Bioinformatics and Biostatistics BB2440: Biostatistics Lecture 1 :Fundamental Probability Timo Koski

ΤK

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Many people associate the word statistics ¹ with diagrams, tables and charts. Here we mean by *statistics* the study of how one draws conclusions from observed data and/or draws correct conclusions under uncertainty and describes these conclusions appropriately.



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¹Florence Nightingale invented a form of the pie chart in order to illustrate seasons sources of patient mortality in the military field hospital sheamanaged.

From Data to Knowledge

Data
ightarrow Statistics
ightarrow Information/Knowledge

Statistics is a tool for creating new understanding from a set of numbers





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 Absolute and Relative Frequencies, Histogram, Mean, Variance, Standard Deviation





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- Absolute and Relative Frequencies, Histogram, Mean, Variance, Standard Deviation
- Boxplot





- Absolute and Relative Frequencies, Histogram, Mean, Variance, Standard Deviation
- Boxplot
- Empirical Probability, Theoretical Probability, Meaning of Probability





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- Independent Events





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- Absolute and Relative Frequencies, Histogram, Mean, Variance, Standard Deviation
- Boxplot
- Empirical Probability, Theoretical Probability, Meaning of Probability
- Rules of Probability
- Independent Events
- Binomial Probability Distribution





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Statistics

We check 35 boxes of matches and check in each case how many matches it contains.

Obviously this might be of importance for a producer of matches in industrial scale, but you might think that this is of no interest for you (counting matches in box is of no special hobby of mine either). This is, however, not the point. This is a simple example of observed data, quite clear for everyone, and to introduce the concepts of probability.







Observed Data

We obtain 35 boxes of matches and check in each case how many matches it contains. Result:





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Frequencies

The table above can be summarized by the table of frequencies

- absolute frequency f_i for the observed values
- relative frequency $p_i = f_i/n$. (i = 1 lowest class, i = 2 next class e.t.c..) <math>n = number of data (=35).





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Table of frequencies

Table: Frequencies for the observed numbers of matches.

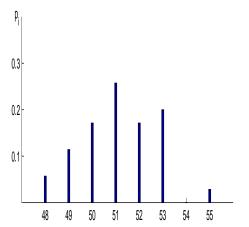
Class	Absolute frequency	Relative frequency (%)	
i	f_i	100 p _i	
48	2	5.7	
49	4	11.4	
50	6	17.1	
51	9	25.7	
52	6	17.1	
53	7	20.0	
54	0	0.0	
55	1	2.9	
Sum	35	100.0	





Histogram

The data becomes easier to grasp if we plot the relative frequencies p_i in *histogram*.

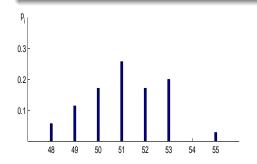






Histogram for relative frequencies

A histogram is a bar graph in which the horizontal scale represents classes of data values and the vertical scale represents the relative frequencies. The heights of the bars correspond to the relative frequency values.







A Measure of Central Tendency: The Mean

Let x_1, \ldots, x_n denote the data to be analysed. As a measure of central tendency one often uses the arithmetic mean

$$\bar{x} = \frac{1}{n} \sum_{i=1}^{n} x_i = \frac{x_1 + x_2 + \dots + x_n}{n}.$$

In the matchbox example we find $\sum_{i=1}^{35} x_i = 1789$ and $\bar{x} = 1789/35 = 51.1143$. This gives us the typical value of the data.





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Measure of dispersion: Variance, Standard deviation

A measure of dispersion refers to how closely the data cluster around the measure of central tendency. Here we consider the variance

$$s^{2} = \frac{1}{n} \sum_{j=1}^{n} (x_{j} - \bar{x})^{2}$$

standard deviation

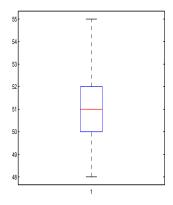
$$s = \sqrt{\frac{1}{n} \sum_{j=1}^{n} (x_j - \bar{x})^2},$$

We take the square root to get back to the original units. In matchbox example we get the standard deviation $s \approx 1.62$.



