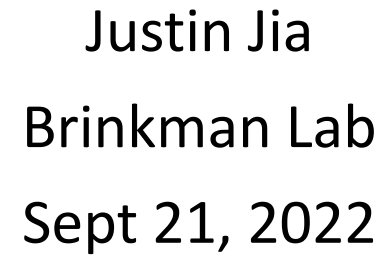
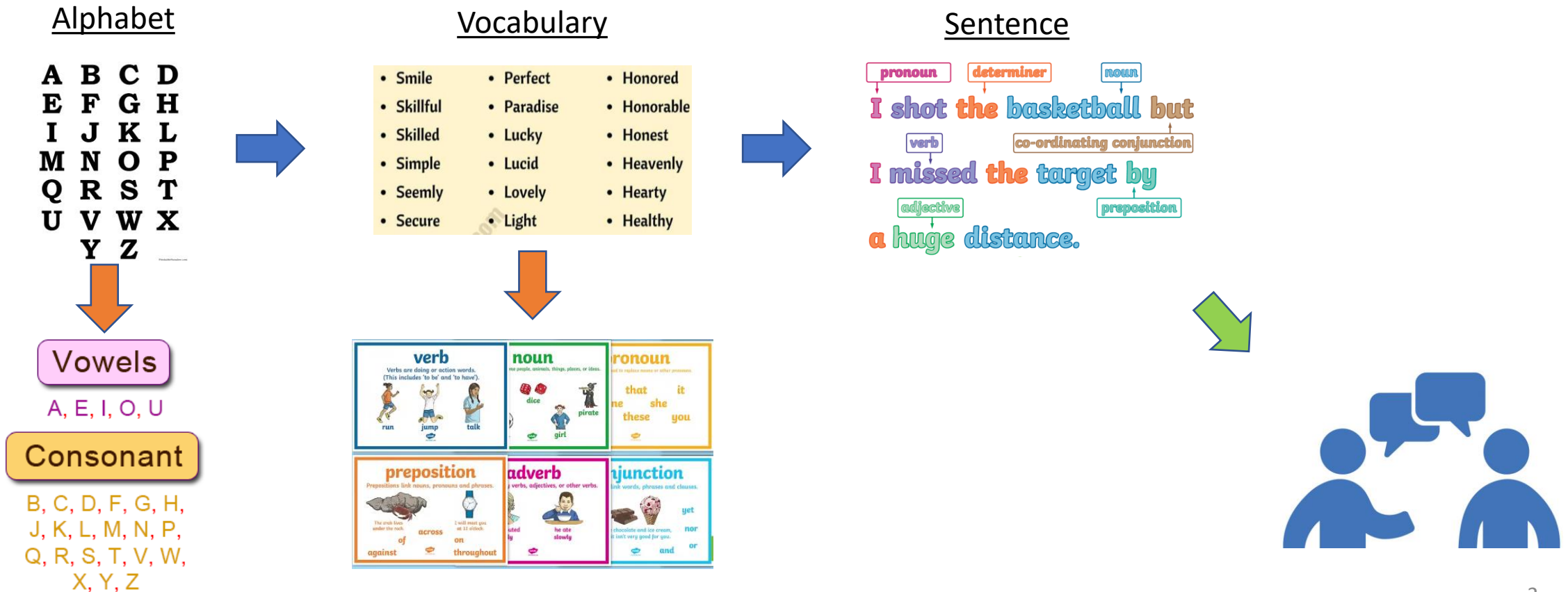


# Can Machines Understand DNA Like English?



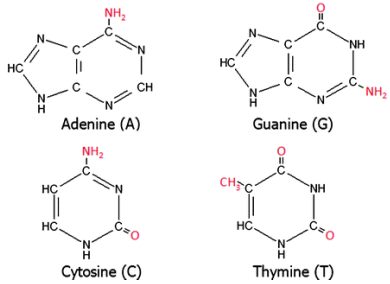
# Natural Language

- Any language that has evolved naturally in humans through use and repetition without conscious planning or premeditation



