

# Attention and Transformer Models for Genomics

BMI/CS 776

[www.biostat.wisc.edu/bmi776/](http://www.biostat.wisc.edu/bmi776/)

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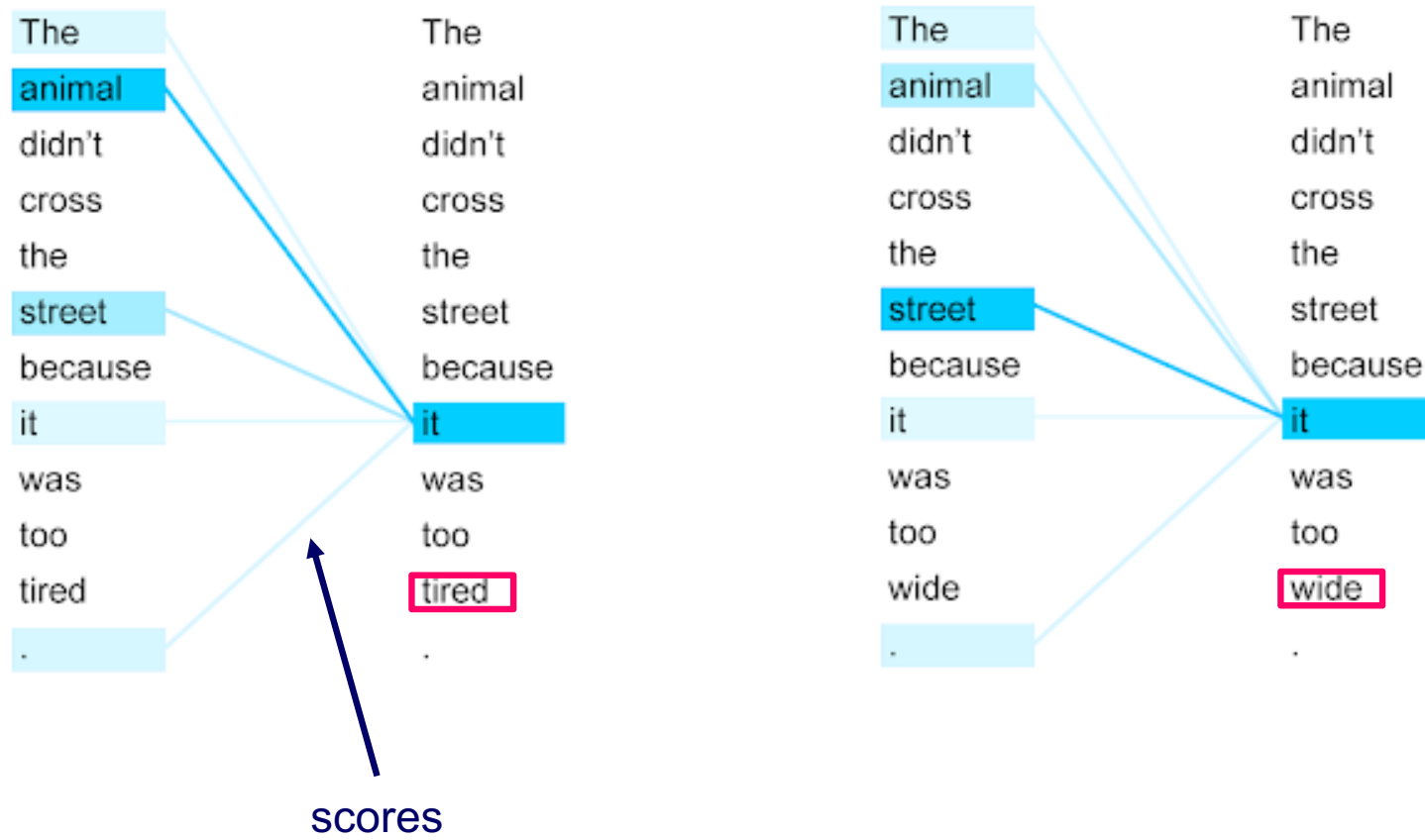
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*Thanks to Noah Cohen Kalafut for slides!*