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BOXPLOT for the matchbox data



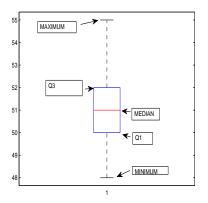
51	52	49	51	52	51	53
52	48	52	50	53	49	50
51	53	51	52	50	51	53
53	55	50	49	53	50	51
51	52	48	53	50	49	51





BOXPLOT a.k.a box-and-whisker diagram

A **boxplot** is a graph of a data set that consists of a line from the minimum value to the maximum value and a box with lines drawn at the first quartile Q_1 , the median and the third quartile Q_3

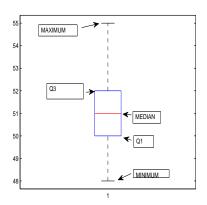






BOXPLOT

first quartile Q_1 = separates the bottom 25% of the sorted values, the median = midpoint, 50 % below, the third quartile Q_3 = separates the bottom 75% of the sorted values







New York Times 2009: Big Data

statistics ... uses powerful computers and sophisticated mathematical models to hunt for meaningful patterns and insights in vast troves of data. The applications are as diverse as improving Internet search and online advertising, culling gene sequencing information for cancer research and analyzing sensor and location data to optimize the handling of food shipments.

 $mathematical \ model \leftrightarrow probability$

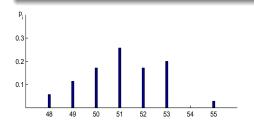




Probability: What is it?

The empirical answer: Let us think of the histogram of the matchbox data again. We assume that nothing changes (in the industrial production of matchboxes).

Then the **probability of the event** " $50 \le$ the number of matches in a box ≤ 53 " is taken as the relative frequency of " $50 \le$ the number of matches in a box ≤ 53 ".



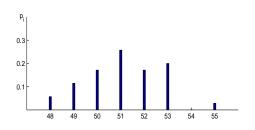




Probability: What is it?

The empirical answer: Let us think of the histogram of the matchbox data again. We assume that nothing changes (in the industrial production of matchboxes). Then the **probability of the event** " $50 \le$ the number of matches in a box ≤ 53 " is

$$p_{50} + p_{51} + p_{52} + p_{53} = 0.171 + 0.257 + 0.171 + 0.20 \approx 0.80$$







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Probability

We write now the probability of **the event** " $50 \le$ the number of matches in a box <53" as

$$Pr(50 \le \text{the number of matches in a box} \le 53) = 0.80$$

Then, of course, the probability of a simple event like for example " the number of matches in a box = 53" is its relative frequency

$$Pr($$
 the number of matches in a box $= 53) = p_{53} = 0.20$





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Probability

The probability of **the event** "50 \leq the number of matches in a box \leq 53 "

$$Pr(50 \le \text{the number of matches in a box } \le 53) = 0.80$$

This is a probability based on past observations. This means, when we assume that nothing changes, that if we receive a new box of mathces and check the number of matches in it, there is 80% chance of the event "50 \leq the number of matches in a box \leq 53" occurring.



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Probability: empirical

The probability

$$Pr(50 \le \text{the number of matches in a box} \le 53) = 0.80$$

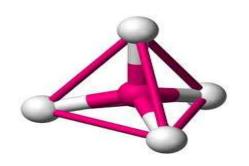
was based on checking 35 boxes. The relative frequencies will change if we observe new boxes, but they will stabilize.





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Probability







To each of the four sides of a tetrahedron there is assigned one of the letters A,T,C,G (nucleotides). All letters are used. We call this the DNA dice. We toss the dice in the air and note the side that it falls on.







