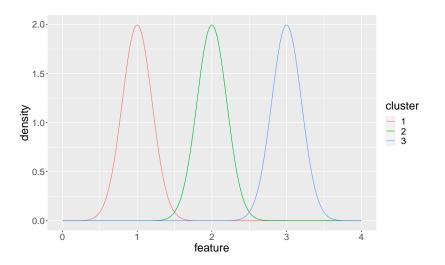
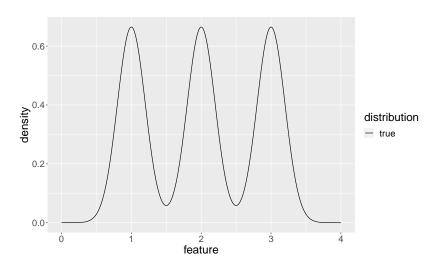
Clustering Model Selection

Toby Dylan Hocking

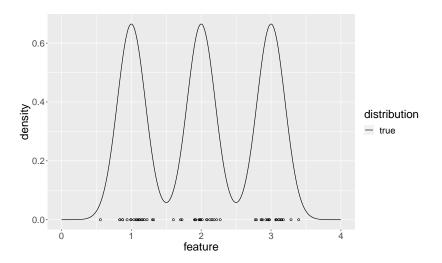
Three normal densities

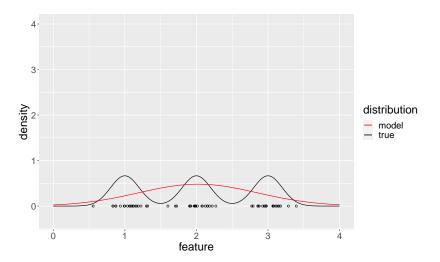


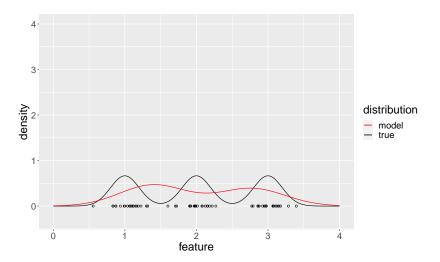
Mixture density

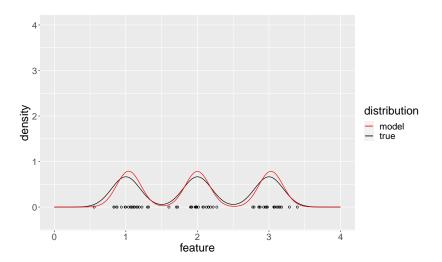


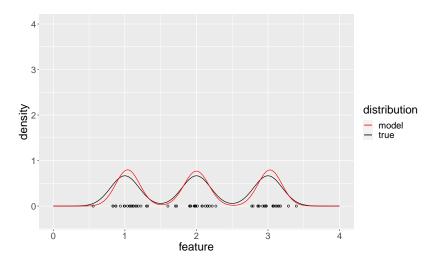
Generate 20 random data from each density