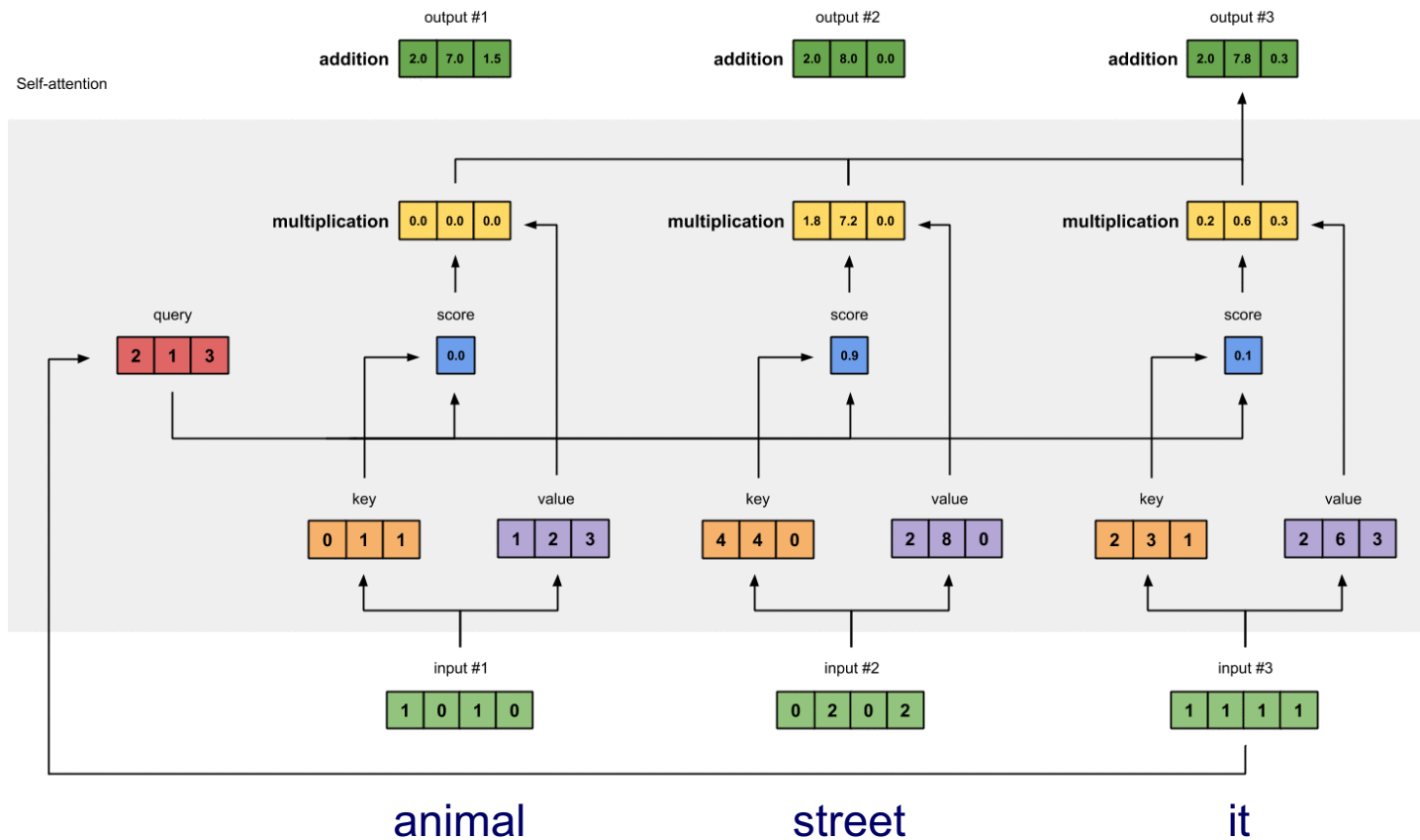
# Goals for lecture

- Attentions
  - Interpretation
  - Self and Cross-Attention Calculation
  - Multi-Head Attention
- Transformer architecture
  - Positional Encoding
  - Generative Output
- Applications to bioinformatics

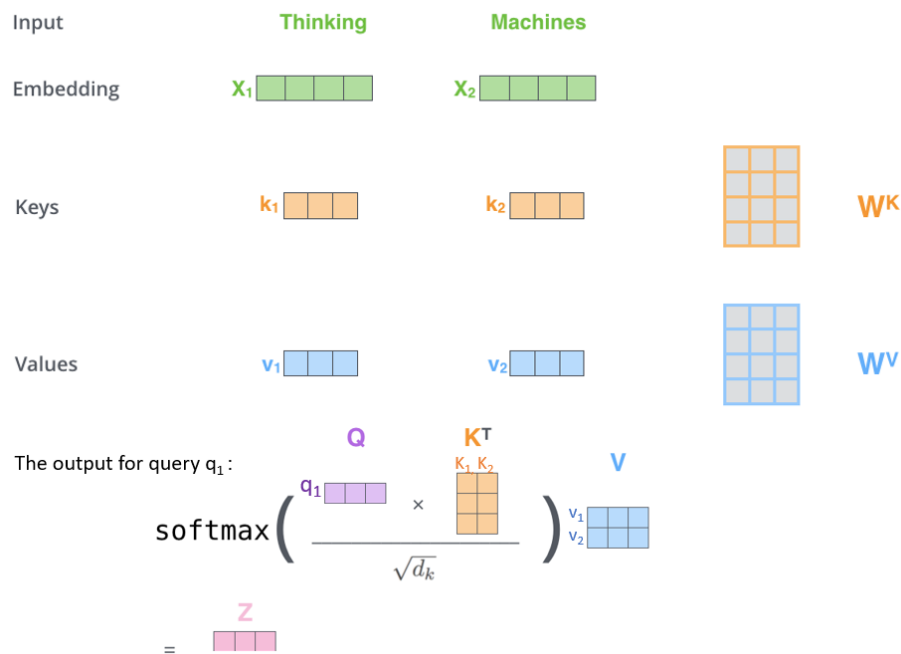
# Attention Interpretation



# Attention Calculation

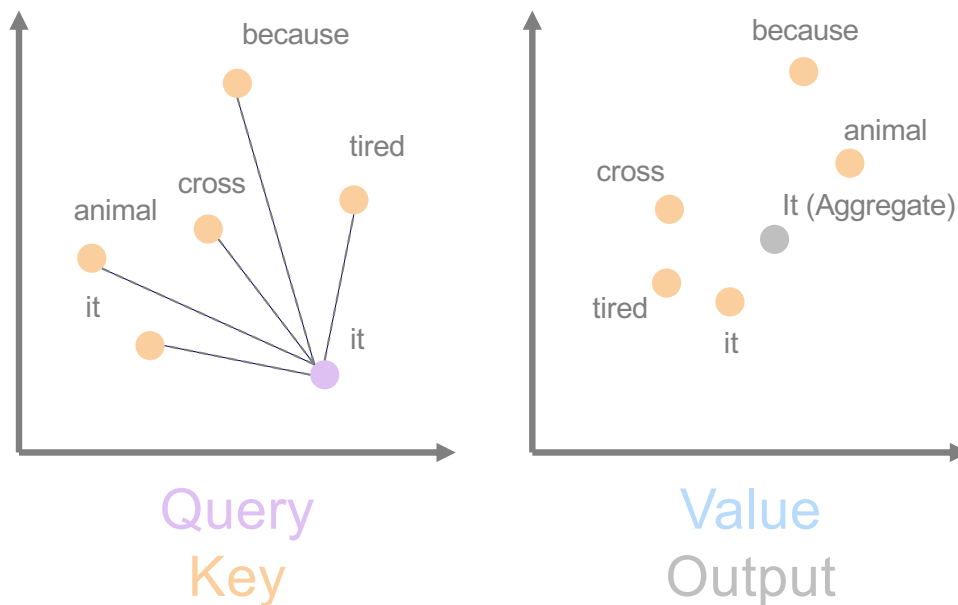


# Attention Calculation



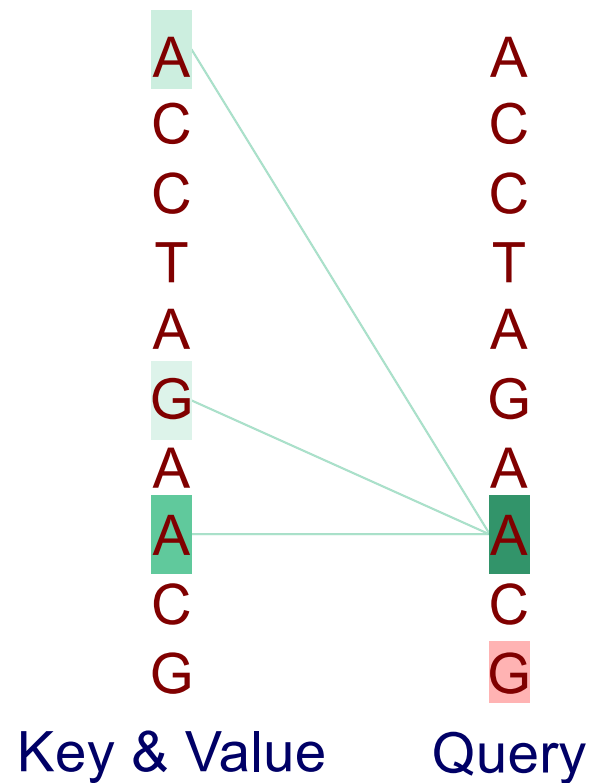
- Embeddings are passed through feed-forward networks to produce a **query** vector, as well as **key** and **value** vectors
- **Value** vectors are summed proportionally to the similarity between their corresponding **keys** and the **query**

