

What is deep learning and how is it used in biology

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What is

- How does deep learning work?
- What types of biological problems is deep learning useful for

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What is machine learning?

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Why use machine learning?

- Automate a manual decision process at larger scale
- Most deep learning doesn't do anything that a human couldn't do given enough time
 - Sort ripe from unripe tomatoes
 - Decide if a cat is in a photo

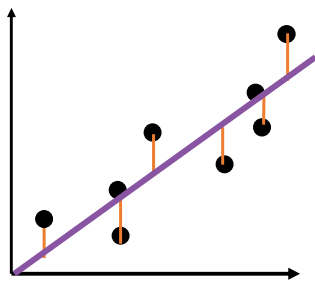
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When do we use machine learning?

- When you have a set of examples with any label
- Classification
 - Data: Tumor biopsies label: malignant vs. benign
 - Data: Protein sequences label membrane vs soluble
- Regression
 - How much fuel will an airplane use?

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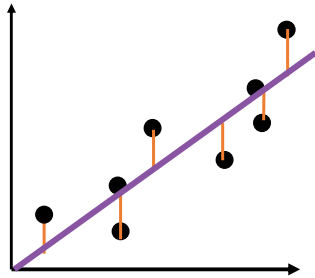
Linear regression is machine learning



- Formula $y = Ax + b$
- A = slope, b = y intercept
- What are the values of A and b that minimize the errors between the line and the points
- Optimization: Find the values of variables by minimizing error to the expected result

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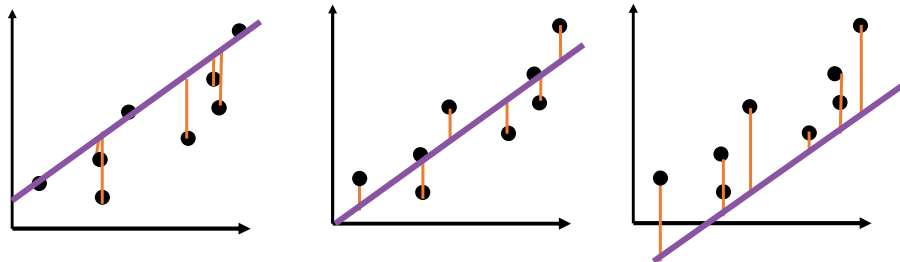
Linear regression is machine learning



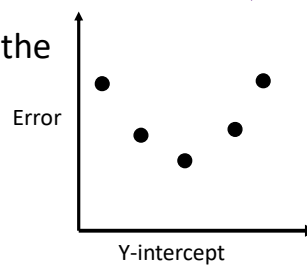
- A person could figure out the best values of $y = Ax + b$ by trying values and checking how large the error is between the line and the point
- However the process of doing can be done automatically by plotting the error as the variable changes

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Linear regression is machine learning

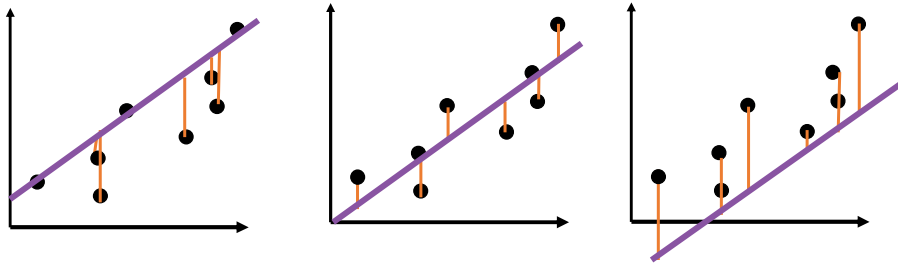


- We can look at the error as the y intercept changes

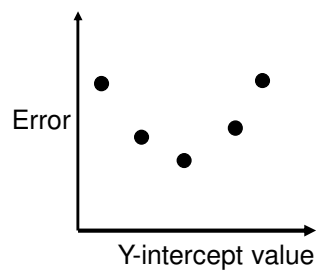


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Linear regression is machine learning

