# CIS 678 Machine Learning

Document & DNA classification

- NLP
  - Machine Translation (MT)
  - Question Answering
  - Document Classification
  - Sentiment Classification
  - Document summarization
- DNA Sequencing
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  - DNA classification

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- Data/Feature encoding
  - One-Hot Encoding
  - Label Encoding

What are the challenges?

- Data/Feature encoding
  - One-Hot Encoding
  - Label Encoding
- NLP/DNA sequencing
  - Tf-idf
  - CountVectorizer

⋖	black	cat	
1	1	1	$d_1$

"A black cat"

4	black	cat	white
1	1	1	0
1	0	1	1

d <sub>1</sub>
$d_2$

"A black cat"

"A white cat"

						ını
4	black		white	įs	as	beautiful
1	1	1	0	0	0	0
1	0	1	1	0	0	0
2	1	2	1	2	2	1

 $d_1$   $d_2$   $d_3$ 

"A black cat"

"A white cat"

"A black cat is as beautiful as a white cat is"

						ul
⋖	black	cat	white	is	as	beautiful
1	1	1	0	0	0	0
1	0	1	1	0	0	0
2	1	2	1	2	2	1

 $d_1$   $d_2$   $d_3$ 

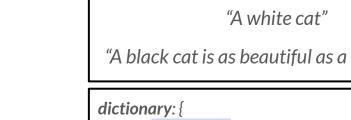
Corpus

"A black cat"

"A white cat"

"A black cat is as beautiful as a white cat is"

						iul	
⋖	black	cat	white	is	as	beautifu	
1	1	1	0	0	0	0	$d_1$
1	0	1	1	0	0	0	$d_2$
2	1	2	1	2	2	1	$d_3$



```
"A black cat is as beautiful as a white cat is"
      "a", "is", "as",
      "cat", "black",
      "white", "beautiful"
```

Corpus

"A black cat"

						ıul
4	black	cat	white	is	as	beautiful
1	1	1	0	0	0	0
1	0	1	1	0	0	0
2	1	2	1	2	2	1

 $d_1$   $d_2$   $d_3$ 

Corpus

"A black cat"

"A white cat"

"A black cat is as beautiful as a white cat is"

**Unigram:** ["a", "is", "as", "cat", "black", "white", "beautiful"]

						iul	
⋖	black	cat	white	is	as	beautiful	
1	1	1	0	0	0	0	$d_1$
1	0	1	1	0	0	0	$d_2$
2	1	2	1	2	2	1	$d_3$

### Corpus

"A black cat"

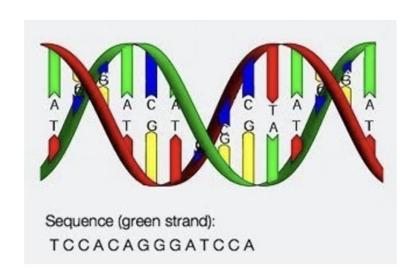
"A white cat"

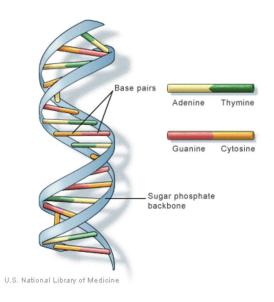
"A black cat is as beautiful as a white cat is"

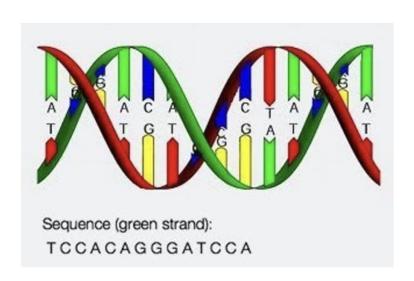
Unigram: ["a", "is", "as", "cat", "black", "white", "beautiful"] bigram: [("a", "cat"), ("cat", "a"), ("black", "cat"),

("cat", "black"), ("beautiful", "cat"), ("cat",

"beautiful"]



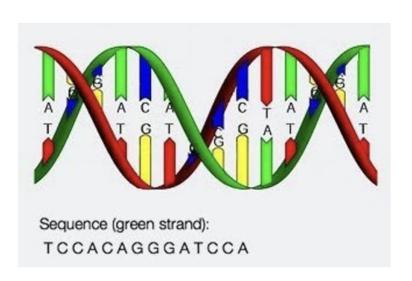




#### ENST00000435737.5

390

### ENST00000419127.5



#### ENST00000435737.5

390

#### ENST00000419127.5

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390

### ENST00000419127.5

### k-mer counting!

DNA sequence as a "language", known as k-mer counting

```
[9] def getKmers(sequence, size=6):
    return [sequence[x:x+size].lower() for x in range(len(sequence) - size + 1)]

[62] mySeq = 'GTGCCCAGGTT'
    getKmers(mySeq, size=5)

['gtgcc', 'tgccc', 'gccca', 'cccag', 'ccagg', 'caggt', 'aggtt']
```

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# Notebook presentation!