# The Language of DNA

## Alphabet



Purines/Pyrimidines

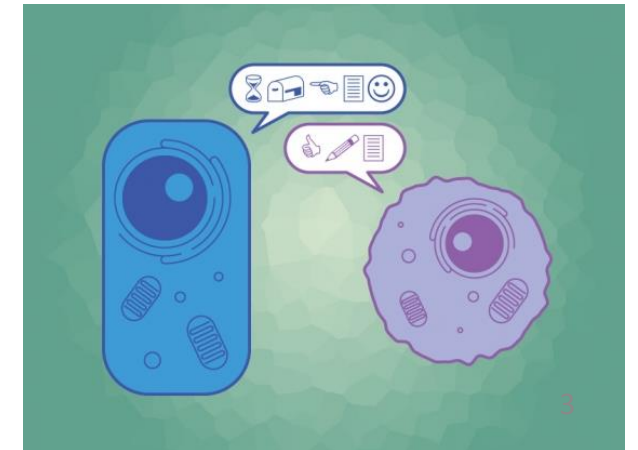
## Vocabulary

AGTTGA  
AGTTGA  
GTTGAG  
GTTGAG  
TTGAGT  
TGAGTT  
GAGTTG



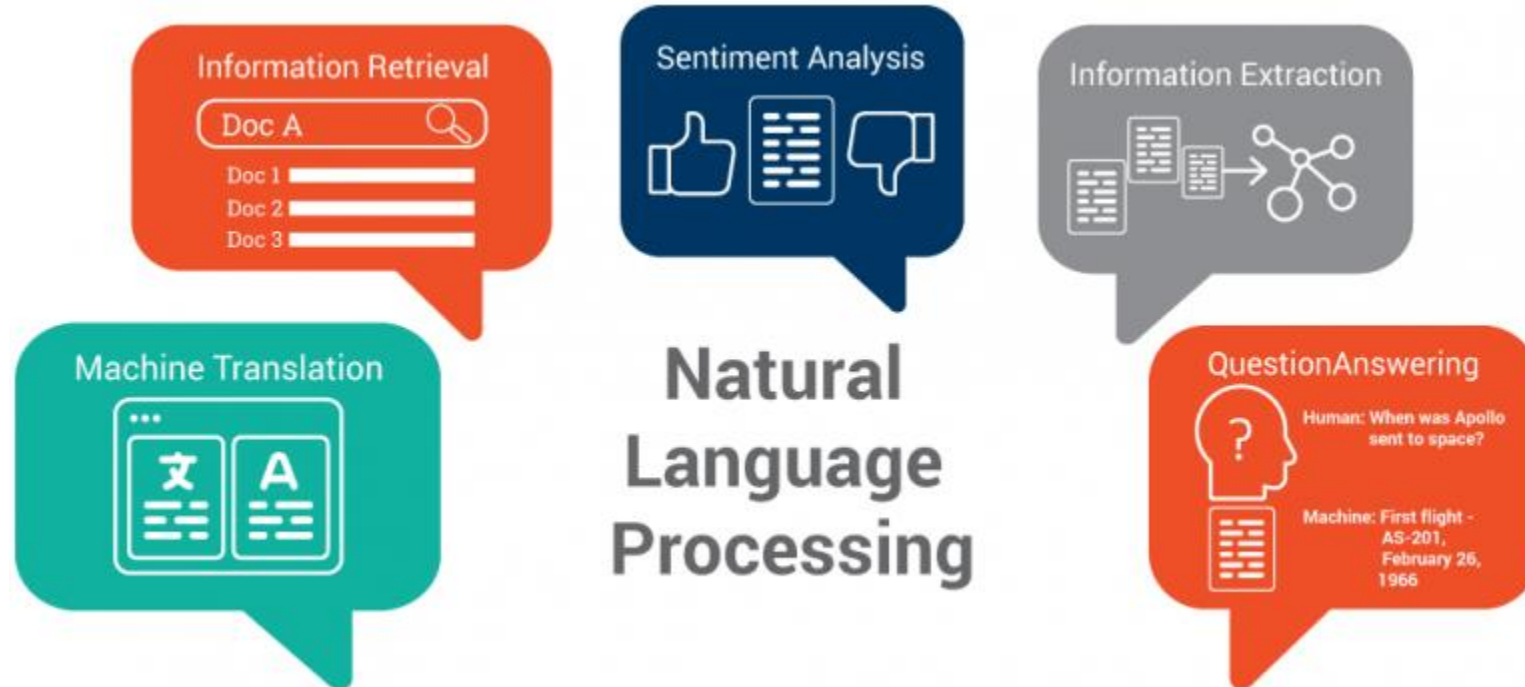
Recognition sites  
Promoters  
Start codons  
Etc...

