

I. Pen-and-paper

①

EM clustering algorithm

↳ E-step (I) → expectation

↳ M-step (II) → maximization

①.

$$P(y_2, y_3 | c_k = 1) = \mathcal{N}(x | \mu_k, \Sigma_k) = \frac{1}{(2\pi)^{D/2}} \cdot \frac{1}{|\Sigma_k|^{1/2}} \cdot \exp\left(-\frac{1}{2} \cdot (x - \mu_k)^T \Sigma_k^{-1} \cdot (x - \mu_k)\right)$$

$$P(c_k = 1, x_\eta) = \pi_k \cdot P(y_2, y_3 | c_k = 1) \cdot P(y_1 | c_k = 1)$$

$$P(x_\eta) = \sum_{k=1}^K P(c_k = 1, x_\eta)$$

$$\gamma(c_{\eta k}) = P(c_k = 1 | x_\eta) = \frac{P(c_k = 1, x_\eta)}{P(x_\eta)}$$

②.

$$N_k = \sum_{\eta=1}^N \gamma(c_{\eta k})$$

$$\mu_k = \frac{1}{N_k} \cdot \sum_{\eta=1}^N \gamma(c_{\eta k}) \cdot x_\eta$$

$$\Sigma_k = \frac{1}{N_k} \cdot \sum_{\eta=1}^N \gamma(c_{\eta k}) \cdot (x_\eta - \mu_k) \cdot (x_\eta - \mu_k)^T$$

$$\pi_k = P(c_k = 1) = \frac{N_k}{N}$$

1. Expectation

Given observations $x = \left\{ \begin{pmatrix} 1 \\ 0.6 \\ 0.1 \end{pmatrix}, \begin{pmatrix} 0 \\ 0.4 \\ 0.8 \end{pmatrix}, \begin{pmatrix} 0 \\ 0.2 \\ 0.5 \end{pmatrix}, \begin{pmatrix} 0 \\ 0.4 \\ -0.1 \end{pmatrix} \right\}$:

$$\gamma(c_{11}) = P(c_1 | x_1)$$

$$\gamma(c_{12}) = P(c_2 | x_1)$$

$$\gamma(c_{41}) = P(c_1 | x_4)$$

$$\gamma(c_{42}) = P(c_2 | x_4)$$

$$\gamma(c_{21}) = P(c_1 | x_2)$$

$$\gamma(c_{22}) = P(c_2 | x_2)$$

$$\gamma(c_{31}) = P(c_1 | x_3)$$

$$\gamma(c_{32}) = P(c_2 | x_3)$$

$$P(c_1, x_1) = \underbrace{\pi_1}_{0.5} \cdot P(y_2 = 0.6, y_3 = 0.1 | c_1 = 1) \cdot \underbrace{P(y_1 = 1 | c_1 = 1)}_{0.3}$$

Bernoulli
distribution

$$P(X=x) = \begin{cases} p & \text{for } x=1 \\ 1-p & \text{for } x=0 \end{cases}$$

$$P(y_2 = 0.6, y_3 = 0.1 | c_1 = 1) = \frac{1}{2\pi} \cdot \frac{1}{|\Sigma_1|^{1/2}} \cdot \exp\left(-\frac{1}{2} [-0.4 - 0.9] \Sigma_1^{-1} \begin{bmatrix} -0.4 \\ -0.9 \end{bmatrix}\right)$$

$$\begin{bmatrix} 0.6 \\ 0.1 \end{bmatrix} - \begin{bmatrix} 1 \\ 1 \end{bmatrix} = \begin{bmatrix} -0.4 \\ -0.9 \end{bmatrix} \quad |\Sigma_1|^{1/2} = 1.93649 \quad \Sigma_1^{-1} = \frac{1}{3.75} \begin{bmatrix} 2 & -0.5 \\ -0.5 & 2 \end{bmatrix} = \begin{bmatrix} 0.53333 & -0.13333 \\ -0.13333 & 0.53333 \end{bmatrix}$$

