
Final

Just pick a W and Run the blue subsection

Constructing W's

To get a letter sequence go to <https://www.ncbi.nlm.nih.gov/nuccore/>
pick a gene from GenBank
go to FASTA
Copy + paste letter seq into data subsection

```
StringReplace[ToString[{StringReplace["ACGTAGTCAATT",  
  {"A" → "0,", "C" → "1,", "G" → "2,", "T" → "3,"}]}], ",}" → "}"]  
{0,1,2,3,0,2,3,1,0,0,3,3}
```

```
ToString["ACGTAGTCAATT"]  
ToString[ACGTAGTCAATT]  
ACGTAGTCAATT // ToString  
StringQ[ACGTAGTCAATT // ToString]  
StringQ[ACGTAGTCAATT]  
ACGTAGTCAATT  
ACGTAGTCAATT  
ACGTAGTCAATT  
  
True  
  
False
```

```
sample = "ACGTAGTCAATT" // ToString;  
StringReplace[ToString[{StringReplace[ToString[sample],  
  {"A" → "0,", "C" → "1,", "G" → "2,", "T" → "3,"}]}], ",}" → "}"]  
{0,1,2,3,0,2,3,1,0,0,3,3}
```

```
lettersample = {ACGTAGTCAATT} // ToString;
```

```
LetterDNAtoNum[Sample_] := ToExpression[StringReplace[ToString[  
  {StringReplace[StringReplace[ToString[{Sample}], {" " → "", " " → "", "{" → "",  
    "}" → "", "(" → "", ")" → "", "[" → "", "]" → "", ";" → "", ":" → "", "_" → "",  
    "+" → "", "&" → "", "/" → "", "." → "", "RowBox" → "", "Null" → ""}],  
    {"0" → "0,", "1" → "1,", "2" → "2,", "3" → "3,", "A" → "0,", "C" → "1,",  
    "G" → "2,", "T" → "3,", "a" → "0,", "c" → "1,", "g" → "2,", "t" → "3,"}]}]  
],  
  ",}" →  
  "}"]]
```

```

LetterDNAToNum[lettersample]
{{0, 1, 2, 3, 0, 2, 3, 1, 0, 0, 3, 3}}

numgenesample = LetterDNAToNum[lettersample];

Flatten[numgenesample][[3]]
2

lettersample = {ACGTAGTCAATT} // ToString;
LetterDNAToNum[Sample_] := ToExpression[StringReplace[ToString[
  {StringReplace[StringReplace[ToString[{Sample}], {" " → "", " " → "", "{" → "",
    "}" → "", "(" → "", ")" → "", "[" → "", "]" → "", ";" → "", ":" → "", "_" → "",
    "+" → "", "&" → "", "/" → "", "." → "", "RowBox" → "", "Null" → ""}],
  {"0" → "0,", "1" → "1,", "2" → "2,", "3" → "3,", "A" → "0,", "C" → "1,",
    "G" → "2,", "T" → "3,", "a" → "0,", "c" → "1,", "g" → "2,", "t" → "3,"}]]]
], ",," → "}]"]
numgenesample = LetterDNAToNum[lettersample];
Flatten[numgenesample]
{0, 1, 2, 3, 0, 2, 3, 1, 0, 0, 3, 3}

```

```
basepairs = ToString[Input["Paste the base pair sequence (ex: AAGCTATGG) here"]];
```

```

Wgenesample = StringJoin[ToString[Input["What Gene is this?"]], " gene"]
(*Lets us know which gene we're dealing with,
used in pdf coding later, so be sure to name it *)

```

BRCA2 mRNA Wolf gene

```

lettersample = {ACGTAGTCAATT} // ToString;

LetterDNAToNum[Sample_] := ToExpression[StringReplace[ToString[
  {StringReplace[StringReplace[ToString[{Sample}], {" " → "", " " → "", "{" → "",
    "}" → "", "(" → "", ")" → "", "[" → "", "]" → "", ";" → "", ":" → "", "_" → "",
    "+" → "", "&" → "", "/" → "", "." → "", "RowBox" → "", "Null" → ""}],
  {"0" → "0,", "1" → "1,", "2" → "2,", "3" → "3,", "A" → "0,", "C" → "1,",
    "G" → "2,", "T" → "3,", "a" → "0,", "c" → "1,", "g" → "2,", "t" → "3,"}]]]
], ",," → "}]"]
numgenesample = LetterDNAToNum[lettersample];
Export[StringReplace["GENE_genesample.txt", "GENE_gene" → Wgenesample],
  Flatten[numgenesample]]
wolf genesample.txt

```

Test of method of Construction of W

H1Avec =

```
{ {3, 0, 2, 2, 1, 3, 2, 1, 2, 3, 3, 2, 2, 2, 2, 1, 1, 3, 3, 3, 3, 3, 3, 1, 2, 1, 0, 3, 1, 1, 3, 2,
  1, 3, 3, 1, 2, 3, 1, 0, 2, 2, 3, 3, 3, 0, 3, 0, 1, 1, 0, 1, 3, 3, 3, 0, 3, 3, 3, 2, 2, 3, 2, 3,
  2, 1, 3, 2, 3, 2, 3, 3, 0, 2, 3, 1, 0, 1, 1, 0, 3, 2, 3, 1, 3, 2, 0, 0, 0, 1, 0, 2, 3, 2, 1, 1,
  3, 1, 1, 1, 2, 1, 1, 1, 1, 1, 2, 1, 1, 2, 1, 3, 3, 1, 3, 2, 1, 3, 2, 1, 3, 1, 1, 3, 2, 0,
  2, 0, 0, 0, 1, 1, 3, 3, 3, 0, 2, 1, 3, 2, 2, 1, 0, 0, 2, 0, 0, 2, 2, 1, 0, 0, 0, 2, 0, 0,
  0, 1, 1, 3, 2, 1, 3, 0, 0, 2, 2, 1, 3, 2, 1, 0, 2, 1, 0, 2, 1, 1, 3, 1, 1, 0, 0, 2, 0, 0,
  0, 0, 0, 0, 1, 1, 1, 2, 1, 3, 2, 2, 1, 1, 1, 3, 3, 1, 1, 2, 3, 2, 3, 1, 0, 2, 0, 2, 1, 3,
  2, 0, 3, 1, 2, 3, 2, 1, 0, 2, 2, 1, 3, 2, 1, 3, 3, 1, 1, 3, 1, 1, 3, 1, 3, 0, 0, 2, 2, 0,
  2, 1, 2, 3, 2, 2, 3, 2, 2, 3, 2, 3, 2, 3, 1, 2, 3, 3, 2, 2, 1, 0, 2, 1, 3, 1, 3, 3, 0, 0,
  0, 0, 0, 2, 2, 1, 2, 1, 3, 2, 2, 1, 2, 2, 1, 1, 2, 1, 0, 2, 2, 1, 3, 0, 1, 2, 0, 1, 2, 3,
  2, 2, 0, 2, 0, 0, 2, 0, 0, 1, 0, 0, 1, 0, 2, 1, 1, 2, 1, 0, 3, 3, 0, 0, 2, 1, 3, 2, 2, 2,
  1, 0, 3, 3, 0, 0, 2, 0, 2, 1, 1, 3, 2, 2, 3, 0, 0, 2, 1, 0, 0, 2, 2, 2, 0, 0, 1, 2, 3, 3,
  2, 2, 3, 2, 1, 0, 2, 0, 1, 0, 0, 0, 2, 2, 2, 3, 0, 1, 1, 2, 2, 0, 2, 1, 1, 3, 1, 2, 2, 2,
  3, 3, 1, 1, 3, 3, 1, 0, 0, 2, 1, 3, 1, 0, 0, 1, 0, 0, 2, 0, 0, 2, 2, 1, 2, 3, 1, 1, 3, 1,
  1, 2, 3, 2, 2, 0, 0, 0, 1, 1, 0, 0, 2, 1, 1, 1, 2, 2, 1, 2, 1, 1, 3, 1, 0, 0, 0, 2, 2, 3,
  2, 2, 1, 3, 0, 1, 0, 0, 0, 0, 0, 1, 3, 0, 0, 2, 2, 1, 0, 0, 1, 2, 2, 2, 3, 2, 1, 0, 3, 1,
  3, 0, 0, 0, 0, 0, 2, 1, 3, 1, 0, 0, 0, 0, 0, 2, 2, 1, 1, 0, 1, 2, 2, 2, 2, 2, 1, 3, 0, 2,
  1, 0, 0, 0, 0, 0, 2, 0, 2, 1, 2, 3, 1, 0, 0, 2, 0, 1, 3, 1, 1, 2, 0, 0, 0, 0, 0, 2, 2, 1,
  3, 0, 0, 0, 0, 0, 2, 1, 1, 3, 2, 1, 2, 2, 1, 0, 0, 1, 0, 0, 2, 2, 0, 0, 0, 3, 1, 1, 3, 1,
  1, 0, 0, 2, 0, 0, 3, 1, 1, 0, 0, 0, 0, 0, 0, 1, 1, 1, 0, 0, 0, 0, 1, 3, 2, 3, 0, 0, 0,
  2, 1, 1, 1, 0, 0, 2, 0, 0, 0, 2, 3, 0, 2, 1, 3, 0, 0, 0, 0, 2, 1, 1, 1, 3, 2, 1, 3, 0,
  0, 0, 2, 1, 3, 0, 0, 2, 2, 1, 3, 2, 3, 0, 0, 0, 0, 1, 1, 1, 0, 0, 2, 2, 1, 2, 2, 1, 1,
  0, 0, 2, 2, 1, 3, 0, 2, 2, 2, 3, 2, 0, 1, 2, 0, 0, 2, 1, 1, 0, 0, 0, 2, 0, 1, 3, 2, 1,
  1, 0, 0, 0, 1, 1, 1, 0, 0, 2, 0, 0, 0, 2, 1, 2, 2, 1, 0, 1, 1, 1, 0, 0, 2, 0, 0, 0, 0,
  0, 2, 3, 0, 0, 0, 3, 3, 1, 0, 2, 3, 3, 0, 2, 0, 0, 2, 3, 3, 3, 1, 3, 3, 1, 3, 0, 2, 3,
  0, 0, 1, 1, 1, 0, 0, 1, 2, 2, 1, 3, 1, 3, 3, 3, 3, 0, 0, 2, 0, 2, 1, 1, 0, 1, 1, 3, 0} };
```

```

H1AW =
{{3, 0, 2, 2, 1, 3, 2, 1, 2, 3, 3, 2, 2, 2, 2, 1, 1, 3, 3, 3, 3, 3, 3, 3, 1, 2, 1, 0, 3, 1, 1, 3},
 {2, 1, 3, 3, 1, 2, 3, 1, 0, 2, 2, 3, 3, 3, 0, 3, 0, 1, 1, 0, 1, 3, 3, 3, 0, 3, 3, 3, 2, 2, 3, 2},
 {3, 2, 1, 3, 2, 3, 2, 3, 3, 0, 2, 3, 1, 0, 1, 1, 0, 3, 2, 3, 1, 3, 2, 0, 0, 0, 1, 0, 2, 3, 2, 1},
 {1, 3, 1, 1, 1, 2, 1, 1, 1, 1, 1, 2, 1, 1, 2, 1, 3, 3, 1, 3, 2, 1, 3, 2, 1, 3, 1, 1, 3, 2, 0, 2},
 {0, 0, 0, 1, 1, 3, 3, 3, 0, 2, 1, 3, 2, 2, 1, 0, 0, 2, 0, 0, 2, 2, 1, 0, 0, 0, 2, 0, 0, 0, 1, 1},
 {3, 2, 1, 3, 0, 0, 2, 2, 1, 3, 2, 1, 0, 2, 1, 0, 2, 1, 1, 3, 1, 1, 0, 0, 2, 0, 0, 0, 0, 0, 1},
 {1, 1, 2, 1, 3, 2, 2, 1, 1, 1, 3, 3, 1, 1, 2, 3, 2, 3, 1, 0, 2, 0, 2, 1, 3, 2, 0, 3, 1, 2, 3, 2},
 {1, 0, 2, 2, 1, 3, 2, 1, 3, 3, 1, 1, 3, 1, 1, 3, 1, 3, 0, 0, 2, 2, 0, 2, 1, 2, 3, 2, 2, 3, 2, 2},
 {3, 2, 3, 2, 3, 1, 2, 3, 3, 2, 2, 1, 0, 2, 1, 3, 1, 3, 3, 0, 0, 0, 0, 0, 2, 2, 1, 2, 1, 3, 2, 2},
 {1, 2, 2, 1, 1, 2, 1, 0, 2, 2, 1, 3, 0, 1, 2, 0, 1, 2, 3, 2, 2, 0, 2, 0, 0, 2, 0, 0, 1, 0, 0, 1},
 {0, 2, 1, 1, 2, 1, 0, 3, 3, 0, 0, 2, 1, 3, 2, 2, 2, 1, 0, 3, 3, 0, 0, 2, 0, 2, 1, 1, 3, 2, 2, 3},
 {0, 0, 2, 1, 0, 0, 2, 2, 2, 0, 0, 1, 2, 3, 3, 2, 2, 3, 2, 1, 0, 2, 0, 1, 0, 0, 0, 2, 2, 2, 3, 0},
 {1, 1, 2, 2, 0, 2, 1, 1, 3, 1, 2, 2, 2, 3, 3, 1, 1, 3, 3, 1, 0, 0, 2, 1, 3, 1, 0, 0, 1, 0, 0, 2},
 {0, 0, 2, 2, 1, 2, 3, 1, 1, 3, 1, 1, 2, 3, 2, 2, 0, 0, 0, 1, 1, 0, 0, 2, 1, 1, 1, 2, 2, 1, 2, 1},
 {1, 3, 1, 0, 0, 0, 2, 2, 3, 2, 2, 1, 3, 0, 1, 0, 0, 0, 0, 0, 1, 3, 0, 0, 2, 2, 1, 0, 0, 1, 2, 2},
 {2, 3, 2, 1, 0, 3, 1, 3, 0, 0, 0, 0, 0, 2, 1, 3, 1, 0, 0, 0, 0, 0, 2, 2, 1, 1, 0, 1, 2, 2, 2, 2},
 {2, 1, 3, 0, 2, 1, 0, 0, 0, 0, 0, 2, 0, 2, 1, 2, 3, 1, 0, 0, 2, 0, 1, 3, 1, 1, 2, 0, 0, 0, 0, 0},
 {2, 2, 1, 3, 0, 0, 0, 0, 0, 2, 1, 1, 3, 2, 1, 2, 2, 1, 0, 0, 1, 0, 0, 2, 2, 0, 0, 0, 3, 1, 1, 3},
 {1, 1, 0, 0, 2, 0, 0, 3, 1, 1, 0, 0, 0, 0, 0, 0, 0, 1, 1, 1, 0, 0, 0, 0, 1, 3, 2, 3, 0, 0, 0, 2},
 {1, 1, 1, 0, 0, 2, 0, 0, 0, 2, 3, 0, 2, 1, 3, 0, 0, 0, 0, 2, 1, 1, 1, 3, 2, 1, 3, 0, 0, 0, 2, 1},
 {3, 0, 0, 2, 2, 1, 3, 2, 3, 0, 0, 0, 0, 1, 1, 1, 0, 0, 2, 2, 1, 2, 2, 1, 1, 0, 0, 2, 2, 1, 3, 0},
 {2, 2, 2, 3, 2, 0, 1, 2, 0, 0, 2, 1, 1, 0, 0, 0, 2, 0, 1, 3, 2, 1, 1, 0, 0, 0, 1, 1, 1, 0, 0, 2},
 {0, 0, 0, 2, 1, 2, 2, 1, 0, 1, 1, 1, 0, 0, 2, 0, 0, 0, 0, 0, 2, 3, 0, 0, 0, 3, 3, 1, 0, 2, 3, 3},
 {0, 2, 0, 0, 2, 3, 3, 3, 1, 3, 3, 1, 3, 0, 2, 3, 0, 0, 1, 1, 1, 0, 0, 1, 2, 2, 1, 3, 1, 3, 3, 3},
 {3, 0, 0, 2, 0, 2, 1, 1, 0, 1, 1, 3, 0, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4},
 {4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4},
 {4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4},
 {4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4},
 {4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4},
 {4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4},
 {4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4},
 {4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4}};

