Probability and Random Processes

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Probability Theory

Defintion 1.0.1 (Sample Space, Ω). *The set of all possible outcomes of an experiment.*

1.1 Discrete Random Variables

- A <u>discrete random variable</u> associates a real number with each outcome in Ω , e.g., X = i is a random variable if the throw of a die is i for i = 1, 2, ..., 6.
- Note that X^2 is still a random variable and in general, a function of a random variable is also a random variable.
- Probability mass function is the set $\{a, P_X(a)\}$ where $a \in \mathcal{A}$ and \mathcal{A} is the set of all possible values taken by X.

Example 1.1.1. Chess match between Alice and Bob. The first to win a game wins the match and conversely, the match is drawn if there are 10 consecutive draws.

$$P(Alice\ wins\ a\ game) = 0.3$$
 $P(Bob\ wins\ a\ game) = 0.4$ $P(Draw) = 0.3$

1. What is the PMF of the duration of a match?

Let L denote the duration of the match. Suppose L = 10, then

$$P_L(10) = (0.3)^9.$$

Otherwise, if L = k < 10,

$$P_L(k) = (0.3)^{k-1}(0.7)$$
 for $k = 1, 2, ..., 9$.

2. Calculate $P(Alice\ wins\ the\ match)$.

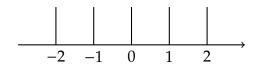
$$P(Alice wins the match) = (0.3) + (0.3)(0.3) + \dots + (0.3)^{9}(0.3)$$
$$= \sum_{k=0}^{9} (0.3)^{k}(0.3)$$

3. Assuming no limit to the number of games played, find $P(Alice\ wins\ the\ match)$ again.

(a)
$$P(Alice\ wins\ the\ match) = \sum_{k=0}^{\infty} (0.3)^k (0.3) = (0.3) \sum_{k=0}^{\infty} (0.3)^k = \frac{0.3}{1-0.7} = \frac{3}{7}$$

• <u>Functions of a RV:</u>

Let
$$P_X(x) = 0.2$$
 $x = -2, -1, 0, 1, 2$.



Information Theory

Defintion 2.0.1 (Entropy).

$$H(X) = \mathbb{E}\left[-\log_2 p(x)\right] = \sum_{x \in X} -p(x)\log_2 p(x)$$

If $X \sim B(p)$, then

$$H(X) = p - \log_2 p + (1 - p) \log_2 (1 - p) \triangleq h(p)$$

called the binary entropy function.

Defintion 2.0.2 (Joint Entropy).

$$H(X,Y) = \mathrm{E}\left[-\log_2 p(x,y)\right] = \sum_x \sum_y -p_{x,y}(x,y)\log_2 p(x,y)$$

If X, Y are independent, then H(X, Y) = H(X) + H(Y).

Proof.

$$H(X,Y) = E \left[-\log_2 p(x,y) \right]$$

$$= E \left[-\log_2 p(x) p(y) \right]$$

$$= E \left[-\log_2 p(x) - \log_2 p(y) \right]$$

$$= E \left[-\log_2 p(x) \right] + E \left[-\log_2 p(y) \right]$$

$$= H(X) + H(Y)$$

Defintion 2.0.3 (Conditional Entropy).

$$H(Y \mid X) = \mathbb{E}_{XY} \left[-\log_2 p(y \mid x) \right] H(X, Y) - H(X)$$

Defintion 2.0.4 (Mutual Information).

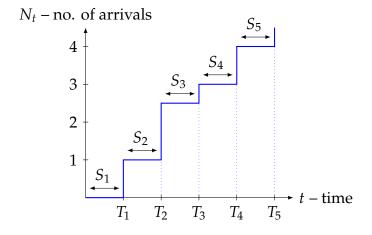
$$I(X;Y) \triangleq H(X) - H(X \mid Y)$$
$$= H(Y) - H(Y \mid X)$$
$$= H(X) + H(Y) - H(X,Y)$$

Theorem 2.0.5 (Asymptotic Equipartition). *If* $X_1, X_2, ..., X_n$ *are iid* $\sim p(X)$, then

$$\frac{-\log_2 p(X_1, X_2, \dots, X_n)}{n} \stackrel{p}{\longrightarrow} H(X).$$

Poisson Proceess

A <u>Poisson process</u> is the continuous time analog of "coin flipping" (or Bernoulli) processes. This makes it a good model for arrival processes: photons hitting a detector, packets in a network, number of emails per hour, etc.