## Sentence

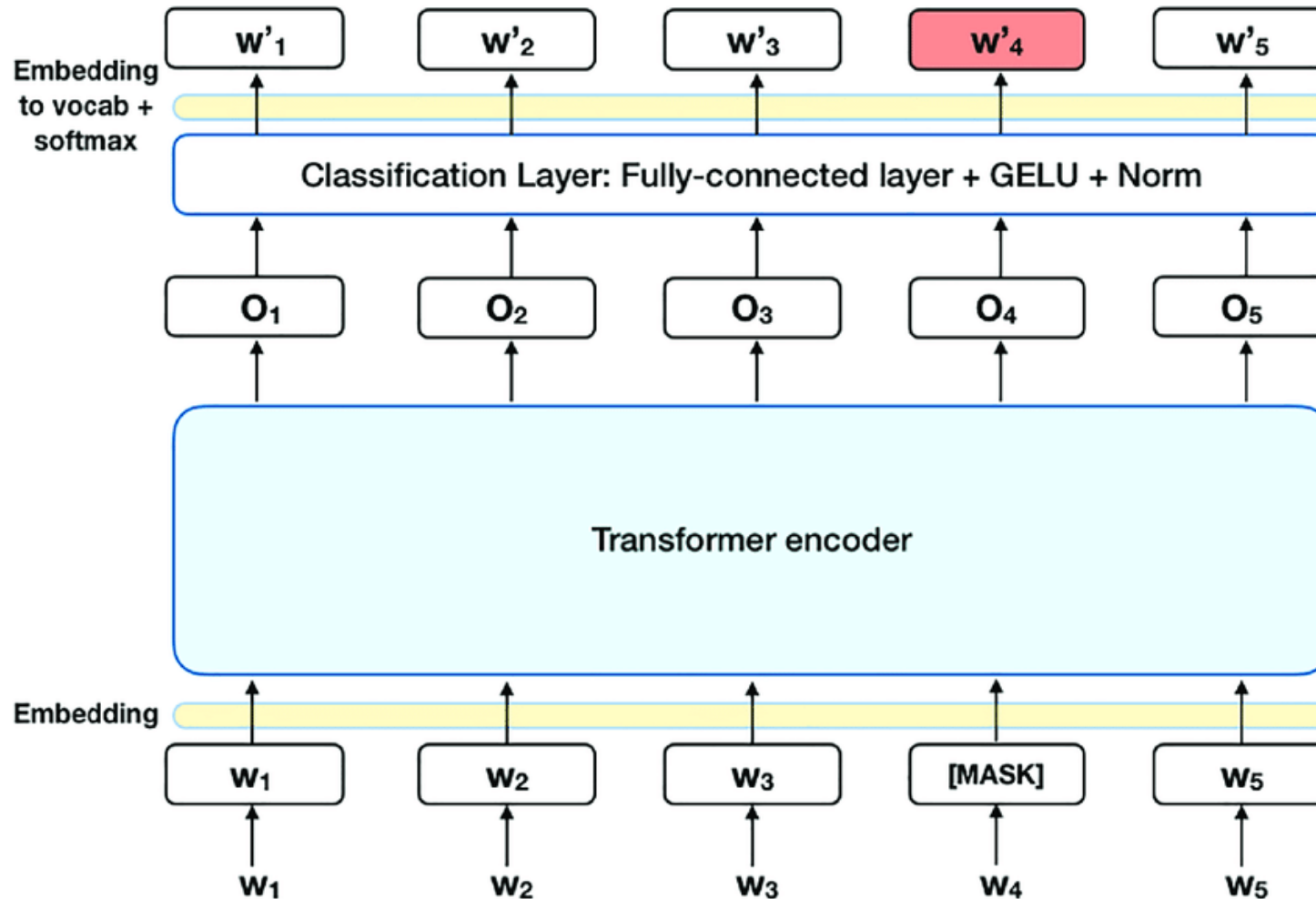


# Natural Language Processing (NLP)

- Programming computers to process and analyze large amounts of natural language data.
- The goal is a computer capable of "understanding" the contents of documents, including the contextual nuances of the language within them.