```

```
numgenesample = H1Avec;
```

```

M = numgenesample;
lengthvec[M_] := Length[M[[1, All]]]
For[npow = 1, npow < 1000, npow++, If[lengthvec[M] < (2^(npow)), Break[]];
  FilledSize = 2^(npow + 1)];
Filler[M_] := Table[4, {i, 1, FilledSize - lengthvec[M]}]
FilledVec[M_] := Join[Flatten[M], Filler[M]]
lengthvec[M_] := Length[M[[1, All]]]
Filler[M_] := Table[4, {i, 1, FilledSize - lengthvec[M]}]
FilledVec[M_] := Join[Flatten[M], Filler[M]]
For[npow = 1, npow < 1000, npow++, If[lengthvec[M] ≤ (2^npow), Break[]]];
(* gives npow such that 2^npow > lengthvec[M] > 2^(npow - 1) *)
FilledSize = 2^npow;
FilledM = FilledVec[M];

```

```

npow
Length[FilledM] (*Should be an (even) power of 2*)
numrowsW =  $\sqrt{\text{Length[FilledM]}}$ 
10
1024
32

(*Table[FilledM[[i]],{i,1,numrowsW}]
Table[FilledM[[i]],{i,33,2*(numrowsW)}]
Table[FilledM[[i]],{i,(2*(numrowsW))+1,3*(numrowsW)}]*)

testWW = Table[Table[FilledM[[i]],
  {i, ((j - 1) * (numrowsW) + 1), (j * (numrowsW))}], {j, 1, numrowsW}];

Print["so this construction is valid if and only if the following: ",
H1AW - testWW // Union // Flatten // Union ,
" is exactly {0}, if not something is wrong"]

```

so this construction is valid if and only if the following:
 {0} is exactly {0}, if not something is wrong

W for Wolf BRCA2 mRNA

Data

Source: GenBank: AB043895.5

```

(*lettersample={}/ToString;*)

lettersample = {AAAGAAGGTCGGCGGAGGCGGAGGCGGAGCTGCTGGGGCTTGGCGCTCTGGAAGTCGTCCAGCCGCGGG ×
TCGCCGAGGAAAGGAGCCTGCGGGTCAGCTTTCTGGCCGAAGTGCCGGCGGAATTTGTAGCCGTCTCC ×
GGCCAAAAAGAGCGGCACCTCGGAAGGCGAGTTATTTACCAAGCACTGGAGTAATATTGTAGATAAAAAT ×
GCCTGTTGGATGCAAGAGAGGCCAACATTTTTTGAATTTTAAAGACGCGGTGCAATCAAGCAGATTTA ×
GGACCAATAAGCCTTAATTGGTTGAAGAACTTTCTTTAGAAGCTCCACCCTATAATTCTGAACCCACAG ×
AAGAATCTGGTTATAAAATCAGCTATGAACCAACCTATTTAAACACCACAAAGGAAACCTTATAATCA ×
GTTGGCTTCAACTCCAATAGTATTACAGAGGCAATATACCAACAATCTCCTTTAAAGAATTAGATAAA ×
TACAGATTAGATTCAGGAAAGGATATTACCGATAGTAAACATAAAAGTTGTTGCACAAATGAAGCTAAAA ×
TGGATCGAGCAAAATGATGTTACCAGCCACCTCTAAATTCTTATCTTAGTGAAAGTCCTGTTCTACGATG ×
TACACATGTAACACCACAGAGAGAAAAAGTCGGTGGTATGTGGAAGCTTATTCATACACCGAAGCTTATG ×
AAGGGTCAGACACCAAAACGCATTTCTGAAAGCTAGGAGCTGAGGTGGATTCTGATATGTCTTGGTCAA ×
GTTCTTTAGCCACACCACCAACCCTTAGTTCTACTGTGCTAATAGTCAGAGATGAAGAAGTATCTGCAGC ×
TGTAATTCCTAATGACACTACTGTATTTTTTAAAGCTGTTTTTCTAACCATGATGAAAGCTGAAGAAA ×
AATGATCGGTTTATCCCTTGTGGTCCAGGCAAAGAAAACAAAATCAAAGGGAAGCTAAAAGTCAAAGTT ×
TGGGGAATTCATTTGGTAAAGTAAATAGCACCAAAGACCATTTGTAAAGTCTACACCAAATGTCCTAGA ×
GGATGAAGTACATGAAAAAGTTCTAGATGTTTCTGAAGAAGAAGATAGTTTTTCATTATGTGTTCTCTAAA ×
TATAAAACAAGAAATCTACAAAAATAAACTAGCAAACTAGGAAAAATATTTTAAATGAGACAAAAA ×
CCAGTGAATGTGAAGAAGCTAAAAAGCAAATGAAAGAAAATAAACATTCATTGGTATCTGAAATGGAACC ×
AAATGACAGTCATCCATTAGATTGGAATGTAACACATGAGAAGCCCTTTGGGAATGGAAGTACAAAAATC ×
TCCAAGGAAATTGTACTGTCTTCAGCCTCTGGATGTTCTGACCTAACCTCTCAAGTCTAAATGGAGCTC ×

```

AGATGGAGAAAAACACCTCTATTGCATACTTCTTATGACCAAAATAATTGAGAAAAAGACCTCATAATCAC ×
AGATAAAGAATGCACCAACTTCATTACTTTGGAAAATCTTGGCCACAGATTTCAAATGTACCAAAGTAT ×
TCAGAGAAGACGTTAAATGAGGAAATAGTAGTAAATAAGATAAACGAAGGGCAGTGTCTTGAATCTCATG ×
AAGATTCCGTTGTTTCGGTAAAGCAAGCAATATATGAACTACTTTAATAGCTTCTCCACTTCAGGGTAT ×
CAGAAAGTCTATATTCAGGATAAGAGAATCACCTGAAGGGATGTCCAATGCAATGTTCTCAAATAATATG ×
ACTAATCCAACTTTAAAGAACCTGAAGCCTCTGAAAGTGGATTGGAAAAACATACTATTTGCTCTCAGA ×
AAGAGGATTCTTTATGTACAAGTTCAATTGATGATGGAAGCTGGCCAGCAACTATCAAACATACTTCTGT ×
AGCTTTGAAGAATTTAGGTTTAAATATCTAGTTTGAAAAAGAAAAAGTTTATTTACGTTATAAAT ×
GATGAAACATCTAATCAAGGCCGTGAAAACACAGAAAGACCAAGAGTCAAGACTAATTAACCTTTGACCC ×
AATTTGAAGCAAATGCTTTTGAAGGACCCCTGACATTTACAAATGCTGATTGAGGTTTATTGCATTCTTC ×
TTCCATCAAAAAAAGTGTTCAGAAATGACTCAGAAAAACCAGCTTTGTCTTTAACCAGCTCTTTTGGG ×
ACAATTCTGAGAAAAGTTTCCAGTAATGGAGCCAGTTCTCCTAATAATAAAATAATATCTCAGGATCCTG ×
ATTATAAAGAAGCAAAAATTAATAAGAAAAAATTGGAGTCATTTATAACCACAGAAACTGATTGTCTGTC ×
ATCCCTGCAGGAAAAACATTGGGAAGATGATGCAAAAAACAAAGAGTTTCAGATATAAAAGAAAAAGTC ×
TTGCCTACAGTAAGTCACCTCCTGTGCCACATTGAGAAGTGAAGGTAGTGATATTCATTTTCACTCTC ×
CAGAAAGCTTTTTCATTGACTGTGATAATACCAGTCTGTTAACTCCTAGCTCTAGGGATTCTCCATCAAG ×
CCTAGTTGTGATGTCTAGAGGAAAAAGAAATCATATAAAATATCAGAGAACTAAAATGTAAGAATCATGAA ×
ACTGTTTTTGAATTAACCAAAAATATTCCCATGGAAGAAGTCAAGACATACATGTTTTAAATGCAGATT ×
CTAAAAATGCTAAACTGTTGTCAACTGAAAAACATATAACAGTAGCATCATCTTCAGTAAAGGTTCAAGT ×
CAACCAAAATGCAATCTCACCACAATCCAAAAAGACCAAAAAGAACTACTTTAATTTCAAAAAATACT ×
GTTAATCCAACTCTGAAGAATTTTCCAGATGATGAAAATAATTTTGTCTTAAAGATAACTAATGAAA ×
GTAATACTCCTGTTTTAGGAAATACTAAGGAACTACATGATTCAAACCTCTGTTGTGTAAGAGATTCTGT ×
TCCTAAGAAGTCTACCATGGTAGTATGTACAGACCTGGATGACAAACAAACAGCCAAAGTGTGATTATG ×
AAAGATTGTTATTCATCAAGCATAGATGATCTTACAGAAAGGAACAGAAGTACCATAAAGCAACAATAA ×
AAATGACTCTAGATCAAGATTCAAAATCAGACATTACCTCAGATATAGTTAGGAAATCAAATGGAAACAG ×
TGATTATATGGATAATTGGGCAAGACTGTCTGATCCAATTTCAAATCACAGTTTTGAAATGGCTTCAAA ×
ACAGCTTCTAATAAAGAGATAAACTCTCTGAAAACAACATTAGGAAAAGTAAAATGCTTTTCAAAGATA ×
TTGAGGAACATTATCCTACTAAGTCTAGCATGTCTTGAATTTGTAATACTTCATCATTAGAAAAGTCAAAA ×
GAAACCAAGCAAATCTCATGCACTTGATCCACAGTCAATTAATATCATATCTGGGTTTGTGCAGAATAGC ×
ACATATGTTTTCTGATAGTGAAAGTGGTCACACAGCTCCTCCAACTTTATCTTTAAAGCAAGATTTTGATT ×
CAAAATCGTAATTTAACTCCTAGTCAAAAGGCAGAAATTACAGAACTTTCTACTATTTTGAAGAATCAGG ×
AAGCCAGTTTGAATTTACACAGTTTAGAAAACCAAGCCACATAATACAGAAAAATCCATTTGAAATGCCT ×
GAAAACAGCTGACTATCTTGAATAGCACTTCTAAGGAATGGAAAGATGATGATCTTCATCTCACAACTA ×
ATGCTCCATCTATCAGTCAGGTAGATAGCAAGAAATCTGAAGGTATAATTGGAGGTAAGCAGAAGTTTGC ×
TTGCTTGTCAAGAACCAGCTGTAAACAGAAGTGCTTCTGGCTATTCAACAGATAAAAATGAAGTGGAGTTT ×
AGAGGCTTTTATTCTGCTCGTGGCACAAAAGTGAATGTTGGTAGTGAAGCATTGCAAAAAGCTAAGAAAC ×
TGTTTCAGTGACCTTGAGAATATCAATGAGGAACTTCTGTAGAAGTAGATAGAAGTTTCTCCTCAAGCAA ×
ATACAATGATTCTGTCTCAATGATTGAGATAGAAGATTGTAATGATAAAAAATTTAAATGAGAAAAATAAT ×
AAATGCGCGCTAATACTACAAAATAATATTGAAATGACTACTGACATTTTGTGGAAGAATATACTGAAA ×
GTTACAGGAGAAAATACAGAAAATGAAGGTAACCAATGTACTGACGCTGGTAGAAAATCTTGTAACCTCAGA ×
ATCTGATGGCAGTGATTCAAGTAAAAATGATACAGTTTATATTCATGAAGAAGAAAATGGCTTGCCCTGT ×
ATTGATCAGCACACATAGATCTGAAATTATTTAGCCAGTTTATGAAGGAGGGGAACACTCAAATTAAG ×
AAGGTTTGTGAGATTTAACTGTTTGAAGTTATGAAAGCTGAAGAAACATCTCATGTTACTATGTCAAA ×
TAAACAGCAGTTAACAGCTAATACGGGGCAAAACATAAAAGATTTTGACACTTTTTATTTATCCTTTCAG ×
ACTGCAAGCAGAAAAAATAAAGGCTCTCCAAAGAGTCATTAATAAAGCTAGAAGTCTCCTTAATCAAA ×
AATGGACAGAAGAAGAATTAATAACTTTTCAGATTCTTGAATTCTGAATTACTTCTGGCATAGATAT ×
CAAGAAAACAGACATCTCAAATCATGAGGTAAAGAAAATACTGAAAGAAAAGACAAAATAACGAAAGAA ×
AGTGACCTAATTGGTACTGAAAATATATTACTGATCTGCAGCAAGACCAGAAAGTAAAATAAAAAAGA ×
TCAAAGAATCTGCTGTGTTGGGTTTTATACAGCTAGTGGGAAAAAATAGAAATTACAAAGGAATCTTT ×
GGACAAAGTAAAAATCTTTTTGAAGAAAAAGAGCAAGATAATAGTGAAATCACTAATTTTAGCCATCGA ×
GGGGCAAAGATGTCCAAGGACAGAGAAGAATGTAAGATGGGCGTGAATTAGCTTGTGGGACAACTGAAA ×