Each T_i for i = 1, 2, 3, 4, 5 represents an arrival and generally, each arrival time is defined as

$$T_n = \sum_{i=1}^n S_i$$

where the interarrival times $S_1, S_2, \ldots, S_n \stackrel{\text{iid}}{\sim} Exponential(\lambda)$. Thus, every S_i has the probability density function

$$f_{S_i}(t) = \lambda e^{\lambda t}; \ t > 0; \quad i = 1, 2, 3, \dots$$

and cumulative distribution function

$$F_{S_i}(t) = 1 - e^{-\lambda t}; \quad i = 1, 2, 3, \dots$$

Defintion 3.0.1 (Number of Arrivals).

$$N_t = \begin{cases} \max_{n \ge 1} \{n \mid T_n \le t\} & t \ge 0 \\ 0 & t < T_1 \end{cases}$$

• Recall: $Exponential(\lambda)$ is a memoryless RV

1.
$$F_{\tau}(t) = \begin{cases} 1 - e^{-\lambda t} & t \ge 0 \\ 0 & t < 0 \end{cases} \Rightarrow f_{\tau}(t) = \lambda e^{-\lambda t}$$

- 2. $E[\tau] = \frac{1}{\lambda}$ and $Var(\tau) = \frac{1}{\lambda^2}$
- 3. $P(\tau > t + s \mid \tau > s) = P(\tau > t)$
- 4. $P(\tau \le t + \epsilon \mid \tau > t) = \lambda \epsilon + o(\epsilon)$; $\lim_{\epsilon \to 0} \frac{o(\epsilon)}{\epsilon} = 0$

Proof.

$$P(\tau > t + \epsilon \mid \tau > t) = P(\tau > \epsilon)$$

$$= e^{-\lambda \epsilon}$$

$$= 1 - \lambda \epsilon + o(\epsilon)$$

$$\therefore P(\tau \le t + \epsilon \mid \tau > t) = 1 - P(\tau > t + \epsilon \mid \tau > t)$$

$$\lambda \epsilon + o(\epsilon)$$

- $P(1 \text{ arrival}) = \lambda \epsilon + o(\epsilon)$
- P(2 arrivals) = $1 \lambda \epsilon + o(\epsilon)$
- P(3 arrivals) = $o(\epsilon)$

We can only have one arrival in a particular subinterval with high probability $\lambda \epsilon + o(\epsilon)$. On the other hand, there are zero arrivals with probability $1 - \lambda \epsilon + o(\epsilon)$.

This means that a Poisson process has <u>independent</u> and <u>stationary</u> increments, i.e., for any $0 \le t_1 \le t_2 \le ... \le t_n \le ...$; if you look at the number of arrivals in t_{n+1} and t_n , $\{N_{t_{n+1}} - N_{t_n}\}$ are independent and the distribution depends only on the size of the interval $(t_{n+1} - t_n)$.

Theorem 3.0.2. Let $N_t := number of arrivals in (0, t), then$

$$P(N_t = k) = e^{-\lambda t} \frac{(\lambda t)^k}{k!}; \quad k = 0, 1, 2, \dots$$

Proof.

$$P(N_t = k) = \int_{t_1} \int_{t_2} \dots \int_{t_k} f_{T_1, T_2, \dots, T_k \mid N_t = k}(t_1, t_2, \dots, t_k \mid N_t = k) dt_1 dt_2 \dots dt_k$$

We constrain the region of integration by letting $S = 0 \le t_1 \le t_2 \le ... \le t_k \le t \le t_{k+1}$. Shifting focus to the k^{th} dimensional pdf,

$$f_{T_1,...,T_k|N_t=k}(t_1,...,t_k \mid N_t=k) = P(T_1 \in (t_1,t_1+dt_1),...,T_k \in (t_k,t_k+dt_k),T_{k+1}>t)$$

where we leverage the fact that $P(T_{k+1} \in (t_{k+1},t)) = P(T_{k+1}>t)$

$$\Rightarrow P(T_{1} \in (t_{1}, t_{1} + dt_{1})) \times ... \times P(T_{k} \in (t_{k}, t_{k} + dt_{k})) \times P(T_{k+1} > t)$$

$$= P(S_{1} \in (t_{1}, t_{1} + dt_{1})) \times ... \times P(S_{k} \in (t_{k} - t_{k-1}, t_{k} - t_{k-1} + dt_{k})) \times P(S_{k+1} > t - t_{k})$$