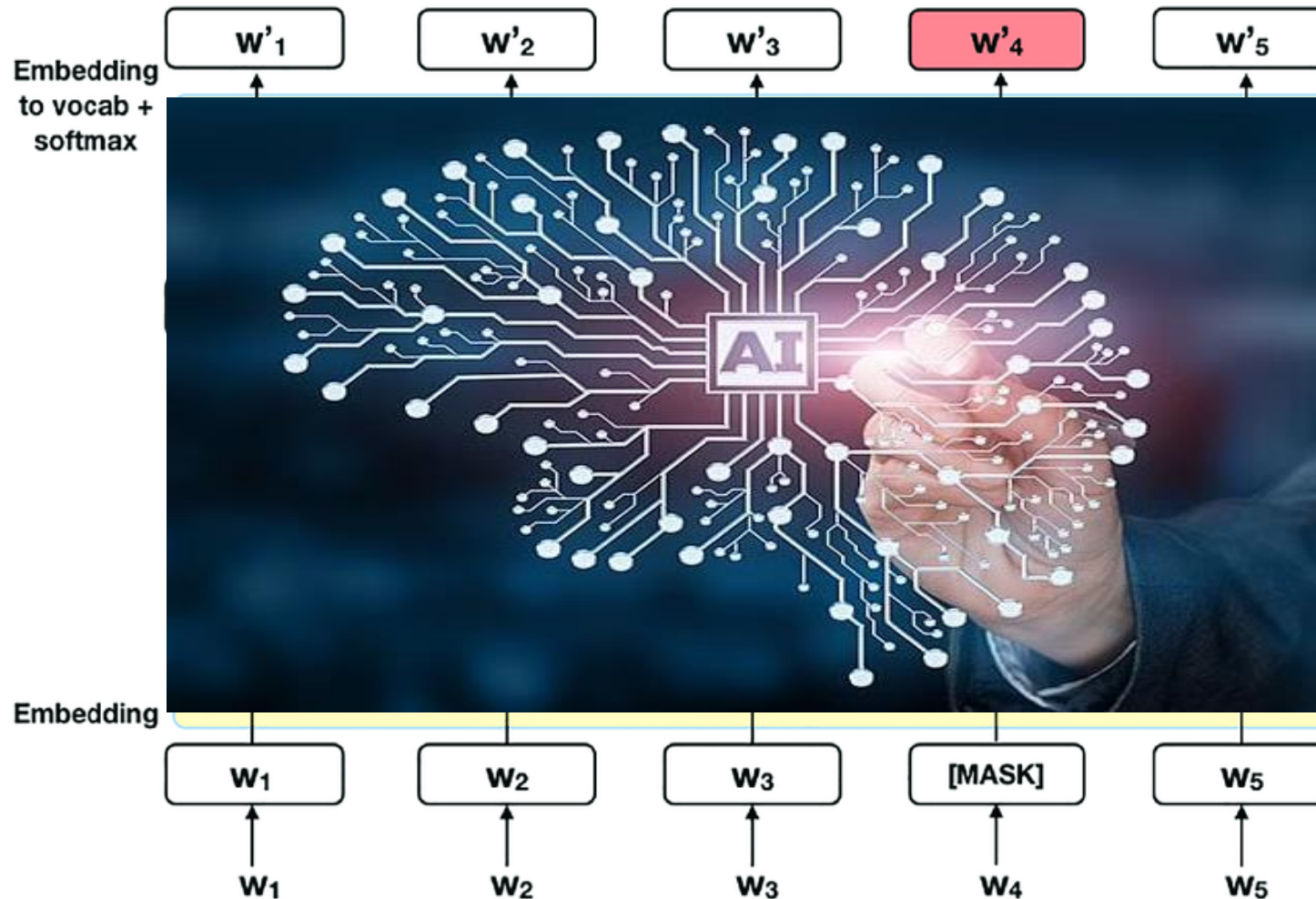


# BERT: Bidirectional Encoder Representation from Transformers



- 2019 Google's research on natural language processing
- Self-learning of natural languages
- Learns contextual relations between words in a text.

