



# 1 simulation

Table 1: Simulation results

Number of clusters	1	2	3	4	5	6	7	8	9	10	PE
<i>Setting 1</i>											
Oracle	<b>100</b>	0	0	0	0	0	0	0	0	0	1
Gap	<b>92</b>	0	1	1	1	2	1	0	1	1	$8.0 \pm 28.3$
Gaussian-Mix	<b>100</b>	0	0	0	0	0	0	0	0	0	$1.0 \pm 0$
CH	—	—	—	—	—	—	—	—	—	—	—
Hartigan	—	—	—	—	—	—	—	—	—	—	—
Jump	<b>0</b>	0	0	0	0	0	1	8	34	57	$102.3 \pm 58.7$
Prediction strength	<b>100</b>	0	0	0	0	0	0	0	0	0	$1.0 \pm 0$
Stability	—	—	—	—	—	—	—	—	—	—	—
Gabriel CV	<b>100</b>	0	0	0	0	0	0	0	0	0	$1.0 \pm 0$
Wold CV	<b>100</b>	0	0	0	0	0	0	0	0	0	$1.0 \pm 0$
<i>Setting 2</i>											
Oracle	0	<b>100</b>	0	0	0	0	0	0	0	0	1
Gap	0	<b>98</b>	1	1	0	0	0	0	0	0	$1.2 \pm 1.3$
Gaussian-Mix	0	<b>100</b>	0	0	0	0	0	0	0	0	$1.0 \pm 0$
CH	0	<b>100</b>	0	0	0	0	0	0	0	0	$1.0 \pm 0$
Hartigan	0	<b>0</b>	6	16	21	14	12	10	8	13	$28.1 \pm 50.8$
Jump	0	<b>70</b>	0	0	0	0	0	2	7	21	$15.5 \pm 55.4$
Prediction strength	0	<b>100</b>	0	0	0	0	0	0	0	0	$1.0 \pm 0$
Stability	0	<b>100</b>	0	0	0	0	0	0	0	0	$1.0 \pm 0$
Gabriel CV	0	<b>99</b>	1	0	0	0	0	0	0	0	$1.1 \pm 1.0$
Wold CV	0	<b>66</b>	3	14	11	4	1	0	1	0	$6.0 \pm 10.6$
<i>Setting 3</i>											
Oracle	0	0	0	<b>100</b>	0	0	0	0	0	0	1
Gap	0	0	0	<b>69</b>	28	3	0	0	0	0	$1.3 \pm 0.5$
Gaussian-Mix	85	2	3	<b>3</b>	1	0	4	2	0	0	$35.9 \pm 13.7$
CH	0	36	28	<b>36</b>	0	0	0	0	0	0	$12.1 \pm 9.9$
Hartigan	0	0	13	<b>75</b>	11	0	1	0	0	0	$2.5 \pm 3.5$
Jump	0	0	0	<b>0</b>	0	0	0	0	14	86	$5.6 \pm 0.4$
Prediction strength	56	0	0	<b>44</b>	0	0	0	0	0	0	$23.5 \pm 20.6$
Stability	0	0	0	<b>33</b>	59	8	0	0	0	0	$1.7 \pm 0.5$
Gabriel CV	0	0	0	<b>100</b>	0	0	0	0	0	0	$1.0 \pm 0$
Wold CV	0	0	0	<b>100</b>	0	0	0	0	0	0	$1.0 \pm 0$
Number of clusters	$\leq 6$	7	8	9	10	11	12	13	14	15	
<i>Setting 4</i>											
Oracle	0	0	0	0	<b>100</b>	0	0	0	0	0	1
Gap	0	0	0	0	<b>12</b>	39	23	16	7	3	$1.5 \pm 0.4$
Gaussian-Mix	71	6	4	2	<b>4</b>	4	0	2	3	4	$22.2 \pm 13.5$
CH	84	6	6	4	<b>0</b>	0	0	0	0	0	$20.8 \pm 8.9$
Hartigan	13	4	13	21	<b>17</b>	12	5	9	1	5	$4.4 \pm 4.8$
Jump	0	0	0	20	<b>80</b>	0	0	0	0	0	$1.4 \pm 0.9$
Prediction strength	100	0	0	0	<b>0</b>	0	0	0	0	0	$35.2 \pm 4.4$
Stability	0	0	0	0	<b>0</b>	3	24	29	28	16	$2.0 \pm 0.3$
Gabriel CV	0	0	0	0	<b>100</b>	0	0	0	0	0	$1.0 \pm 0$
Wold CV	0	0	0	<del>0</del>	<b>100</b>	0	0	0	0	0	$1.0 \pm 0$

In setting 1, “—” means the method can not be used with parameter  $k = 1$ .

