

# Universal Law of G on

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- Let's say I eat this leaf, and I feel sick.

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- What other leaves might I avoid?

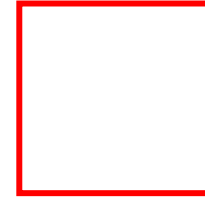


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- What other leaves might I avoid?

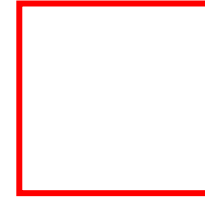


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- There is a “consequential region” of different possible objects that satisfy the property of being sickness-inducing leaves.
- I have generalized the property to other leaves.



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- The region basically tells you how generalizable a property is
- A bigger region will mean there are more objects that share that property

# Property: being able to reach things on a high shelf

The consequential region (spread of objects that have this property) might be easy to estimate here



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The celebrities would be considered objects that may/may not be in the region.

# Property: being able to reach things on a high shelf

The consequential region (spread of objects that have this property) might be easy to estimate here

But sometimes it might not be obvious.

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The celebrities would be considered objects that may/may not be in the region.



# Which animals have the T-9 hormone?



- Let's say you know canaries have T-9 hormones.

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# Which animals have the T-9 hormone?



- Let's say you know canaries have T-9 hormones.
- What other animals might have them?

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# Which animals have the T-9 hormone?

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# Which animals have the T-9 hormone?

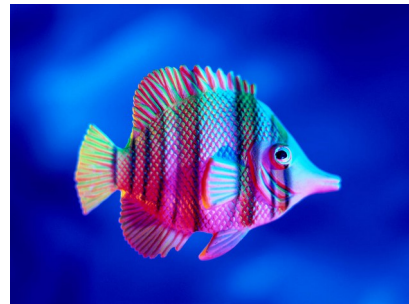
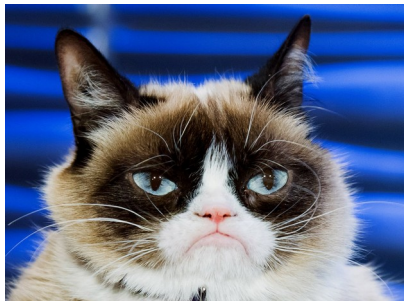


- Let's say you know canaries have T-9 hormones.
- What other animals might have them? It's not obvious which objects (animals) are in the region R (which has the property "T-9 hormone"). The spread of the collection of the number of objects/animals that share the property, is not obvious.

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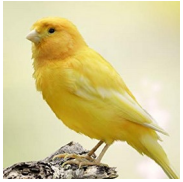
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2. [10pts, HELP] What is the probability of getting  $x=1$  for regions containing  $x=0$ ?

You can think of it as:



- A canary ( $x=0$ ) has the T-9 hormone
- The canary is in a region  $R$  where all objects in  $R$  share the property of having a T-9 hormone
- You don't know the exact region  $R$
- But there are different regions,  $R$ , and say the canary ( $x=0$ ) is in all of them.

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- What's the probability that a robin ( $x=1$ ) has a T-9 hormone?
- What's the probability that a robin is in the  $R$  of objects that have the property of having a T-9 hormone (the region to which canary,  $x=0$ , belongs)?

0

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X axis: different  
objects that are  
arranged by just one  
feature.

Y axis: probability  $y$  is  
in a region given that  
 $x$  is in a region

A canary has a T-9 hormone ( $X=0$  is in the region)

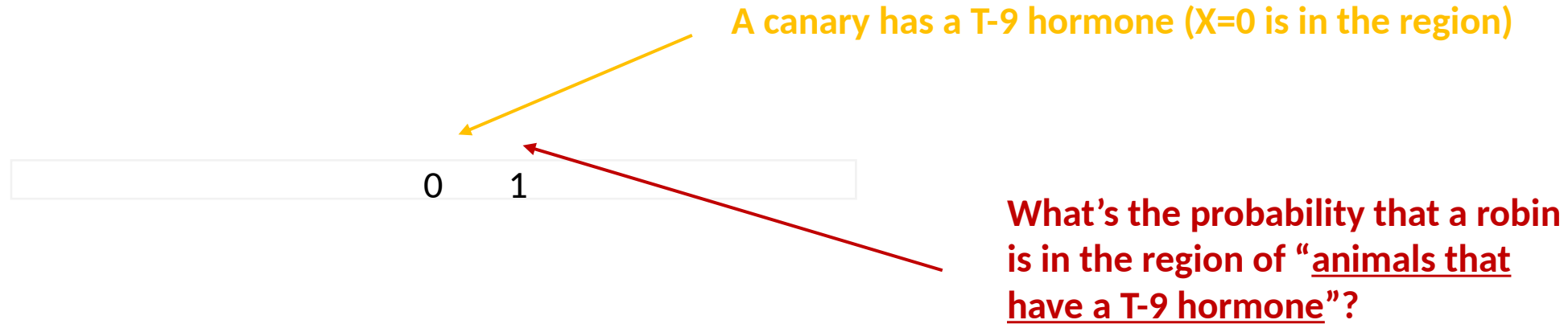


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(What's the probability that  $X=1$  is in the region given that  $X=0$  is in the region?)