And thus $P(y_2=0.6, y_3=0.1 | c_1=1) = \frac{1}{2\pi} \cdot \frac{1}{1.93649} \cdot \exp(-0.21067) \simeq$
 $\simeq 0.06658$

We can now compute $P(c_1, x_1) = 0.5 \times 0.06658 \times 0.3 = 9.987 \times 10^{-3}$

$P(c_2, x_1) = \underbrace{\pi_2}_{0.5} \cdot P(y_2=0.6, y_3=0.1 | c_2=1) \cdot \underbrace{P(y_1=1 | c_2=1)}_{0.7}$

$P(y_2=0.6, y_3=0.1 | c_2=1) = \frac{1}{2\pi} \cdot \frac{1}{1.11803} \exp\left(-\frac{1}{2} \begin{bmatrix} 1.2 & -0.8 \\ -0.8 & 1.2 \end{bmatrix} \begin{bmatrix} 0.6 \\ 0.1 \end{bmatrix}\right) = *$

$\begin{bmatrix} 0.6 \\ 0.1 \end{bmatrix} - \begin{bmatrix} 0 \\ 0 \end{bmatrix} = \begin{bmatrix} 0.6 \\ 0.1 \end{bmatrix} \quad |\Sigma_2|^{\frac{1}{2}} = 1.11803 \quad \Sigma_2^{-1} = \begin{bmatrix} 1.2 & -0.8 \\ -0.8 & 1.2 \end{bmatrix}$

$* = 0.11962$

$P(c_2, x_1) = 0.5 \times 0.11962 \times 0.7 = 0.04187$

$P(c_1, x_2) = \underbrace{\pi_1}_{0.5} \cdot P(y_2=-0.4, y_3=0.8 | c_1=1) \cdot \underbrace{P(y_1=0 | c_1=1)}_{0.7}$

$P(y_2=-0.4, y_3=0.8 | c_1=1) = \frac{1}{2\pi} \cdot \frac{1}{1.93649} \cdot \exp\left(-\frac{1}{2} \begin{bmatrix} -1.4 & -0.2 \end{bmatrix} \begin{bmatrix} 0.53333 & -0.13333 \\ 0.17333 & 0.53333 \end{bmatrix} \begin{bmatrix} -1.4 \\ -0.2 \end{bmatrix}\right)$

$\begin{bmatrix} -0.4 \\ 0.8 \end{bmatrix} - \begin{bmatrix} 1 \\ 1 \end{bmatrix} = \begin{bmatrix} -1.4 \\ -0.2 \end{bmatrix}$

$** = 0.05005$

$P(c_1, x_2) = 0.5 \times 0.05005 \times 0.7 = 0.01752$

$P(c_2, x_2) = \underbrace{\pi_2}_{0.5} \cdot P(y_2=-0.4, y_3=0.8 | c_2=1) \cdot \underbrace{P(y_1=0 | c_2=1)}_{0.3}$

$P(y_2=-0.4, y_3=0.8 | c_2=1) = \frac{1}{2\pi} \cdot \frac{1}{1.11803} \cdot \exp\left(-\frac{1}{2} \begin{bmatrix} -0.4 & 0.8 \end{bmatrix} \begin{bmatrix} 1.2 & -0.8 \\ -0.8 & 1.2 \end{bmatrix} \begin{bmatrix} -0.4 \\ 0.8 \end{bmatrix}\right) =$

$= 0.06819$

$P(c_2, x_2) = 0.5 \times 0.06819 \times 0.3 = 0.01023$

$$P(C_1, x_3) = \underbrace{\pi_1}_{0.5} \cdot P(y_2=0.2, y_3=0.5 | C_1=1) \cdot \underbrace{P(y_1=0 | C_1=1)}_{0.7}$$

$$P(y_2=0.2, y_3=0.5 | C_1=1) = \frac{1}{2\pi} \cdot \frac{1}{1.93649} \cdot \exp\left(-\frac{1}{2} \begin{bmatrix} 0.8 & -0.5 \end{bmatrix} \begin{bmatrix} 0.53333 & -0.13333 \\ 0.17333 & 0.53333 \end{bmatrix} \begin{bmatrix} -0.8 \\ -0.5 \end{bmatrix}\right) =$$

