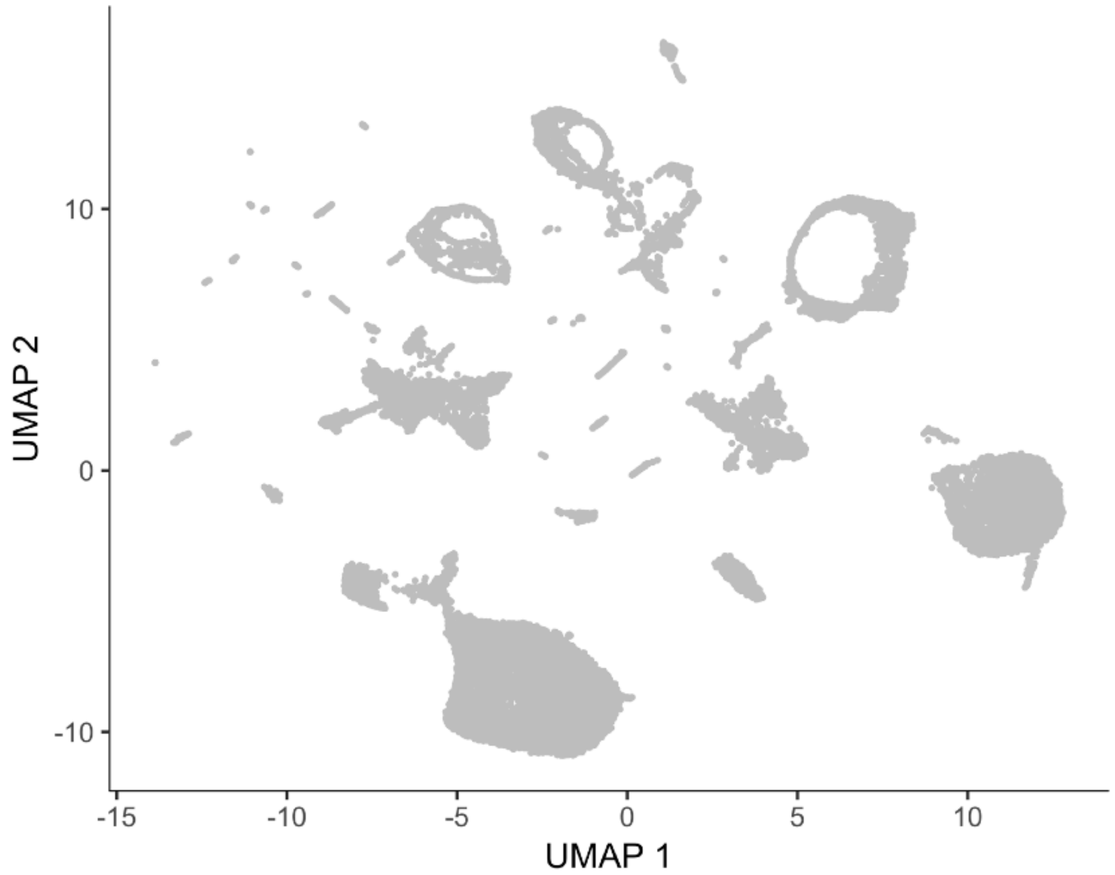

Pre-process
Reduce dimension

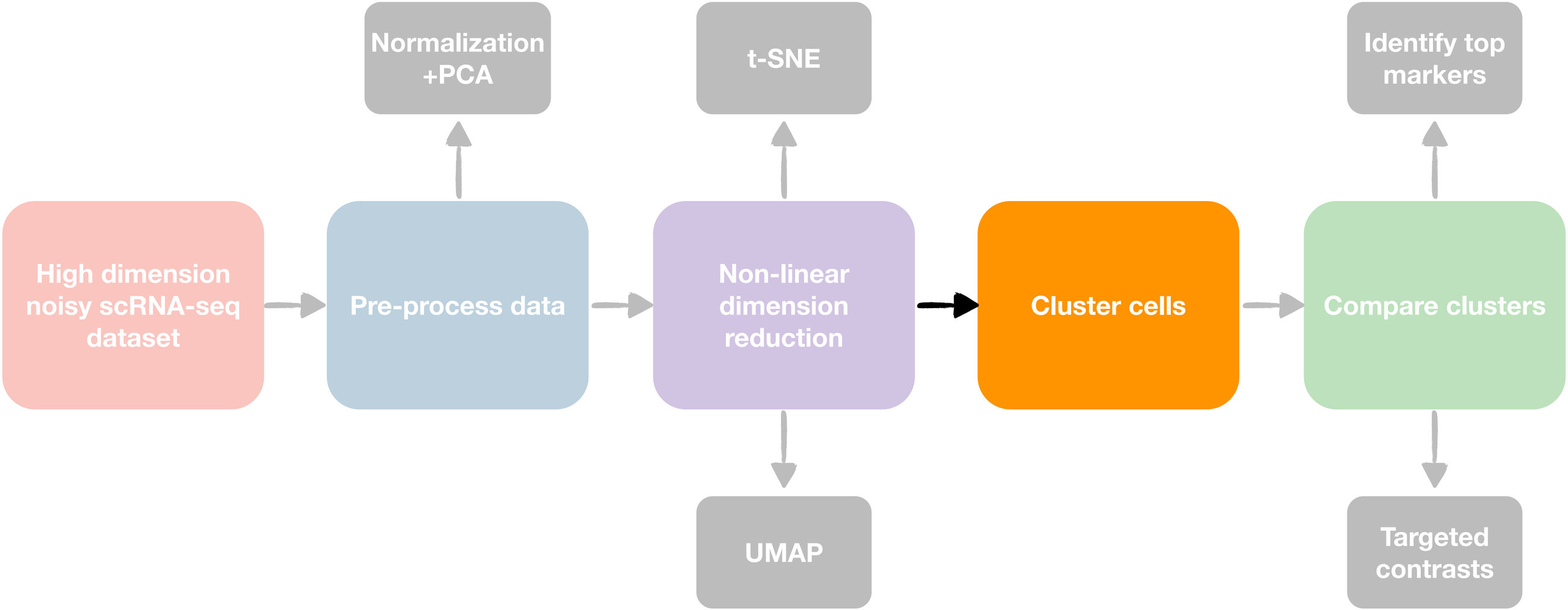

10
0
-10
UMAP 2
-15
-10
-5
0
5
10
UMAP 1

Pre-process

Reduce dimension

Reduce dimensions




```
cds <- reduce_dimension(cds)
plot_cells(cds)
```

Cluster cells