Course > Section 2: Machine ... > 2.2: Conditional Pro... > Comprehension Ch...

## Comprehension Check: Conditional Probabilities Part 1

Q1

1/1 point (graded)

In a previous module, we covered Bayes' theorem and the Bayesian paradigm. Conditional probabilities are a fundamental part of this previous covered rule.

$$P(A|B) = P(B|A) \frac{P(A)}{P(B)}$$

We first review a simple example to go over conditional probabilities.

Assume a patient comes into the doctor's office to test whether they have a particular disease.

- The test is positive 85% of the time when tested on a patient with the disease (high sensitivity): P(test + | disease) = 0.85
- The test is negative 90% of the time when tested on a healthy patient (high specificity): P(test-|heathy) = 0.90
- The disease is prevalent in about 2% of the community: P(disease) = 0.02

Using Bayes' theorem, calculate the probability that you have the disease if the test is positive.

0.1478261 **✓ Answer**: 0.1478261

## Explanation

$$P(\text{disease}|\text{test+}) = P(\text{test+}|\text{disease}) \times \frac{P(\text{disease})}{P(\text{test+})} = \frac{P(\text{test+}|\text{disease}) P(\text{disease})}{P(\text{test+}|\text{disease}) P(\text{disease}) + P(\text{test+}|\text{healthy}) P(\text{healthy})]} = \frac{0.85 \times 0.02}{0.85 \times 0.02 + 0.1 \times 0.9}$$

Submit

You have used 1 of 2 attempts

**1** Answers are displayed within the problem

The following 4 questions (Q2-Q5) all relate to implementing this calculation using R.

We have a hypothetical population of 1 million individuals with the following conditional probabilities as described below:

- The test is positive 85% of the time when tested on a patient with the disease (high sensitivity): P(test + | disease) = 0.85
- The test is negative 90% of the time when tested on a healthy patient (high specificity): P(test-|heathy) = 0.90
- The disease is prevalent in about 2% of the community: P(disease) = 0.02

Here is some sample code to get you started:

```
set.seed(1)
disease <- sample(c(0,1), size=le6, replace=TRUE, prob=c(0.98,0.02))
test <- rep(NA, le6)
test[disease==0] <- sample(c(0,1), size=sum(disease==0), replace=TRUE, prob=c(0.90,0.10))
test[disease==1] <- sample(c(0,1), size=sum(disease==1), replace=TRUE, prob=c(0.15, 0.85))</pre>
```

What is the probability that a test is positive? 0.114509 ✓ Answer: 0.114509 0.114509 Explanation The probability of a positive test can be calculated using mean(test). You have used 1 of 5 attempts Submit Answers are displayed within the problem Q3 1/1 point (graded) What is the probability that an individual has the disease if the test is negative? 0.003461356 Answer: 0.003461356 0.003461356 Explanation The probability of having the disease given a negative test can be calculated using <code>[mean(disease[test==0])]</code>. Submit You have used 1 of 5 attempts **1** Answers are displayed within the problem Q4 1/1 point (graded) What is the probability that you have the disease if the test is positive? Remember: calculate the conditional probability the disease is positive assuming a positive test. 0.1471762 **✓ Answer:** 0.1471762 0.1471762 Explanation The probability of having the disease given a positive test can be calculated using [mean(disease[test==1]==1)]. You have used 1 of 5 attempts Submit **1** Answers are displayed within the problem Q5 1/1 point (graded) If the test is positive, what is the relative risk of having the disease? First calculate the probability of having the disease given a positive test, then normalize it against the disease prevalence. 7.389106 **✓ Answer:** 7.389106 7.389106 **Explanation** The relative risk can be calculated using mean(disease[test==1]==1)/mean(disease==1). You have used 1 of 5 attempts Submit

Answers are displayed within the problem

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