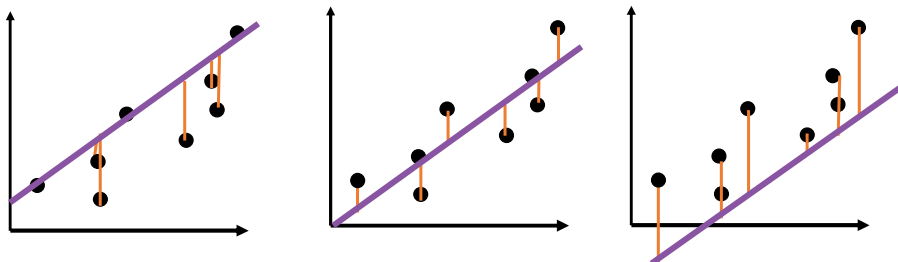


- We can look at the error as we vary value for the y intercept

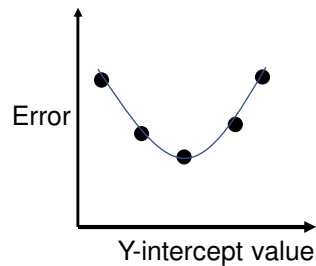


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Linear regression is machine learning

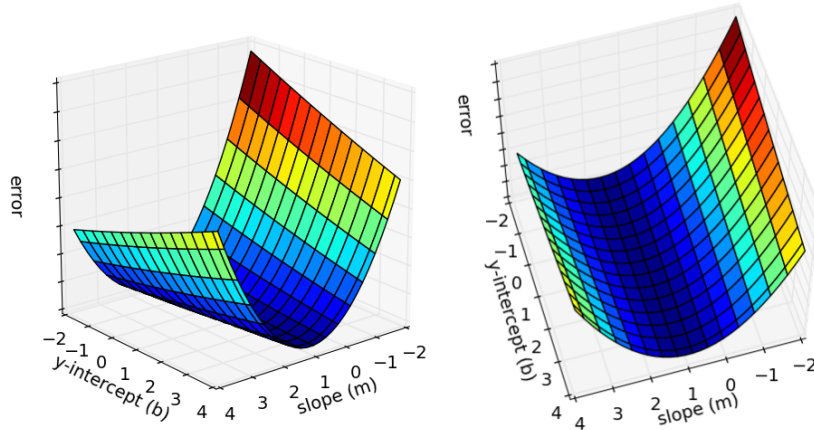


- With the relationship between y-intercept and error, we can find the value of y-intercept that minimizes the error



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Importantly, both slope and intercept can be optimized simultaneously



Imagine rolling a ball into this plane, seeing where it lands, and taking the values of slope and y-intercept at that point

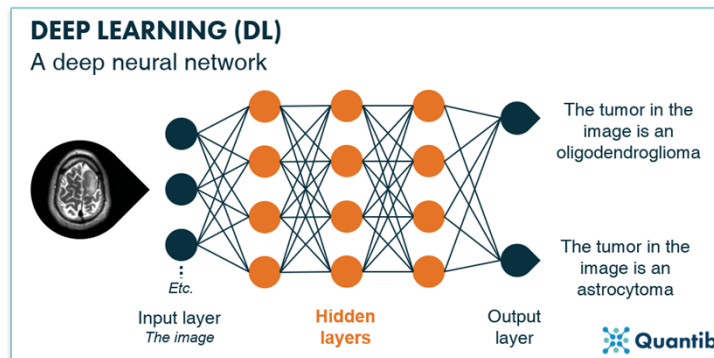
This process isn't limited to just two variable at a time, many variables can be optimized at one

<https://spin.atomicobject.com/2014/06/24/gradient-descent-linear-regression/>

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Deep learning is subset of machine learning

- Deep learning refers specifically to deep neural network architecture
- Architecture = how a model is set up

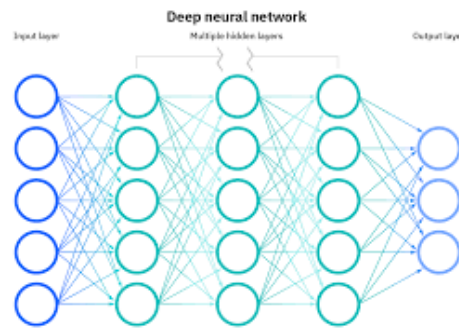


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Imagery around deep learning is unhelpful



What does this mean?



What are the dots?
What are the line?
How does this predict?