TAACAACTACCCAGAGTATGAAGAACTCACAGTTCTCTAGAGAAGAAAAAATTGTTTCTAATGAGAT ×
 TGCAGCCTTAAGACCCAGGCTCTTAAGTGATAATTTATACAAACAACTGAAAATCTTAAATATCAGAT ×
 CATGCCTCTCAGAAAGTTGATGTACATGAAAATACAGAAAAAGAAACAGCAAAAAAGCCTACAATGTATA ×
 CAAATCAATCCACTTATTCTAGCCATTGAAAACCTACCTTTAACATTTTACACAGGACACGGAAGAAAAAT ×
 TTCTGTGAGTGAGGCTTCACTATTTGAAGCAAAAAAATGGCTTAGAGAAGGAGAATGGGATGATCAATCA ×
 GAAAGAATAAATGCTGCCAAGGTTAACTGCTTAAAGAATATCCTGATGATTACGTAGAAAATCCTTCAT ×
 GTGGAAATAGTTCAAATAGTGCCATAACTGAAAATGACAAAAATCATCTCTCTGAAAAACAAGGCTCAAC ×
 TTATTTAAGTAATAGTACCATGTCTAACAGCTATTCATACCATCCTGGCTTTTGTCTATTCTAGTGAAGTG ×
 TATAATAAATCAGAATATCTTTCAAGAAGTAAATTTGATAATTTGGTATTGAACCAGTAATAAAGAATA ×
 TTAGAGAGAGAAAAACATTGGTTTTCTGAAATAATGTCCCCTGGAAGAGAAGCAGACACAGACCCACA ×
 AAGTGTAATGAAGATATTTGTGTTGAGAACTTGCGACTAACTCTTCATGCAAAAATAAAAATACAGCC ×
 ATTAAGTGGCCATATCTGACTCAAATAATTTAATACAATTCAAAAGTTGAATTCTGATTCAAATAATT ×
 CTGTACCTGCATACAGTACAGTAAATAGTAAAGAGTCTTTGTTGCACACCAGACAAAAGTGACAGAGGG ×
 GTTTACAGACAACTGCAGCATGGTAACTAAACAAAACACCAAGAGTAAATCAGACACTTGCCATGCAGAA ×
 ATTGTGGCAGATTATCCTAAGGCACTGGATGATTAGAGGCTATTTTTCTAACTCTCTGGGTGCTATAG ×
 AATGTTACCTTCACATAAGGTTTTTGTGACATTCAAAGTGAACAACTTCACAACTTAACCAAAGTAT ×
 GTCTGGATTGGAGAAAGTTTCTGAAACACCACCTTGTGAGATTAATTCAAAAACTTCTGATAGATGTGAA ×
 CTTCTAGGGGGAAGCTTCCCAAGTCAGTCTTTACACAAATGCATGTGGGATTTTTAGCACAGCAAGTG ×
 GAAAATCTGTACAAGTATCAGATGCTGCAATACAAAAGGCAAGAGAGGTGTTTTCTAAGCTAGAAGATAG ×
 TGCCAAGCAACTCTTTCTGAAGTATCACTTAAAGATAATGAAGAACATTAGAAAAGTTCACAAATGAA ×
 GAAAATACTGTGATATATACCTCCCAAATTTACTATCATCTGCTTTCTCTGGATTTAGGACAGCAAGTG ×
 GGAACAAGTTCAGTTTCTGAAAGTGCTTATGCAAAGTTAAGGGAATGTTAGAAGAATTCAATCTGAT ×
 CAGAACTGAAAGTTGTCTTCAGCATTCTACTTCTAGACAAGATGTATCAAAAATGCCTCCTCCCTCT ×
 TGTATTGGTAAGAGAACCCAGAACACTCCAGAACTCCAAATTGGATAAAGCCTGCAATAAAGAATTTA ×
 GATTATCAAGTAACTGTAAACATCAGAGTGGTCTTCAGAAAATCATCACTCTATTAAAGTTTCTCCATG ×
 TCCCTCTCAATTGAAGCGAGACAAACCACAGTTGCTAGTCGGAAGCAAAGGATCACTTGTTGAGAACATT ×
 CATCCTTTGGGAAAAGAACAAGCTTTACCTAAAAATATAAAAACAGAGATTGGGAAAGCTGAACTTTTTC ×
 CTAATCTTCTGTGAAAACAAATATAGAATTTTGTCTACTTACTCCAAGGATCCAGAAAACTATTTTGA ×
 AACAGAAACCGTAGAGATTGCCAAAGCTTTTATGGAAGATGGTGAGCTGACAGATTCCGAACTGCTAAGT ×
 CATGCCAAACACTTTGTTTTTACATGCCAAAACACTAAGGAAATGGTTTTGTTAAATCAAGAATTGGAA ×
 AAAGAAGAGGAGATGCACTTGTCTCAGTTGGAGAACCCCAATTAAGAACTTGTAAATGAATTCGA ×
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 AGAAGCTTGTTTATGCATCATATTTCTTTAGAGCCAAATTTCTGTGGACCTTTTCGCACAACTGAGGAAC ×
 GGCAAGAAATACAGAAATCCAAATTTCACTGCACCTGGTCAAGAAATTTTGCCTAAATCTCATTTTTATGA ×
 ACACCTGGCTTCAGAAAAATCTTCAAGTAATTTATCAGTTTCACGGCAACCATTTTGTATGGTTCCTGCC ×
 ACAGGAAATGAAAAAAGGAGACACTTGATTGCTCCAGGCAACCAAGTGAAGTCTTTGTCCCACCTTTTA ×
 AAACTAAATCACAATTTTACAGAGATGAGCAGTGCATTAGCAAGAATACTAAATTGGAAAAAACAAACA ×
 AAACTCCAAAGACATAGATGAATTTGGCTCTGGTGATAGTGAAGAAAAATATTAATGACAGTGGAATCCAT ×
 CAGCTTAAGAAAAATAACTCCAATCAAGCAGCAACTATAATATTCACAAAGAATGAAAAAGAACCTTTAG ×
 ATTTAATTACAAATCTTCAGAACGCCAGAGATATACAGGATATGCGGATTAAGAAAGAAACAAAGGCAGCA ×
 TATTTTTCCACAGCCAGGTAGTCTGTATCTTGCAAAAACCTCCACTTTGCCTAGAATCTCTCTGAGAGAA ×
 GCAGTAGAAGGCCGAGTCCCCTCTGCATGTTCTCATAAACAGCTCTATATGTATGGTGTTCCAAACATT ×
 GTGTAATAAATAACAGCAAAAATGCAGAGTCTTTTCAGTTTCATGCTCAGGATTATTTTGGTAAGGAAGG ×
 CCTATGGTCTGGAGAAGGAATACAATTGGCTGATGGTGGATGGCTCATACCCTCCAATGATGGAAAGATT ×
 GGAAGAAGAAGATTTTATAGGGCTCTGTGTGACACCCAGGTGTGGATCCAAATTGTATTTCTAGAGTTT ×
 GGGTATATAATCACTATAGATGGATTATATGGAATTTGGCAGCCATGGAATTTGCCTTTCTTAAGGAATT ×
 TGCTAATAGGTGTCTAAGTCCAGAAAGAGTGCTTCTTCAACTAAAATACAGATATGATGTGGAAATTGAT ×
 AAAAGCAGAAGATCAGCTATAAAGAAGATAATGGAAGGGATGACACAGCTGCAAAAACACTTGTCTCT ×
 GTATTTCTGAAATCATTTCTGCAAGTGCAGATATATCTGAACTTCTAGTAGTAAACTAGTAGTGTGGG ×
 TACCAAAAAAGTGGGCATTATTGAGCTCACAGATGGGTGGTATGCTATTAAGGCCAGTTAGACCCTCCC ×
 CTCTTAGCTCTCGTAAAGAACGGGAGATTGACTGTGGGTGAGAAGATCACTATTCTGGAGCAGAACTGG ×

```

TAGGCTCTCCTGATGCCTGCACACCACTTGAAGCCCCAGAATCTCTTATGTTAAAGATTTCTGCTAACAG ×
TACTCGTCCTGCTTGTGTTATACCAAACCTGGATTCTCTCCTGATCCTAGACCTTTCCCTCTCCCTTG ×
TCATCACTTTTCAGTGATGGAGGAAATGTTGGTTGTGTTGATGTAGTTGTTCAAAGAGCATACCCAATAC ×
AGTGGATGGAGAGGACCCCATCTGGATTATGCATATTTGCAATGAAAGAGAGGAAGAAAAGGAAGCAAC ×
AAAAATATGCAGAAATCCAACAAAAGAACTAGAAGTTTTATTCAATAAAATTCAAGCAGAATTTGAAAAG ×
AATGATGAAAAATATAACAAAGCAGTGTATACCATCATGTGCATTAACAAGACAGCAGATCTGTGCTCTGC ×
AAGATGGTGCAGAGCTTTATGAAGCAGTGACAAATGCACCAGACCCAAGTGACCTGGAGGGTTATTTTAG ×
TGAAGAGCAGTTAAGAGCCTTGAATAATCACAGACAGATGTTGAATGATAAGAAGCAAGCACAGATCCAG ×
TTAGAATTCAAGAAGGCTATGGAATCTGCTGAGCAAGGAGAACAAATCTACCAAGGGATGTTACAAC TG ×
TGTGGAAGTTACGTATCATAAGCTACAGGAAAAAAGAAAAAGATTCAGTTACATTGAGTATCTGGCGTCC ×
ATCACCAGATTTATATTCCTGTTAATAGAAGGAAAGAGATACAGAATCTATCATCTTGACAGCATCACAA ×
TCTAAAAGTAAATCTGGAAGGCAACACACAGCTAACAGCAACAAAGAAAACCTCAGTACCAGCAACTAC ×
CAGCATCAGATGAAATCCTATCCCAAGTTTATCAGCCAAGGGAACCCCTTTACTTCAACAAACTGTTGGA ×
TCCGGACTTCCAACCACCTTGTTCTGAGGTGGACCTAATAGGATTTGTAGTTTCTGTTGTGAAAAAATA ×
GGTCTTGCTCCTGTGGTCTATTTGTGATGAATGCCATAATTTATTGGCAATAAAGTTCTGGACTGATT ×
TTAATGAAGACATTATTAACCTTACACATTAATTGCTGCAAGCAACCTCCAGTGGCGACCAGAAGCCAA ×
ATCAGGAATTCCTACTTTATTTGCTGGAGATTTTTCCAGTTTTCTGCCAGTCCAAAGGAGGAGCATTTT ×
CAAGAGACATTCCACAAAATGAAAAATACTGTTGAGAATATTGGTATGTTTTACAATGATGCAGAAAACA ×
AACTTGTGCATATACTTAATGCAAATGATCCCAAGTTGTCCACCCGACTAAAGACTATGCTTCAGAGCC ×
ACACACAGCTCAAATAGTCCTTGGCATAGGAAATAAATTTCTGATGTCTTCTCCAATAATGAGATGAAT ×
TATCAGAGTCCTTTATCACTTTGTAAGCCAAAAGAGAAGTCTGTCCCATACTGGATCAACCCAAATGA ×
CTTCAAAGTCTTATTGTAAGAGGAGAAAGAGATGGATGACCCAAAACCTGCAAAAAGAGAAGAGCCTT ×
GGACTTTTTGAGTAGAGTGCCTTTACCTCCATCTGTGAGTCCCATTGTGACATTTGTTTCTCCAGCTGCA ×
CAGAAGGCATTTAGCCACCACGGAGTTGCGGCACCAAAATATGAAACACTGATGAAGAAAGAGTTGAATT ×
CTCCACAGATGACTCCACGTAAATTTAATGACCTTTCCCTTTTGGAAAGTGATTCAATAGCAGACGAAGA ×
ACTCGCAATGATAAACACCCAAGCCCTTTTGTGGGTTACCAGGAGAACATCAACTTGTGTCTGTGAGT ×
GACTCTACCAGGACTGCTCCACGAGCTCAAAAGATTATCTTGGACTGAAAAGGCATTCTACTGCACCCG ×
GGGTGAGAGGACCCGAGAGCCCCAGGCCCTGCACCAGGAAGCGGGAGCCCCGTGTACAGAACACAAGTGA ×
TCTGAAAAGGACATCTCTGAGACTGCAGAGGCAACAAACACAAAAATGACAATGAATTGGTGACTGACTC ×
AACCTTTCCAATGTGTGGAACACAGCCTCAACCTGTATGTCAAGATGTGCATAATGAGACAAGAAAGA ×
CCACATCCCAAATCTCCTGTGTGCTTGTCTATCTTAGGAAACCTGGCCTATCTCTGTACTGGTCGGTGTA ×
CTTTATTTTCAGTTATGTGTCTGAAAATTGTGTATTTATTAGCTAATCAGGAAAAAAAATCTCCTTTAAAC ×
TCTTATGATTGGATATGATCAAGTATATTTACAAAGTAAACACACTTTTTCTTTAAATTGTGTCCCTAA ×
TTAAATGAAAGTAGGTTTCAAAGTACTGTTATTTTGAAGTCTGTTAGTTCCTTTTGGTGACTTGGTTTTG ×
TTTTGTTTTTCGGAGGTAACCTACTATGAACCAAGTTTCTTAATAAACGTGTTGGTTCTCTTATAGTTG ×
TATCCTGATCAAAAGTCAGGAGGAGTAAGGAACAAACAGCAGTGCTCTCTCTGGACCAGTTCTTTAACCT ×
TACGTCAGCATAAGTGCAAGAAAAACAGAATCCTCAATGTGATTCTTTTATGATTCTAGTGTGATTGC ×
TGAATTATTTCAATTAATAAAATTCAAATGCTTTTAAAAAAGAAAAAAGAAAAAAGAAAAAAGAAAAAAG // ToString;