$$= 0.06837$$

$$P(C_1, x_3) = 0.5 \times 0.06837 \times 0.7 = 0.02393$$

$$P(C_2, x_3) = \underbrace{\pi_2}_{0.5} \cdot P(y_2=0.2, y_3=0.5 | C_2=1) \cdot \underbrace{P(y_1=0 | C_2=1)}_{0.3}$$

$$P(y_2=0.2, y_3=0.5 | C_2=1) = \frac{1}{2\pi} \cdot \frac{1}{1.11103} \cdot \exp\left(-\frac{1}{2} \begin{bmatrix} 0.2 & 0.5 \end{bmatrix} \begin{bmatrix} 1.2 & -0.8 \\ -0.8 & 1.2 \end{bmatrix} \begin{bmatrix} 0.2 \\ 0.5 \end{bmatrix}\right) =$$

$$= 0.12958$$

$$P(C_2, x_3) = 0.5 \times 0.12958 \times 0.3 = 0.01944$$

$$P(C_1, x_4) = \underbrace{\pi_1}_{0.5} \cdot P(y_2=0.4, y_3=-0.1 | C_1=1) \cdot \underbrace{P(y_1=1 | C_1=1)}_{0.3}$$

$$P(y_2=0.4, y_3=-0.1 | C_1=1) = \frac{1}{2\pi} \cdot \frac{1}{1.93649} \cdot \exp\left(-\frac{1}{2} \begin{bmatrix} 0.6 & -1.1 \end{bmatrix} \begin{bmatrix} 0.53333 & -0.13333 \\ 0.17333 & 0.53333 \end{bmatrix} \begin{bmatrix} -0.4 \\ -1.1 \end{bmatrix}\right) =$$

$$= 0.05905$$

$$P(C_1, x_4) = 0.5 \times 0.05905 \times 0.3 = 8.8575 \times 10^{-3}$$

$$P(C_2, x_4) = \underbrace{\pi_2}_{0.5} \cdot P(y_2=0.4, y_3=-0.1 | C_2=1) \cdot \underbrace{P(y_1=1 | C_2=1)}_{0.7}$$

$$P(y_2=0.4, y_3=-0.1 | C_2=1) = \frac{1}{2\pi} \cdot \frac{1}{1.11103} \cdot \exp\left(-\frac{1}{2} \begin{bmatrix} 0.4 & -0.1 \end{bmatrix} \begin{bmatrix} 1.2 & -0.8 \\ -0.8 & 1.2 \end{bmatrix} \begin{bmatrix} 0.4 \\ -0.1 \end{bmatrix}\right) =$$

$$= 0.12450$$

$$P(C_2, x_4) = 0.5 \times 0.12450 \times 0.7 = 0.04358$$

Taking the following values into consideration,

$$P(x_1) = 9.987 \times 10^{-3} + 0.04187 = 0.05186$$

$$P(x_2) = 0.01752 + 0.01023 = 0.02775$$

$$P(x_3) = 0.02393 + 0.01944 = 0.04337$$

$$P(x_4) = 8.8575 \times 10^{-3} + 0.04358 = 0.05244$$

We can now compute:

$$\gamma(c_{11}) = P(c_1|x_1) = \frac{P(c_1, x_1)}{P(x_1)} = \frac{9.987 \times 10^{-3}}{0.05186} \approx 0.19258$$

$$\gamma(c_{12}) = P(c_2|x_1) = \frac{P(c_2, x_1)}{P(x_1)} = \frac{0.04187}{0.05186} \approx 0.80737$$