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Neural networks are a series of data transformations

- Training a neural network is figuring out the parameters for the different data transformations

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What neural networks aren't

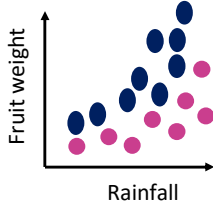


- Not a brain-like or neuron-like structure
- Essentially they are a series of data transformations
- With parameters for the data transformations learned simultaneously
 - Like in the linear regression example

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Contrived example

- A farm with blackberries and raspberries
- They would like a robot to automatically sort berries, but all they have is a scale
- They know that fruit weight varies drastically with rainfall levels

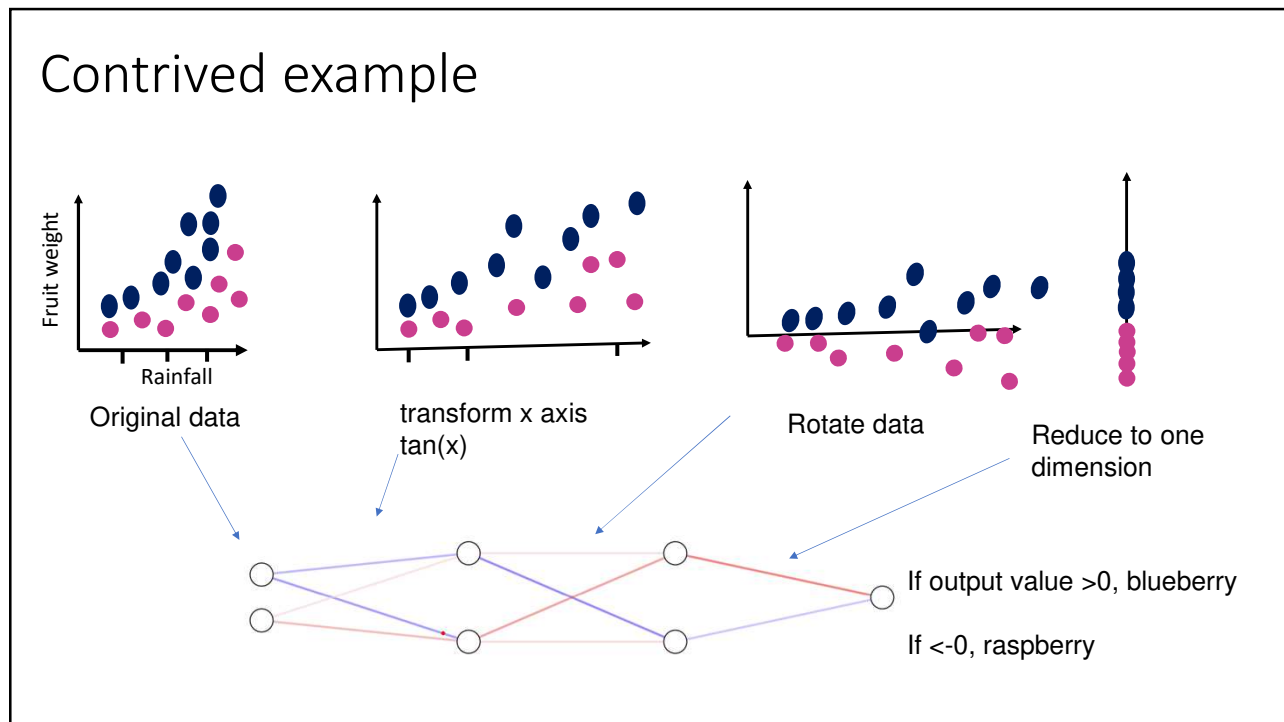


Options

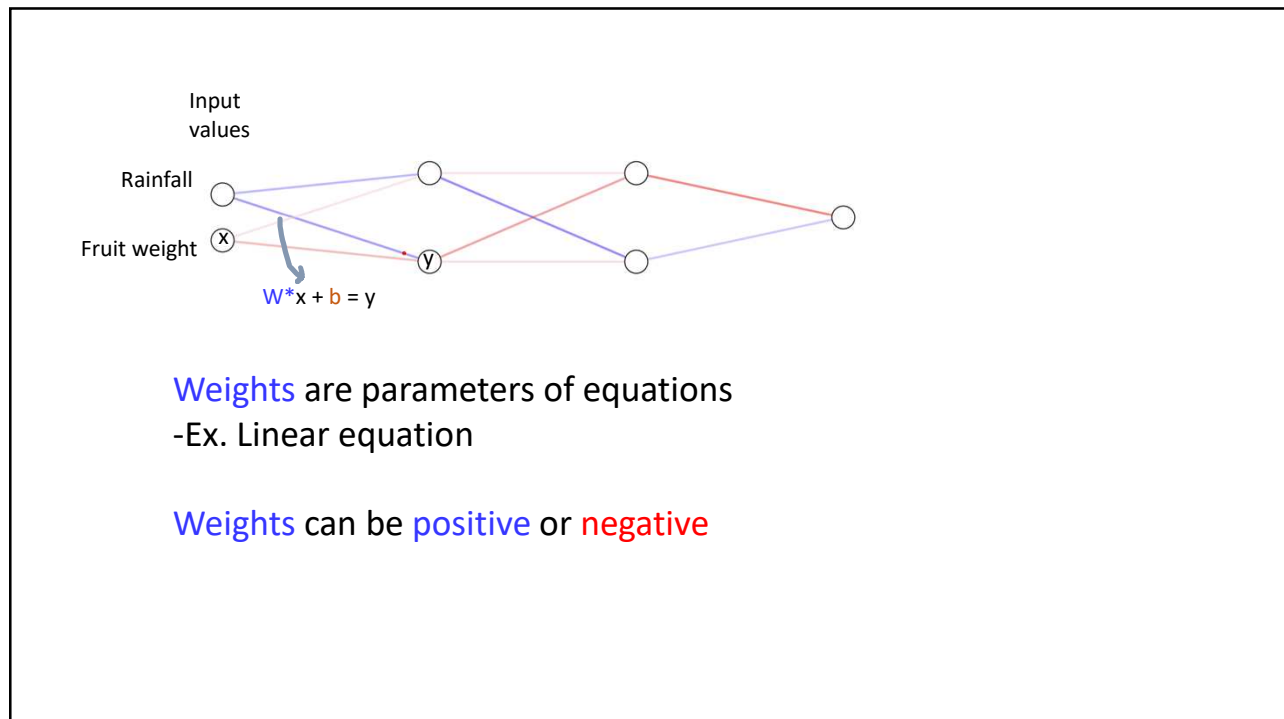
- Make a rule for different rainfall levels
 - Ex. If there's been 3-5 inches of rainfall, anything above 2 grams is a blackberry
- Find an equation that divides the two
- Use a neural network to find data transformations that allow you to cleanly separate between the two berries

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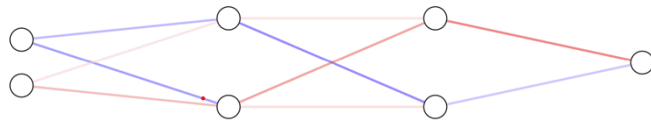
Contrived example



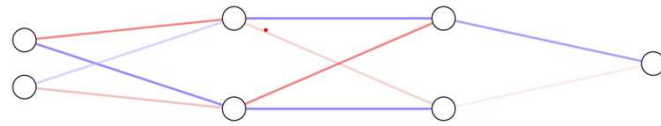
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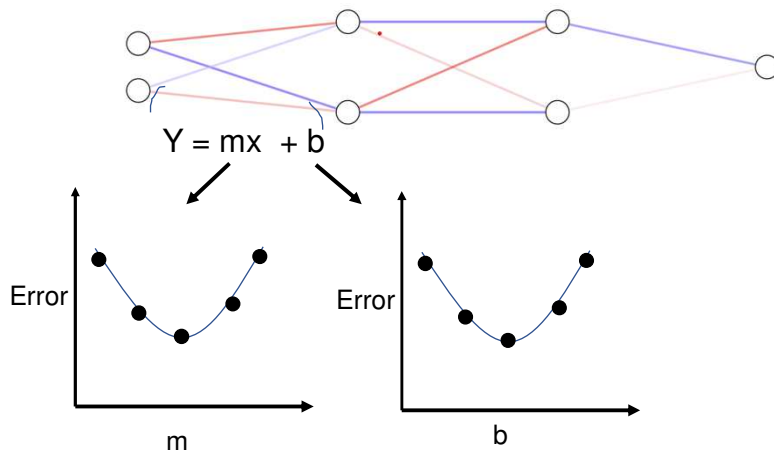


- Neural networks begin with random weights
- Random weights create random output
- We can find good values from the weights because we can measure the error between the produced output and the expected output