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Here, we don't know what the spread of the consequential region will be.

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These black bars are all possible spreads of the CR, generated randomly

(Note that if you are instructed to find the probability of getting  $x=1$  for regions containing  $x=0$ , you need to make sure all your regions contain  $x=0$ )

0 1

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Because we don't know what the true spread of the consequential region (property: animals that have a T-9 hormone) is, we will use all these possible Rs to determine the probability that  $x=1$  (or "robin") is included in that region.

# So how do we actually determine this?

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- This graph can tell you the relative probabilities of the different objects within it

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ry, at 0, has the highest probability.  
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- This graph can tell you the relative probabilities of the different objects within it

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ry, at 0, has the highest probability. Robin, at 1, has a lower probability, and so

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ry, at 0, has the highest probability. Robin, at 1, has a lower probability, and so

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- The y axis of this graph represents the probability that you would sample object x (if you considered all possible CRs).



- This graph can tell you the relative probabilities of the different objects within it

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ry, at 0, has the highest probability. Robin, at 1, has a lower probability, and so

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*In the HW, canary is the only observed object in that property; is in all the regions.*

*So if we sampled objects from all the regions, canary will be sampled the most (every time) whereas robin will be sampled less, and so on.*

0 1

So what does this mean?

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- The y axis of this graph represents the probability that you would sample object x (if you considered all possible CRs).





- You can compute the  $y$  value at any  $x$  using this procedure:

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$$\sum_R P(X|R)P(R)$$

=

$$P(X|A)*P(A)$$

+

$$P(X|B)*P(B)$$

+

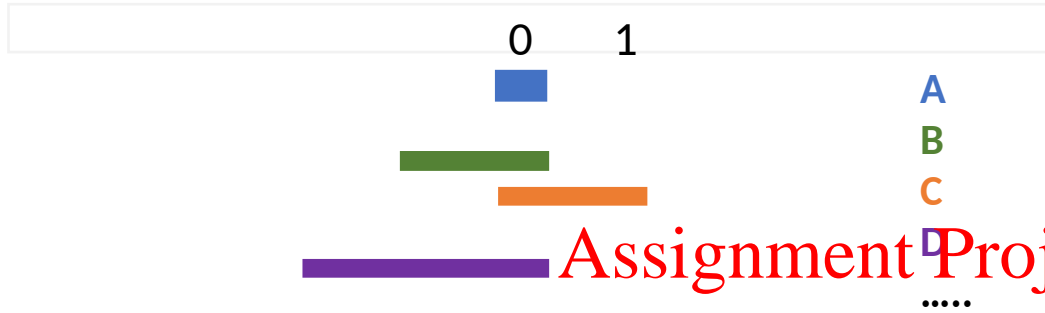
$$P(X|C)*P(C)$$

+

$$P(X|D)*P(D)$$

+

.....



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$$\sum_R P(X|R)P(R)$$

=

$$P(X|A)*P(A)$$

+

$$P(X|B)*P(B)$$

+

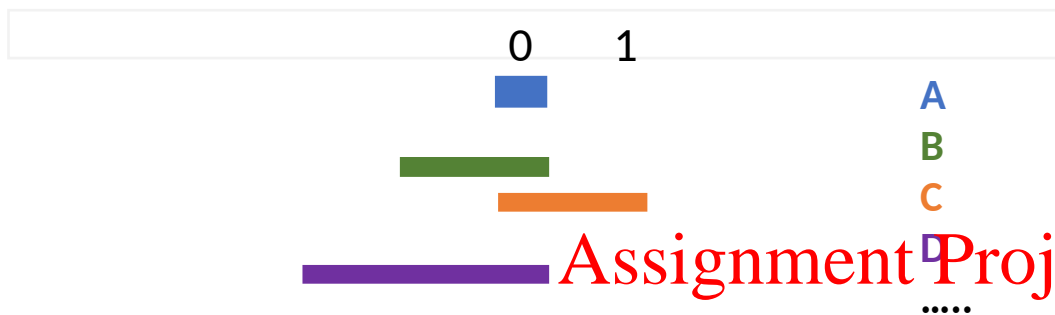
$$P(X|C)*P(C)$$

+

$$P(X|D)*P(D)$$

+

.....



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\*NOTE: If X is not in the consequential region R, then for that specific R,

$$P(X|R)P(R) = 0$$

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-3.2

4.8

All the regions in this assignment should contain  $x=0$ , but how would you check if they contain  $x=1$ ?

