Discrete Distribution Families Binomial(n, p)*binom

successes in n Bernoulli trials.

PMF: $P(X = x) = \binom{n}{x} p^x (1 - p)^{n-x}$

Mean: np, Var: np(1-p)

Geometric(p)*geom

Failures before 1st success.

PMF: $P(X = x) = (1 - p)^x p, x \ge 0$ Mean: (1-p)/p, Var: $(1-p)/p^{\overline{2}}$

NegBin(k, p)*nbinom

Failures before k-th success.

PMF: $P(X = x) = {x+k-1 \choose x} p^k (1-p)^x$ Mean: k(1-p)/p, Var: $k(1-p)/p^2$

 $Poisson(\lambda)*pois$

Counts in interval, rate λ .

PMF: $P(X = x) = \frac{\lambda^x e^{-\lambda}}{x!}$

Continuous Distribution Families

 $\mathbf{Uniform}(a,b)$ - *unif

All values equally likely on [a, b].

PDF: $f(x) = \frac{1}{b-a}, \ a \le x \le b$

Mean: (a+b)/2, Var: $(b-a)^2/12$

For a uniform distribution between a and b,

$$P(X > x) = \frac{b - x}{b - a}, \quad a \le x \le b$$

 $Normal(\mu, \sigma^2)$ - *norm

Bell-shaped, $-\infty < \mu < \infty, \sigma^2 > 0$

PDF: $f(x) = \frac{1}{\sqrt{2\pi\sigma^2}} e^{-(x-\mu)^2/(2\sigma^2)}$

Mean: μ , Var: σ^2

Z score: $\mu = x - Z\sigma$

Lognormal (μ, σ^2) - *lnorm

If $\ln X \sim N(\mu, \sigma^2)$.

PDF: $f(x) = \frac{1}{x\sigma\sqrt{2\pi}}e^{-(\ln x - \mu)^2/(2\sigma^2)}$

Mean: $e^{\mu+\sigma^2/2}$, Var: $(e^{\sigma^2}-1)e^{2\mu+\sigma^2}$ Skewness: $(e^{\sigma^2}+2)\sqrt{e^{\sigma^2}-1}$

Exponential(λ) - *exp

Time between Poisson events, Positive RV, wait time, memoryless

 (λ) is average rate (β) is mean wait time

PDF: $f(x) = \lambda e^{-\lambda x}, x \ge 0$ Mean: $1/\lambda$, Var: $1/\lambda^2$

Skewness: 2

 $Beta(\alpha, \beta)$ - *beta

On [0, 1], Bayesian priors. Uniform distribution special case

 (Γ) function is a generalization of factorial function to non-integer numbers.

PDF: $f(x) = \frac{1}{B(\alpha,\beta)} x^{\alpha-1} (1-x)^{\beta-1}$

Mean: $\alpha/(\alpha+\beta)$, Var: $\frac{\alpha\beta}{(\alpha+\beta)^2(\alpha+\beta+1)}$

Skewness: $\frac{2(\beta-\alpha)\sqrt{\alpha+\beta+1}}{(\alpha+\beta+2)\sqrt{\alpha\beta}}$

Weibull (k, λ) - *weibul

Lifetimes, survival, reliability.

Exponential family (when k = 1), longer you wait Survival Analysis

PDF: $f(x) = \frac{k}{\lambda} (x/\lambda)^{k-1} e^{-(x/\lambda)^k}$

Mean: $\lambda\Gamma(1+1/k)$, Var: $\lambda^2[\Gamma(1+2/k) - \Gamma(1+1/k)^2]$ Skewness: $\frac{\Gamma(1+3/k)\lambda^3 - 3\mu\sigma^2 - \mu^3}{\sigma^3}$

Continuous RV (numerical integration)

 $f_X(x)$ can take on values larger than 1

 $f_X(x)$ cannot give probability, area under the graph does

the density of $f_X(x)$ cannot take negative values even if total area is 1 because probability cannot be negative the support of $f_X(x)$ must be finite $f \leftarrow function(x) 2*x$

EV <- integrate(function(x) x*f(x), 0,1)\$value

Define the PDF function

 $f \leftarrow function(x) 2*x$

prob <- integrate(f, lower = 0.5, upper =</pre> 0.75) \$value # Integrate over [0.5, 0.75] **Conditional Distributions**

Conditional dist is just a segment of marginal dist, then re-normalized to have an area under the curve equal to 1 If X and Y are independent, in continouse case,

$$f_{Y|X}(y) = f_Y(y)$$

This means conditional PDF of Y and X is marginal PDF of Y.

Random Sample

It is independent and identically distributed (iid). Each pair of observations are independent, and each observation comes from the same distribution. MLE

Great way to find estimators. Applied on multi and univariate. Relies on random sample of n observations. Mean, is a estimator for univariate, multivariate, linear regression Steps for MLE

1.(discrete or contin) distribution

2. Find parameters of a theoretical distribution (eg λ in a Poisson distribution)

3.Choose distribution)

4. Play with the parameters for that family of distributions to find the most likely parameters for the corresponding parametric

5. To get estimates, we use the likelihood

function of our observed random sample.)

```
R: Generating Sample and Likelihood
#Generate 30 random samples from Exponential(beta=
set.seed(123)
sample_n30 <- tibble(values = rexp(30, rate = 1/20)</pre>
#Compute likelihood and log-likelihood for candida
exp_values <- tibble(</pre>
  possible_betas = seq(5, 50, 0.5),
  likelihood = map_dbl(1 / possible_betas,
    prod(dexp(sample_n30$values, .))),
  log_likelihood = map_dbl(1 / possible_betas,
      log(prod(dexp(sample_n30$values, .))))
empirical_MLE <- exp_values |>
  arrange(desc(likelihood)) |>
  slice(1)
analytical_MLE <- mean(sample_n30$values)</pre>
#We use the sample mean() function from R!
round(analytical_MLE, 2)
LL <- function(1) log(prod(dexp(sample_n30$values,
rate = 1 / 1)))
optimize(LL, c(5, 50), maximum = TRUE)
stochastic - having some uncertain outcome.
deterministic - an outcome that will be known
with 100% certainty.
Computers cannot actually generate truly
random outcomes.
Instead, they use something called
pseudorandom numbers.
```

The empirical approach (using observed data), resulting in approximate values that improve as the sample size increases (i.e., the frequentist paradigm!).

One pitfall is that neighbouring pairs are not

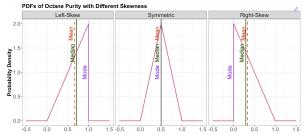
Although this sequence is deterministic,

it behaves like a random sample.

independent of each other.

The Law of Large of Numbers states that, as we increase our sample size n, our empirical mean converges to the true mean we want to estimate.

Skewness



Left-skewed: long tail to the left, mean < Right-skewed: long tail to the right, mean > median < mode median > mode