

1st Step: data pre-processing

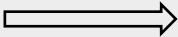
Raw dataset (input)

I_i : individual, $i = 1, \dots, n$

V_j = categorical variable, $j = 1, \dots, m$

	V_1	V_2	...	V_j
I_1				
I_2				
...				
I_i				

Multiple
Correspondence
Analysis

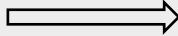


Multiple Correspondence Analysis result

PC_j = Principal Component (continuous variable), $j = 1, \dots, m$

	PC_1	PC_2	...	PC_j
I_1				
I_2				
...				
I_i				

Threshold
Inertia percentage
(at least 50%)



Dataset for Hierarchical clustering (output)

Selected PC , $k < j$

PC_k , $k = 1, \dots, p$

	PC_1	PC_2	...	PC_k
I_1				
I_2				
...				
I_i				

2nd Step: Hierarchical Cluster Analysis

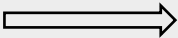
Dataset for Hierarchical clustering (input)

Selected PC , $k < j$

PC_k , $k = 1, \dots, p$

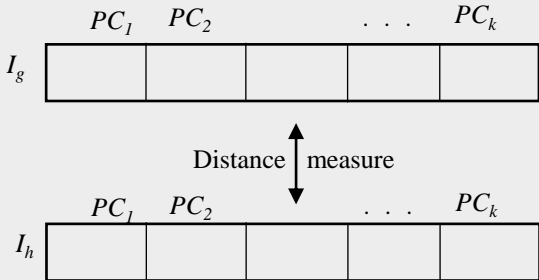
	PC_1	PC_2	...	PC_k
I_1				
I_2				
...				
I_i				

Hierarchical
Cluster Analysis

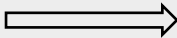


Starting agglomerative process:
distance between p -dimensional
observations

(Euclidian distance)



Ward's algorithm
Agglomeration of
clusters



Hierarchical Cluster Analysis result

Hierarchical tree (output)

