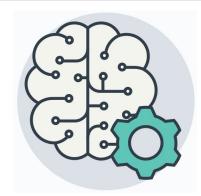
Aprendizado de Máquina

Medidas Estatísticas, Distribuições e Escalonamento de Features



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Medidas estatísticas

$$mean(x) = \frac{\sum_{i=1} x_i}{count(x)}$$

$$variance = \sum_{i=1}^{n} (x_i - mean(x))^2$$

$$covariance = \sum_{i=1}^{n} ((x_i - mean(x)) \times (y_i - mean(y)))$$

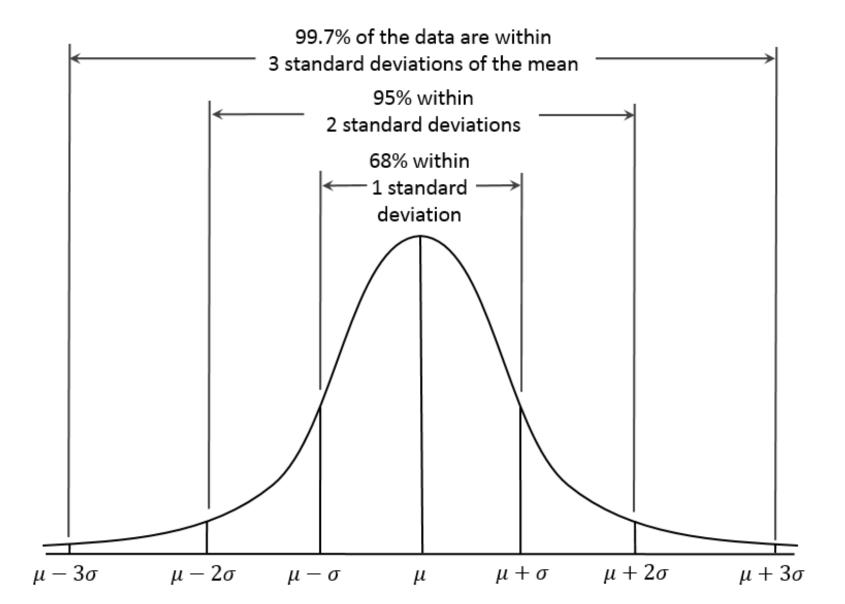
```
# Example of Estimating Mean and Variance
# Calculate the mean value of a list of numbers
def mean(values):
  return sum(values) / float(len(values))
# Calculate the variance of a list of numbers
def variance(values, mean):
  return sum([(x-mean)**2 for x in values])
# calculate mean and variance
dataset = [[1, 1], [2, 3], [4, 3], [3, 2], [5, 5]]
x = [row[0] for row in dataset]
y = [row[1] for row in dataset]
mean x, mean y = mean(x), mean(y)
var x, var y = variance(x, mean x), variance(y,
mean y)
print('x stats: mean=%.3f variance=%.3f' % (mean x,
var x))
print('y stats: mean=%.3f variance=%.3f' % (mean_y,
var vll
```

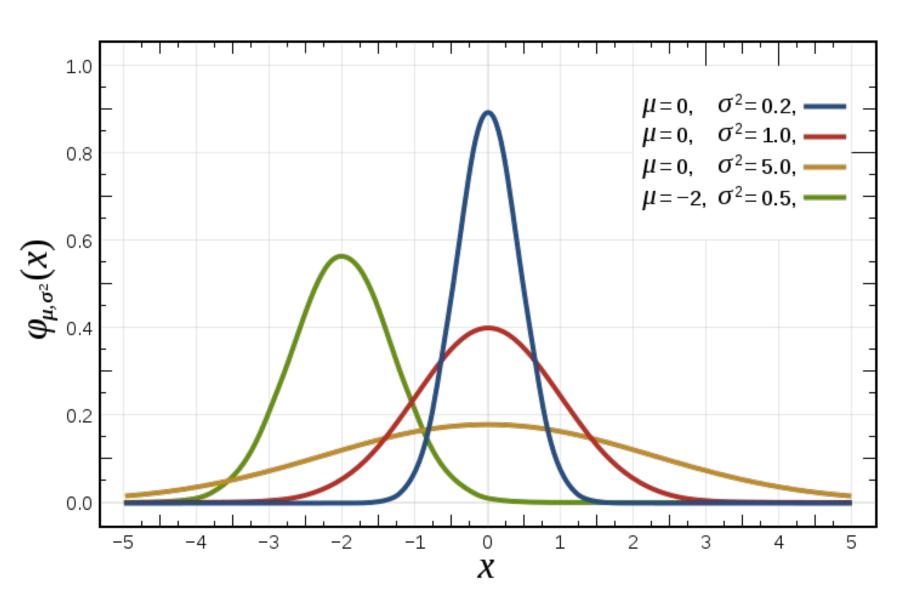
```
# Calculate covariance between x and y
def covariance(x, mean_x, y, mean_y):
    covar = 0.0
    for i in range(len(x)):
        covar += (x[i] - mean_x) * (y[i] -
        mean_y)
    return covar
```