We toss the dice in the air and note the side that side that it falls on. Clearly we can thus produce a **random** DNA sequence. Such a sequence would seem to lack value for bioinformatics. Disregarding that for the moment, we might ask what is the probability of observing the sequence CAAGT in five tosses of the dice, or what is the probability of the event "CAAGT in five tosses".







What is the probability of the event "CAAGT in five tosses"? All the sides of the tetrahedron are of equal area, and the tetrahedron is balanced. The situation is such that we assign the probabilities

$$Pr(A) = Pr(T) = Pr(C) = Pr(G) = \frac{1}{4}$$

to the four possible outcomes of a single toss. In other words, all outcomes of the toss of the dice are equally likely. Note that we have here a very crude model of a DNA sequence and a way to compute the probability of such a sequence.





In addition, we think that the individual tosses do not influence each other. This means that

$$Pr(A) = Pr(T) = Pr(C) = Pr(G) = \frac{1}{4}$$

are the probabilities of the letters at any toss. Hence we get that

$$Pr(CAAGT) = Pr(C) Pr(A) Pr(A) Pr(G) Pr(T)$$
$$= \frac{1}{4} \cdot \frac{1}{4} \cdot \frac{1}{4} \cdot \frac{1}{4} \cdot \frac{1}{4} = \left(\frac{1}{4}\right)^{5}.$$

In probability terms one says that the outcomes at the different tosses are **independent events**.



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The probability

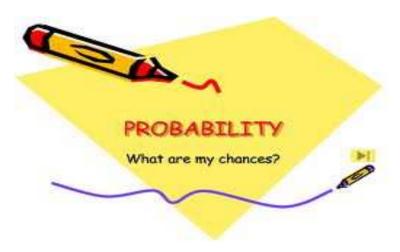
$$Pr(CAAGT) = \frac{1}{4} \cdot \frac{1}{4} \cdot \frac{1}{4} \cdot \frac{1}{4} \cdot \frac{1}{4} = \left(\frac{1}{4}\right)^{5}.$$

was obtained theoretically by reasoning about the symmetry of the tetrahedron and independence of the tosses. Of course, one may toss any given DNA dice, say 10^{10} times, and register the relative frequencies of the letters A,T,C,G. Then one can check whether the theoretical probabilities and the empirical probabilities seem to agree.





Probability





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Probability: a formal statement

We know continue by giving a general formal description and some rules of computation for Pr(A), the probability of an event.

Pr denotes probability

A,B, and C denote specific events.

Pr(A) denotes the probability of event A occurring





Probability: a formal statement

The probability of an event A, Pr(A), has the following properties:

(a)
$$0 \le Pr(A) \le 1$$
;





Probability: a formal statement

The probability of an event A, Pr(A), has the following properties:

- (a) $0 \le Pr(A) \le 1$;
- (b) Pr(S) = 1, if S is an event that is certain to happen.





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Probability: a formal statement

The probability of an event A, Pr(A), has the following properties:

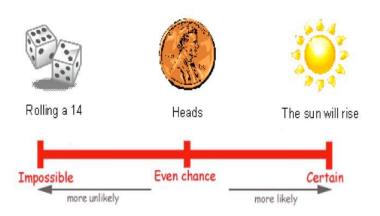
- (a) $0 \le Pr(A) \le 1$;
- (b) Pr(S) = 1, if S is an event that is certain to happen.
 - For example, when tossing the DNA dice once, the event $S = \{A, T, C, G\}$ (= one of the letters A,T,C,G) is certain to happen. (Here we may call A, T, C, G simple events, because they cannot be broken down any further).





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Probability

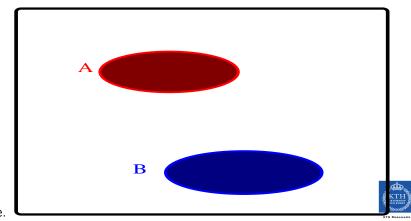






Mutually exclusive events

We say that two events A and B are **mutually exclusive**, if the occurrence of one precludes the occurrence of the other. The diagram below illustrates this. For DNA dice, the events $\{A, T\}$, $\{C\}$ are mutually



exclusive.