```

Process

```
SpecialNote = " ";
```

```

Wgenesample = "BRCA2 mRNA Wolf gene"
(*Lets us know which gene we're dealing with,
used in pdf coding later, so be sure to name it *)

```

```
BRCA2 mRNA Wolf gene
```



```

LetterDNAToNum[Sample_] := ToExpression[StringReplace[ToString[
  {StringReplace[StringReplace[ToString[{Sample}], {" " → "", " " → "", "{" → "",
    "}" → "", "(" → "", ")" → "", "[" → "", "]" → "", ";" → "", ":" → "", "_" → "",
    "+" → "", "&" → "", "/" → "", "." → "", "RowBox" → "", "Null" → ""}],
  {"0" → "0,", "1" → "1,", "2" → "2,", "3" → "3,", "A" → "0,", "C" → "1,",
    "G" → "2,", "T" → "3,", "a" → "0,", "c" → "1,", "g" → "2,", "t" → "3,"}]]
], ",,"} → "}]"]
numgenesample = LetterDNAToNum[lettersample];
Export[StringReplace["GENE_genesample.txt", "GENE_gene" → Wgenesample],
  Flatten[numgenesample]]
BRCA2 mRNA Wolf genesample.txt

lengthofgeneitself = Length[Flatten[numgenesample]]
(*To make sure no base pairs are left out *)
11190

```

Construction of W

Can compare to W constructen in Python file **W_hat_construction.py** if we want

```

lettersample = {basepairs} // ToString;
LetterDNAToNum[Sample_] := ToExpression[StringReplace[ToString[
  {StringReplace[StringReplace[ToString[{Sample}], {" " → "", " " → "", "{" → "",
    "}" → "", "(" → "", ")" → "", "[" → "", "]" → "", ";" → "", ":" → "", "_" → "",
    "+" → "", "&" → "", "/" → "", "." → "", "RowBox" → "", "Null" → "", "
    " → "", "
    " → ""}],
  {"0" → "0,", "1" → "1,", "2" → "2,", "3" → "3,", "A" → "0,", "C" → "1,",
    "G" → "2,", "T" → "3,", "a" → "0,", "c" → "1,", "g" → "2,", "t" → "3,"}]]
], ",,"} → "}]"]

numgenesample = LetterDNAToNum[lettersample];
lengthofgeneitself = Length[Flatten[numgenesample]];
M = numgenesample;

For[npow = 1, npow < 1000, npow++, If[Length[M] < (2^(npow)), Break[]];
  FilledSize = 2^(npow + 1)];
Filler[vecvar1_] := Table[4, {i, 1, FilledSize - lengthofgeneitself}]
FilledVec[vecvar2_] := Join[Flatten[vecvar2], Filler[vecvar2]]

Filler[vecvar4_] := Table[4, {i, 1, FilledSize - lengthofgeneitself}]
FilledVec[vecvar5_] := Join[Flatten[vecvar5], Filler[vecvar5]]
For[npow = 1, npow < 1000, npow++, If[lengthofgeneitself ≤ (2^npow), Break[]]];
(* gives npow such that 2^npow > lengthofgeneitself > 2^(npow - 1) *)
FilledSize = 2^npow;
FilledM = FilledVec[M];
numrowsW =  $\sqrt{\text{Length[FilledM]}}$ ;

```

```

W = Table[Table[FilledM[[i]],
  {i, ((j - 1) * (numrowsW) + 1), (j * (numrowsW))}], {j, 1, numrowsW}];

(*numgenesample=LetterDNAtoNum[lettersample];
lengthofgeneitself=Length[Flatten[numgenesample]];
M=numgenesample;
lengthvec[M_]:=Length[M[[1,All]]]
For[npow=1,npow<1000,npow++,If[lengthvec[M]<(2^(npow)),Break[]];
  FilledSize=2^(npow+1)];
Filler[M_]:=Table[4,{i,1,FilledSize-lengthvec[M]}]
FilledVec[M_]:=Join[Flatten[M],Filler[M]]
lengthvec[M_]:=Length[M[[1,All]]]
Filler[M_]:=Table[4,{i,1,FilledSize-lengthvec[M]}]
FilledVec[M_]:=Join[Flatten[M],Filler[M]]
For[npow=1,npow<1000,npow++,If[lengthvec[M]≤(2^npow),Break[]]];
(* gives npow such that 2^npow > lengthvec[M] > 2^(npow - 1) *)
FilledSize=2^npow;
FilledM=FilledVec[M];
numrowsW=√Length[FilledM];*)

```

W for Human isolate NA19240 chromosome 9 genomic scaffold

Is a “genomic scaffold” a single gene?

Data

Source: GenBank: KZ268583.1

```
ln[ ]:= (*lettersample={ } //ToString;*)
```

```

ln[ ]:= lettersample = {TCTTAAGCCAGTCTTTTATTTATATTTTGATTCTGTTTTGTGGAGGACATGCTTCCTGAATTTTAGGAA ×
GAGGCCAAAAGGTTTGTCAAGTTTTACCTGATACATTGGATTAAATTATCTAATGTACACACTTTTAA ×
TTTAAGTCTAGGGGCGACTGTCTACTCTTGATTTTGATAGTATTATTTTCTTAACATCCAAGTCCATC ×
TTCATCTATTTGTATAAGATCAATAAAAAATATATTTGTCCAGAACCTGCTTTGGCGGAGTTACTTCTTT ×
CTAAGTAGTAGAGGTAGCAGTTGAGACATGAGCTGGGTTCTGGGTGAGTTTAGAGGGCTGGGCGACATTC ×
CTCCTTTTGGTCTGTATGACTGAATGAATGCAGTTCTTGCTGTCTCGCTCCTCTCCTTAACACATTGAGC ×
CATTGCAGCAGATAAGAAGGAATAATCTTGATCTGCCATTGAGGTGGAACACATGTTCTCTCAACCACA ×
CCCATAGGTTGTACTCACACTCGGCCAGAAGGTATCCTGTCAATGATATGGAGATGTATCTATCTATCTA ×
GATAGATATCTACTTTGGTTTATGCTCTCTGGTTGCCCGCAAATTATCTCCTTAAATTGAATATCAAAAG ×
AGAGCTTGGTGATGGCAGTGTTATAAAATCCTCAAAATGCAGCACCCATACCCAGAGGAATTTGTAGATT ×
CTGAGATTCTAATTCAGATACCAAATATATAAAAGGGGAATTGGTCATTGAGGGTTGCTAGGCTCTTTG ×
TTGAGCATATATGCTCTTTTCATGACTTCGAAATTATTTTAAAAATCTAAACTTTTTCTCAGTGTGCTGC ×
AAGATGATTTGATTTGAATGCATAAGCAGTAATTCTCCCTAAGATTTGTACAATATATTTGCTCTGACA ×
AGCGATAGCCAGCAACTCACTTCACAGCAATTTACAGCGTTTCCATGATAAGTTGAATTATTTTAACTA ×
GACTCTCTTTGCCTTAATAAAAAATGAAGAAGCAATATACTTGTCTAATTAGGTTCAAAAGTTGGCAG ×
TCTCTCTCTGGAAAGAATAGTAAACTTTTCAGCGGCCTAATATGCATGTATAAACACACAAACACACA ×
CACACACACACACACACACAAGCACTATTCATAATATTTAAAGCACATTCTGTTCTATGACTTCATTT ×
GTCTAGCACAAAATAAAACGATCTCAGTATATGTCAAGTATCAATTTTTTCGTATGGCCAATTATAGATA ×
TTTTATTTTTTAAAGATTAGAGTGTTCTTGAAGCTCTTTATATTTCTTTGTCAATGAACTAAACATTGGC ×
AAATATGTAGGGTTTCCACATAAGAACATTATTAACATCAAAATAGAAAGCTGGTGGTAGCAATAATGA ×