# Attention Calculation

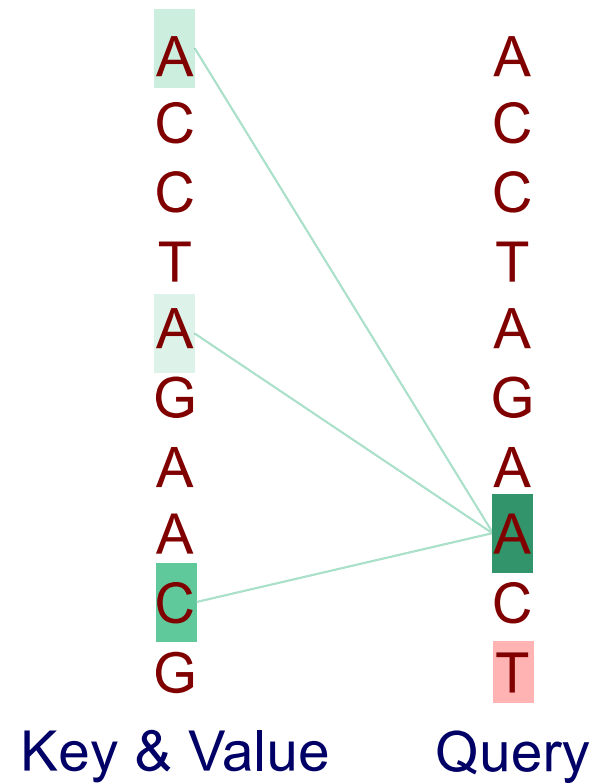


- **Key** and **query** vectors can be thought of as sharing a latent space
- The distance between the **query** and **keys** then determines the final **output** in **value** space

# Types of Attention

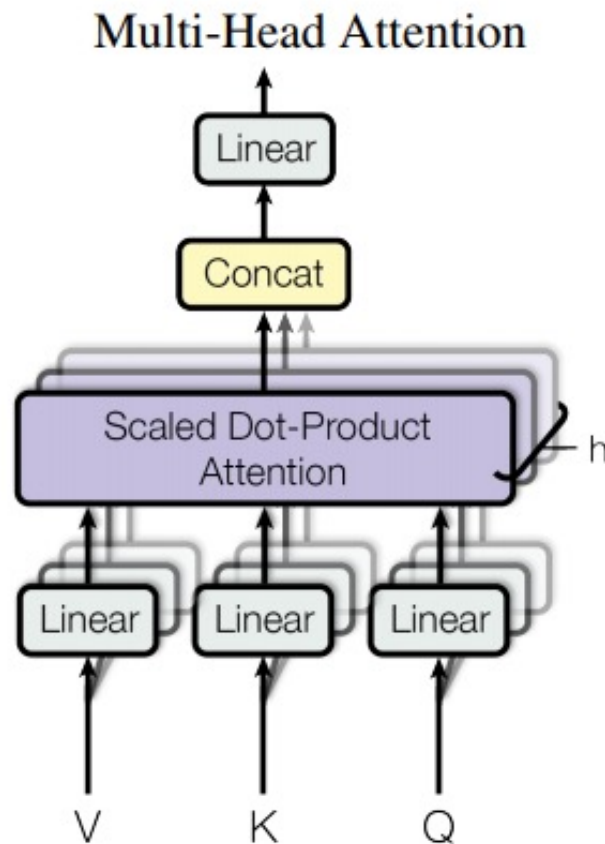


Self-Attention



Cross-Attention

# Multi-Head Attention



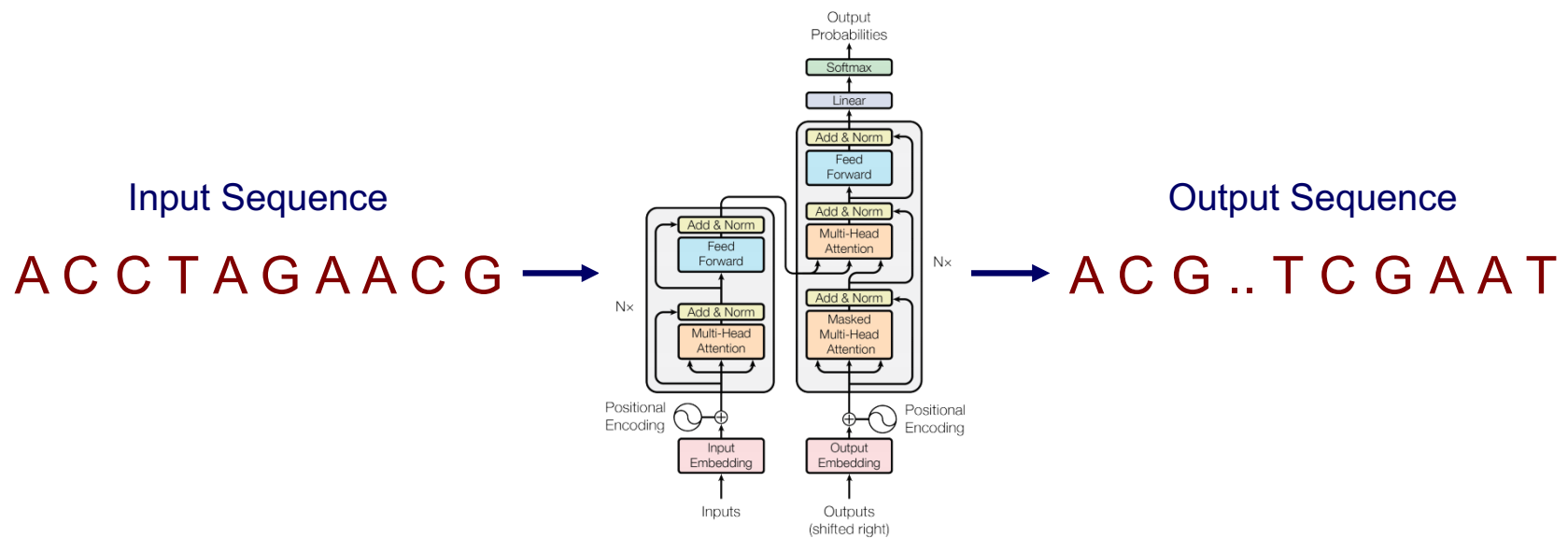
Each attention module is known as a 'head'

Multi-head attention involves aggregating multiple heads, usually culminating in average pooling

This allows each head to prioritize differing features of the sequence (e.g. noun-noun relationships, noun-verb relationships)



# Transformer Outline



# Example: Generative Text



Find the bug with this code:

```
...  
for (var i = 0; i < 5; i++) {  
  setTimeout(() => console.log(i), 1000)  
}  
...
```



The code contains a bug that will cause it to always print the number 5 to the console, rather than the expected sequence of numbers from 0 to 4. This is because the `setTimeout` function creates a closure that captures the value of the `i` variable, but the `i` variable is changed before the callback function is executed.