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A or B

The event A or B means that A occurs or both A and B occur or B occurs.





Probability: a formal statement of the additive law

The event A or B means that A occurs or both A and B occur or B occurs.

(c) if A and B are mutually exclusive events then we have the **additive** law

$$Pr(A \text{ or } B) = Pr(A) + Pr(B).$$





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Example

The DNA dice again, the events $A_1 = \{A, T\}$, $A_2 = \{C\}$ are mutually exclusive. Also the events $\{A\}$ and $\{T\}$ are mutually exclusive. Thus the additive law together with the theoretical probability model from the above gives

$$Pr(A_1) = Pr(\{A, T\}) = Pr(\{A\} \text{ or } \{T\}) = Pr(\{A\}) + Pr(\{T\})$$

$$= \frac{1}{4} + \frac{1}{4} = \frac{1}{2}$$

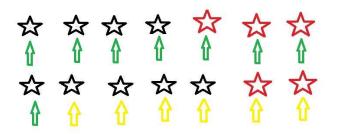
The additive law gives also

$$Pr(A_1 \text{ or } A_2) = Pr(A_1) + Pr(A_2) = \frac{1}{2} + \frac{1}{4} = \frac{3}{4}$$



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Probabilities play a very big role in genetics. In the figure we have 14 (abstract) peas: nine have white flowers, five have red flowers, six have yellow pods, eight have green pods.

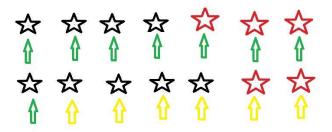






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What is Pr(green pods or white flowers)?







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• green pods or white flowers means green pods or white flowers or both



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- green pods or white flowers means green pods or white flowers or both
- it is wrong to add to the 8 peas with green pods the 9 peas with white flowers, because then you have counted 5 of the peas twice.







- green pods or white flowers means green pods or white flowers or both
- it is wrong to add to the 8 peas with green pods the 9 peas with white flowers, because then you have counted 5 of the peas twice.
- $Pr(\text{green pods or white flowers }) = \frac{12}{14} = \frac{6}{7}$





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The General Rule

(d)

$$Pr(A \text{ or } B) = Pr(A) + Pr(B) - Pr(A \text{ and } B)$$

In the example above

Pr(green pods or white flowers)

$$= Pr(\text{green pods }) + Pr(\text{ white flowers }) - Pr(\text{green pods and white flowers })$$

$$=\frac{8}{14}+\frac{9}{14}-\frac{5}{14}=\frac{12}{14}=\frac{6}{7}$$





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Complementary event A*

"A does not occur" = A^* .

Example

$$S = \{A, T, C, G\}$$

$$A = \{A, T\}. A^* = \{C, G\}.$$





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Probability of the Complementary event A^*

$$Pr(A^*) = 1 - Pr(A).$$





Probability of the Complementary event A*

$$Pr(A^*) = 1 - Pr(A).$$

The DNA dice again, the event $A = \{A, T, C\}, A^* = \{G\}$ we have by the additive law

$$Pr(A) = Pr(\{A\}) + Pr(\{T\}) + Pr(\{C\}) = \frac{3}{4}$$

and

$$Pr(A^*) = 1 - Pr(A) = \frac{1}{4}$$

but this agrees with what we have by our model for the dice, namely that $Pr(\{G\}) = \frac{1}{4}$.



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Probability: independent events

We say that A and B are **independent** events if

$$Pr(A \text{ and } B) = Pr(A) \cdot Pr(B).$$



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Probability

So far our examples been dealing with probability empirically as a relative frequency or by the rule 'simple events have an equal chance of occurring' (e.g., $Pr(A) = Pr(T) = Pr(C) = Pr(G) = \frac{1}{4}$). Next we check probabilities of different sort.