x stats: mean=3.000 variance=10.000

y stats: mean=2.800 variance=8.800

Covariance: 8.000

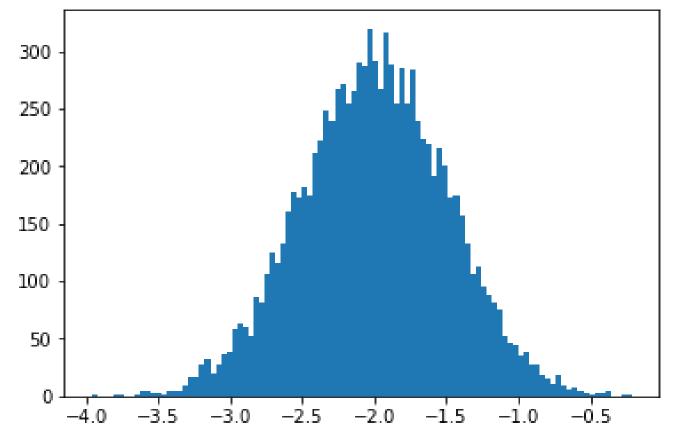




```
np.random.normal(size=10000)
plt.hist(s, bins=100);
300
250
200
150
100
50
```

```
np.random.normal(size=10000)
plt.hist(s, bins=100);
300
250
200
150
100
50
```

```
s = np.random.normal(loc=-2, scale=0.5, size=10000)
plt.hist(s, bins=100);
```



loc : Mean ("centre") of the distribution.

scale: Standard deviation (spread or "width") of the distribution.

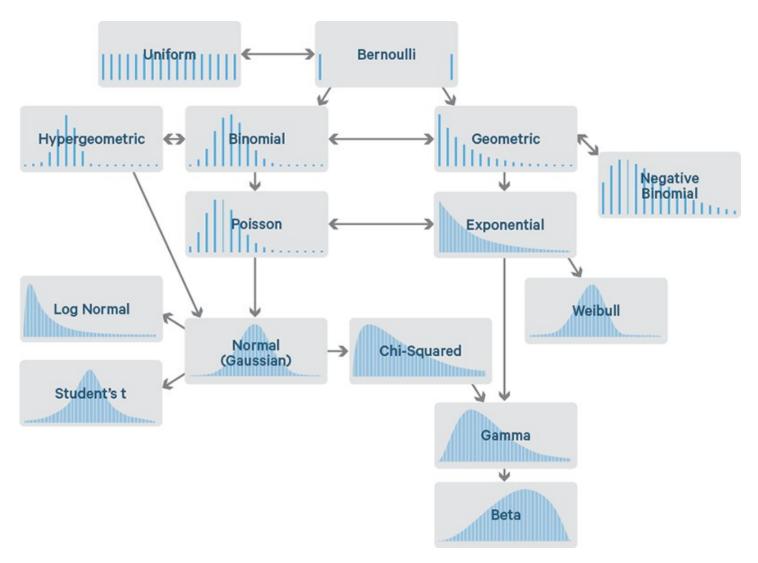
Distribuições

```
beta (a, b[, size])
binomial (n, p[, size])
chisquare (df[, size])
dirichlet (alpha[, size])
exponential ([scale, size])
f (dfnum, dfden[, size])
gamma (shape[, scale, size])
geometric (p[, size])
qumbel ([loc, scale, size])
hypergeometric (ngood, nbad, nsample[, size])
laplace ([loc, scale, size])
logistic ([loc, scale, size])
lognormal ([mean, sigma, size])
logseries (p[, size])
multinomial (n, pvals[, size])
multivariate_normal (mean, cov[, size, ...)
negative_binomial (n, p[, size])
noncentral_chisquare (df, nonc[, size])
noncentral_f (dfnum, dfden, nonc[, size])
normal ([loc, scale, size])
```

numpy.random

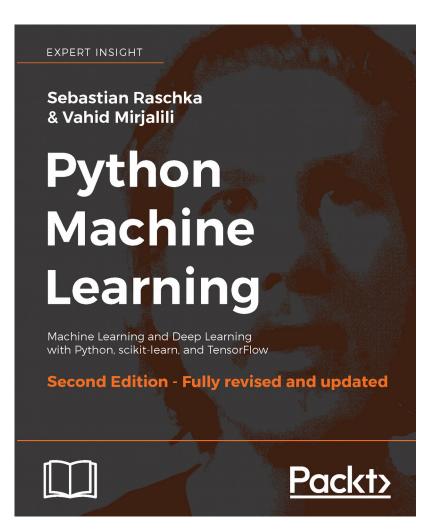
```
pareto (a[, size])
poisson ([lam, size])
power (a[, size])
rayleigh ([scale, size])
standard_cauchy ([size])
standard_exponential ([size])
standard_gamma (shape[, size])
standard_normal ([size])
standard_t (df[, size])
triangular (left, mode, right[, size])
uniform ([low, high, size])
vonmises (mu, kappa[, size])
wald (mean, scale[, size])
weibull (a[, size])
zipf (a[, size])
```