# BERT: Bidirectional Encoder Representation from Transformers



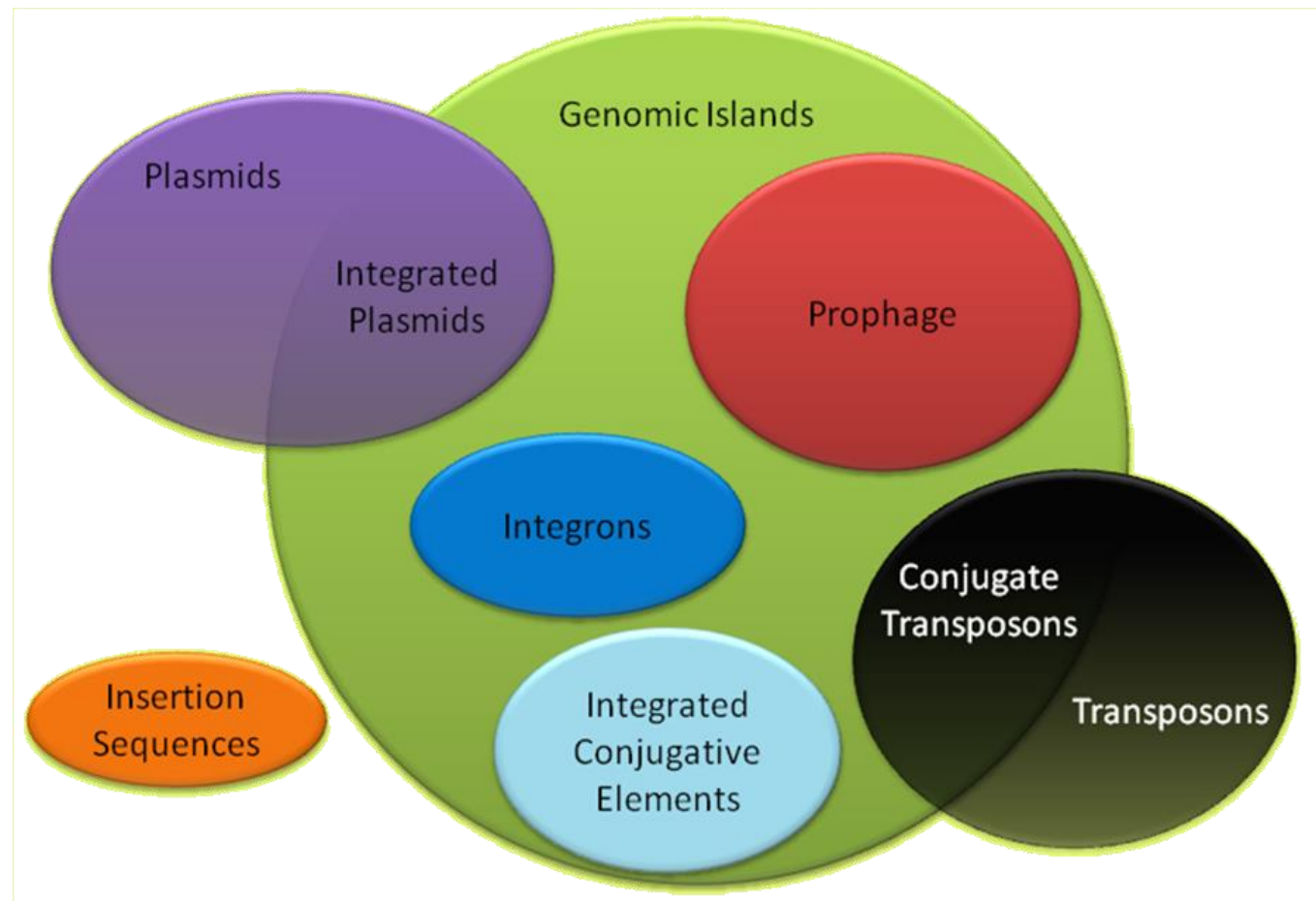
- 2019 Google's research on natural language processing
- Self-learning of natural languages
- Learns contextual relations between words in a text.