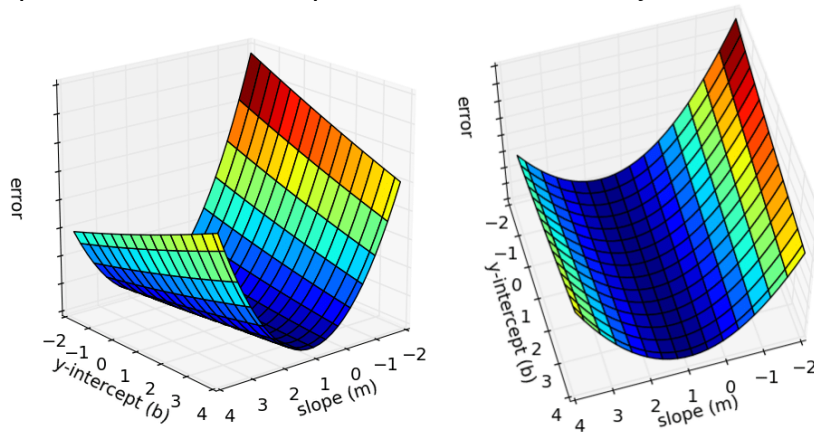
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Every variable has its own optimization relative to the error



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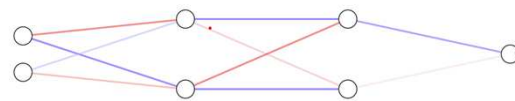
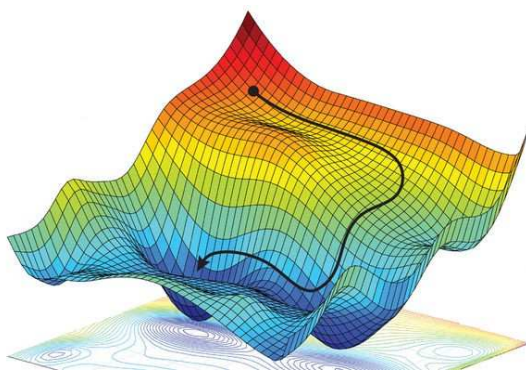
Multiple parameters can be optimized simultaneously



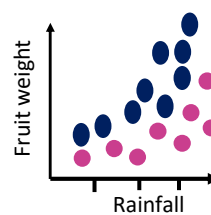
<https://spin.atomicobject.com/2014/06/24/gradient-descent-linear-regression/>

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Gradient descent allows the whole network to optimized simultaneously



Finding the values for each of the data transformations that separate blueberries and raspberries



Source: <https://reconsider.news/2018/05/09/ai-researchers-allege-machine-learning-alchemy/>

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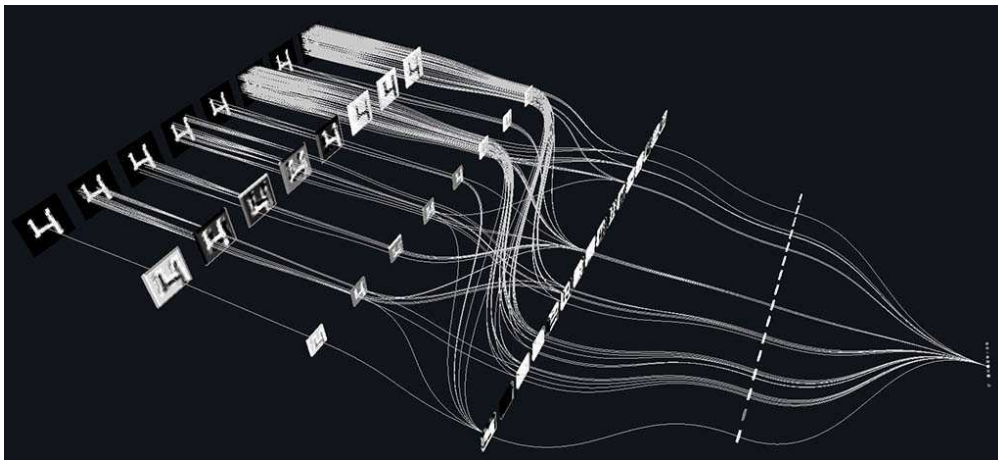
Applications of neural networks in biology

- Something a human could do, but it takes too much time to do it more than a few times
- Particularly used in image analysis
 - A person can look at a few hundred images, a train classifier can look through many more

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Convolution is a common step in image processing neural networks

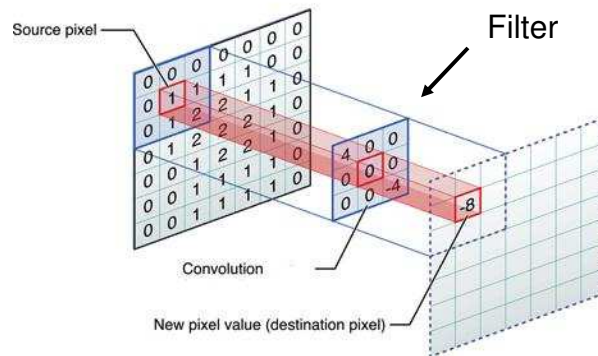
In a convolutional neural networks, the original image is modified with multiple filters to create richer inputs for the model to learn from