Distribuições



http://blog.cloudera.com/blog/2015/12/common-probability-distributions-the-data-scientists-crib-sheet/

Main Reference



Python Machine Learning

Chapter 4 - Building Good Training Sets – Data Preprocessing

Bringing features onto the same scale

Feature Scaling

$$x_{std}^{(i)} = \frac{x^{(i)} - \mu_x}{\sigma_x}$$

$$x_{norm}^{(i)} = \frac{x^{(i)} - \mathbf{x}_{min}}{\mathbf{x}_{max} - \mathbf{x}_{min}}$$

standardizatio n min-max scaling ("normalization")

	input	standardized	normalized
0	0	-1.46385	0.0
1	1	-0.87831	0.2
2	2	-0.29277	0.4
3	3	0.29277	0.6
4	4	0.87831	0.8
5	5	1.46385	1.0

Standardization

$$\text{standardized_value}_i = \frac{\sum_{i=1}^{n} (value_i - mean)}{stdev}$$

- Standardization is a rescaling technique that refers to centering the distribution of the data on the value 0 and the standard deviation to the value 1.
- The mean and the standard deviation summarize a normal distribution.
- Standardization is a scaling technique that assumes your data conforms to a normal distribution.
- If a given data attribute is normal or close to normal, this is probably the scaling method to use.

Normalization

scaled value =
$$\frac{value - min}{max - min}$$

- Normalization can refer to different techniques depending on context.
- Here, we use normalization to refer to rescaling an input variable to the range between 0 and 1.
- Normalization is a scaling technique that does not assume any specific distribution.
- If your data is not normally distributed, consider normalizing it prior to applying your machine learning algorithm.

Normalization

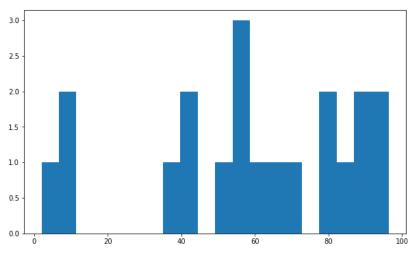
```
np.random.seed(0)
x = np.random.rand(20)
x = (x * 100).round(2)
x = np.resize(x, (20,
1))
```

```
[[ 54.88]
  71.52]
  60.281
  54.491
  42.371
  64.591
  43.761
  89.181
  96.371
  38.341
  79.171
  52.891
  56.8 1
  92.561
   7.1 1
   8.71
   2.021
  83.26]
  77.821
 [ 87. ]]
```

Normalization

```
x norm = normalize(x)
```

plt.hist(x, bins=20)



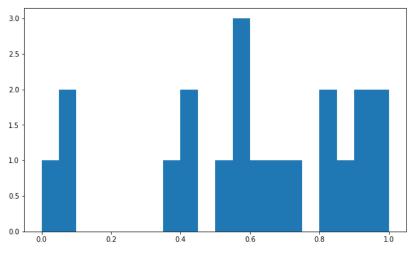
```
x

mean: 58.16,

std: 27.59,

min: 2.02,

max: 96.37
```



```
x_norm
---
mean: 0.59,
std: 0.29,
min: 0.0,
max: 1.0
```

```
0.560254371,
 0.73661897],
 0.61748808],
[ 0.55612083],
[ 0.42766296],
 0.663169051,
 0.44239534],
 0.92379438],
 0.38494966],
 0.817700051,
 0.539162691,
 0.58060413],
 0.959618441,
 0.053842081,
 0.0709062 1.
 0.
 0.861049281,
 0.80339163],
[ 0.90068892]]
```

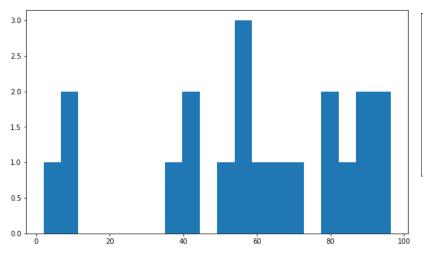
Standardization

```
x_{std}^{(i)} = \frac{x^{(i)} - \mu_x}{\sigma_x}
```

Standardization

```
x \text{ std} = \text{standardize}(x)
```

plt.hist(x, bins=20)



x mean: 58.16, std: 27.59, min: 2.02,

max: 96.37

```
3.0 - 2.5 - 2.0 -1.5 -1.0 -0.5 0.0 0.5 10 15
```

```
x_std
---
mean: 0.0,
std: 1.0,
min: -2.03,
max: 1.38
```

```
[[-0.11870903],
  0.48434953],
  0.076995071,
[-0.13284322],
 [-0.5720902]
  0.233195931,
 [-0.52171451],
  1.12437442],
  1.384950811,
 [-0.71814345],
  0.761597 1.
 [-0.19082962],
 [-0.04912535],
  1.246870691,
 [-1.85032791],
 [-1.79197909],
 [-2.03443473],
  0.909824741,
  0.71267098],
 [ 1.04536795]]
```

