# Harvard Medical School Test

### Introduction

"We developed a test for a disease. The false negative rate is zero ("sensivity"), the false positive rate is  $\frac{5}{100}$  ("specifity") and the incidence of the disease in the population is  $\frac{1}{1000}$ ."

"What is the probability that a randomly selected person has the disease?"

This question was posed at the Harvard Medical School to staff and students alike (see [1]), but similar questions are surely part of many Statistics introductory courses as well.

### Sampling

We solve the problem by sampling.

We denote with "0" a person in the population which does not have the disease under question. We denote with "1" a person which does have the disease.

Define the size of the population, the *prevalence* of the disease in the population and the *false positive* rate of the test:

Generate the population to draw from:

```
in[5]:= Clear[prev, pop];
prev = prevalence * popSize;
pop = Append[Table[i - i, {i, 1, popSize - prev}],
Table[j - j + 1, {j, 1, prev}]] // Flatten;
```

Define a predicate for the result of the test, person  $\in \{0,1\}$ , 'True' if the test(!) delivers a positive test result, 'False' otherwise:

```
In[8]= Clear[testQ];
testQ[person_] :=
Module[
{falsePositive = First[RandomSample[Range[100], 1]],
result = Indeterminate},
If[person == 1, result = True];
If[person == 0,
Which[
falsePositive <= 5, result = True,
falsePositive > 5, result = False]];
Return[result]]
```

Define a predicate whether a person *really* has the disease:

```
In[10]= Clear[diseaseQ];
diseaseQ[person_] :=
Module[
{result = Indeterminate},
If[person == 0, result = False]; If[person == 1, result = True];
Return[result]]
```

Do a Monte Carlo simulation to find an estimate of the probability.

```
In[12]:= Timing[Module[
```

```
{
        numIter = 100000, (* How many trials? *)
        patient, (* The sample drawn *)
        patientRes, (* Disease status of sample *)
        testRes, (* The test results of sample *)
        countDis, (* sample member with disease *)
        countPos(* sample member withpos. test *)
       },
       patient = Flatten[Table[RandomSample[pop, 1], {numIter}]];
       patientRes = diseaseQ /@ patient; testRes = testQ /@ patient;
       countDis = Count[patientRes, True];
       countPos = Count[testRes, True];
       Print["H = People with disease: ", countDis];
       Print["E = People tested 'positive': ", countPos];
       Print["P(H|E): ", N[countDis/countPos, 2]]]]
     H = People with disease: 101
     E = People tested 'positive': 5157
     P(H|E): 0.020
Out[12]= {1.548235, Null}
```

### Bayes' Rule

It is instructive to solve for this conditional probability analytically.

To do this, we need to apply Bayes' rule:

$$P(H \black) = \frac{P(E \black) H) * P(H)}{P(E)}, P(H), P(E) > 0$$
(1)

H denotes the hypothesis and E the corresponding evidence. Note that this is the form as derived by Thomas Bayes himself.

#### First Implementation

Let's write a function which implements this rule.

```
In[13]:= Clear[bayesRule];
bayesRule[prior_, likelihood_, evidence_] :=
Module[{}, (likelihood * prior) / evidence]
What are the arguments for our problem?
H = "The person has the disease.",
E = "The test for the person is positive.",
P(H) = 1/(1000),
```

 $P(\neg H) = \frac{999}{1000}$ P(E)H) = 1 [because the test finds the disease always (= 1- false neg. rate)],  $P(E \square H) = \frac{5}{100}$  ( = false positive rate),  $P(E) = P(E)H) *P(H) + P(E) H)*P(H) = 1*\frac{1}{1000} + \frac{50}{1000} * \frac{999}{1000}$ In[15]:= Clear [PH, PnotH, PEH, PEnotH, PE]; PH = 1 / 1000;PnotH = 999 / 1000;PEH = 1;PEnotH = 5 / 100;PE = PEH \* PH + PEnotH \* PnotH; In[21]:= Clear[PHE]; PHE = bayesRule[PH, PEH, PE] 20 Out[22]= 1019 In[23]:= **N[PHE]** Out[23] = 0.0196271

#### Second Implementation

We follow the approach as given in stackoverflow:

```
In[24]:= Clear[conditionalProb];
                                                                       \texttt{conditionalProb[pC_, pTC_, pTNC_] /; (0 < pC < 1) \&\& (0 < pTC \le 1) \&\& (0 < pTNC \le 1) := \texttt{conditionalProb[pC_, pTC_, pTNC_] /; (0 < pC < 1) \&\& (0 < pTNC \le 1) := \texttt{conditionalProb[pC_, pTC_, pTNC_] /; (0 < pC < 1) &\& (0 < pTC \le 1) \&\& (0 < pTNC \le 1) := \texttt{conditionalProb[pC_, pTC_, pTNC_] /; (0 < pC < 1) &\& (0 < pTC \le 1) &\& (0 < pTNC \le 1) &\& (0 < pTNC
                                                                                       Module[{}, (pTC * pC) / ({pTC, pTNC}, {pC, 1 - pC})]
```

```
In[26]:= conditionalProb[1/1000, 1, 5/100]
```

20 Out[26]= 1019

#### Third Implementation

We follow the second approach as given in stackoverflow:

```
In[27]:= Remove [P];
     Unprotect@Intersection;
     Intersection[A_Symbol, B_Symbol] := {A, B}
     Intersection[A_Not, B_Symbol] := {A, B}
     Intersection[A_Symbol, B_Not] := {A, B}
     P[Int_List /; Length@Int == 2] := P[Int[[2]] & Int[[1]]] P[Int[[1]]]
      (*//P(B) given knowledge of P(A)//*)
     P[B_, A_] := If[NumericQ@B, B, P[B & A] P[A] + P[B & Not@A] P[Not@A]]
     P[Not@B_, A_: 1] := If[NumericQ@A, 1 - P[B], 1 - P[B, A]]
     P[A_ & B_] := P[A \cap B] / P[B, A]
     P[Not@A_ & B_] := 1 - P[A & B];
In[37]:= P[H] = 1 / 1000
Out[37] = \frac{1}{1000}
In[38]:= P[EV & H] = 1
```

Out[38]= 1

```
In[39]:= \mathbf{P}[\mathbf{Ev} \approx \neg \mathbf{H}] = 5 / 100
Out[39]= \frac{1}{20}
In[40]:= \mathbf{P}[\mathbf{H} \approx \mathbf{Ev}]
Out[40]= \frac{20}{1019}
In[41]:= \mathbf{N}[\%]
Out[41]= 0.0196271
```

## Bibliography

1. Casscells, Ward and Schoenberger, Arno and Graboys, Thomas B., Interpretation by Physicians of Clinical Laboratory Results. New England Journal of Medicine, 1978.