$$\gamma(c_{21}) = \frac{0.07752}{0.02775} \approx 0.63135$$

$$\gamma(c_{22}) = \frac{0.01023}{0.02775} \approx 0.36865$$

$$\gamma(c_{31}) = \frac{0.02393}{0.04337} \approx 0.55176$$

$$\gamma(c_{32}) = \frac{0.01944}{0.04337} \approx 0.44824$$

$$\gamma(c_{41}) = \frac{8.8575 \times 10^{-3}}{0.05244} \approx 0.16891$$

$$\gamma(c_{42}) = \frac{0.04358}{0.05244} \approx 0.83105$$

2. Maximization

We evaluate

$$N_k = \sum_{n=1}^N \gamma(c_{nk})$$

$$N_1 = 0.19258 + 0.63135 + 0.55176 + 0.16891 = 1.5446$$

$$N_2 = 0.80737 + 0.36865 + 0.44824 + 0.83105 = 2.45531$$

We determine the mean values (and $P(y_1=1|c_1)$)

$$\mu_k = \frac{1}{N_k} \cdot \sum_{n=1}^N \gamma(c_{nk}) \cdot x_n$$

$$\mu_1 = \frac{1}{1.5446} \left(0.19258 \cdot \begin{pmatrix} 0.6 \\ 0.1 \end{pmatrix} + 0.63135 \cdot \begin{pmatrix} 0 \\ -0.4 \end{pmatrix} + 0.55176 \cdot \begin{pmatrix} 0 \\ 0.2 \end{pmatrix} + 0.16891 \cdot \begin{pmatrix} 1 \\ 0.4 \end{pmatrix} \right) = \begin{bmatrix} 0.23403 \\ 0.02649 \\ 0.50714 \end{bmatrix} \rightarrow \begin{matrix} P(y_1=1|c_1) = 0.23403 \\ \mu_1 = \begin{bmatrix} 0.02649 \\ 0.50714 \end{bmatrix} \end{matrix}$$

$$\mu_2 = \frac{1}{2.45531} \left(0.80737 \cdot \begin{pmatrix} 0.6 \\ 0.1 \end{pmatrix} + 0.36865 \cdot \begin{pmatrix} 0 \\ -0.4 \end{pmatrix} + 0.44824 \cdot \begin{pmatrix} 0 \\ 0.2 \end{pmatrix} + 0.83105 \cdot \begin{pmatrix} 1 \\ 0.4 \end{pmatrix} \right) = \begin{bmatrix} 0.66730 \\ 0.30914 \\ 0.21043 \end{bmatrix} \rightarrow \begin{matrix} P(y_1=1|c_2) = 0.66730 \\ \mu_2 = \begin{bmatrix} 0.30914 \\ 0.21043 \end{bmatrix} \end{matrix}$$

and the new covariance matrix

$$\Sigma_k = \frac{1}{N_k} \cdot \sum_{n=1}^N r(c_{nk}) \cdot (x_n - \mu_k) \cdot (x_n - \mu_k)^T$$

$$\Sigma_1 = \frac{1}{1.5446} \left(0.19258 \cdot \begin{pmatrix} 0.6 - 0.02649 \\ 0.1 - 0.50714 \end{pmatrix} \begin{pmatrix} 0.6 - 0.02649 & 0.1 - 0.50714 \end{pmatrix} + \right.$$

$$+ 0.63135 \cdot \begin{pmatrix} -0.4 - 0.02649 \\ 0.8 - 0.50714 \end{pmatrix} \begin{pmatrix} -0.4 - 0.02649 & 0.8 - 0.50714 \end{pmatrix} +$$

$$+ 0.55176 \cdot \begin{pmatrix} 0.2 - 0.02649 \\ 0.5 - 0.50714 \end{pmatrix} \begin{pmatrix} 0.2 - 0.02649 & 0.5 - 0.50714 \end{pmatrix} +$$

$$+ 0.16891 \cdot \begin{pmatrix} 0.4 - 0.02649 \\ -0.1 - 0.50714 \end{pmatrix} \begin{pmatrix} 0.4 - 0.02649 & -0.1 - 0.50714 \end{pmatrix} \Bigg) =$$

$$= \begin{bmatrix} 0.14137 & -0.10541 \\ -0.10541 & 0.09605 \end{bmatrix}$$

and

$$\Sigma_2 = \begin{bmatrix} 0.10829 & -0.08865 \\ -0.08865 & 0.10412 \end{bmatrix}$$