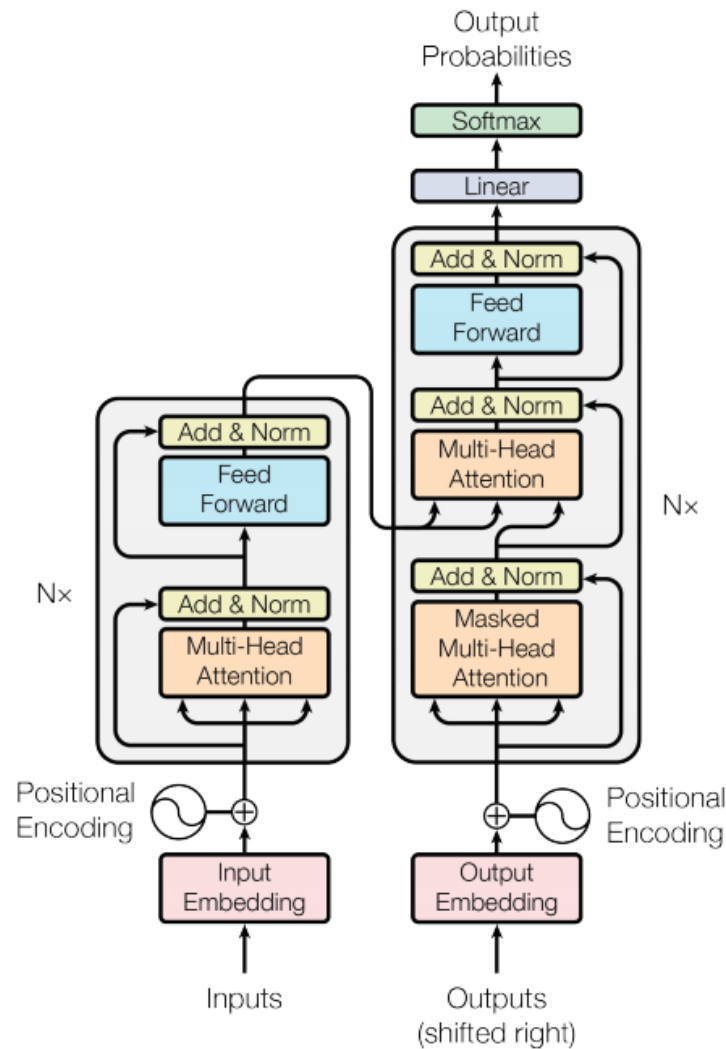


To fix this bug, the code can be modified to create a new variable in the loop that is passed to the `setTimeout` callback function, rather than using the loop variable directly. This will ensure that each callback function captures the correct value of the loop variable, and the expected output will be produced. Here is an example of how the code could be fixed:

```
Copy code  
for (var i = 0; i < 5; i++) {  
  // Create a new variable and pass its value to setTimeout  
  const value = i;  
  setTimeout(() => console.log(value), 1000)  
}
```

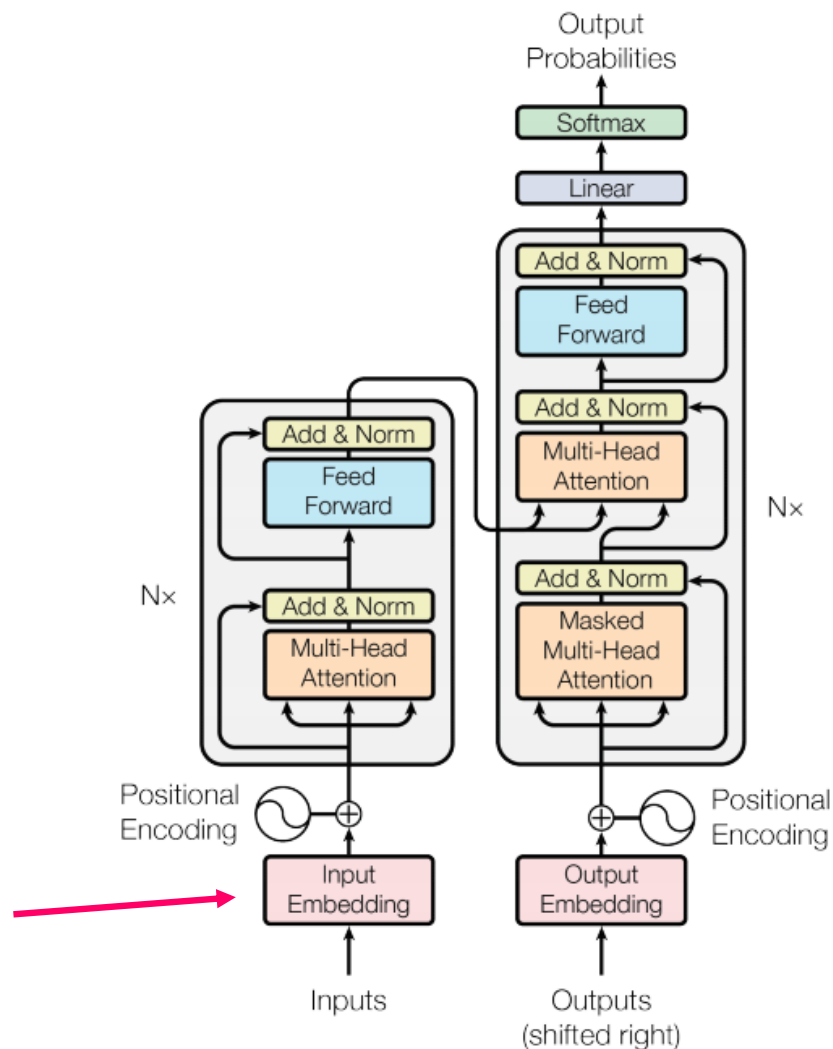
This code will print the numbers 0 to 4 to the console, as expected. Alternatively, the `let` keyword can be used in place of `var` to declare the loop variable, which will automatically create a new variable for each iteration of the loop and avoid the need to create a new variable manually. This is a common pattern for avoiding closure-related bugs in JavaScript.