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Suppose you take a multiple Choice test with 10 questions, and each question has 5 answer choices (a,b,c,d,e), what is the probability you get exactly 4 questions correct just by guessing? $P(Success) = \frac{1}{5}$ $P(x) = {n \choose x} p^x e^{n-x}$ $P(Failure) = \frac{4}{5}$



《四》《圖》《意》《意》 [2]

Probability: series of independent events

We discuss finally the **binomial distribution** which shows the probabilities of different outcomes for a series of independent random events, each of which can have only one of two values, e.g., success and failure. Usually one presents here the toss of a coin, with outcomes heads and tails. I shall talk about tosses of a thumbtack.

Fair coin: $Pr(\text{heads}) = Pr(\text{tails}) = \frac{1}{2}$





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Tossing of a Thumbtack



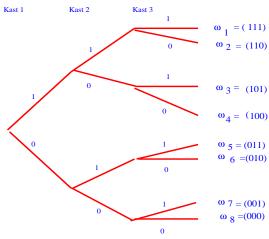
Toss a thumbtack. If it falls on

its point as in the picture above, we say that the event a digital 'one' (1), occurs. We say that att the event a digital zero (0) occurs if the thumbtack lands on its head.





Three tosses of a thumbtack

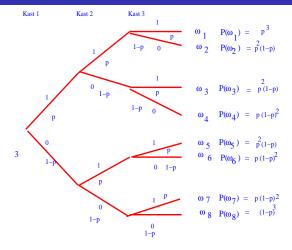


Number of possible outcomes=



 2^3

Three tosses of a thumbtack: probabilities by independence

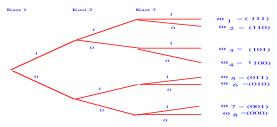


$$Pr(\text{ zero }(0)) = 1 - p, Pr(\text{ one }(1)) = p, 0$$



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Three tosses of a thumbtack: probability of the number of ones



We consider the probabilities of the number of ones $\omega_1 \to 3$, ω_2 , ω_3 , $\omega_5 \to 2$ ω_4 , ω_6 , $\omega_7 \to 1$, $\omega_8 \to 0$.





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Three tosses of a thumbtack: probability of the number of ones

$$Pr(3) = Pr(\omega_{1}) = p^{3},$$

$$Pr(2) = Pr(\omega_{2}) + Pr(\omega_{3}) + Pr(\omega_{5}) = 3p^{2}(1 - p)$$

$$Pr(1) = Pr(\omega_{4}) + Pr(\omega_{6}) + Pr(\omega_{7}) = 3p(1 - p)^{2}$$

$$Pr(0) = Pr(\omega_{8}) = (1 - p)^{3}$$





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Three tosses of a thumbtack: probability of the number of ones

We rewrite using the binomial coefficients:

$$Pr(3) = p^{3} = {3 \choose 3} p^{3} (1 - p)^{0}$$

$$Pr(2) = 3p^{2} (1 - p) = {3 \choose 2} p^{2} (1 - p)$$

$$Pr(1) = 3p(1 - p)^{2} = {3 \choose 1} p(1 - p)^{2}$$

$$Pr(0) = {3 \choose 0} (1 - p)^{3}$$

or with a single formula

$$Pr(k) = {3 \choose k} p^k (1-p)^{3-k}, \quad , k = 0, 1, 2, 3.$$



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Binomial coefficients

$$\binom{n}{k} = \frac{n!}{k!(n-k)!}$$

reads as 'n choose k',

$$n! = n \cdot (n-1) \dots 2 \cdot 1$$
 n factorial





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Binomial distribution

Definition

$$Pr(k) = \binom{n}{k} p^k (1-p)^{n-k}$$
, for $k = 0, 1, ..., n$.

This is called the binomial probability distribution with parameters n and p. The **mean** of this distribution is np and the **variance** is np(1-p).





Binomial distribution

The binomial distribution shows the probabilities of different outcomes for a series of random events, each of which can have only one of two values.