and the new mixing parameter is

$$\pi_k = p(c_k = 1) = \frac{N_k}{N}$$

$$\pi_1 = p(c_1 = 1) = \frac{1.5446}{4} \simeq 0.38615$$

$$\pi_2 = p(c_2 = 1) = \frac{2.45531}{4} \simeq 0.61383$$

②

$$x_{\text{new}} = \begin{pmatrix} 1 \\ 0.3 \\ 0.7 \end{pmatrix} \quad \boxed{\text{posteriors: } p(c_1 | x_{\text{new}}) \text{ and } p(c_2 | x_{\text{new}})}$$

$$p(c_1, x_{\text{new}}) = \underbrace{\pi_1}_{0.38615} \cdot p(y_2=0.3, y_3=0.7 | c_1=1) \cdot \underbrace{p(y_1=1 | c_1=1)}_{0.23403}$$

$$P(y_2=0.3, y_3=0.7 | c_1=1) = \frac{1}{2\pi} \frac{1}{|\Sigma_1|^{\frac{1}{2}}} \exp\left(-\frac{1}{2}(x-\mu_1)^T \Sigma_1^{-1} \cdot (x-\mu_1)\right) =$$

$$|\Sigma_1|^{\frac{1}{2}} = 0.04967 \quad \begin{bmatrix} 0.3 \\ 0.7 \end{bmatrix} - \begin{bmatrix} 0.02649 \\ 0.50714 \end{bmatrix} = \begin{bmatrix} 0.27351 \\ 0.19286 \end{bmatrix}$$

$$\Sigma_1^{-1} = \begin{bmatrix} 38.92887 & 42.72246 \\ 42.72246 & 57.27698 \end{bmatrix}$$

$$* = \frac{1}{2\pi} \cdot \frac{1}{0.04967} \cdot \exp(-4.77524) \simeq 0.02703$$

$$\text{and thus } P(c_1, x_{\text{new}}) = 0.38615 \times 0.02703 \times 0.23403 = \\ = 2.44272 \times 10^{-3}$$

$$P(c_2, x_{\text{new}}) = \underbrace{\pi_2}_{0.61383} \cdot P(y_2=0.3, y_3=0.7 | c_2=1) \cdot \underbrace{P(y_1=1 | c_2=1)}_{0.66730}$$

$$P(y_2=0.3, y_3=0.7 | c_2=1) = \frac{1}{2\pi} \frac{1}{|\Sigma_2|^{\frac{1}{2}}} \exp\left(-\frac{1}{2}(x-\mu_2)^T \Sigma_2^{-1} \cdot (x-\mu_2)\right) = \\ = \frac{1}{2\pi} \cdot \frac{1}{0.05845} \cdot \exp(-3.68380) \simeq 0.06842$$

$$|\Sigma_2|^{\frac{1}{2}} \simeq 0.05845 \quad \begin{bmatrix} 0.3 \\ 0.7 \end{bmatrix} - \begin{bmatrix} 0.30914 \\ 0.21043 \end{bmatrix} = \begin{bmatrix} -9.14 \times 10^{-3} \\ 0.48957 \end{bmatrix}$$

$$\Sigma_2^{-1} = \begin{bmatrix} 30.47713 & 25.94888 \\ 25.94888 & 31.69774 \end{bmatrix}$$

$$\text{and thus } P(c_2, x_{\text{new}}) = 0.61383 \times 0.06842 \times 0.66730 \simeq \\ \simeq 0.02803$$

$$P(x_{\text{new}}) = P(c_1, x_{\text{new}}) + P(c_2, x_{\text{new}}) = \\ = 2.44272 \times 10^{-3} + 0.02803 \simeq 0.03047$$

$$P(c_1 | x_{\text{new}}) = \frac{2.44272 \times 10^{-3}}{0.03047} \simeq 0.08017$$

$$P(c_2 | x_{\text{new}}) = \frac{0.02803}{0.03047} \simeq 0.91992$$

③

under a ML assumption, we can assign each observation to a cluster by computing $\arg\max_k P(x|c_k)$:

$$P(x_1|c_1) = \frac{P(c_1, x_1)}{P(c_1)} = \frac{9.987 \times 10^{-3}}{0.38615} = 0.02586$$

$$P(x_1|c_2) = \frac{0.04187}{0.61383} = 0.06821$$

→ x_1 should be assigned to c_2

$$P(x_2|c_1) = \frac{0.01752}{0.38615} = 0.04537$$

$$P(x_2|c_2) = \frac{0.01023}{0.61383} = 0.01667$$

→ x_2 should be assigned to c_1

$$P(x_3|c_1) = \frac{0.02393}{0.38615} = 0.06197$$

$$P(x_3|c_2) = \frac{0.01944}{0.61383} = 0.03167$$

→ x_3 should be assigned to c_1

$$P(x_4|c_1) = \frac{8.8575 \times 10^{-3}}{0.38615} = 0.02294$$

$$P(x_4|c_2) = \frac{0.04358}{0.61383} = 0.07100$$

→ x_4 should be assigned to c_2

thus, clusters = $\{c_1 = \{x_2, x_3\}, c_2 = \{x_1, x_4\}\}$

Silhouette score (for both clusters)

$$S(x) = \frac{b(x) - a(x)}{\max\{a(x), b(x)\}}, \quad a(i) = \frac{1}{\text{size of cluster} - 1} \sum_{\substack{j \text{ in} \\ \text{the same} \\ \text{cluster as} \\ i}} d(i, j),$$

$$b(i) = \min_{\substack{c \neq \text{cluster} \\ \text{of } i}} \left(\frac{1}{\text{size of } c} \sum_{\substack{j \text{ in} \\ \text{cluster} \\ c}} d(i, j) \right)$$

manhattan distance

$$d(x_1, x_2) = |1-0| + |0.6-(-0.4)| + |0.1-0.8| = 2.7$$

$$d(x_1, x_3) = 1.8$$

$$d(x_1, x_4) = 0.4$$

$$d(x_2, x_3) = 0.9$$

$$d(x_2, x_4) = 2.7$$

$$d(x_3, x_4) = 1.8$$

$$S(c_1) = \sum_{x_i \in c_1} S(x_i)$$

$$S(x_2) = \frac{b(x_2) - a(x_2)}{\max\{a(x_2), b(x_2)\}} = 1 - \frac{a(x_2)}{b(x_2)} = 1 - \frac{0.9}{2.7} = \frac{2}{3}$$

$$a(x_2) = 0.9$$

$$b(x_2) = \frac{1}{2}(2.7 + 2.7) = 2.7$$

$$S(x_3) = 1 - \frac{a(x_3)}{b(x_3)} = 1 - \frac{0.9}{1.8} = \frac{1}{2}$$

$$a(x_3) = 0.9$$

$$b(x_3) = \frac{1}{2}(1.8 + 1.8) = 1.8$$

$$S(c_1) = S(x_2) + S(x_3) = \frac{2}{3} + \frac{1}{2} = \frac{7}{6} = 1.1(6)$$

$$S(c_2) = \sum_{x_i \in c_2} S(x_i)$$

$$S(x_1) = 1 - \frac{a(x_1)}{b(x_1)} = 1 - \frac{0.4}{2.25} = \frac{37}{45}$$

$$a(x_1) = 0.4$$

$$b(x_1) = \frac{1}{2}(2.7 + 1.8) = 2.25$$

$$S(x_4) = 1 - \frac{a(x_4)}{b(x_4)} = 1 - \frac{0.4}{2.25} = \frac{37}{45}$$

$$a(x_4) = 0.4$$

$$b(x_4) = \frac{1}{2}(2.7 + 1.8) = 2.25$$

$$S(c_2) = S(x_1) + S(x_4) = 2 \times \frac{37}{45} = 1.6(4)$$

④ Given the fact that purity is the number of correctly matched class and cluster labels divided by the number of total datapoints and that

$$\frac{1}{4} \sum_{k=1}^2 (\arg \max (|c_k \cap g_j|)) = 0.75$$

we can infer that $0.75 \times 4 = 3$ is the maximum number $\sum_{k=1}^2 (\arg \max (|c_k \cap g_j|))$ can be.

since $\text{clusters} = \{c_1 = \{x_2, x_3\}, c_2 = \{x_1, x_4\}\}$,
 $\arg \max (|c_k \cap g_j|) \in \{1, 2\}$.