<https://medium.com/analytics-vidhya/understanding-convolution-operations-in-cnn-1914045816d4>

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Convolution moves a filter over an input image to create a new image



<https://medium.com/@bdhuma/6-basic-things-to-know-about-convolution-daef5e1bc411>

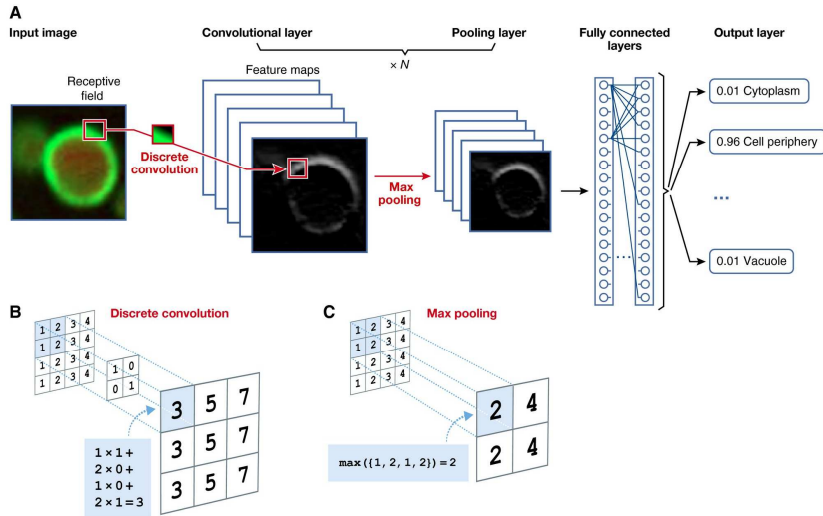
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Different types of filters

<i>Original</i>	<i>Gaussian Blur</i>	<i>Sharpen</i>	<i>Edge Detection</i>
$\begin{bmatrix} 0 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 0 \end{bmatrix}$	$\frac{1}{16} \begin{bmatrix} 1 & 2 & 1 \\ 2 & 4 & 2 \\ 1 & 2 & 1 \end{bmatrix}$	$\begin{bmatrix} 0 & -1 & 0 \\ -1 & 5 & -1 \\ 0 & -1 & 0 \end{bmatrix}$	$\begin{bmatrix} -1 & -1 & -1 \\ -1 & 8 & -1 \\ -1 & -1 & -1 \end{bmatrix}$
			

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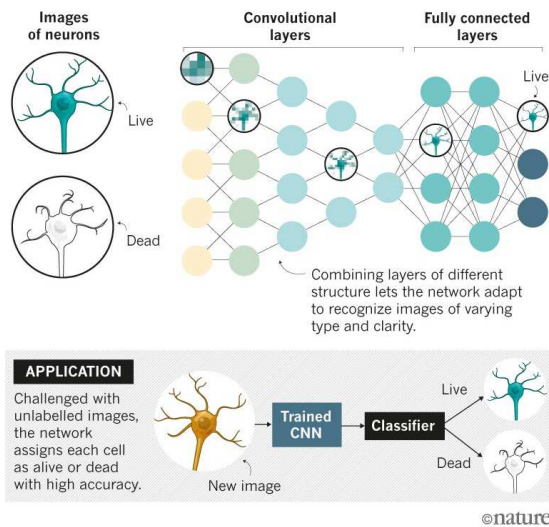
Classifying protein localization



Angermueller, 2016, MSB

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Classifying living/dead neurons



- Network to separate images of live and dead neurons
- Convolutional neural networks create richer representations for the fully connected layers

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Natural language processing

- By just looking at all the text in Wikipedia, a natural language model can accurately describe English, even without prior knowledge about grammar/parts of speech/etc.
 - Many differently styles of computer model for doing this
- Give these same models every protein sequence, and you get a model that describes “protein language”

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Why model protein sequence as a language?

- Sequences of tokens, where token meaning depends on context
- Language: “I drank a cold strawberry smoothie”
- Protein: “M E C E C A H N S H Q”

* To format protein sequences for language models, all you need to do is put a space between each amino acid

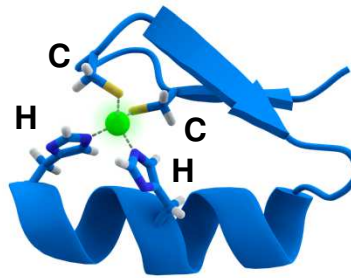
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Why model protein sequence as a language?