# Transformer Architecture



Encoder Decoder

# Input Embedding

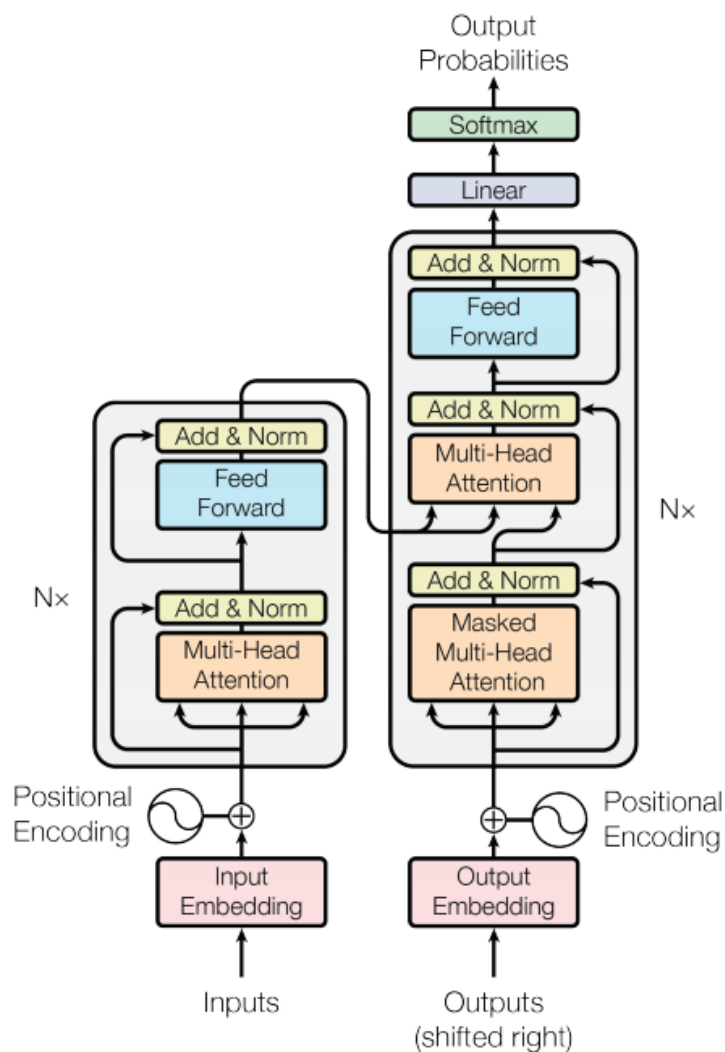


Input Sequence  
A C C T A G A A C G

Learned  
Embeddings

0 4 4 5 0 8 0 0 0 8  
2 8 8 8 2 3 2 2 8 3  
5 5 5 2 5 0 5 5 5 0  
3 8 8 8 3 6 3 3 8 6

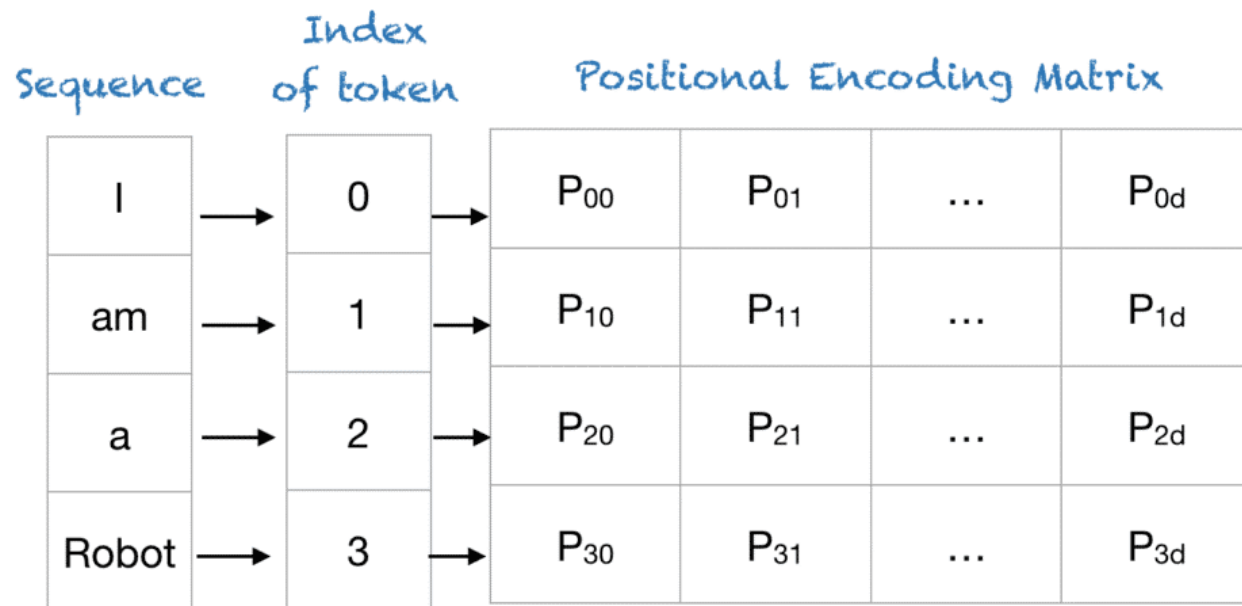
# Positional Encoding



Add positional representations to word embeddings

- Allows the network to consider word proximity

# Positional Encoding



Positional Encoding Matrix for the sequence 'I am a robot'

# Positional Encoding

$$P(k, 2i) = \sin\left(\frac{k}{n^{2i/d}}\right)$$

$$P(k, 2i + 1) = \cos\left(\frac{k}{n^{2i/d}}\right)$$

Positional Encoding Matrix with  $d=4$ ,  $n=100$

Sequence	Index of token, $k$	$i=0$	$i=0$	$i=1$	$i=1$
I	0	$P_{00}=\sin(0)$ = 0	$P_{01}=\cos(0)$ = 1	$P_{02}=\sin(0)$ = 0	$P_{03}=\cos(0)$ = 1
am	1	$P_{10}=\sin(1/1)$ = 0.84	$P_{11}=\cos(1/1)$ = 0.54	$P_{12}=\sin(1/10)$ = 0.10	$P_{13}=\cos(1/10)$ = 1.0
a	2	$P_{20}=\sin(2/1)$ = 0.91	$P_{21}=\cos(2/1)$ = -0.42	$P_{22}=\sin(2/10)$ = 0.20	$P_{23}=\cos(2/10)$ = 0.98
Robot	3	$P_{30}=\sin(3/1)$ = 0.14	$P_{31}=\cos(3/1)$ = -0.99	$P_{32}=\sin(3/10)$ = 0.30	$P_{33}=\cos(3/10)$ = 0.96

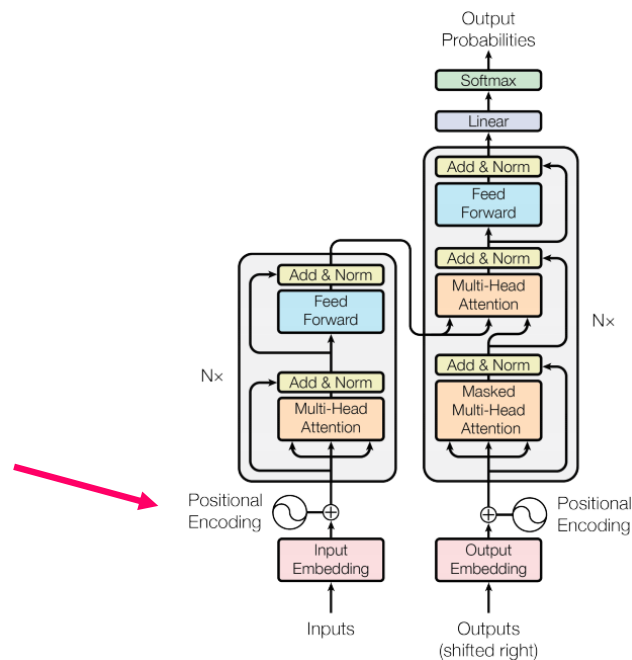
Positional Encoding Matrix for the sequence 'I am a robot'

