```
Machine Learning Home work I
                                                                                                   How F. Ourleya - 504241526
     Q1-A) P(21 X=0, Y=0) = P(X=0, Y=0, 2)
                                                                          P(X=0, Y=0)
      ->P(X=0,Y=0,Z=1): X=0, Y=0, Z=1 is 0.06 (from table)
->P(X=0,Y=0,Z=2): X=0, Y=0, Z=2 1s 0.09
          0,06+0,09 = 0,15 P(X=QY=0) = 0,75
        P(21x=0, Y=0) : for 2=1:
[2-1] P(2-1/X=0, Y=0) = P(X=0) Y=0, Z=1)
                                                                                                            = 0.06 = 0.4
                                                                     P(x=0, 4=0)
  (2=2] P(2=21 X=0, Y=0) = P(X=0, Y=0, Z=2) = 0,09 = 0,6
                                                                                P(x=0, Y=0)
           therefore;
                                              P(21X=0, Y=0) = { 0,4 \ 1/2=1 }
     Q1-B) Mognetizing over X, we must find P(Z; Y) values for each X.
             \rho(2,4) = \sum \rho(x,4,2) = 1
    Q=1,Y=-1) => P(Z=1,Y=-1) = P(X=0, Y=-1,Z=1) + P(X-1, Y=-1,Z=1) = 0,06+0,06
    (2=1, Y=0) =) P(X=0, Y=0, Z=1) + P(X=1, Y=0, Z=1) = 0.06 + 0.02 = 0.08
    (2=1, y=1) => P(x=0, Y=1, Z=1) + P(x=0, Y=1, Z=1) = 0.08 + 0.02 = 0.10
   (2-2, 4=-1) =) P(x=0, 4=1, 2=2) + P(x=1, 4=-1, 2=2) = 0.09+0.24=0.33
   (2-2, Y=0) => P(X=0, Y=0, Z=2) + P(X=1, Y=0, Z=2) = 0,09+0,08=0,17
   (2=2, Y=1) =) p(X=0, Y=1, 2=2) + p(X=1, Y=1, 2=2) = 0112,+0,01 = 0,20
     P(4) = \( \langle (\frac{1}{2}, \frac{1}{2}) - \frac{1}{2} + \frac{1}{2}
      -> 2=1,4=-1 P(2=1,4=-1) =0,72, P(2=1).P(4=-1) = 0,12 7 0,3-0,4
          Since P(2=1,4=-1) # P(2-1-). P(4-1) 2 and 4 are not independent, they are depost
```

Q2-) Ducusion questions => -> What hoppers if folds are different for different models? If the folds differ for each model. Models will be trained and vollaged on different data points. This reduces the reliability of comparison If models will work on differt data subset loss and error calculations are herd to comment because of inconsistency. Each model performance ore directly related to data samples. Therefore if the Holds are different compaison is less rearingfull - what happens if the folds ore obtained without shuffling! Without shuffling dater may not be rondonly distributed accross folds. Therefore biosed Justrubston will appear in training and validation sets. It different tolds how patterns because of the nature of dedouset. This case may couse overfitting or underfitting, depends on the fold plts Q5-A) $\Rightarrow P(x)2) = \int P(x)dx \quad \Rightarrow p(x) = \int \exp(-|x|) ||x| = x, \text{ for } x > 0;$ $P(x)(2) = \int_{\frac{1}{2}} exp(-x) dx$ $\rightarrow S \exp(-x) dx = -\exp(-x)$ $P(x)2 = 0 - \left(\frac{1}{2} \exp(-2)\right) = \frac{1}{2} \exp(-2)$ So, P(x)2)81.01353 = 0,0677 % 6,77 chance of x to be >2

Machine Leaning Homewal DE 3.) A.) with virus => %90 accoracy + /10%2 of population sick without virus => % 20 occuracy actually sick, tested positive -> 9 P(Postive | Sict) = 0.9 P(Negative | Healty) = 0.9 P(Sick | Positive) P(Healthy) = 1- P(Sick) = 0.98 P(Sick)=0.02 P (Positive | Sick) = 0.9 P(Positive | Healthy) = 0.1 P(Sick | Positive) = P(Positive | Sick). P(sick) P(Positive) P(Positive Trick) P(sick) P(Sick | Positie) = 0.9.0.02 = 0.018 = 0.155 0.116 = 0.116 = 0.155P (Portle Healthy) . P (Helly) 0.9. 0.02+01.09 P(Positive) = 0.018+ 0.038 = 0.116 % 15 is a relatively low probability so there is a high chance of false posities due to the %2 of population, Hest is untrustable