```

TTGGGAACACAGAGTCTCTACTCAACGTTCTAGTTCTGCCATACCATAACTTTGTGATCTCAGGAAATAT ×
 CTCTCCATGTTGTCTCATCTCAAAGTATAGTTCTGTCAATTTTCAATAAGAGCTTTTGTCTAATTATGAAG ×
 TACTAGTTAGTGTAACCATTATTTTGAGCTTCATGTAATCAAGAACACATGGACTCCACTTGCAAAACA ×
 TCGAAAATGTAGTAGGGATTGGGGGCATAAAGCAACACTTTAAAATGTGTAAGACAATGAGTAAGCAAC ×
 AAAGTGTCCAATTTTTAGGGGAAAGTTGCATACGTTAGGAAAAGGCAGGATTAAGTAACAGAGAATTTG ×
 AATGATAACTGGCCAATTGGTGTCAATTTACAATTGCAAGTCATACAAATGAAGTTTTCTGTTTTAAAGAG ×
 AAAAGGAGTTATTTAGAATGGGTCAACCTATTGGGGAAGCAATGTAGTTAGAAACAATGCCCAAAACCAT ×
 GTAAGCAAATGCTCTGTAGAGCACACCCCTGCAATGCTGCCATTGTGAGGCCAAGTCTCTCCTCGTCTTG ×
 GTACTGAGCCCTCCGTTCTGCCTCCATCATTGCCACTGTAGCTGCCACAAAATGACCCCTCAACCACCGC ×
 TGCCCAAGGAACAAAGAAAGAATTCTGTCTTCCGCGCTCTCAGATCAATTTCCAACATCAGGTGAGCCTT ×
 TGATGGGCACTATTCAAGTTCCTTATCCCTGAAATAGATGCAGTAAAAATATAGAAATTGCCTATGTGTT ×
 TCCCAATAAGACACATATGGAAGCCTGTTTTCCACAACAGGAAGGAGTTTGACAATGGGTGTTCAAAG ×
 GAACAATATTCCTGTAAACCGTACTTTGCCCATATGAAGAAAAGCAATAAGGATTATTTAGTAAATAGA ×
 CATGGAAACTCATCCAGGGTTGGCTGATGAGAAGCTGGTTAGCAAGGGGGTCTGCCTTCAGTTAGGACAA ×
 GGTTTGTGCTTCCACGGGTTCTCTCCACAGCAGGAGGGATGCAAACTTCCCTTTCCTCCCCTGCACCTA ×
 CCCTCAAATGGCCAGAGGTCTTCAGGTGCTAGAATTTCTCAATTAATGCTGCACAAAATATCAGACAGC ×
 CTTGACTGTACAGTCTGTTCTCATGAAGCTAGTCTCTGCTCACTACATAAAAACAGGAGAGTAAGAACAA ×
 GGGTGTTTAACGCTACCTTAGCTCAAACAAGTTTCTCTCTGTATTATGCCAAGAACCTGGGAACCAAGTGC ×
 ATCTGCTGCTTTCCTTCTTGGATTCTAGCCAGACAAAAGAGGCAAGGGGCATTTCTTCAGAGGCCTTG ×
 AGCTTCACTACACAATGACCCAGGCTCTACATGCACCCTCTTTATATATTTCTACCTTGAAAAAAAATTT ×
 TTATATAATATTAATAATATATATTTTTATATAATAAACACTTTTTTAATAGATAGATATAGATATACAT ×
 AGATAAAGATCTCTAGTCAGCCTTTTTTAAGGCTGGGCTGATCGCAGTGCCTCAAACTATAATCCCAGC ×
 ACTTTGGGAGGCCAAGGTGGCCAGATCTCTTGAGTCCAGGAGTTGGAGATCACCCAGGGCAACATGGTGA ×
 AACCCCATCTTCACAAAAATTAGCTAGTATGGTATTATGCACCTGCAGTCCCTGCTACTCAGGAGGCTGA ×
 GGTGGGAGAATTGCTTGAGCACAGTATGTGAAGGCTTCAGTGAGCTCTAATCACATGACTGCACTCCATC ×
 TTGGGTGACAAAGTGAGACCCTCTCTCAAAAATAAAATAAAATAAAAGGCTACCACCATACTCAGAGAT ×
 AAGTGTGTCAAGGTATATTTGAGCTATCCTTCTATATTCTATTTGGTAAAAAAAAAAAAAGGCAAGA ×
 ACTCTTCTCATTCTAGATTTTTGTATTAATTAGACATTTGAAGTTTATAGCAGAAGAGCTATAATCACTC ×
 TATAGACCAGATAGTGCAACAGATATCAATGCTTTTTAAAAGTATAGAAGGTTATTAGAAATTTTTTAA ×
 ACTACTTATAGGTATATATGTATCTAATTGAAGTATCAAATGCAAGTAAGATCATTTCTTAGCGTGTGA ×
 AATCCACTCAATTTATTAATAATTTTTCTAATATCTATTACAATAATATTTCTTAATTAGTTAACAAAAG ×
 AGGAGTTTTAAGACATTTGTTTATATGTACTTACTAGATTCAAACCTCGATTCCACTATTTTCAGAACTCA ×
 TACTCTGAGACAAGTCTTTTTTATGTAAGTATGTTTCTGCCTATATTAAGACAGATATGTCAATTT ×
 TGCTAGTCATGCTGTTCCAAAGCTCTCCATCCTGATTATTTTTCGGTTTGTCTAGCAGTCATTGAGAGA ×
 CTTACTTATATTCAAATTTCTCTAGGTTTAAACATTCGTGTATGTCTTGTGTGGTTTTGTCTATTTTT ×
 GCTGTATATAATTTAAGACATTTTATTGACATACACATGCAGAAAAGTACAATGATTAATATGATAGCT ×
 TGATTAATGAACACATGTATTTGCTTATAGCCATGTACGAAAATAGAACATTATTAATAATAGTGATAC ×
 CTCTCCTGCCCCCTTCCAAACACTAACCTCATCCTCAATAGTAACAGATTTTTTTTATCATAATTTGGT ×
 CTACTTTCAAATTTTTATTAATAAATCGGAGTATCTACTCTAAGTCTATGTTTATTTTATTGTTGTTAT ×
 TTTACTTGTAGTATTTATCTGCTAATGGACATGGTAGATTGAAGACGGCTACATACACATTTTTTAATTA ×
 ATAGATTTTTTGTAGCACTTTGTGGCTCATGCCTTAATCCCATCACTTTGGGAGGCTGAGGTGCGTGGAT ×
 CATGAGGTCAAGGATCCAGACAATCCTGGCTAACGTGGTAAACCCCTTCTCTACTAAAATACAAAAAA ×
 TTAGCTGATAGATAACATCAAGATAACATCTGAGTCTTAGCTGCACTGAGTCAAGCCTACTTACATCTT ×
 TGTCTTCTGCTGCACTTTTCTTCCACATACCGTCCAGGAATGCCAAGCTCCGTTGGCTTCTACCCCA ×
 TTTCCACTATTGTTCCCTGCCACCGCGGCTTTTTGCCGCTACCGCCGCGGGTTTTTGCCTCCGCTGCT ×
 TTTTGCCACCGCCGCGCGGCTTTTTACCCCAACGCTGGGGCTTTTTGCGGCTCTTTGACCCACCACCG ×
 GGGCTTGTGCTCTTTTTGCACCCGCCGCGTGCCTTTTTGCCCCGCCGCTGCGGCTTTTCCCCCGC ×
 CTCACGGCTTTCTGCCCCACCGCTGTGGCTTTTCATCACTACTACCGGCTTTTGGCCCCGCCACTG ×
 CGGGTTTCTCCACCGCGGTTTTTGCCCCGCCGCGTGGCTTTTACCCCGCGCCATGGCATTTTTG ×
 CCCACGTCACCGTGGCTTTTTGAGCTTTTTGCCGCTGCGGCTTTTGGCCCTGAAGCACGGCTTTTTG ×
 CCCTTGGCGCGTGGCTTTTTGCCCCGCCGCGGCTTTTTGCGGCTTTTTGCCCCGCCAATACGGC ×

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TTTTTGCCGCTGCGGCTCTTTGCCTCCGAAGCCACGGGTTTTTACCCCCACCGCCACGGGTTTTTGCCCC ×
CGTGGCTTTTTGCTGCCACGGCTTTTTGCCCCGCGCCATGGCTTTTTGCTCCTGCCGCTGAGGCTTTT ×
TGCCGCGCAGTTTTTGCCCCGCGCGCTGCTTTTTGTGGCTTTTTGCCCCGCGAGCTTTCCCCCCC ×
GTCGCGCGGGTTTTGTGGTTTTTTGCACCCGCTCCCGCTGCTTTTTGCCCCGCCACTACGGCTTTTT ×
GCCGCTTGTTTTTTGCCCCGAAGCCACAGCTTTTTGCCCTCGCCGCGCGGCTTTTTGTGGCATTTT ×
ACTCTCAGCCACGGTGGCTTTTTGCCCCTGCCACGGCTTTTTATCCCTATCGCCGTCGCCATGGCTTTTT ×
GCAGTCACAGCTTTTTATCCCCACCGCGCGGCTTTTTGGGCCACCACCGCGGCTTTTCAACCGCGGC ×
TTTTTGCCCCCACCACGTGGCTTTTGGCTCCTGCCACTGAGGCTTTTTGCTGCCGCGGTTTTTGCCCT ×
GCCGCGCTGCTTTTTGCGGCTTTTGGCCCCGCCAGGGCGGCTTTTTGCCCCGTGAATTTTCCCCC ×
GTCTAAGCGGGATTTGTGATTTTTTTGCCCCGCTCCTGCTGCTTTTTGCCCCGCCGCGCAGCTTT ×
TTGCCCCACTGCTACGGTTTTTTGCCACAGTTGCTTTTTGCCCCGAAGCCACAGCTTTTACCCTCGAC ×
ACCGCGGCTTTTTGTGGCTTTTGCCTCGCGCTGAGGCTTTTTGCCGCGCGGTGTTGTTCCCGCCA ×
CCGCTGCTTTTTGCGGCTTTTGGCCCCCACCGCCACGGCTTTTTGCCCGCCACTACGGCTTTTTGCCG ×
CTGCAGCTTTTTGCCCCGAAGCCACGGCTTTTTGCCCTCACCCTGCGGCTTTTTGCACCCACAGCCGG ×
GGCTATTTACCCCCGCTTCCACGGCTTTTTGCCACCGCTGTTTTTCCCCACCACCGCGGTTTTTGCC ×
CCCGCGCGCGTGGGTTTTTGCCGCTGAGGCTTTTTTTAAACGCCACCAGCGCGGCTTTTAGTCCCCGC ×
CACCGCGGTTTTTGCTCCGCTGCCGAGGCTTTTTGTCCCCACCGCTTTGCTTTTTGTAGGTTTTCGC ×
CCCCGTGCTGCGAGGTTTTTTGCCCCCTGCCACCACGGCTTTTTCCCCAGCGCGCGGATTTTTGTGT ×
TTTTTTTCCCCGCTCCCGCTGCTTTTTGCCCCCTCAGTGGCTTTTTACCCCCCTCAGCGCGGCTTTT ×
TGCTCCACCGCCACGTCTTCTCAACGCCACCGTGGCTTTTTGCCCTGCCGCCACGCTTTTTGCC ×
GCCGTGGCTTTTTGCCCCGCTGCTTTTGAACCTTAATTTCACTTGAAATCTGACTTCCCACTGCCATG ×
CAACCTAACATATTTGTATGTTAGACTCCGGGAATTAGGACATGAACATTTCTGGGAGGCCATTATTTG ×
TCTACAACAGACATAATCTATTTACCTGAAGATTAAAGTGATCTTTATTTTTCTGCCTCTCTTTCTTAAT ×
TTTTTTTTAAATAATATGGATTGTAGTAAAGAGAAAGAAAGAAAGAAAGAAAGGAAGGAAGGAAGAAAG ×
AGAAAGAAAGGAGGAAATGAGAGAAGGGAGGGAGGGAGGAAGGGAGAAAGGCAGGAAGGAGAGAAAAAG ×
AAAGCAAGAACTCAAGAAAGAAAGAGAAAGAAAGTGAGAAAAGAAAGGAAGGAGGAAGAGAGAATGGTAAA ×
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GCATTCTCATAAACCTGTTTGTGATGTGTGTA CTCACTAACAGAGTTGAACCTTTCTTTTGATAGAGCA ×
GTTTTGAAACACTCTTTTGTAGAATCTGCAAGTGGATATTTTGATAGCTTTGACGGTTTCGTTGGAAAC ×
GGAAATATCTTCATATAAAATCTAGACAGAAGCATTCTCAGAAACATCTCTGTGATGTTTGCATTCAAGA ×
CACATAGTTGAACTTTCTCTTTACTGTAACAGCTTTGAAACACTGATTTTGTAGTATCTGGAAGTGGACA ×
TTTCATGTGCATTGGGGCTATATTGAAAAAGGAAATATCTCCCATAAAAATTGACAGAAACATTCTCA ×
GAAACTAGTTTGTGATGTGTGTA CTCACTAACACAGTTGAACTTTCTTTTGATAGAGTAGTTTGAAG ×
CAGTCTTTTGTAGAATCTGCAAGTGGATATTTGGATACCTTTGAGGATTTGTTGCAAAACAGGAATATC ×
TTCAGGTAAATCTACACAGAAGCATTCTCAGAAACATCTTTGGATGTTTGCATTCAAGTAACAGTGTT ×
GAACATTCCCTTTCTAGAGCAGGTTTGAACACTCTTTTGTAAATATCTGGAAGTGGACATTTGGATCG ×
CTTTGAGGTCTGTGGTGAAGAAAGTACCTTGGCATAAAAACTAGACAGAACCATTCTCATAAACTTG ×
TTTGTGATGTGTGTA CTCACTAACAGAGTTGAACCTTTCTTTTGATAGAACAGTTTGAACAATCTTTT ×
GGTAGAATCTACAAGAGGATATTTGGATAGCTTTGAGTATTTTCATCGGAAACGGGAATATGTTTATATAA ×
AATCTAGACAGAAGCATTATCAGAAACATCTTTGTGATGTTGCAATCAAGTCACAGATTTGAACATTCC ×
CTTTTCATAGAGCAGGTTTGAACACTCTTTTGTAGTATCTGGAAGTGGACATTTGGATCACTTTGAGGC ×
CTATGGTGTAAAAGGAAATATCTTCGCATAAAAACTAGACAGAAGCATTCTCACAACCTTGTCTGTGATA ×
TGTGTA CTCACTAACAGAGGTGAACCTTTCTTTTGATAGAGCAGTTTGAACACTCTTTTGGAGAAT ×
CTGCAAGTGGGATTTTGGATAGCTTTGAGGATTTGTTGGAACGGGAATATCCTAATACAAATCTAGA ×
AAGAAGCATTCTCAGAAACATATCTGTGATGTTTGCATTCAAGTCACAGGGGTGAATATTCCTTTTCTG ×
GAGCAGGTTTGAAGAACTGATTTTGTGGTATCTGGAAGTGGACATTTGAGCGCACTGTGGCCTTTATTG ×
AAAAAGGAAATATCTCCCATAAAACTAGACAGAAATATTCTCAGAACTACTTTGTGATGTGTGTA CT ×
CACTATCAGAGTTGAACCTTTCTTTTCATAGAGCAGTTTGAACACACTTATTCTAGAATCTGCAAGT ×
GGATATTTGGATAGCTTTGAGGAATTCGGTGGAAACGCGAATATCTTCATATAAAATCTAGACAGAAGCA ×
TTCTCAGAAACATCTTTGGGATGTTTGCCTTCACATCAGACAGTTGAACATTCCCTTTTCATAGGGCAGG ×
TTGAAATACTCTTTTGTAGTATCTGGAAGTGGACATTTGGAACGGTATCAGGCCTATGGTTAAAAAGGA ×
AATATCTCCCATAAAAACAAGACAGAAGCACTCTCAGAACTTATTTCTGATATCTGTCCTCACTTAAC ×
GGACATGAACCTTTCTTTTATAGAGCACTTTTGAACACTCTTTTGTAGTATCTGCAAGTGGATATTT ×
GGATGGCTTTGAGGATTTGTTGGAACGGGAATATCTTCATATAAAATCTAGACAGAAGCATTCTGAGA ×
AACTACTTTGTGATGTTTGCATAAAAGACACAGAGTTGAACATTCCCTGTCTAGAGCAGGTTTGAACA ×
ATCTTTTGTAGTATCTGGAAGTGGACACTTCGAACGCTTTGAGGCCTATGGTTAAAAAGGAAACATCT ×
CTCATATAAAGAAGACAGAAGCATTCTCAGAACTTATTTGTGATGTGTGCTAACTAACAGACTTGA ×
AACTGTCTTGATACAGCAGTTTTCAAACACTCTTTTTCTAGAATCTGCAAGTGGACATTTGGATAGTTTT ×
GAGGATTTGTTGCAACGGGATTACATATAAAAAAGTAGACAGCAGCATTCTCAGAACTTCTTGTGATG ×
TTTGCATTCAAGTCACAGAGTTGAACATTCCCTGTCTAGAGCAGGTTTGAACAATCTTTTGTAGTAT ×
CTGGAAGTGGACACTTCGAACGCTTTGAGGCCTATGGTTAAAAACGAAATATCTTCATATAAACAAGA ×
CAGAAGCATTCTCAGAAATTTATTTGTGATGTGTGCTCACTAACAGACTTGAACCTGCCTTTTAATA ×
CAGCAGTTTGAACACTCTTTTGTAGAATCTGCAAGTGGACATTTGGACAGCTTTGAGGATTTGTTG ×
GAAACTGGATTACATATAAAAGTAGACAGTAGAATCTCAGAACTTTTGTGATGTTTGCATTCAAGT ×
CACAGAGTTGAACATACCCTTTTCATAGAGCAGGTTTGATACACTCTTTTGTAGTATCTGGAAGTGGACA ×
TTTGGAGCGCTTTGTGGCCTACGGTAAAAAGGAAAGTACCCTCCCATAAAAACAACATAGAAGCAATCTC ×
AGAAACTTGTTTATGTGTATCTACTCAACGAACAGTGTGCAACTTTCTATTGATAGAGCAGGTGTGAA ×
ACACTCTTTTGTGAATCTGCAGGTGCATACTTGGATAGAATTGAGGATTTGTTGGAAGGGATTAC ×

```
TTATAAAAAGTAGACTGCAGCATTCTCAGAAACTTCTTTGTGATGTTTGCATTCAAGTCACAGAGTTGAA ×
CATTCCCTTTTCATAGAGCAGGTTTGAAACACTCTTTTGTAGTATCTGGAAGTGGACATTTCCAACGCTT ×
TCAGGCCTACGGAGAAAAAGGATATATCTTCCCATAAAAACAAGACAGAAGCATTCTCAGAAACTTATTT ×
GTGATGTGTCTCCTCAACTAACGGACTTGAACCTTTCTTTTATAGAGCACTTTTGAAACACTCTTTTGTG ×
TACTATCTGCAAGTGGATAGTTGGATGGCTTTGACGATTTCTGTTGAAACGGGAATATCTTCCTATAAAA ×
TCTACACAGAAGCATTCTCAGAAACTTCTTTGTGATGTTTGCATTCAAGTCACAGAGTTGAACATTCCCT ×
GTCATAGAGCAGGTTTGAAACAATCTTTTTTTAGTATCTGGAAGTGGACACTTCGAACGCTTTCAGGCCT ×
ATGGTTAAAAAGGAAAT} // ToString;
```