Definition

$$Pr(k) = \binom{n}{k} p^k (1-p)^{n-k}$$
, for $k = 0, 1, ..., n$.







BINOMIAL PROBABILITY FUNCTION / BINOMIAL DISTRIBUTION

Requirements:

- 1) a repeated identical independent trials.
- 2) Two outcomes (success/facture)
- 3) P(Success) = p, P(Failure) = q, p+q=1

Then
$$P(x)$$
, the probability that there will
be exactly x successes in n trials is: $\frac{n!}{p(x)} = \binom{n}{x} p^x q^{-x}$; $\binom{n}{x} = nCr \frac{(n-r)!r!}{(n-r)!r!}$



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Alignment of Sequences & series of random events

We wish to compare two sequences ${\bf x}$ and ${\bf y}$ with 15 nucleotides in each.

$$\mathbf{x} = \begin{matrix} \mathbf{G} \stackrel{\downarrow}{A} & T \stackrel{\downarrow}{A} & A & G \stackrel{\downarrow}{C} & C \stackrel{\downarrow}{C} & C \stackrel{\downarrow}{C} & T \stackrel{\downarrow}{G} & T \stackrel{\downarrow}{C} & T \\ C & A & A & A & T & C & C & C & A & G & T & C & T \end{matrix}$$

We say that we have a **match**, if the paired nucleotides are the same in both sequences. We have eleven matches indicated by \downarrow .





Alignment of Sequences & Binomial Distribution

$$\mathbf{x} = \begin{matrix} G \stackrel{\downarrow}{A} T \stackrel{\downarrow}{A} \stackrel{\downarrow}{A} G \stackrel{\downarrow}{C} \stackrel{\downarrow}{$$

If we are willing to assume that the nucleotides are random (DNA dice!) and independent, and that the probabilities are the same at each site, then the probabilities of the number of matches (=successes) are a binomial distribution with parameters n=15, $p=\frac{1}{4}$. Then we can compute how probable or likely it is to get 11 matches in a sequence of 15 nucleotides. This is a case of the question of significance in sequence alignment.





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Rare Event Rule

If, under a given assumption, the probability of an observed event is extremely small (\approx 0), we conclude that the assumption is likely not correct.

In the example with alignment of sequences, the probability of 11 matches in two sequences of 15 nucleotides under the assumption of independent tosses of the DNA dice is 1.0297e-04. (A computation done using a Matlab function for the binomial probability with n=15, $p=\frac{1}{4}$, k=11).

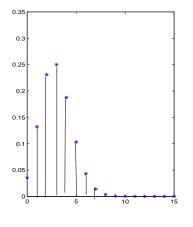
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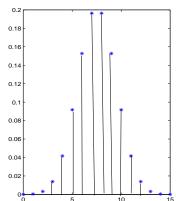




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Binomial distributions with n = 15 and p = 0.2, p = 0.5







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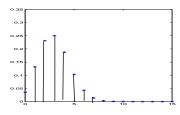


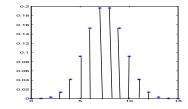
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Binomial distributions with n = 15 and p = 0.2, p = 0.5

For p = 0.2 the distribution is skewed, mean is $15 \cdot 0.2 = 3$, with variance $15 \cdot 0.2 \cdot 0.8 = 2.4$. For p = 0.5 the distribution is symmetric around its mean $15 \cdot 0.5 = 7.5$, with variance $15 \cdot 0.5 \cdot 0.5 = 3.75$.









Binomial distribution

Note that

$$\sum_{k=0}^{n} \Pr(k) = \sum_{k=0}^{n} \binom{n}{k} p^{k} (1-p)^{n-k} = (p + (1-p))^{n} = 1$$

by the binomial formula

$$(a+b)^n = \sum_{k=0}^n \binom{n}{k} a^k b^{n-k}$$





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Thank You!



"How do you want it—the crystal mumbo-jumbo or statistical probability?"

