

# SEIR Model

**About:** SEIR stands for susceptible (S), exposed (E), infected (I), and resistant (R). It simplifies mathematical modeling of infectious diseases. The population considered in this model is categorized in S, E, I and R categories. The flow pattern is depicted by the order of the labels of each compartment. The parameters of the flow of control are:

**alpha and beta:** control the rate of flow of people through the pattern

**sigma:** controls the rate of flow from exposed to infected

**gamma:** controls the rate of flow from infected to resistant

**mu:** mortality ate of those unaffected by the disease

**nu:** controls the rate of flow from susceptible to resistant by vaccinating to avoid getting infected

**Algorithms used:** According to the national institutes of health, this model uses a two-step diagnostic algorithm.

The algorithm is as follows:

1] The first diagnostic (denoted dx1) is assumed to have sensitivity  $s_1$  and specificity  $sp_1$ . The second diagnostic (denoted dx2) is assumed to have sensitivity  $s_2$  and specificity  $sp_2$ .

2] It follows that the overall sensitivity of the algorithm (in which the tests are applied sequentially, and a positive overall result requires a positive result on both tests) is  $s_1 \cdot s_2$ , and the overall specificity is  $sp_1 + (1 - sp_1) \cdot sp_2$ .

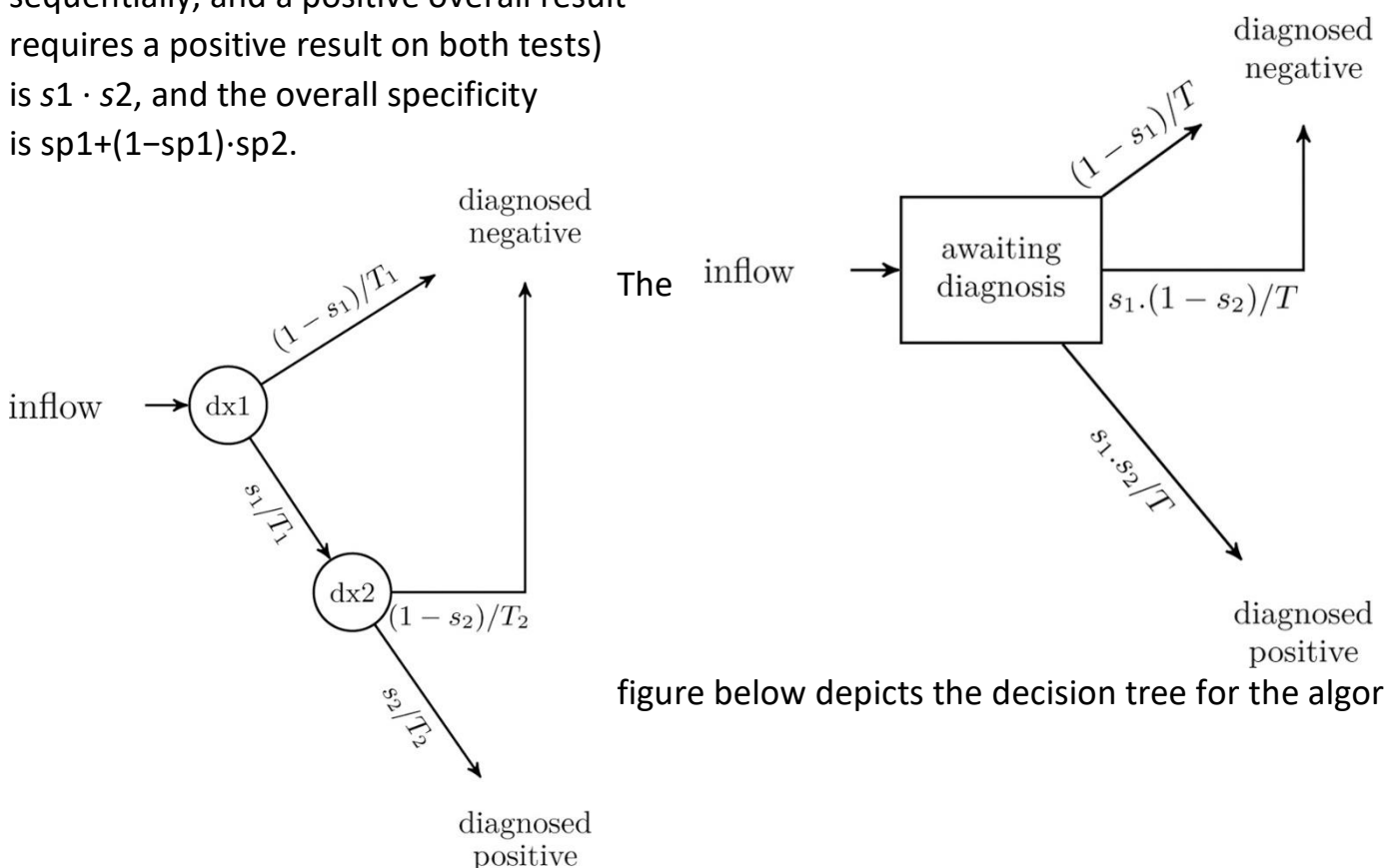


figure below depicts the decision tree for the algorithm:

References: <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC6711160/>  
<http://www.public.asu.edu/~hnesse/classes/seir.html>