Process

```
In[ ]:= SpecialNote =
  " Not sure if this is a single gene or multiple. Also not sure if this is necessarily
    a set of whole genes or if it's just the DNA sequence corresponding
    to a particular chromosomal structure. In that case it might
    includes only parts of some genes (not the whole genes)";
```

```
In[ ]:= Wgenesample = "Human chromosome9 scaffold gene"
  (*Lets us know which gene we're dealing with,
  used in pdf coding later, so be sure to name it *)
```

```
Out[ ]:= Human chromosome9 scaffold gene
```

```
In[ ]:= LetterDNAToNum[Sample_] := ToExpression[StringReplace[ToString[
  {StringReplace[StringReplace[ToString[{Sample}], {" " → "", " " → "", "{" → "",
    "}" → "", "(" → "", ")" → "", "[" → "", "]" → "", ";" → "", ":" → "", "_" → "",
    "+" → "", "&" → "", "/" → "", "." → "", "RowBox" → "", "Null" → ""}],
  {"0" → "0,", "1" → "1,", "2" → "2,", "3" → "3,", "A" → "0,", "C" → "1,",
    "G" → "2,", "T" → "3,", "a" → "0,", "c" → "1,", "g" → "2,", "t" → "3,"}]]
  ], ",," → "}]"]
numgenesample = LetterDNAToNum[lettersample];
Export[StringReplace["GENE_genesample.txt", "GENE_gene" → Wgenesample],
  Flatten[numgenesample]]
```

```
Out[ ]:= Human chromosome9 scaffold genesample.txt
```

```
In[ ]:= lengthofgeneitself = Length[Flatten[numgenesample]]
  (*To make sure no base pairs are left out *)
```

```
Out[ ]:= 59027
```

Construction of W

Can compare to W constructen in Python file **W_hat_construction.py** if we want

In[]:=

```

LetterDNAToNum[Sample_] := ToExpression[StringReplace[ToString[
  {StringReplace[StringReplace[ToString[{Sample}], {"," → "", " " → "", "{" → "",
    "}" → "", "(" → "", ")" → "", "[" → "", "]" → "", ";" → "", ":" → "", "_" → "",
    "+" → "", "&" → "", "/" → "", "." → "", "RowBox" → "", "Null" → "", "
    " → "", "
" → ""}],
  {"0" → "0,", "1" → "1,", "2" → "2,", "3" → "3,", "A" → "0,", "C" → "1,",
    "G" → "2,", "T" → "3,", "a" → "0,", "c" → "1,", "g" → "2,", "t" → "3,"}]]],
  ], ",,}" → "}""]

numgenesample = LetterDNAToNum[lettersample];
lengthofgeneitself = Length[Flatten[numgenesample]];
M = numgenesample;

For[npow = 1, npow < 1000, npow++, If[Length[M] < (2^(npow)), Break[]];
  FilledSize = 2^(npow + 1)];
Filler[vecvar1_] := Table[4, {i, 1, FilledSize - lengthofgeneitself}]
FilledVec[vecvar2_] := Join[Flatten[vecvar2], Filler[vecvar2]]

Filler[vecvar4_] := Table[4, {i, 1, FilledSize - lengthofgeneitself}]
FilledVec[vecvar5_] := Join[Flatten[vecvar5], Filler[vecvar5]]
For[npow = 1, npow < 1000, npow++, If[lengthofgeneitself ≤ (2^npow), Break[]]];
(* gives npow such that 2^npow > lengthofgeneitself > 2^(npow - 1) *)
FilledSize = 2^npow;
FilledM = FilledVec[M];
numrowsW =  $\sqrt{\text{Length[FilledM]}}$ ;

```

In[]:=

```

W = Table[Table[FilledM[[i]],
  {i, ((j - 1) * (numrowsW)) + 1}, {j * (numrowsW)}], {j, 1, numrowsW}];

```

In[]:= Dimensions[W]

Out[]:= {256, 256}

In[]:= $\rho = (W.\text{Transpose}[W])$;In[]:= Dimensions[ρ]

Out[]:= {256, 256}

```

In[ ]:= (*numgenesample=LetterDNAToNum[lettersample];
lengthofgeneitself=Length[Flatten[numgenesample]];
M=numgenesample;
lengthvec[M_]:=Length[M[[1,All]]]
For[npow=1,npow<1000,npow++,If[lengthvec[M]<(2^(npow)),Break[]]];
FilledSize=2^(npow+1)];
Filler[M_]:=Table[4,{i,1,FilledSize-lengthvec[M]}]
FilledVec[M_]:=Join[Flatten[M],Filler[M]]
lengthvec[M_]:=Length[M[[1,All]]]
Filler[M_]:=Table[4,{i,1,FilledSize-lengthvec[M]}]
FilledVec[M_]:=Join[Flatten[M],Filler[M]]
For[npow=1,npow<1000,npow++,If[lengthvec[M]≤(2^npow),Break[]]];
(* gives npow such that 2^npow > lengthvec[M] > 2^(npow -1) *)
FilledSize=2^npow;
FilledM=FilledVec[M];
numrowsW=√Length[FilledM];*)

```

W for PCV1 samples

PCV Type 1 samples

PCV1₁ =

```

{{0, 1, 1, 0, 2, 1, 2, 1, 0, 1, 3, 3, 1, 2, 2, 1, 0, 2, 1, 2, 2, 1, 0, 2, 1, 0, 1, 1, 3, 1, 2, 2, 1,
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```

```

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0, 0, 2, 3, 2, 0, 0, 0, 2, 0, 0, 2, 3, 2, 1, 2, 1, 3, 2, 1, 3, 2, 3, 0, 2, 3, 0, 3, 3}};

```

PCV1₂ =

```

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0, 0, 2, 3, 2, 0, 0, 0, 2, 0, 0, 2, 3, 2, 1, 2, 1, 3, 2, 1, 3, 2, 3, 0, 2, 3, 0, 3, 3});

```

Defn of W

```

(*Export["PCV1_1genesample.txt",Flatten[PCV11]]
Export["PCV1_2genesample.txt",Flatten[PCV12]]
Export["PCV1_3genesample.txt",Flatten[PCV13]]
Export["PCV1_4genesample.txt",Flatten[PCV14]]*)
(*^These give the DNA data in the form we need in python *)

```

We constructed the W for $M = \{\text{Join}[PCV1_1[[1]]], PCV1_2[[1]]]\}$; in python

(DensityMatrixREcalc.py)

Only used two genes because this particular choice has length $4096 = 2^{12}$ after filling
i.e. b is even, which helps make the algorithm a bit simpler

```

SpecialNote = " We use 2 samples of PCV1 Genes here
so that we have an even b,  $2^b = \text{Length of the filled data set}$  ";

```

```

lengthofgeneitself = Length[Flatten[{Join[{PCV11[[1]], PCV12[[1]]}]}]]
(*To make sure no base pairs are left out *)

```

3518

```

Wgenesample = "(2 sample) PCV1 gene"
(*Lets us know which gene we're dealing with,
used in pdf coding later, so be sure to name it *)

```

(2 sample) PCV1 gene

```

W = {{0, 0, 1, 1, 1, 1, 0, 0, 2, 2, 1, 1, 2, 2, 1, 1, 0, 0, 1, 1, 3, 3, 3, 3, 1, 1, 2, 2, 2, 2, 1, 1,
0, 0, 2, 2, 1, 1, 2, 2, 2, 2, 1, 1, 0, 0, 2, 2, 1, 1, 0, 0, 1, 1, 1, 1, 3, 3, 1, 1, 2, 2, 2, 2},
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```

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```

[illegible]

[illegible]