# Positional Encoding

Learned Embeddings										Temporal Embeddings									
0	4	4	5	0	8	0	0	0	8	0	6	0	6	0	2	3	8	5	9
2	8	8	8	2	3	2	2	8	3	2	2	0	9	7	4	9	1	2	7
5	5	5	2	5	0	5	5	5	0	4	7	1	5	3	3	7	6	3	1
3	8	8	8	3	6	3	3	8	6	3	5	7	1	2	5	3	0	3	9

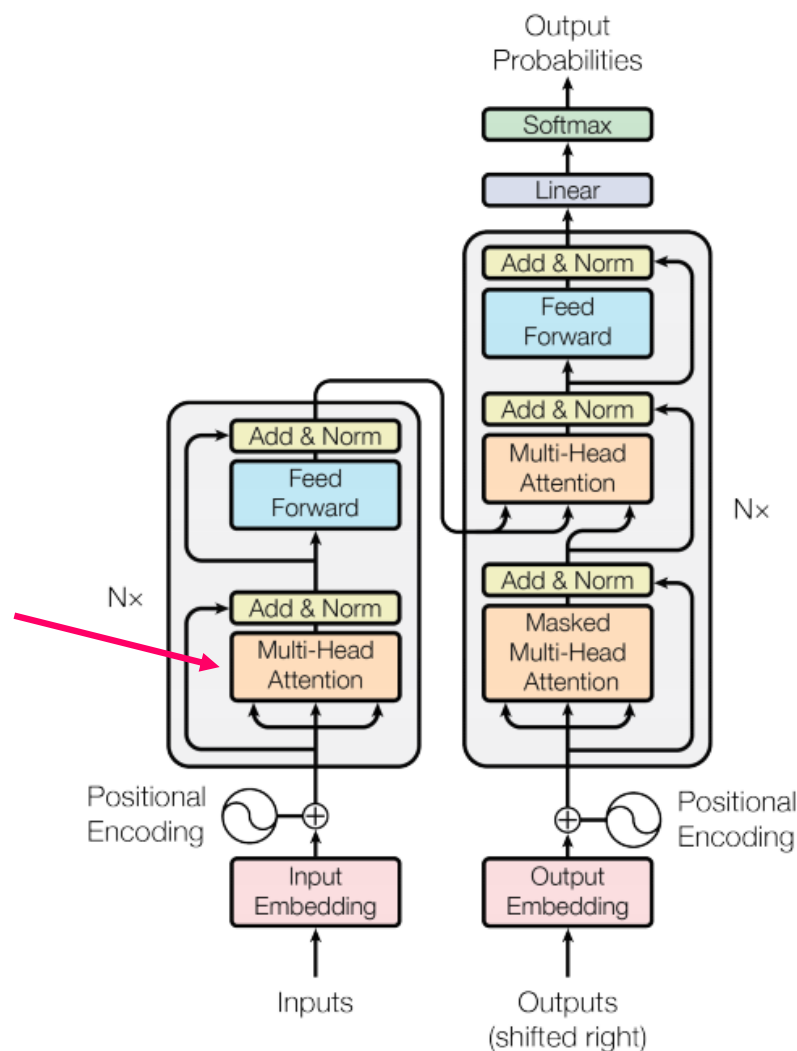
Positional Encoding

Addition





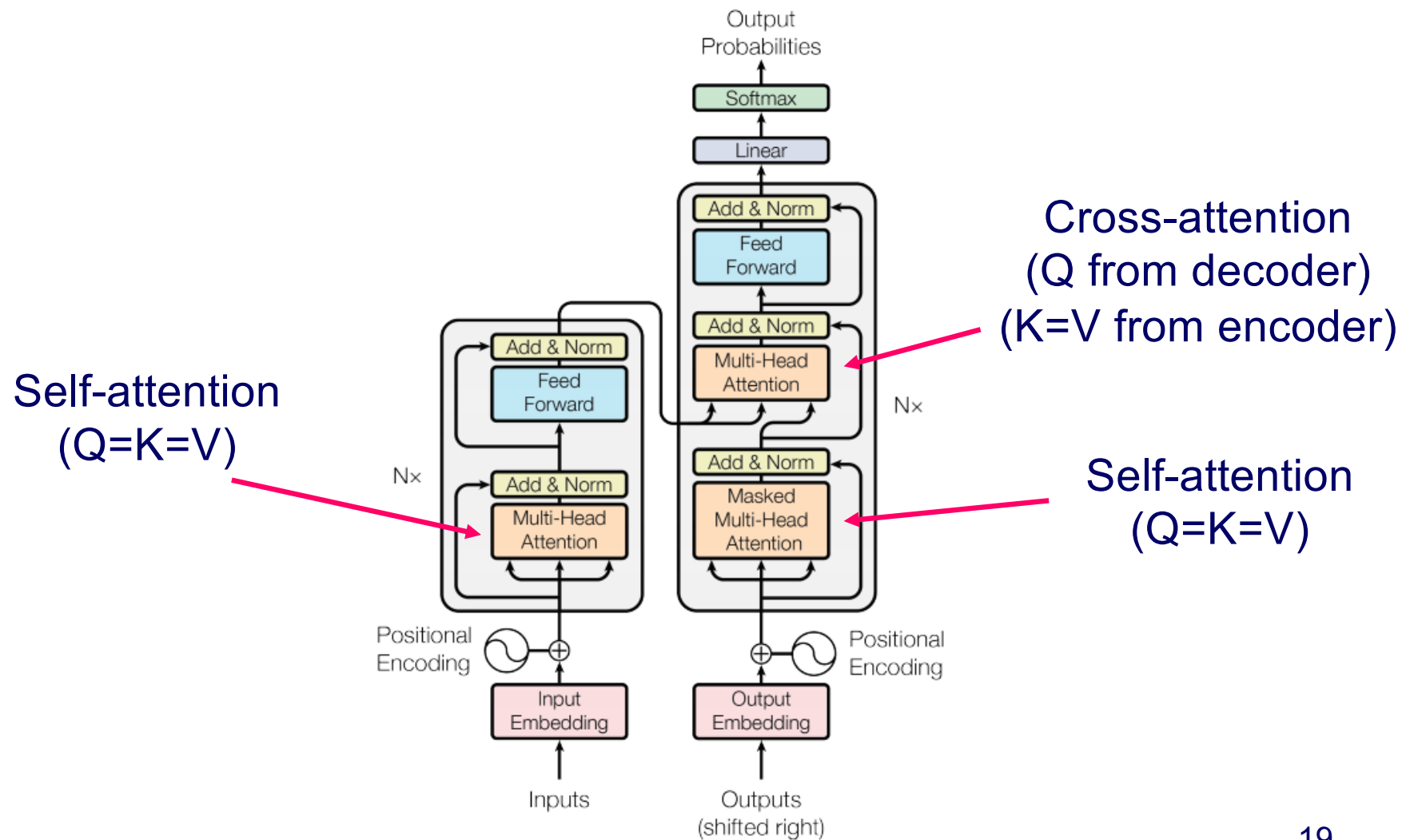
# Attention Modules



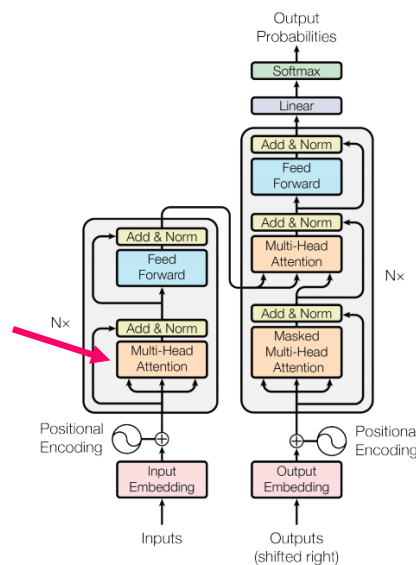
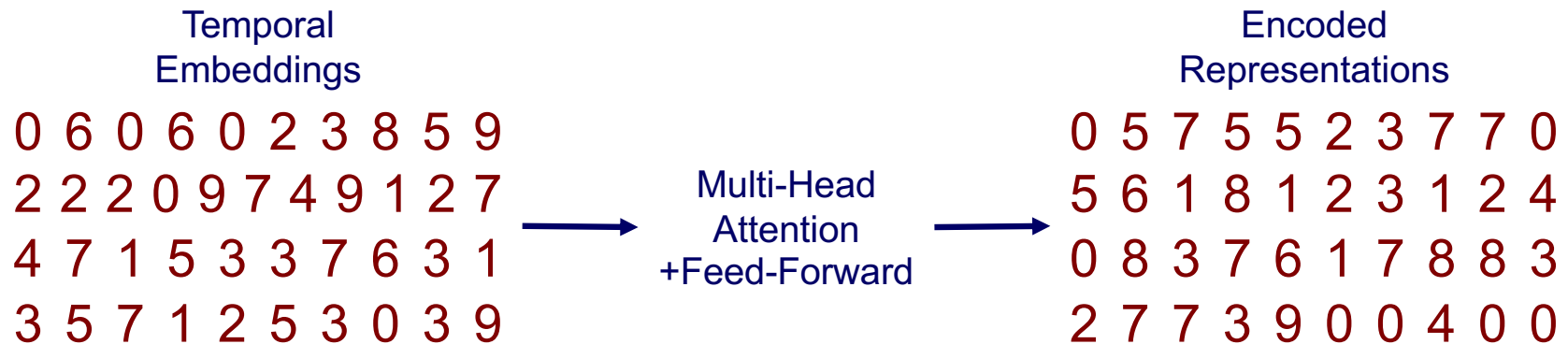
Prioritize bases relative to each other

- This is the primary mechanism which allows transformers to work
- Essentially adds context to existing embeddings