Thus, $\sum_{k=1}^2 (\arg \max (|c_k \cap g_j|))$ can only be 3 if

$$\sum_{k=1}^2 (\arg \max (|c_k \cap g_j|)) = \arg \max_{c_1} (1, 1) + \arg \max_{c_2} (0, 2) =$$

↑ number of observations per class

$= 2 + 1 = 3$ (there's also the possibility of swapping these values, but what matters is that two observations of one cluster are part of the same class and the other two of the remaining cluster are part of different classes.)

Taking this into consideration, we can conclude that the number of possible classes is ≥ 2 .

II. Programming and critical analysis

1. Code:

```
import pandas as pd
from scipy.io.arff import loadarff

import warnings
warnings.filterwarnings("ignore")

data = loadarff('column_diagnosis.arff')
df = pd.DataFrame(data[0])
df['class'] = df['class'].str.decode('utf-8')
df.head()

from sklearn import metrics, cluster
from sklearn.preprocessing import MinMaxScaler
import numpy as np

def purity_score(y_true, y_pred):
    confusion_matrix = metrics.cluster.contingency_matrix(y_true, y_pred)
    return np.sum(np.amax(confusion_matrix, axis=0)) /
np.sum(confusion_matrix)

X = df.drop("class", axis=1)
y = df["class"]
X_normalized = MinMaxScaler().fit_transform(X)
kmeans_list = []

for k in [2, 3, 4, 5]:
    print("k =", k)
    kmeans = cluster.KMeans(n_clusters=k, random_state=0).fit(X_normalized)
    kmeans_list.append(kmeans)
    print("\tSilhouette Score =", metrics.silhouette_score(X_normalized,
kmeans.labels_))
    print("\tPurity =", purity_score(y, kmeans.labels_))

k = 2
    Silhouette Score = 0.36081773371884557
    Purity = 0.6290322580645161

k = 3
    Silhouette Score = 0.29579055730002257
    Purity = 0.667741935483871

k = 4
    Silhouette Score = 0.2686566721650703
```

```
Purity = 0.6612903225806451
```

```
k = 5
```

```
Silhouette Score = 0.24328260038805272
```

```
Purity = 0.6741935483870968
```

2.

i. Code:

```
from sklearn.decomposition import PCA
```

```
learnt_pca = PCA(n_components=2)
```

```
learnt_pca.fit(X_normalized)
```

```
X_trans = learnt_pca.transform(X_normalized)
```

```
top2_explained_variance = learnt_pca.explained_variance_ratio_
```

```
print(top2_explained_variance[0]+top2_explained_variance[1])
```

```
0.77137397434354
```

ii. Code:

```
labels = list(df.columns)
```

```
labels.remove("class")
```

```
weights_component1 = list(abs(learnt_pca.components_[0]))
```

```
weights_component2 = list(abs(learnt_pca.components_[1]))
```

```
input_vars_by_relevance1= [x for _, x in sorted(zip(weights_component1, labels), key=lambda pair: pair[0])]
```

```
input_vars_by_relevance2= [x for _, x in sorted(zip(weights_component2, labels), key=lambda pair: pair[0])]
```

```
print("Most relevant input variables by component (in ascending order of importance):")
```

```
print("- Component 1:", input_vars_by_relevance1)
```

```
print("- Component 2:", input_vars_by_relevance2)
```

```
Most relevant input variables by component (in ascending order of importance):
```

```
- Component 1: ['pelvic_radius', 'degree_spondylolisthesis', 'sacral_slope', 'pelvic_tilt', 'lumbar_lordosis_angle', 'pelvic_incidence']
```

```
- Component 2: ['degree_spondylolisthesis', 'lumbar_lordosis_angle', 'pelvic_incidence', 'sacral_slope', 'pelvic_radius', 'pelvic_tilt']
```