W for H1A

Define a 1x781 vector H1Avec which contains all the base pair info of the H1A gene

```
H1Avec =
{{3, 0, 2, 2, 1, 3, 2, 1, 2, 3, 3, 2, 2, 2, 2, 1, 1, 3, 3, 3, 3, 3, 3, 3, 1, 2, 1, 0, 3, 1, 1, 3, 2,
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  3, 1, 1, 1, 2, 1, 1, 1, 1, 1, 2, 1, 1, 2, 1, 3, 3, 1, 3, 2, 1, 3, 2, 1, 3, 1, 1, 3, 2, 0,
  2, 0, 0, 0, 1, 1, 3, 3, 3, 0, 2, 1, 3, 2, 2, 1, 0, 0, 2, 0, 0, 2, 2, 1, 0, 0, 0, 2, 0, 0,
  0, 1, 1, 3, 2, 1, 3, 0, 0, 2, 2, 1, 3, 2, 1, 0, 2, 1, 0, 2, 1, 1, 3, 1, 1, 0, 0, 2, 0, 0,
  0, 0, 0, 0, 1, 1, 1, 2, 1, 3, 2, 2, 1, 1, 1, 3, 3, 1, 1, 2, 3, 2, 3, 1, 0, 2, 0, 2, 1, 3,
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  2, 1, 2, 3, 2, 2, 3, 2, 2, 3, 2, 3, 1, 2, 3, 3, 2, 2, 1, 0, 2, 1, 3, 1, 3, 3, 0, 0,
  0, 0, 0, 2, 2, 1, 2, 1, 3, 2, 2, 1, 2, 2, 1, 1, 2, 1, 0, 2, 2, 1, 3, 0, 1, 2, 0, 1, 2, 3,
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  3, 0, 0, 0, 0, 0, 2, 1, 3, 1, 0, 0, 0, 0, 0, 2, 2, 1, 1, 0, 1, 2, 2, 2, 2, 2, 1, 3, 0, 2,
  1, 0, 0, 0, 0, 0, 2, 0, 2, 1, 2, 3, 1, 0, 0, 2, 0, 1, 3, 1, 1, 2, 0, 0, 0, 0, 0, 2, 2, 1,
  3, 0, 0, 0, 0, 0, 2, 1, 1, 3, 2, 1, 2, 2, 1, 0, 0, 1, 0, 0, 2, 2, 0, 0, 0, 3, 1, 1, 3, 1,
  1, 0, 0, 2, 0, 0, 3, 1, 1, 0, 0, 0, 0, 0, 0, 1, 1, 1, 0, 0, 0, 0, 1, 3, 2, 3, 0, 0, 0,
  2, 1, 1, 1, 0, 0, 2, 0, 0, 0, 2, 3, 0, 2, 1, 3, 0, 0, 0, 0, 2, 1, 1, 1, 3, 2, 1, 3, 0,
  0, 0, 2, 1, 3, 0, 0, 2, 2, 1, 3, 2, 3, 0, 0, 0, 0, 1, 1, 1, 0, 0, 2, 2, 1, 2, 2, 1, 1,
  0, 0, 2, 2, 1, 3, 0, 2, 2, 2, 3, 2, 0, 1, 2, 0, 0, 2, 1, 1, 0, 0, 0, 2, 0, 1, 3, 2, 1,
  1, 0, 0, 0, 1, 1, 1, 0, 0, 2, 0, 0, 0, 2, 1, 2, 2, 1, 0, 1, 1, 1, 0, 0, 2, 0, 0, 0, 0,
  0, 2, 3, 0, 0, 0, 3, 3, 1, 0, 2, 3, 3, 0, 2, 0, 0, 2, 3, 3, 3, 1, 3, 3, 1, 3, 0, 2, 3,
  0, 0, 1, 1, 1, 0, 0, 1, 2, 2, 1, 3, 1, 3, 3, 3, 3, 0, 0, 2, 0, 2, 1, 1, 0, 1, 1, 3, 0}};
```

```
Export["H1A_genesample.txt", Flatten[H1Avec]]
```

H1A_genesample.txt

```
SpecialNote = " ";
```

```
lengthofgeneitself = Length[Flatten[H1Avec]] (*To make sure no base pairs are left out *)
```

781

```
Wgenesample = "Human H1A gene"
(*Lets us know which gene we're dealing with,
used in pdf coding later, so be sure to name it *)
```

Human H1A gene

```
W = {{3, 0, 2, 2, 1, 3, 2, 1, 2, 3, 3, 2, 2, 2, 2, 1, 1, 3, 3, 3, 3, 3, 3, 3, 1, 2, 1, 0, 3, 1, 1, 3},
      {2, 1, 3, 3, 1, 2, 3, 1, 0, 2, 2, 3, 3, 3, 0, 3, 0, 1, 1, 0, 1, 3, 3, 3, 0, 3, 3, 3, 2, 2, 3, 2},
      {3, 2, 1, 3, 2, 3, 2, 3, 3, 0, 2, 3, 1, 0, 1, 1, 0, 3, 2, 3, 1, 3, 2, 0, 0, 0, 1, 0, 2, 3, 2, 1},
      {1, 3, 1, 1, 1, 2, 1, 1, 1, 1, 1, 2, 1, 1, 2, 1, 3, 3, 1, 3, 2, 1, 3, 2, 1, 3, 1, 1, 3, 2, 0, 2},
      {0, 0, 0, 1, 1, 3, 3, 3, 0, 2, 1, 3, 2, 2, 1, 0, 0, 2, 0, 0, 2, 2, 1, 0, 0, 0, 2, 0, 0, 0, 1, 1},
      {3, 2, 1, 3, 0, 0, 2, 2, 1, 3, 2, 1, 0, 2, 1, 0, 2, 1, 1, 3, 1, 1, 0, 0, 2, 0, 0, 0, 0, 0, 1},
      {1, 1, 2, 1, 3, 2, 2, 1, 1, 1, 3, 3, 1, 1, 2, 3, 2, 3, 1, 0, 2, 0, 2, 1, 3, 2, 0, 3, 1, 2, 3, 2},
      {1, 0, 2, 2, 1, 3, 2, 1, 3, 3, 1, 1, 3, 1, 1, 3, 1, 3, 0, 0, 2, 2, 0, 2, 1, 2, 3, 2, 2, 3, 2, 2},
      {3, 2, 3, 2, 3, 1, 2, 3, 3, 2, 2, 1, 0, 2, 1, 3, 1, 3, 3, 0, 0, 0, 0, 0, 2, 2, 1, 2, 1, 3, 2, 2},
      {1, 2, 2, 1, 1, 2, 1, 0, 2, 2, 1, 3, 0, 1, 2, 0, 1, 2, 3, 2, 2, 0, 2, 0, 0, 2, 0, 0, 1, 0, 0, 1},
      {0, 2, 1, 1, 2, 1, 0, 3, 3, 0, 0, 2, 1, 3, 2, 2, 2, 1, 0, 3, 3, 0, 0, 2, 0, 2, 1, 1, 3, 2, 2, 3},
      {0, 0, 2, 1, 0, 0, 2, 2, 2, 0, 0, 1, 2, 3, 3, 2, 2, 3, 2, 1, 0, 2, 0, 1, 0, 0, 0, 2, 2, 2, 3, 0},
      {1, 1, 2, 2, 0, 2, 1, 1, 3, 1, 2, 2, 2, 3, 3, 1, 1, 3, 3, 1, 0, 0, 2, 1, 3, 1, 0, 0, 1, 0, 0, 2},
      {0, 0, 2, 2, 1, 2, 3, 1, 1, 3, 1, 1, 2, 3, 2, 2, 0, 0, 0, 1, 1, 0, 0, 2, 1, 1, 1, 2, 2, 1, 2, 1},
      {1, 3, 1, 0, 0, 0, 2, 2, 3, 2, 2, 1, 3, 0, 1, 0, 0, 0, 0, 0, 1, 3, 0, 0, 2, 2, 1, 0, 0, 1, 2, 2},
      {2, 3, 2, 1, 0, 3, 1, 3, 0, 0, 0, 0, 0, 2, 1, 3, 1, 0, 0, 0, 0, 0, 2, 2, 1, 1, 0, 1, 2, 2, 2, 2},
      {2, 1, 3, 0, 2, 1, 0, 0, 0, 0, 0, 2, 0, 2, 1, 2, 3, 1, 0, 0, 2, 0, 1, 3, 1, 1, 2, 0, 0, 0, 0, 0},
      {2, 2, 1, 3, 0, 0, 0, 0, 0, 2, 1, 1, 3, 2, 1, 2, 2, 1, 0, 0, 1, 0, 0, 2, 2, 0, 0, 0, 3, 1, 1, 3},
      {1, 1, 0, 0, 2, 0, 0, 3, 1, 1, 0, 0, 0, 0, 0, 0, 0, 1, 1, 1, 0, 0, 0, 0, 1, 3, 2, 3, 0, 0, 0, 2},
      {1, 1, 1, 0, 0, 2, 0, 0, 0, 2, 3, 0, 2, 1, 3, 0, 0, 0, 0, 2, 1, 1, 1, 3, 2, 1, 3, 0, 0, 0, 2, 1},
      {3, 0, 0, 2, 2, 1, 3, 2, 3, 0, 0, 0, 0, 1, 1, 1, 0, 0, 2, 2, 1, 2, 2, 1, 1, 0, 0, 2, 2, 1, 3, 0},
      {2, 2, 2, 3, 2, 0, 1, 2, 0, 0, 2, 1, 1, 0, 0, 0, 2, 0, 1, 3, 2, 1, 1, 0, 0, 0, 1, 1, 1, 0, 0, 2},
      {0, 0, 0, 2, 1, 2, 2, 1, 0, 1, 1, 1, 0, 0, 2, 0, 0, 0, 0, 0, 2, 3, 0, 0, 0, 0, 3, 3, 1, 0, 2, 3},
      {0, 2, 0, 0, 2, 3, 3, 3, 1, 3, 3, 1, 3, 0, 2, 3, 0, 0, 1, 1, 1, 0, 0, 1, 2, 2, 1, 3, 1, 3, 3, 3},
      {3, 0, 0, 2, 0, 2, 1, 1, 0, 1, 1, 3, 0, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4},
      {4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4},
      {4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4},
      {4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4},
      {4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4},
      {4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4},
      {4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4},
      {4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4}};
```

```
Length[W]
```

```
Length[W[[1]]]
```

```
Print["This should be 1024 for H1A: ", Length[W] * Length[W[[1]]]]
```

```
32
```

```
32
```

```
This should be 1024 for H1A: 1024
```