Number of clusters	1	2	3	4	5	6	7	8	9	10	PE
<i>Setting 5</i>											
Oracle	0	0	0	<b>100</b>	0	0	0	0	0	0	1
Gap	0	0	0	<b>66</b>	25	7	1	1	0	0	$1.8 \pm 1.3$
Gaussian-Mix	0	0	0	<b>56</b>	36	6	2	0	0	0	$2.0 \pm 1.3$
CH	0	5	25	<b>53</b>	14	2	1	0	0	0	$10.3 \pm 17.0$
Hartigan	0	0	24	<b>61</b>	5	6	3	1	0	0	$7.3 \pm 11.5$
Jump	0	0	0	<b>73</b>	0	0	1	0	5	21	$3.1 \pm 3.5$
Prediction strength	74	8	5	<b>13</b>	0	0	0	0	0	0	$114.5 \pm 67.8$
Stability	0	4	5	<b>23</b>	36	23	7	2	0	0	$5.7 \pm 10.2$
Gabriel CV	0	0	0	<b>100</b>	0	0	0	0	0	0	$1.0 \pm 0.0$
Wold CV	0	0	0	<b>99</b>	1	0	0	0	0	0	$1.0 \pm 0.1$
<i>Setting 6</i>											
Oracle	0	0	<b>100</b>	0	0	0	0	0	0	0	1
Gap	0	0	<b>10</b>	8	13	11	12	11	17	18	$9.8 \pm 5.9$
Gaussian-Mix	0	0	<b>88</b>	12	0	0	0	0	0	0	$1.4 \pm 1.3$
CH	0	17	<b>74</b>	8	1	0	0	0	0	0	$4.0 \pm 6.3$
Hartigan	0	0	<b>87</b>	10	3	0	0	0	0	0	$1.6 \pm 1.6$
Jump	0	0	<b>0</b>	0	0	1	2	13	32	52	$13.7 \pm 5.0$
Prediction strength	19	2	<b>78</b>	1	0	0	0	0	0	0	$10.3 \pm 20.5$
Stability	0	0	<b>24</b>	39	29	7	1	0	0	0	$4.5 \pm 3.0$
Gabriel CV	0	2	<b>97</b>	1	0	0	0	0	0	0	$1.3 \pm 2.1$
Wold CV	0	0	<b>89</b>	9	1	0	1	0	0	0	$1.6 \pm 1.9$

Last column gives the mean and standard deviation of PE for each algorithm.

## 2 Real data application

Table 2: Number of clusters selected by each algorithm

	Congress Voting	Breast Cancer	Sonar
CH-index	2	2	2
Hartigan	3	3	3
Jump	9	9	10
Prediction strength	2	2	1
Bootstrap stability	2	2	10
Gap	10	9	10
Gaussian-Mix	7	5	1
Gabriel	2	2	2
Wold	2	3	10

All the algorithms executed with their default parameter settings with  $k$  ranges from 1 to 10

### 3 CV error with two multivariate normal distributed clusters

#### 3.1 Setup

There are two clusters  $G_1$  and  $G_2$ , where observations from  $G_1$  are distributed as

$$N\left(\begin{pmatrix} \mu_1^X \\ \mu_1^Y \end{pmatrix}, \mathbf{I}\right)$$

and observations from  $G_2$  are distributed as

$$N\left(\begin{pmatrix} -\mu_1^X \\ -\mu_1^Y \end{pmatrix}, \mathbf{I}\right)$$

where  $\mu_1^X > 0$  and  $\mu_1^Y > 0$ . If the true cluster is single cluster  $G$  with

$$P(G = G_1) = P(G = G_2) = 1/2$$

After applying  $K$ -means on  $Y$  axis with  $k = 2$  to the whole data, and assume the observation number  $n \rightarrow \infty$ , we have the estimated center of  $G_1$  be

$$\bar{\mu}_1^Y = 2\varphi(\mu_1^Y) + 2\mu_1^Y \Phi(\mu_1^Y) - \mu_1^Y \quad (1)$$

and the estimated center of  $G_2$  be

$$\bar{\mu}_2^Y = -2\varphi(\mu_1^Y) - 2\mu_1^Y \Phi(\mu_1^Y) + \mu_1^Y \quad (2)$$

where  $\varphi()$  and  $\Phi()$  are the standard normal probability and cumulative distribution function respectively.

### 3.2 CV error with $k = 1$ and $k = 2$

By symmetry, the CV error for points from  $G_1$  is same as the points from  $G_2$ . Because  $P(G = G_1) = P(G = G_2) = 1/2$ , the CV error for  $k = 2$  can be calculated solely from  $G_2$ , that is

$$\begin{aligned}
CV(2) &= E[(Y - \hat{Y})^2], \quad Y \sim N(-\mu_1^Y, 1) \\
&= P(\hat{G} = 2|G = 2) \cdot E[(Y - \hat{Y})^2|\hat{G} = 2] + P(\hat{G} = 1|G = 2) \cdot E[(Y - \hat{Y})^2|\hat{G} = 1] \\
&= P(\hat{G} = 2|G = 2) \cdot E[(Y - \bar{\mu}_2^Y)^2] + P(\hat{G} = 1|G = 2) \cdot E[(Y - \bar{\mu}_1^Y)^2] \\
&= \Phi(\mu_1^X)[var(Y) + (-\mu_1^Y - \bar{\mu}_2^Y)^2] + [1 - \Phi(\mu_1^X)][var(Y) + (-\mu_1^Y - \bar{\mu}_1^Y)^2] \\
&= \Phi(\mu_1^X)[1 + (\mu_1^Y + \bar{\mu}_2^Y)^2] + [1 - \Phi(\mu_1^X)][1 + (\mu_1^Y + \bar{\mu}_1^Y)^2] \\
\bar{\mu}_2^Y = -\bar{\mu}_1^Y &= 1 + (\mu_1^Y + \bar{\mu}_1^Y)^2 - 4\Phi(\mu_1^X)\mu_1^Y\bar{\mu}_1^Y
\end{aligned}$$

where  $\bar{\mu}_1^Y$  is given in equation (1).

When  $k = 1$ , the result is straight forward since the estimated center will be  $(0, 0)$ , so

$$CV(1) = E[Y^2] = 1 + (\mu_1^Y)^2$$

where  $Y \sim N(-\mu_1^Y, 1)$  or  $Y \sim N(\mu_1^Y, 1)$