Solution 1:

See R code

Solution 2:

```
a) k = 3
```

2 circles and 1 triangle, so our point is also a circle

```
## Warning in library(package, lib.loc = lib.loc, character.only = TRUE, logical.return =
TRUE, : there is no package called 'ggplot2'
## Error in qplot(x, y, shape = z, size = I(6), col = z): could not find function "qplot"
```

b) k = 5

3 circles and 3 triangles, we have to specify beforehand what to do in case of a tie

```
## Warning in library(package, lib.loc = lib.loc, character.only = TRUE, logical.return =
TRUE, : there is no package called 'ggplot2'
## Error in qplot(x, y, shape = z, size = I(6), col = z): could not find function "qplot"
```

c) k = 7

3 circles and 4 triangles, so our point is also a triangle

```
## Warning in library(package, lib.loc = lib.loc, character.only = TRUE, logical.return =
TRUE, : there is no package called 'ggplot2'
## Error in qplot(x, y, shape = z, size = I(6), col = z): could not find function "qplot"
```

a) When using the naive Bayes classifier, the features $x := (x_{\text{Color}}, x_{\text{Form}}, x_{\text{Origin}})$ given the category $y \in \{\text{yes}, \text{no}\}$ are assumed to be conditionally independent of each other, s.t.

$$p((x_{\text{Color}}, x_{\text{Form}}, x_{\text{Origin}})|y=k) = p(x_{\text{Color}}|y=k) \cdot p(x_{\text{Form}}|y=k) \cdot p(x_{\text{Origin}}|y=k).$$

For the posterior probabilities $\pi_k(x)$ it holds that

$$\pi_k(x) \propto \underbrace{\pi_k \cdot p(x_{\text{Color}}|y=k) \cdot p(x_{\text{Form}}|y=k) \cdot p(x_{\text{Origin}}|y=k)}_{=:\alpha_k(x)}$$

$$\iff \exists c \in \mathbb{R} : \pi_k(x) = c \cdot \alpha_k(x),$$

where π_k is the prior probability of class k. From this and since the posterior probabilities need to sum up to 1, it holds that

$$1 = c \cdot \alpha_{\text{yes}}(x) + c \cdot \alpha_{\text{no}}(x)$$
$$\iff c = \frac{1}{\alpha_{\text{ves}}(x) + \alpha_{\text{no}}(x)}.$$

This means in order to compute $\pi_{\text{yes}}(x)$ the scores $\alpha_{\text{yes}}(x)$ and $\alpha_{\text{no}}(x)$ are needed.

Now we want to compute for a new fruit the posterior probability $\hat{\pi}_{yes}((yellow, round, imported))$.

Note that we do not know the true prior probability and the true conditional densities. Here -since the target and the features are categorical- we can estimate them with the relative frequencies encountered in the data, s.t.

$$\begin{split} \hat{\alpha}_{\text{yes}}(x) &= \hat{\pi}_{yes} \cdot \hat{p}(\text{yellow}|y = \text{yes}) \cdot \hat{p}(\text{round}|y = \text{yes}) \cdot \hat{p}(\text{imported}|y = \text{yes}) \\ &= \hat{\mathbb{P}}(y = \text{yes}) \cdot \hat{\mathbb{P}}(x_{\text{Color}} = \text{yellow}|y = \text{yes}) \cdot \hat{\mathbb{P}}(x_{\text{Form}} = \text{round}|y = \text{yes}) \cdot \hat{\mathbb{P}}(x_{\text{Origin}} = \text{imported}|y = \text{yes}) \\ &= \frac{3}{8} \cdot \frac{1}{3} \cdot \frac{1}{3} \cdot 1 = \frac{1}{24} \approx 0.042, \\ \hat{\alpha}_{\text{no}}(x) &= \hat{\pi}_{no} \cdot \hat{p}(\text{yellow}|y = \text{no}) \cdot \hat{p}(\text{round}|y = \text{no}) \cdot \hat{p}(\text{imported}|y = \text{no}) \\ &= \hat{\mathbb{P}}(y = \text{no}) \cdot \hat{\mathbb{P}}(x_{\text{Color}} = \text{yellow}|y = \text{no}) \cdot \hat{\mathbb{P}}(x_{\text{Form}} = \text{round}|y = \text{no}) \cdot \hat{\mathbb{P}}(x_{\text{Origin}} = \text{imported}|y = \text{no}) \\ &= \frac{5}{8} \cdot \frac{2}{5} \cdot \frac{3}{5} \cdot \frac{2}{5} = \frac{3}{50} = 0.06. \end{split}$$

At this stage we can already see that the predicted label is "no", since $\hat{\alpha}_{\text{no}}(x) = 0.06 > \frac{1}{24} = \hat{\alpha}_{\text{yes}}(x)$. With this we can calculate the posterior probability

$$\hat{\pi}_{\text{yes}}(x) = \frac{\hat{\alpha}_{\text{yes}}(x)}{\hat{\alpha}_{\text{ves}}(x) + \hat{\alpha}_{\text{no}}(x)} \approx 0.41.$$

Corresponding R-Code:

```
df_banana <- data.frame(
   Color = as.factor(
        c("yellow", "yellow", "brown", "brown", "green", "green", "red")),
   Form = as.factor(
        c("oblong", "round", "oblong", "oblong", "round", "oblong", "round")),
   Origin = as.factor(
        c("imported", "domestic", "imported", "imported", "domestic", "imported",
        "domestic", "imported")),
   Banana = as.factor(c("yes", "no", "no", "yes", "no", "yes", "no", "no"))
)

new_fruit <- data.frame(Color = "yellow", Form = "round", Origin = "imported", Banana = NA)
df_banana <- rbind(df_banana, new_fruit)</pre>
```

b) For the distribution of a numerical feature given the the category we need to specify a probability distribution with continuous support. For example, for the information $x_{\rm Length}$ we could assume that $p(x_{\rm Length}|y={\rm yes}) \sim \mathcal{N}(\mu_{\rm yes},\sigma_{\rm yes}^2)$ and $p(x_{\rm Length}|y={\rm no}) \sim \mathcal{N}(\mu_{\rm no},\sigma_{\rm no}^2)$. (To estimate these normal distributions one would need to estimate their parameters $\mu_{\rm yes},\mu_{\rm no},\sigma_{\rm yes}^2,\sigma_{\rm no}^2$ on the data respectively)