- There are specific relationships between tokens

subject-verb modifier
 “I drank a cold strawberry smoothie”

C2H2 zinc-finger motif
 “M E C E C A H N S H Q”



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Why model protein sequence as a language?

- A sentence can be represented as a **long numeric vector** ↗
- The difference between any two sentences can be measured by comparing their vectors



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Why model protein sequence as a language?

- Some positions in the sequence can be changed without changing the meaning of the sentence too much

“I drank a cold strawberry smoothie”
 “I sipped a chilled strawberry smoothie”

“MECECAHNSHE”
 “MECECAHNSHQ”

Highly similar

Highly similar

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Why model protein sequence as a language?

- Some positions in the sequence can be changed without changing the meaning of the sentence too much

“I drank a cold strawberry smoothie”
 “I drank a cold computer smoothie”

“MECECAHNSHE”
 “MECECAHNSHP”

Change in meaning

Change in meaning

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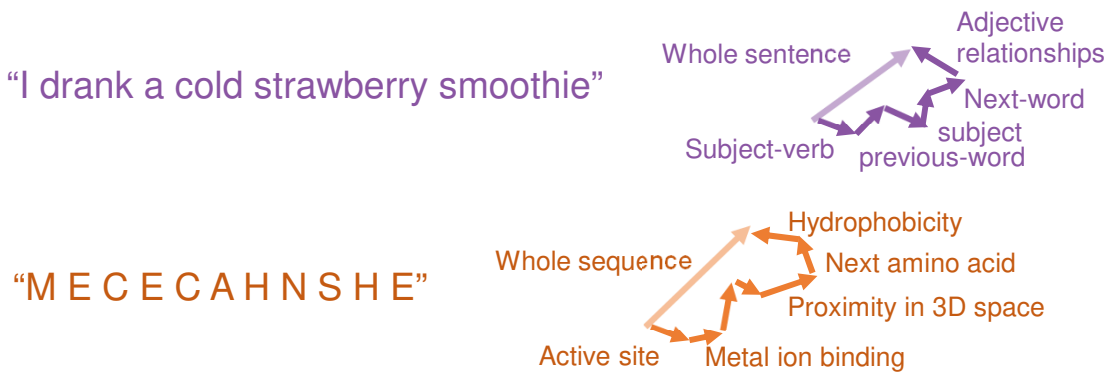
Each amino acid has its own meaning vector

*The sentence vector is the mean of all word vectors



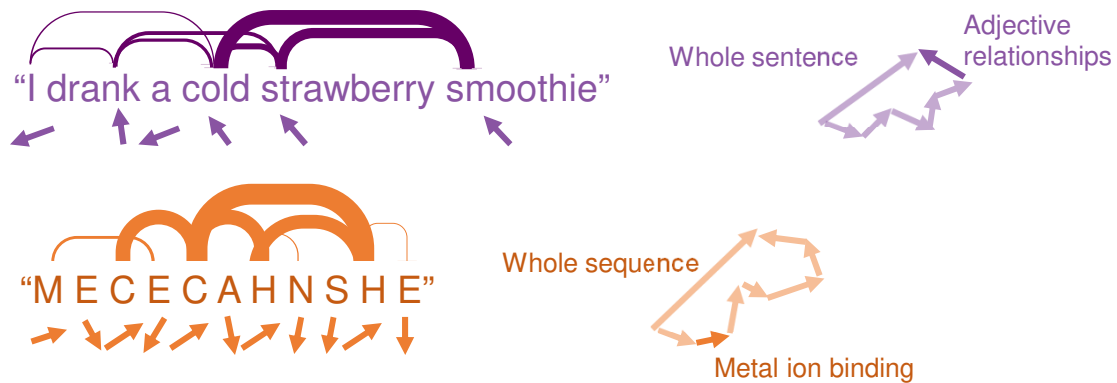
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Vectors are composed of many smaller vectors, which capture different relationships



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A calculation between word vectors overlays a network onto the sequences