3. Code and graphs:

```
import matplotlib.pyplot as plt

X_2d = X_trans
cluster_labels = kmeans_list[1].labels_

class_labels = df['class'].unique()

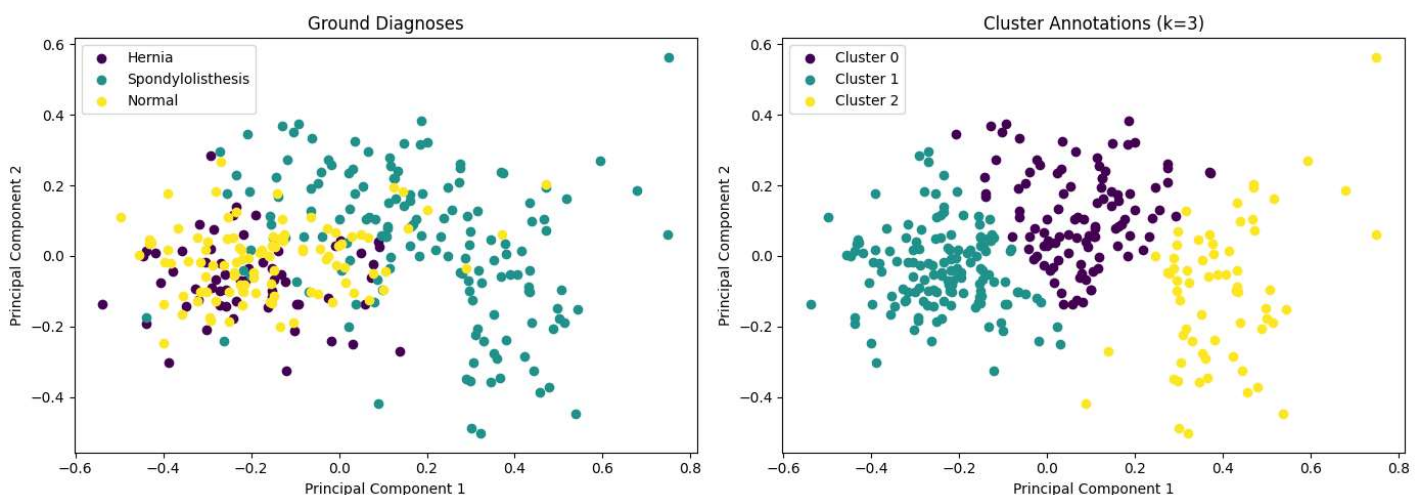
colors = plt.cm.viridis(np.linspace(0, 1, len(class_labels)))

fig, ax = plt.subplots(1, 2, figsize=(14, 5))

ax[0].set_title('Ground Diagnoses')
ax[0].set_xlabel('Principal Component 1')
ax[0].set_ylabel('Principal Component 2')
for i, label in enumerate(class_labels):
    indices = df['class'] == label
    ax[0].scatter(X_2d[indices, 0], X_2d[indices, 1], label=label,
color=colors[i])
ax[0].legend()

ax[1].set_title('Cluster Annotations (k=3)')
ax[1].set_xlabel('Principal Component 1')
ax[1].set_ylabel('Principal Component 2')
for i in range(len(class_labels)):
    indices = cluster_labels == i
    ax[1].scatter(X_2d[indices, 0], X_2d[indices, 1], label=f'Cluster {i}',
color=colors[i])
ax[1].legend()

plt.tight_layout()
plt.show()
```



4. We can find natural clusters within the population based on the features used for clustering by applying algorithms like K-means to the dataset. These clusters might highlight trends or associations between people who have similar traits. When it comes to health, it's probable that some clusters correspond to people with comparable diseases or risk factors, while other clusters correspond to people who are generally in good health.

Moreover, clustering can be used to divide the population into various groups, each with a distinct health profile. For instance, people with high-risk health profiles may form one cluster, and people with low-risk profiles may form another. Medical professionals may find this segmentation useful as it enables them to customize interventions or treatments for particular population subgroups.

END