If machines can utilize NLP for human languages, can the same process be used for DNA?



# Genomic Islands (GIs)

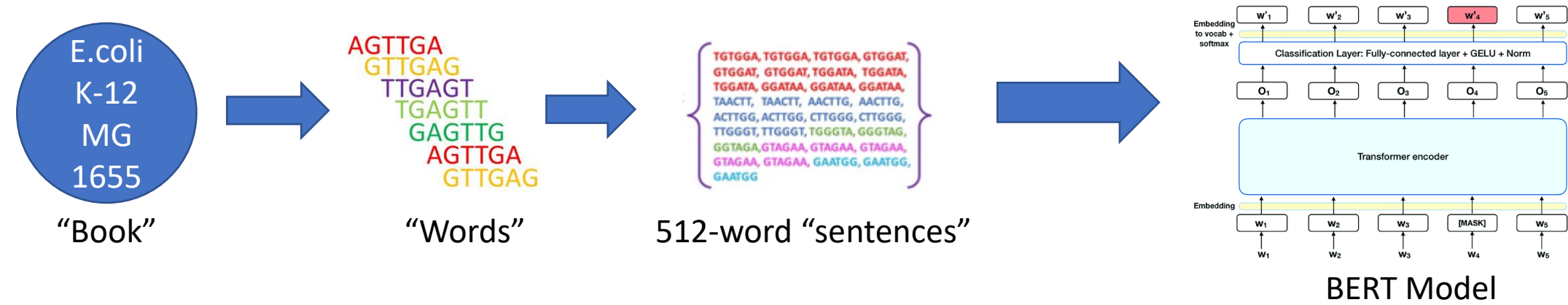
- Large segments (>8KB) of a genome that has evidence of horizontal origins