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Recap

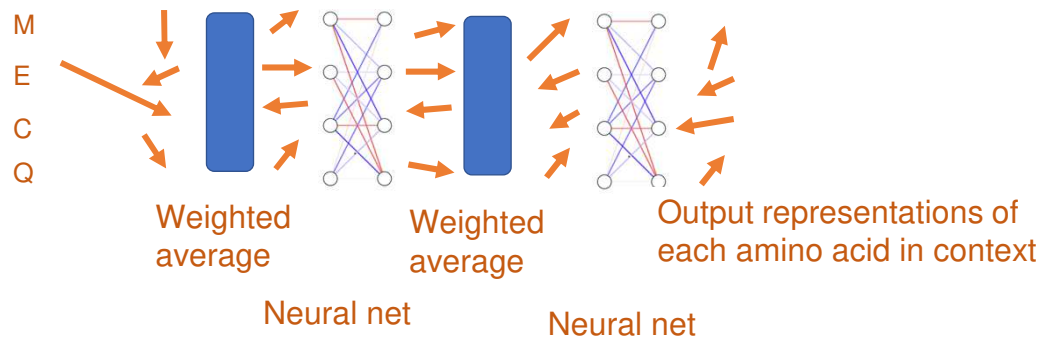
- Each word/amino acid has a long vector that captures its identity
- Averaging these word vectors creates a sentence vector
- Vectors are composed of smaller vectors which capture different properties



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Connecting back to neural networks

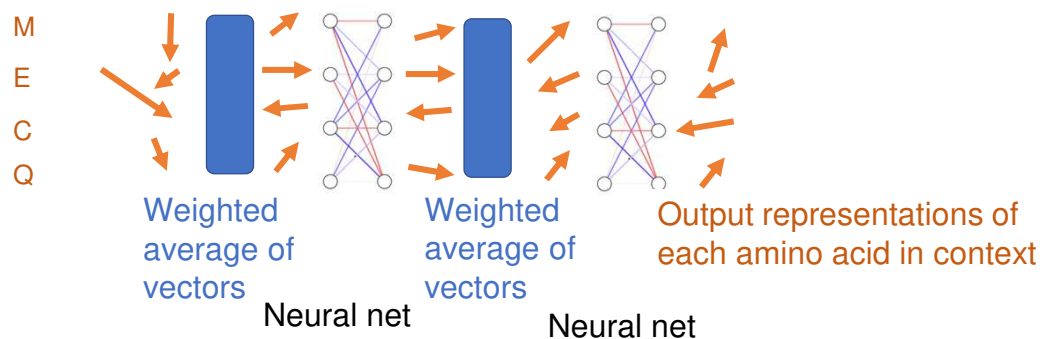
- Weighted average steps alternated with neural networks with trainable weights.
- The weighted average operation is called “attention”



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Connecting back to neural networks

- Each of the twenty amino acids begins with one of twenty vectors
- By mixing information between them, output vectors that represent each amino acid in context are produced



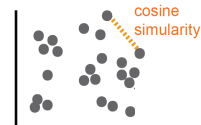
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Semantic shift variant effect prediction

Deep mutational scanning dataset

Measure the similarity between a mutant sequence and wild-type

Sequence	Functional readout	Semantic shift
MECECAHNSHE	Wild-type	0
MEC Q CAHNSHE	Neutral	0.1
MEC P AHNSHE	Deleterious	0.6



Using models trained for sequence similarity should improve performance

Judy Du & Anne Cirincione, Mona Singh lat

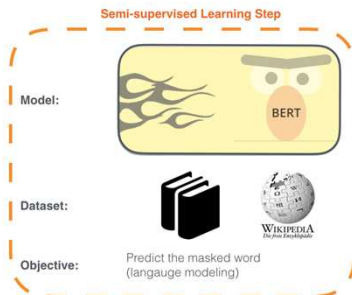
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Modern language models trained on large numbers of sequences can be fine-tuned to produce embeddings that are better suited for particular tasks

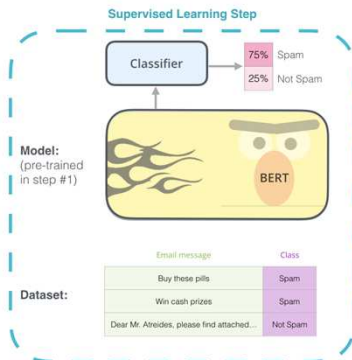
Don't need to create a whole new neural for every problem

1 - **Semi-supervised** training on large amounts of text (books, wikipedia.. etc).

The model is trained on a certain task that enables it to grasp patterns in language. By the end of the training process, BERT has language-processing abilities capable of empowering many models we later need to build and train in a supervised way.



2 - **Supervised** training on a specific task with a labeled dataset.



Jay Alammarr – The illustrated transformer

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