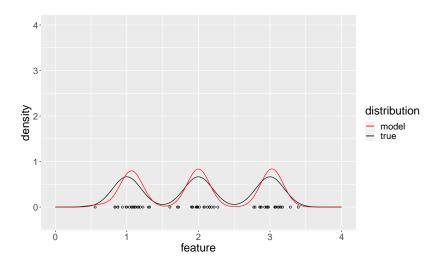


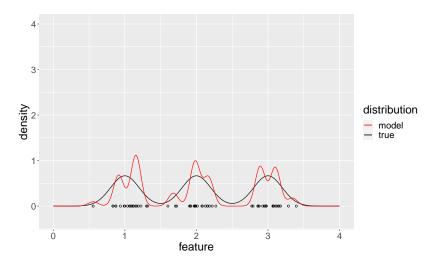


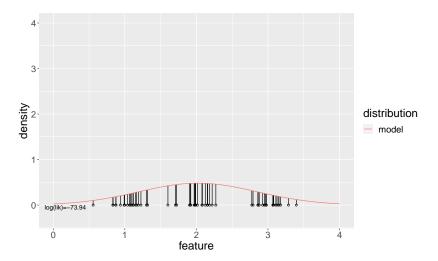


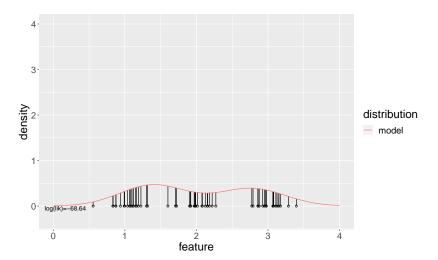


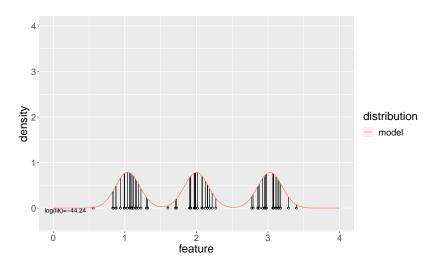


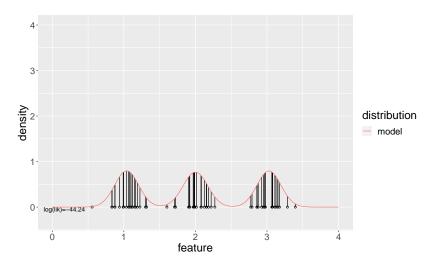


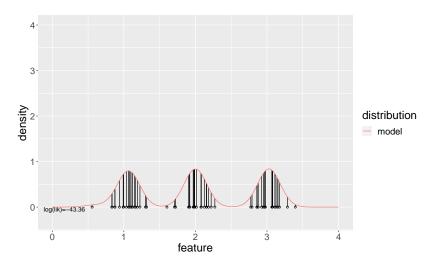


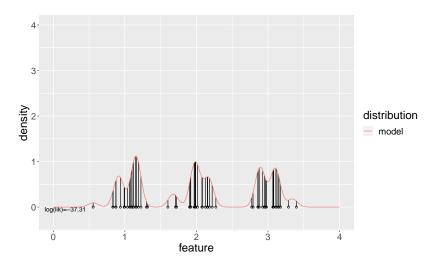




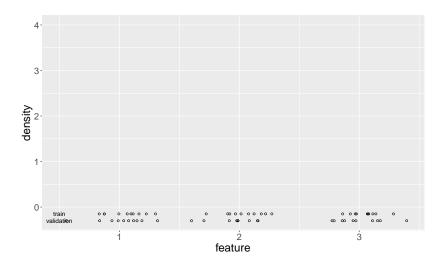


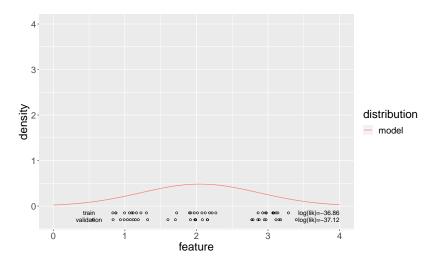


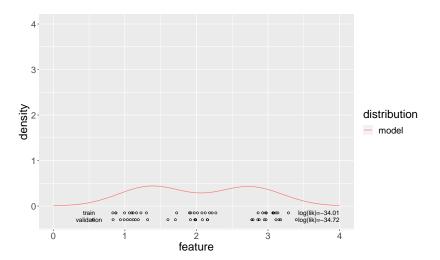


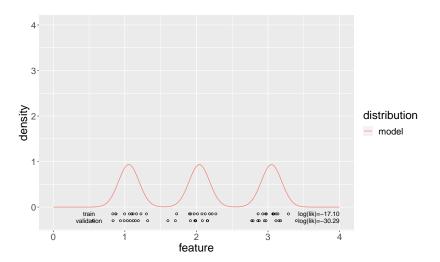


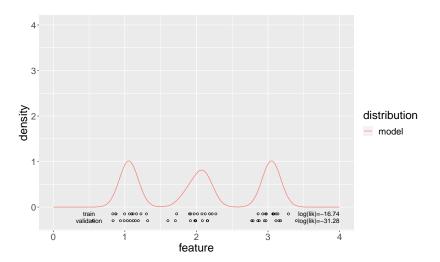
Divide into train and validation

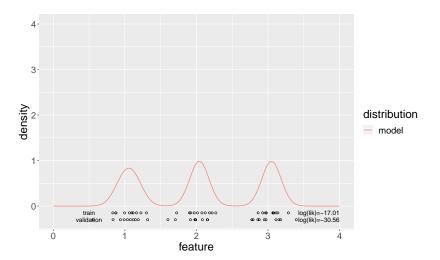


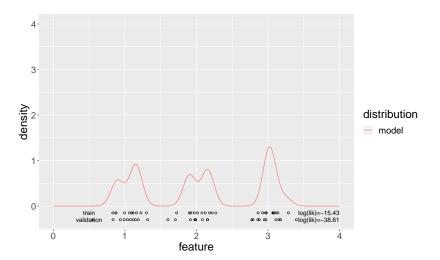