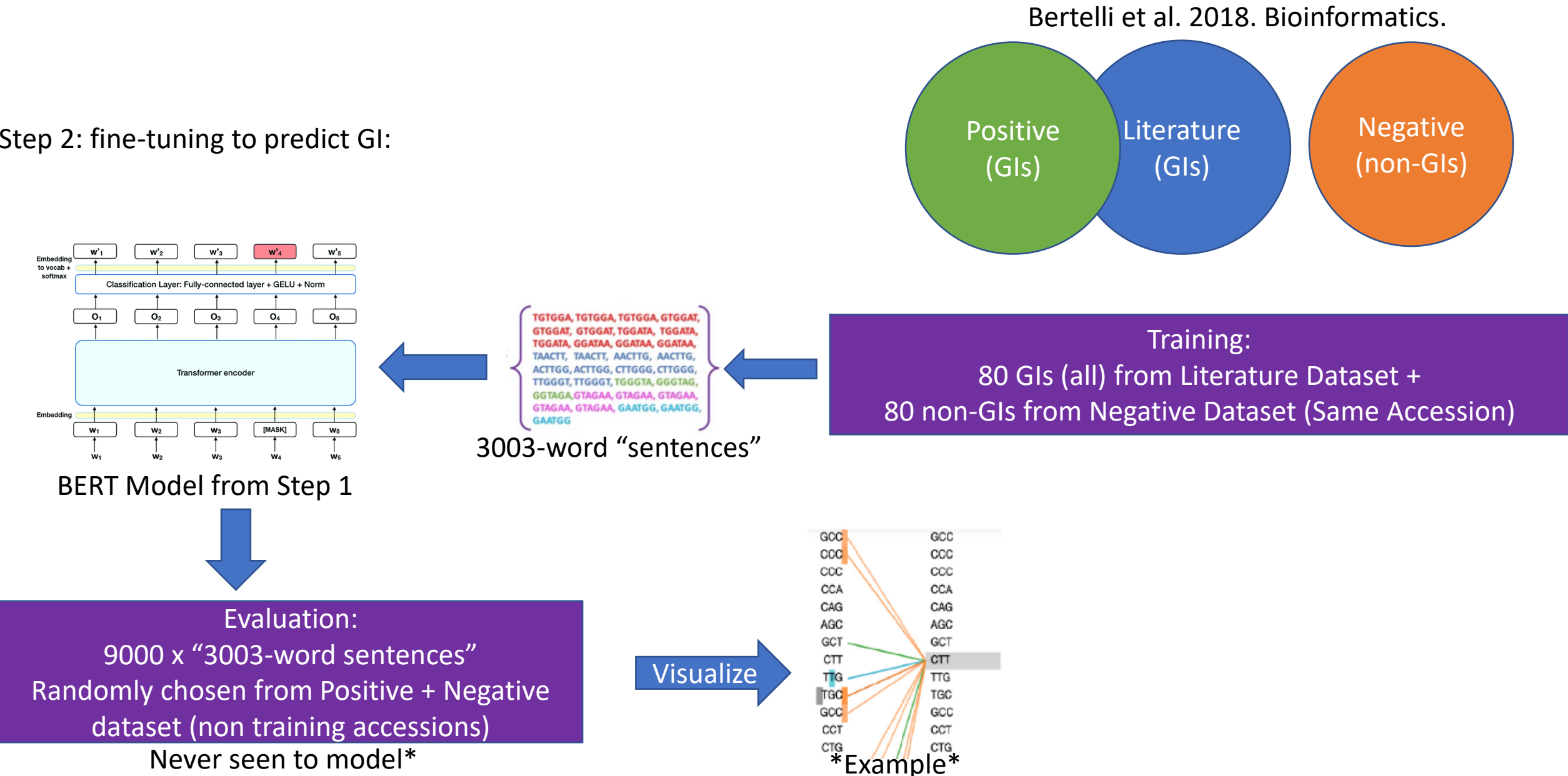
# Teaching the BERT Model the DNA Language

Step 1: Generalized self-learning of Bacterial DNA “words”:



# Predicting Genomic Islands

Step 2: fine-tuning to predict GI:



# Performance - Preliminary

Yes. NLP works on DNA

<i>Method</i>	<i>MCC</i>	<i>F-Score</i>	<i>Accuracy</i>	<i>Precision</i>	<i>Recall</i>
<i>GI-BERT Unoptimized</i>	0.48	0.69	0.71	0.77	0.71
<b><i>GI-BERT Optimized</i></b>	0.67	<b>0.82</b>	0.82	0.82	<b>0.82</b>
<i>IslandViewer4</i>	<b>0.70</b>	0.78	<b>0.89</b>	0.90	0.73
<i>SIGI-HMM</i>	0.35	0.37	0.73	0.92	0.26
<i>IslandPath-DIMOB v1</i>	0.49	0.55	0.77	0.87	0.47
<i>MTGpick</i>	0.32	0.56	0.70	0.55	0.68
<i>ZislandExplorer</i>	0.2	0.23	0.69	0.85	0.18
<i>Islander</i>	0.19	0.20	0.7	<b>0.97</b>	0.14
<i>MSGIP</i>	0.15	0.20	0.68	0.87	0.16

# Why does it work?

- Mostly picking up patterns
  - E.g. repetitive sequences that flank GIs
  - E.g. presence of certain gene sequences in a GI.
- Need in depth evaluation.

# Acknowledgements

- Brinkman lab
- Omar Nassif