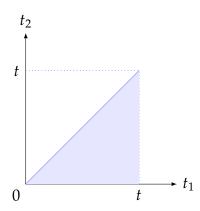
$$= (\lambda e^{-\lambda t_{1}} dt_{1}) ... (\lambda e^{-\lambda (t_{k} - t_{k-1})} dt_{k}) e^{-\lambda (t - t_{k})}$$

$$= \lambda^{k} e^{-\lambda t}$$

The result $\lambda^k e^{-\lambda t}$ is a constant to the k^{th} dimensional integral, so barring any constraints on t_1, t_2, \ldots, t_k , Volume $(S) = t^k$. But, we must respect the fact that $0 \le t_1 \le t_2 \le \ldots \le t_k$. By symmetry, every permutation of (t_1, t_2, \ldots, t_k) has equal volume

Volume
$$(S) = \frac{t^k}{k!} \Rightarrow P(N_t = k) = \frac{(\lambda t)^k e^{-\lambda t}}{k!}; \quad k = 0, 1, 2, \dots$$

- Conditioned on the number of arrivals being k in an interval, the density is uniform
- *Intuition behind symmetry argument:* k = 2



From the image above we know that $P(t_1 \le t_2) + P(t_2 < t_1) = 1$. Thus, $P(t_1 < t_2) = \frac{1}{2} = \frac{1}{2!}$ which generalizes to k dimensions.

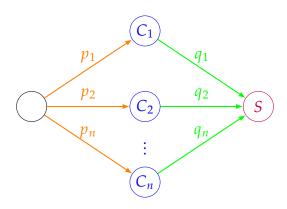
Defintion 3.0.3 (Poisson Merging). *Merging two or more independent Poisson processes yields* $PP(\lambda_1 + \lambda_2 + ... + \lambda_n)$

Defintion 3.0.4 (Poisson Splitting). Let $N \sim PP(\lambda)$ such that N is split into N_1 and N_2 . For each arrival we flip a coin independently with probability H = p. If the coin lands on heads, send the arrival to the N_1 queue otherwise it is sent to the N_2 queue.

Statistical Inference

4.1 Detection & Bayes' Theorem

Consider *N* possible exclusive causes of a particular health symptom.



- Each cause i has a prior probability p_i and it has a probability q_i of causing the observed symptom.
- We want to esimate the posterior probability π_i of cause i given the symptom S,

$$\pi_i = P(C_i \mid S) = \frac{P(C_i \cap S)}{P(S)} = \frac{P(S \mid C_i) P(C_i)}{\sum_j P(S \mid C_j) P(C_j)}.$$

From the diagram above the posterior distribution for cause i can be simplified to

$$\pi_i = \frac{q_i \, p_i}{\sum_j q_j \, p_j}.\tag{4.1}$$

4.2 MAP & MLE

Defintion 4.2.1 (Maximum a posteriori (MAP)).

$$MAP \triangleq \underset{i}{\text{arg max}} \ \pi_i = \underset{i}{\text{arg max}} \ p_i \ q_i$$

Defintion 4.2.2 (Maximum Likelihood Estimation (MLE)).

MLE
$$\triangleq \arg \max_{i} q_i \equiv \text{MAP}$$
 assuming uniform priors $p_i = \frac{1}{N} \forall i$

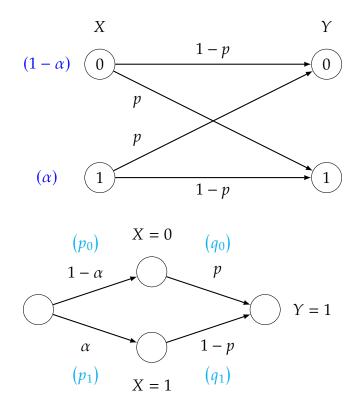
More generally, if X, Y are discrete RVs then

MAP
$$[X \mid Y = y] = \underset{x}{\operatorname{arg max}} P(X = x \mid Y = y)$$

MLE $[X \mid Y = y] = \underset{x}{\operatorname{arg max}} P(Y = y \mid X = x)$

- Called detection because everything is discrete, i.e., detection and classification are synonymous.
- MAP: which cause best explaisn the observed symptom.
- MLE: which cause best generates/induces the observed symptom.

Example 4.2.1 (MAP/MLE Analysis of BSC). *Consider a* BSC (p) *with* $p < \frac{1}{2}$.



$$\Rightarrow \text{MAP}[X \mid Y = 1] = \underset{i \in \{0,1\}}{\text{arg max}} \ p_i q_i.$$

This yields the inequality statement,

$$p_0 q_0 \overset{\hat{X}_1=0}{\leq} p_1 q_1 \overset{\hat{X}_1=1}{\leq} p_1 q_1$$

where \hat{X}_1 is the MAP estimate of X when Y = 1 is 0 or 1. Thus

$$(1-\alpha)p \underset{1}{\overset{0}{\leqslant}} \alpha (1-p)$$