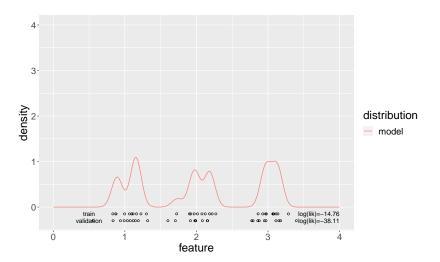


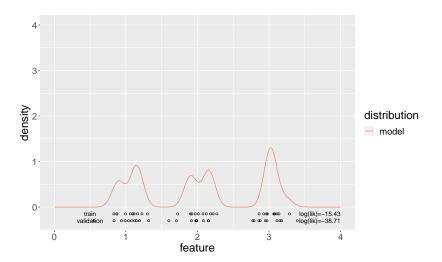


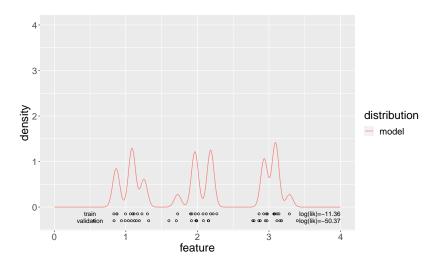


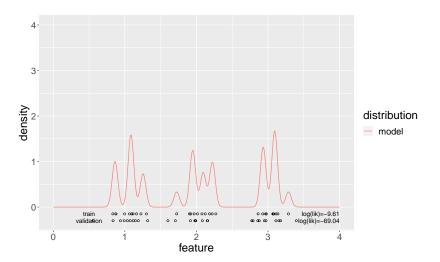




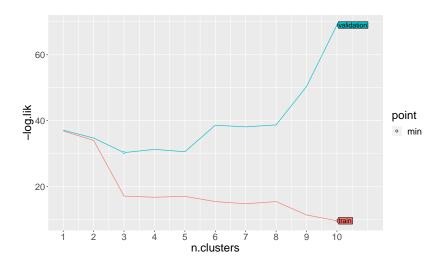




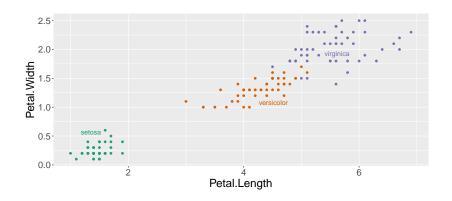




Overall log likelihood plot



Visualize iris data with labels



Visualize iris data without labels

- Let $X = [x_1 \cdots x_n]^{\mathsf{T}} \in \mathbb{R}^{n \times p}$ be the data matrix (input for clustering), where $x_i \in \mathbb{R}^p$ is the input vector for observation i.
- **Example** iris n = 150 observations, p = 2 dimensions.

##		Petal.Width	Petal.Length
##	[1,]	0.2	1.4
##	[2,]	0.2	1.4
##	[3,]	0.2	1.3
##	[4,]	0.2	1.5

