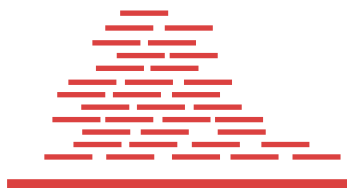
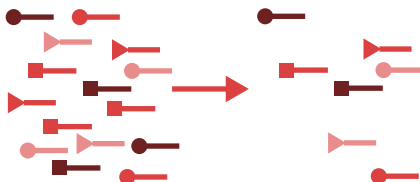


Phase 1 - Data acquisition

Alignment



De-duplication

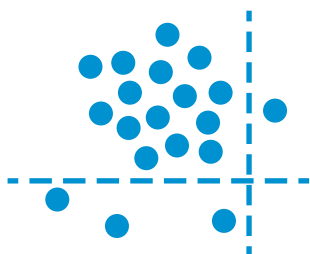


Quantification

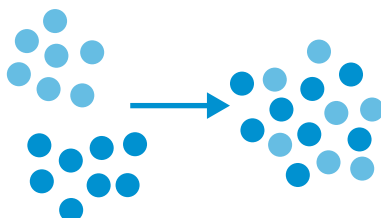


Phase 2 - Data cleaning

Quality control




Normalisation



Imputation

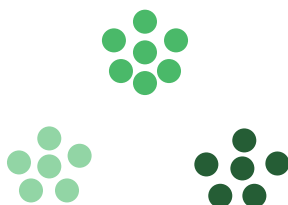
0	1	0	0
0	0	0	0
0	0	2	0
0	1	0	0



1	1	0	1
0	0	0	0
0	1	2	2
0	1	1	0

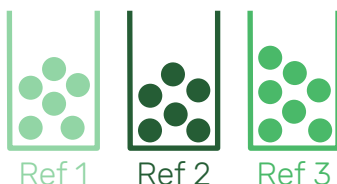
Phase 3 - Cell assignment

Clustering



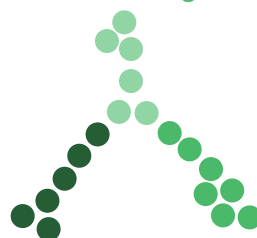
Unknown groups

Classification



Known groups

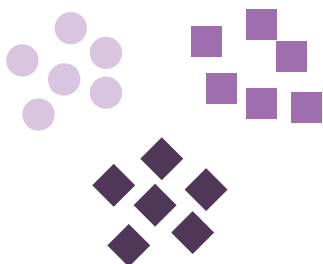
Ordering



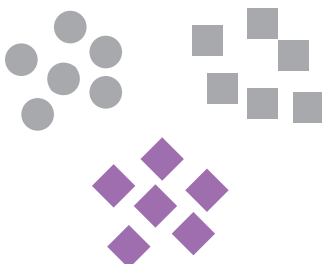
Trajectory

Phase 4 - Gene identification

Differential expression



Marker genes



Expression patterns