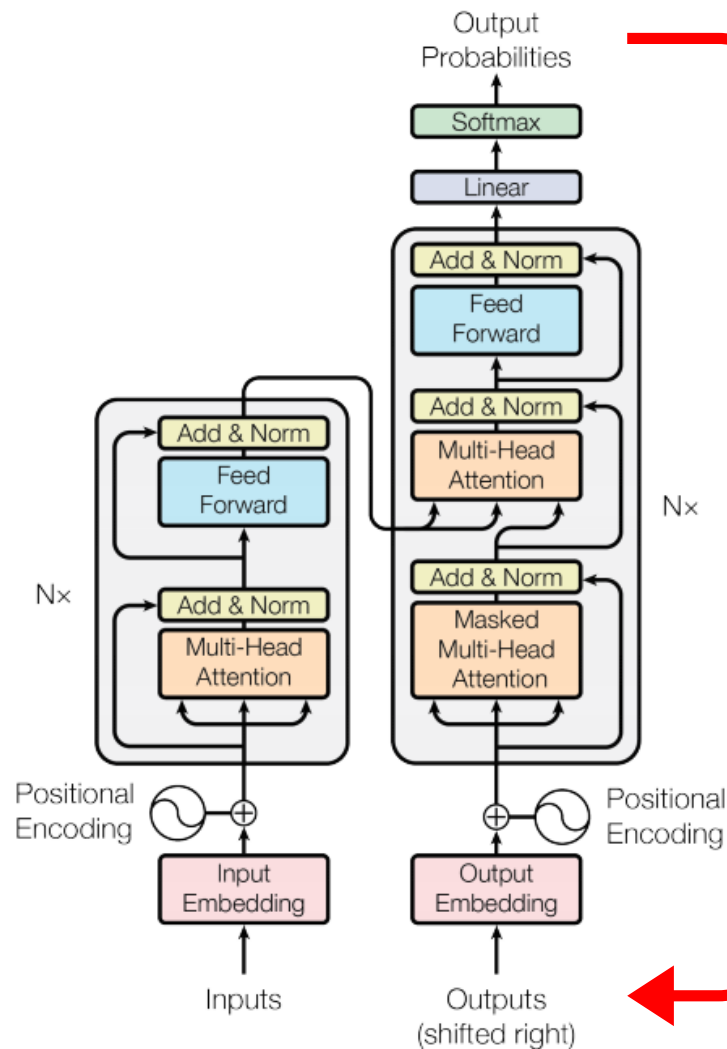
# Self and Cross-Attention



# Transformer Attention



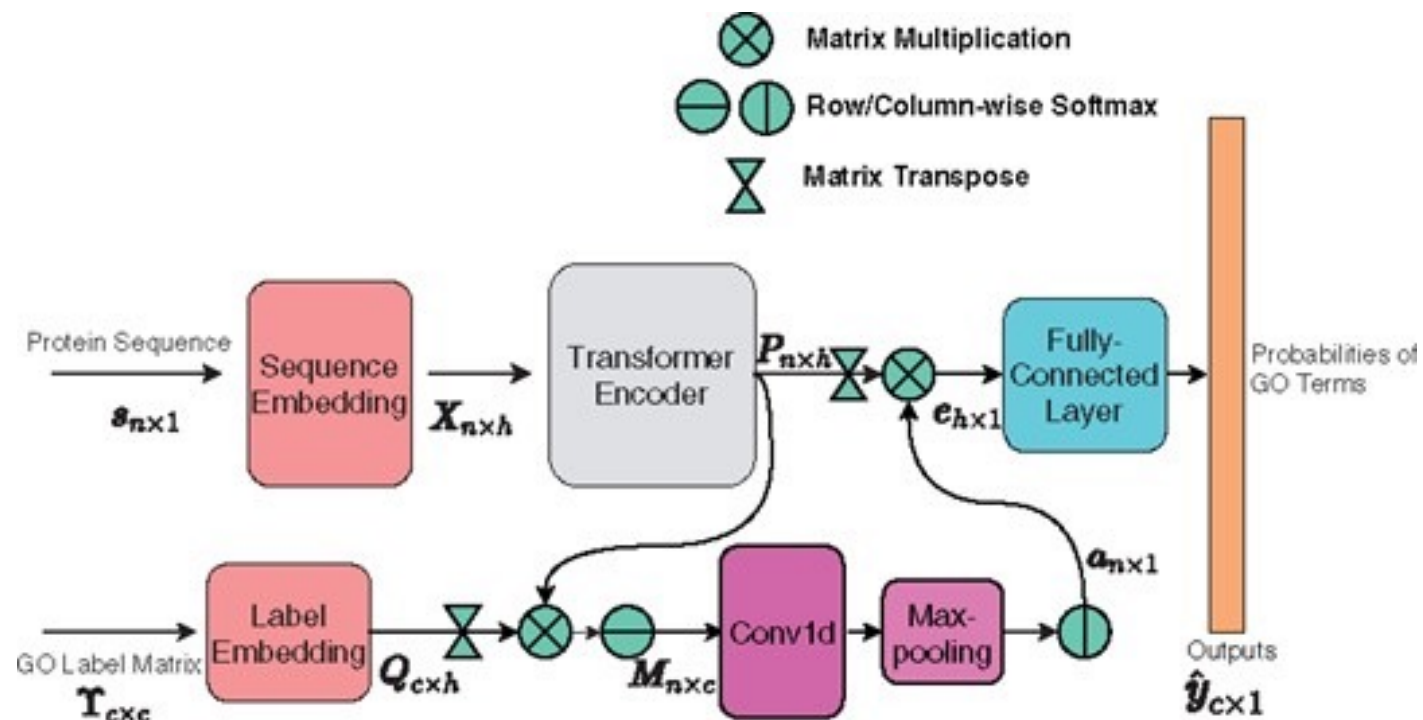
# Transformer Decoder



For generative outputs, repeatedly choose the most likely next element until the end of the sequence

0. <start>
1. <start> A
2. <start> A T
3. <start> A T T
4. <start> A T T G
5. <start> A T T G <end>

# Example: Protein Function Annotation



Transformers have applications other than sequence generation as well

**d**

Legend:  
█ Experiment  
█ Enformer  
█ Basenji2

GENCODE annotation  
CD44  
CD44  
CD44  
CD44  
CD44

DNase/CD14-positive monocyte female  
20 kb

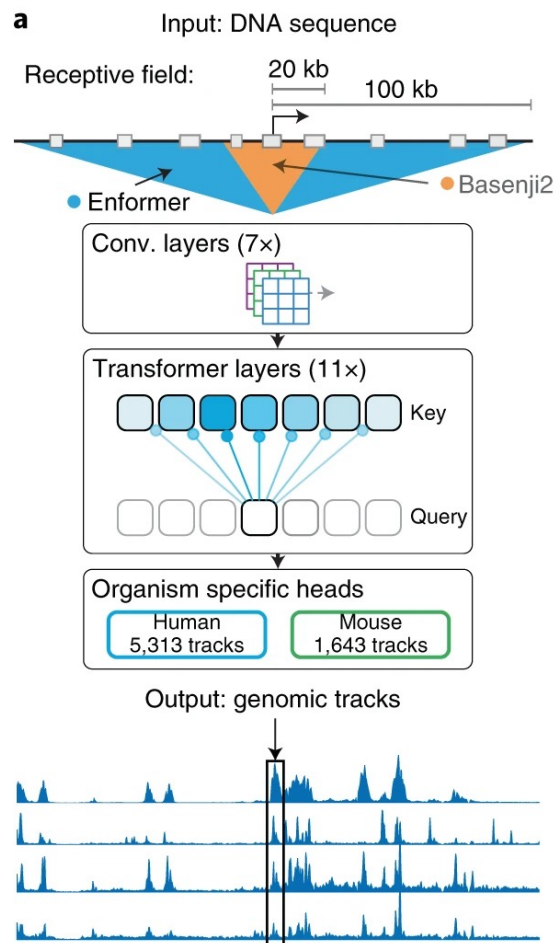
DNase/keratinocyte female

ChIP Histone:H3K27ac/keratinocyte female

CAGE/keratinocyte – epidermal