\*NOTE: If X is not in the consequential region R, then for that specific R,

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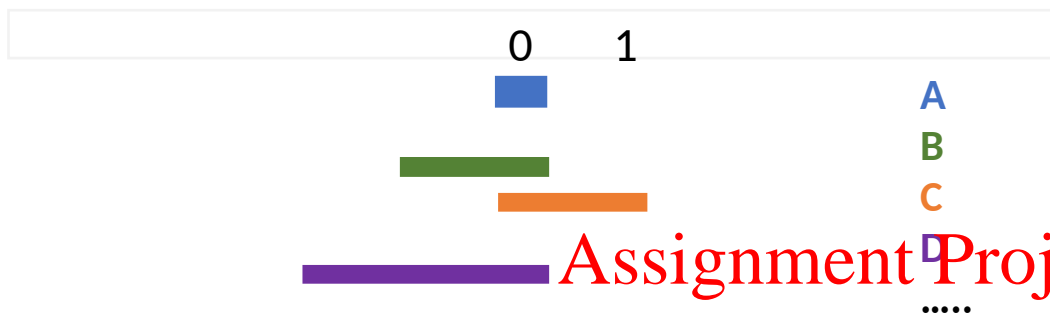
-3.2

4.8

All the regions in this assignment should contain  $x=0$ , but how would you check if they contain  $x=1$ ?

$\text{edge1} \leq (x=1) \leq \text{edge2}$

What's the probability of getting  $x=1$ , for regions containing  $x=0$ ?



$$\sum_R P(X|R)P(R)$$
$$=$$
$$\cancel{P(X|A)*P(A)} + P(X|B)*P(B) + P(X|C)*P(C) + \cancel{P(X|D)*P(D)} + \dots$$

0

0

0

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$$P(X|R)P(R)$$

Probability of  
observing object X in  
the region R

Probability of you  
sampling region R

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(if you put all objects  
that were in R in a  
bag, what would the  
probability of  
sampling this  
particular object X  
be?)

(e.g. if you put all  
regions in a bag, what  
would the probability  
of you sampling that  
particular region R  
be?)



$$\sum_R P(X|R)P(R)$$

probability that your object  $x=1$  (or whatever  $x$  you specify) will have  $x=0$  (or whatever other  $x$  you specify) is the same as the probability that the object that is in all  $R$ .

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Computing this sum is marginalizing over every possible  $R$  there could be (since you don't know the true  $R$ , you see what  $P(X|R)$  would be under any of the possible  $R$ s)

# How to compute $P(X|R)$ and $P(R)$

$P(X|R)$      $1/(\text{length of the } R \text{ you have currently sampled})$     (if  $X$  is in  $R$ , and 0 otherwise)

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$P(R)$      $1/(\text{number of } R\text{s})$     (since every  $R$  has the same chance of being sampled)

# Why do we do this?

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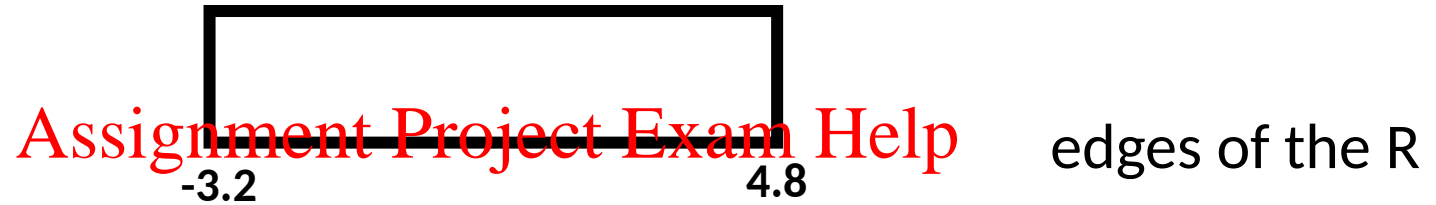
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- You can think of each of these  
s as a uniform distribution

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$$P(X|R)$$



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$$\int_L^R (\text{Consequential Region}) = 1 \quad ; \text{ the area of this rectangle} = 1$$

$$P(X|R)$$

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-3.2

4.8

edges of the R

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Since the length is 8,

$$P(X|R)$$



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Since the length is 8, for length\*height to = 1, height has to be  $1/8$

$$P(X|R)$$



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Hence,  $P(X|R) = 1/(\text{length of } R)$



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# Scaling of axes

- Takes an axis that is written in normal, linear interval form [1,2,3,4....]
- And instead writes it as  $[e^1, e^2, e^3, e^4, e^5]$   
(or  $[10^1, 10^2, 10^3, 10^4, 10^5]$ , which is equivalent because of the change of base formula)
- That changes the spacing (and causes a distortion)  
al is changing a lot, the corresponding spacing on the y axis is just one space, making it seem as if the exponential just decreased by a little, not a lot)
- `plt.yscale`

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