Then P(A,B|C) = P(B,C) = P(B

```
2. [A] OLet F= 12, then ANF= A= NA, Daisin NA ELA
      DLet E = ANF, for YFEC. Then it is known that (ANF) U(ANF, ) = AN(F, UF, ) = A
      (AMEDIN(AMEC) = AM(FINEC) = $ AMEC is the complement of Er in terms of ILA. Since
      FCEC, then (ANF,C) ECA
     BLEE E,= ANFI, Ez= ANFZ for YF, EEC. THEN E, VEZ= (ANF) V (ANE) = AN(F, UE).
       Since FUELEC, then AN(FUE) ECA. E, UELECA
     According to the definition of collection, Ca is a valid collection
     OPA(DA) = PA(DAA) = P(DAA) = P(A) = P(A) = P(A)
     @PALSILA = PLAN >0 as PLA) >0 and PLSNA) >0 for VSEL
    For V Si, Si EC if Si, Si are disjoint other PAC(Si NA) UCS, NA) = PAI(Si US) NA) = PICSI US) NA)
     P[A] = P(S,NA) = P(S,NA) + P(S,NA) [CS,NA) and (S,NA) are disjoint] = P(S,NA) + P(S,NA) P(A)
   = Pa(Sina) + Pa(Sina) According to the definition of probability measure, this measure is valid
  (b) Yes. O P(1)= # of i-values such that NiER = N = 1 Since NiER for Vi
         OPENP(S) >0 as N>0 and the humerator is pleases non-negative
        Tor YS1, Sz ED it S1, Sz are disjoint , then
           P(S,US2)= number of i-values such that K:ES,US2 = # of i-values for MiES, # of i-values for MiES.
                  = # it i-values for ViEs, + # of i-values for ViEs, = P(S,)+P(S)
      According to the definition, it's a valid probability measure
```

```
3. Let f = \begin{cases} 1 \text{ if } food \text{ is } good \\ 0 \text{ otherwise} \end{cases}; S = \begin{cases} 1 \text{ if } good \text{ sleeper} \\ 0 \text{ if } food \text{ is } bad \end{cases}; W = \begin{cases} 1 \text{ if } wake up \\ 0 \text{ otherwise} \end{cases}; S = \begin{cases} 1 \text{ if } good \text{ sleeper} \\ 0 \text{ if } bad \text{ sleeper} \end{cases}

The quality of food and the type of the beby are independent in this case.

Then we get: P(f=0) = 0.1, P(W=1|f=0) = 1, P(S=1) = 0.6, P(f=1) = 0.9, P(S=0) = 0.9

P(W=1|S=1,f=1) = 0.1, P(W=1|S=0,f=1) = 0.8

(a) P(W=1) = P(W=1|f=0) P(f=0) + P(W=1|S=1,f=1) P(S=1) P(S=1)
```

(d) Since $P(f=0|w=1) \neq P(f=0|s=1,w=1)$, these two events are not conditionally independent given that w=1 according to the definition. In the context, given the information that the baby wakes up, it is intuitive to have more certainties on the quality of the food if it is known that the baby is a good sleeper.

```
4. (a) P(x=5) = (\frac{1}{2})^5 = \frac{1}{32}; P(x=4) = (\frac{1}{2})^5 \cdot 2 = \frac{1}{16}; P(x=3) = 2 \cdot (\frac{1}{2})^4 + (\frac{1}{2})^5 = \frac{5}{32}
                  P(X=2) = (\frac{1}{2})^{3} \times 2 + (\frac{1}{2})^{4} \times 2 - (\frac{1}{2})^{5} = \frac{11}{32}
P(X=1) = [-P(X=5) - P(X=4) - P(X=3) - P(X=2) - P(X=0)
= [-\frac{1}{31} - \frac{1}{16} - \frac{5}{32} - \frac{11}{32} - \frac{1}{32} = \frac{3}{8}
```

(b) Code:

```
def p_longest_streak(n, tries):
   count = np.zeros(n+1)
   for i in range(int(tries)):
       sequence = np.random.randint(0,2,size = n)
       currentStreak = 0
       longestStreak = 0
       for j in range(n):
           if sequence[j] == 1:
                currentStreak += 1
               if(currentStreak > longestStreak):
                   longestStreak = currentStreak
               currentStreak = 0
       if(currentStreak > longestStreak):
           longestStreak = currentStreak
       count[longestStreak] += 1
    prob = count/tries
    return prob
   # Write your Monte Carlo code here, n is the length of the sequence and tries is the number
   # of sampled sequences used to produce the estimate of the probability
```

Plot: tries: 1000.0 tries: 5000.0 tries: 10000.0 tries: 10000.0 tries: 10000.0 tries: 5000.0 tries: 5000.0 tries: 5000.0 tries: 10000.0 0.35 0.25 0.20 0.15 0.10 0.20 0.15 0.10 0.05

We can see that the estimated probabilities for each longest streak in a sequence of length 5 have not much difference with the exact probabilities for that

(c)

print("The probability that the longest streak of ones in a Bernoulli iid sequence of length 200 has length 8 or more is ")

print(sum(p_longest_streak(200,1e5)[8:])) # Compute the probability and print it here

The probability that the longest streak of ones in a Bernoulli iid sequence of length 200 has length 8 or more is

The estimated probability that the longest streak of heads has length 8 or more for 200 flips is about 0.32. Therefore, the sequence of 8 ones is not an evidence that the program may not be generating truly random sequences since a randomly generated sequence can have a sequence of 8 or more ones for 200 flips with a high probability(32%).