Chr11:35,082,742–35,197,430

# Example: Enhancer Prediction from DNA Sequence

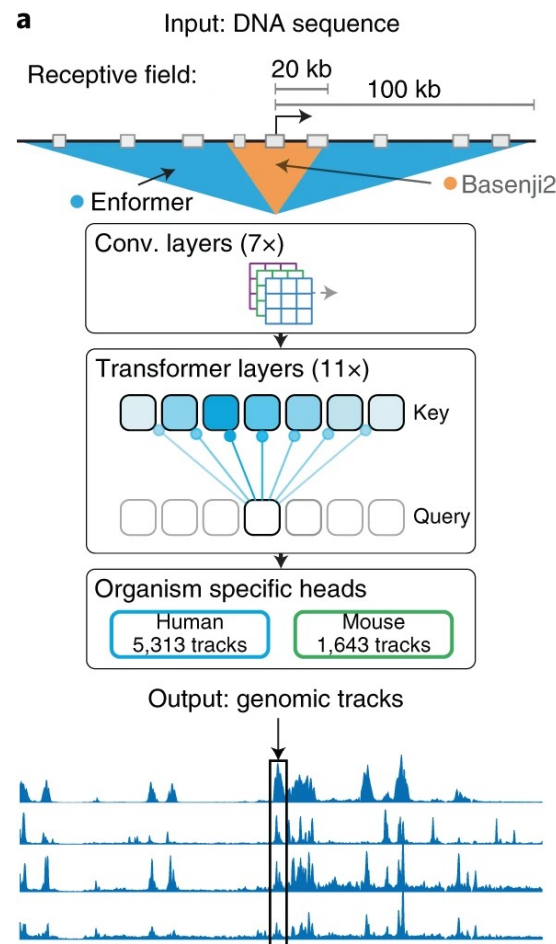


Instead of generating a DNA sequence, genomic tracks (e.g. TF binding, accessibility) can be generated instead

- Convolve 100kb to produce features for each base with attention pooling
- Feed to multiple self-attention blocks (transformer encoder)
- Apply final convolutions to predict tracks for humans or mice

# Example: Enhancer Prediction from DNA Sequence

Transformers allow for broader search regions with fewer computational limits



Predict attribute-correlated locations based on DNA sequence



# Example: Enhancer Prediction from DNA Sequence

After predicting tracks, known enhancers line up with calculated attention scores

CRISPRi Validation

Attention Scores