Run this for output and pdf

```
In[ ]:= EigensofW = Eigenvalues[W] // N
```

```
Out[ ]:= {483.848, 19.8583 + 6.96768 i, 19.8583 - 6.96768 i, -4.00531 + 18.8576 i, -4.00531 - 18.8576 i,
0.780021 + 18.5963 i, 0.780021 - 18.5963 i, -14.8901 + 11.0216 i, -14.8901 - 11.0216 i,
-17.4194 + 4.58851 i, -17.4194 - 4.58851 i, 13.727 + 11.6623 i, 13.727 - 11.6623 i,
5.50844 + 17.0659 i, 5.50844 - 17.0659 i, 17.907, 16.802 + 5.88984 i, 16.802 - 5.88984 i,
-17.6938 + 1.52994 i, -17.6938 - 1.52994 i, 2.7767 + 17.3978 i, 2.7767 - 17.3978 i,
-15.8433 + 6.40316 i, -15.8433 - 6.40316 i, -10.4457 + 13.3045 i, -10.4457 - 13.3045 i,
11.7951 + 11.9015 i, 11.7951 - 11.9015 i, 8.85116 + 14.2089 i, 8.85116 - 14.2089 i,
-4.3147 + 16.16 i, -4.3147 - 16.16 i, -3.34967 + 16.183 i, -3.34967 - 16.183 i,
-11.2547 + 11.7013 i, -11.2547 - 11.7013 i, -8.99223 + 13.3767 i, -8.99223 - 13.3767 i,
13.0153 + 9.36275 i, 13.0153 - 9.36275 i, 15.8232 + 1.46337 i, 15.8232 - 1.46337 i,
14.2088 + 7.0725 i, 14.2088 - 7.0725 i, 5.87228 + 14.5199 i, 5.87228 - 14.5199 i,
-12.3893 + 9.41411 i, -12.3893 - 9.41411 i, -14.9914 + 3.96472 i, -14.9914 - 3.96472 i,
-2.08969 + 15.229 i, -2.08969 - 15.229 i, 14.5249 + 4.83292 i, 14.5249 - 4.83292 i, -15.1298,
-4.34439 + 14.2896 i, -4.34439 - 14.2896 i, 1.26689 + 14.7232 i, 1.26689 - 14.7232 i,
14.7278, -14.6933, 10.356 + 10.331 i, 10.356 - 10.331 i, -6.1602 + 13.1693 i,
-6.1602 - 13.1693 i, 8.80256 + 11.5068 i, 8.80256 - 11.5068 i, 4.07596 + 13.747 i,
4.07596 - 13.747 i, -11.9787 + 7.8384 i, -11.9787 - 7.8384 i, 4.78932 + 13.2623 i,
4.78932 - 13.2623 i, -7.12071 + 12.1164 i, -7.12071 - 12.1164 i, 13.8338 + 0.49743 i,
13.8338 - 0.49743 i, 12.1535 + 6.27809 i, 12.1535 - 6.27809 i, -10.567 + 8.64602 i,
-10.567 - 8.64602 i, -12.7852 + 4.75438 i, -12.7852 - 4.75438 i, -13.3918,
-12.9044 + 3.56866 i, -12.9044 - 3.56866 i, 10.5261 + 8.1339 i, 10.5261 - 8.1339 i,
11.0537 + 7.32294 i, 11.0537 - 7.32294 i, -8.04821 + 10.4704 i, -8.04821 - 10.4704 i,
-2.78767 + 12.7516 i, -2.78767 - 12.7516 i, 12.9311, 2.04162 + 12.7079 i, 2.04162 - 12.7079 i,
-0.586609 + 12.7444 i, -0.586609 - 12.7444 i, -11.9498 + 4.14973 i, -11.9498 - 4.14973 i,
8.13711 + 9.52736 i, 8.13711 - 9.52736 i, -8.5541 + 9.0017 i, -8.5541 - 9.0017 i,
11.9562 + 2.80519 i, 11.9562 - 2.80519 i, -5.41773 + 10.9764 i, -5.41773 - 10.9764 i,
-10.3646 + 6.09143 i, -10.3646 - 6.09143 i, 6.88196 + 9.72759 i, 6.88196 - 9.72759 i,
-3.73646 + 11.2323 i, -3.73646 - 11.2323 i, -2.39128 + 11.5907 i, -2.39128 - 11.5907 i,
-2.90984 + 11.3502 i, -2.90984 - 11.3502 i, -11.6004, 5.48515 + 10.1858 i,
5.48515 - 10.1858 i, -0.961992 + 11.3353 i, -0.961992 - 11.3353 i, 8.53358 + 7.47957 i,
8.53358 - 7.47957 i, 11.194, 10.7642 + 2.96244 i, 10.7642 - 2.96244 i, 2.70555 + 10.786 i,
2.70555 - 10.786 i, -10.4185 + 2.96466 i, -10.4185 - 2.96466 i, -8.50841 + 6.68741 i,
-8.50841 - 6.68741 i, 9.78122 + 4.44744 i, 9.78122 - 4.44744 i, 6.97967 + 8.07633 i,
6.97967 - 8.07633 i, 3.83441 + 9.86985 i, 3.83441 - 9.86985 i, -6.23061 + 8.37987 i,
-6.23061 - 8.37987 i, -10.1229 + 1.89246 i, -10.1229 - 1.89246 i, 10.1699 + 1.11161 i,
10.1699 - 1.11161 i, 0.48782 + 10.1037 i, 0.48782 - 10.1037 i, -3.91686 + 9.20553 i,
-3.91686 - 9.20553 i, -8.74962 + 4.26766 i, -8.74962 - 4.26766 i, 1.98637 + 9.31578 i,
1.98637 - 9.31578 i, -5.85892 + 7.403 i, -5.85892 - 7.403 i, 5.99014 + 7.18589 i,
5.99014 - 7.18589 i, 7.61044 + 5.33771 i, 7.61044 - 5.33771 i, -7.25101 + 5.51437 i,
-7.25101 - 5.51437 i, -8.32108 + 3.00497 i, -8.32108 - 3.00497 i, 8.11617 + 2.96488 i,
8.11617 - 2.96488 i, -6.10965 + 6.00601 i, -6.10965 - 6.00601 i, 6.52477 + 5.54903 i,
6.52477 - 5.54903 i, -3.38787 + 7.49355 i, -3.38787 - 7.49355 i, -0.343758 + 7.88408 i,
-0.343758 - 7.88408 i, -7.86774, 7.62836 + 0.690443 i, 7.62836 - 0.690443 i,
-7.25496 + 1.26918 i, -7.25496 - 1.26918 i, 6.80626 + 2.08624 i, 6.80626 - 2.08624 i,
-5.04238 + 5.02122 i, -5.04238 - 5.02122 i, 6.20929 + 3.09539 i, 6.20929 - 3.09539 i,
-6.42482 + 2.48272 i, -6.42482 - 2.48272 i, 2.83363 + 6.23469 i, 2.83363 - 6.23469 i,
```



```

Wgenesample
ρ = (W.Transpose[W]); (* ρ as inner product *)
rhoEigens = Sort[DeleteCases[Eigenvalues[ρ] // N, 0.], Greater];
(*DeleteCases Removes 0's from the set of Eigenvalues,
Sort puts the list in order of greatest to least *)
set =  $\frac{\text{rhoEigens}}{\text{Total[rhoEigens]}}$ ;
(* This is the set of nonzero normalized eigenvalues in order of greatest to least *)
n = Length[set];
 $H[\alpha_] := \frac{1}{1-\alpha} \text{Log}[\text{Sum}[(\text{set}[[i]])^\alpha, \{i, 1, n\}]] // N$ 
H0 = Log[n] // N; (* H0 = Hartley Entropy*)
H1 = -Sum[(set[[i]]) (Log[set[[i]]]), {i, 1, n}] // N; (* H1 = Shannon Entropy*)
H2onward = Table[H[a], {a, 2, 20}] // N; (* H2 onward *)
RenyiEntropyofEigenvalues = Join[{H0}, {H1}, H2onward];

```

Out[]:= Human chromosome9 scaffold gene

In[]:= Abs[rhoEigens]

Out[]:= {266.021., 3806.06, 3389.28, 2450.78, 1912.6, 1773.95, 1645.93, 1562.73, 1421.38, 1393.77, 1338.46, 1279.79, 1258.53, 1237.62, 1202.43, 1147.59, 1134.25, 1117.5, 1091.67, 1042.43, 1004.17, 993.133, 975.255, 952.125, 932.025, 923.146, 894.656, 871.752, 856.101, 848.158, 837.193, 827.084, 802.064, 789.353, 782.585, 764.194, 759.974, 744.507, 733.538, 727.8, 709.079, 696.459, 683.526, 669.329, 660.707, 650.239, 648.121, 630.997, 621.876, 620.126, 610.721, 605.601, 594.97, 588.609, 581.376, 562.353, 553.969, 548.08, 530.752, 520.77, 515.185, 504.421, 498.906, 491.608, 481.472, 473.282, 471.475, 462.801, 458.669, 448.371, 444.36, 434.536, 429.254, 421.526, 414.627, 404.69, 402.189, 392.587, 385.249, 376.745, 374.051, 361.052, 357.922, 353.906, 346.076, 340.597, 337.628, 331.13, 326.501, 319.903, 317.057, 314.908, 308.495, 307.951, 298.837, 298.009, 293.027, 286.488, 281.229, 274.199, 272.686, 267.677, 261.447, 255.358, 244.996, 242.571, 233.2, 231.124, 229.491, 225.681, 224.051, 221.124, 218.349, 216.522, 211.058, 204.307, 201.943, 195.608, 191.972, 190.154, 187.058, 185.029, 181.413, 175.733, 174.211, 168.877, 167.461, 164.878, 158.908, 158.374, 155.961, 154.325, 148.703, 145.967, 145.18, 142.788, 142.085, 139.663, 135.557, 129.041, 125.922, 122.917, 121.442, 120.44, 117.932, 115.184, 112.394, 110.399, 107.631, 105.408, 103.149, 99.939, 98.7483, 94.5576, 92.5514, 89.7287, 88.9743, 85.5292, 84.2914, 82.9885, 80.7292, 79.1281, 76.2596, 72.9578, 71.1938, 68.4218, 66.5738, 64.591, 63.4024, 62.9802, 60.479, 59.559, 57.5124, 54.9998, 53.9055, 52.8349, 50.2557, 49.1445, 47.7769, 45.0571, 44.2219, 41.9247, 41.4539, 40.8924, 40.0632, 37.5205, 36.0087, 35.5347, 34.1811, 32.6806, 32.5248, 29.3879, 28.2356, 27.6005, 26.9967, 24.4664, 23.2797, 22.5026, 21.8369, 21.2764, 19.8888, 18.8483, 17.3046, 15.8729, 15.5208, 14.2935, 13.7374, 13.1778, 12.4535, 12.0886, 11.3875, 10.6453, 9.54106, 8.9693, 8.82021, 8.41967, 7.65915, 6.62035, 5.99868, 5.08291, 4.55845, 4.44252, 4.31448, 3.77024, 2.98685, 2.77102, 2.42774, 2.00658, 1.73301, 1.60463, 1.11771, 0.830514}

In[]:= RenyiEntropyofEigenvalues

Out[]:= {5.44674, 2.52786, 0.82095, 0.617115, 0.548553, 0.514268, 0.493697, 0.479984, 0.470188, 0.462841, 0.457127, 0.452556, 0.448816, 0.445699, 0.443062, 0.440801, 0.438842, 0.437128, 0.435615, 0.434271, 0.433068}

In[]:=

```

button =
  Button["Click here for output and pdf", Print[Style[Wgenesample, Black, Bold, 28]] ×

```

```

Print[Style["The ", Blue, Italic, 18], Style[Wgenesample, Black, Italic, 18],
  Style[" has ", Blue, Italic, 18], Style[lengthofgeneitself, Black, Italic, 18],
  Style[" base pairs ", Blue, Italic, 18]] ×
If[StringLength[SpecialNote] > 3, Print[Style["(Special Note): ", Black, Bold, 16],
  Style[SpecialNote, Black, Italic, 12]], Print[" "]] ×
Print[Style["W is a ", Blue, Italic, 18], Style[Length[W], Black, Italic, 18],
  Style[" by ", Blue, Italic, 18], Style[Length[W[[1]]], Black, Italic, 18],
  Style[" matrix with ", Blue, Italic, 18],
  Style[Length[W] * Length[W[[1]]], Black, Italic, 18],
  Style[" = 2^b elements", Blue, Italic, 18], Style[" for b = ", Blue, Italic, 18],
  Style[Log[2, Length[W] * Length[W[[1]]]], Black, Italic, 18]] ×
If[(Length[W] * Length[W[[1]])] == (Length[W])^2,
  Print[Style["(If statement safecheck): ", Black, Bold, 12],
    Style[Length[W], Black, Italic, 12], Style[" times ", Red, Italic, 12],
    Style[Length[W[[1]]], Black, Italic, 12],
    Style[" equals ", Red, Italic, 12], Style[(Length[W]^2), Black, Italic, 12],
    Style[" W is of the right size, you may proceed ", Red, Italic, 12]],
  Print[Style["(If statement safecheck): ", Black, Bold, 12],
    Style["Warning!!!", Red, Italic, 28],
    Style[" W is of wrong size, STOP and check W ", Red, Italic, 12]]] ×
Print["The number of nonzero eigenvalues is = ", Length[rhoEigens]] ×
Do[Print["The i-th Eigenvalue " $\lambda_i$ , " is = ", (rhoEigens)[[i]],
  {i, 1, Length[rhoEigens]}] ×
Print[Graphics[ListPlot[rhoEigens // N, AxesLabel → {Style["i", Medium, Bold],
  Style[" $\lambda_i$ ", Medium, Bold]}, PlotLabel → "Eigenvalue PLOT"]]] ×
Print[Graphics[ListLogPlot[rhoEigens // N, AxesLabel → {Style["i", Medium, Bold],
  Style["Log[ $\lambda_i$ ]", Medium, Bold]}, PlotLabel → "Eigenvalue Log PLOT"]]] ×
Print["Zooming in on the Log Plot so as to Exclude the first
  eigenvalue gives the following plot:"] ×
Print[Graphics[ListLogPlot[Table[{i, rhoEigens[[i]]}, {i, 2, Length[rhoEigens]}],
  AxesLabel → {Style["i", Medium, Bold], Style["Log[ $\lambda_i$ ]", Medium, Bold]},
  PlotRange → {{10, 2 * rhoEigens[[2]]}}, PlotStyle → Red,
  PlotLabel → Style["Logplot of Eigenvalues, excluding  $\lambda_1$ ", Red, Bold, 16]]] ×
Print["The approximate linearity of the above plot tells us
  that the eigenvalues decrease exponentially. If it's
  nowhere near linear try adjusting the plot range. "]
×
Print[" "] ×
Print[Style[
  " ", 18]] ×
Print[" "]
×
Print[Style["The First normalized eigenvector is: ", Blue, Italic, 18],
  Style[set[[1]], Blue, Italic, 18]] ×
Print[Style["The Second normalized eigenvector is: ", Blue, Italic, 18],
  Style[set[[2]], Blue, Italic, 18]] ×
Print[Style["The Last (n-th) normalized (nonzero) eigenvector is: ",
  Blue, Italic, 18], Style[set[[n]], Blue, Italic, 16]] ×
If[Total[set] == 1, Print[Style["(If statement safecheck): ", Black, Bold, 12],
  Style["Total[set] = ", Red, Italic, 12], Style[Total[set], Black, Italic, 12],

```

```

Style[" = 1, so the Eigenvalue set is properly normalized", Red, Italic, 12]],
Print[Style["(If statement safecheck): ", Black, Bold, 12],
Style["Warning!!!", Red, Italic, 28], Style[" Total[set] = ", Red, Italic, 12],
Style[Total[set], Black, Italic, 12], Style[" ≠ 1, ", Red, Italic, 12],
Style[" so the Eigenvalue set is NOT properly normalized.", Red, Italic, 12],
Style[" This will render the entropies invalid. Fix it. ", Red, Italic, 12]]]
×
Print[" "] ×
Print[Style[
" ", 18]] ×
Print[" "]
×
Do[Print["The  $\alpha$ -th Renyi Entropy  $H_\alpha \rightarrow H_{i-1}$ , " is = ",
RenyiEntropyofEigenvalues[[i]]], {i, 1, Length[RenyiEntropyofEigenvalues]}] ×
Print[Graphics[Show[
ListPlot[RenyiEntropyofEigenvalues, PlotRange → All,
AxesLabel → {Style[" $\alpha$ ", Large, Bold], Style[" $H_\alpha$ ", Large, Bold]}],
ListLinePlot[RenyiEntropyofEigenvalues, PlotStyle → {Red, Thin}]
]]] ×
Export["rhoEigenEntropies.pdf", EvaluationNotebook[]] ×
NotebookSave[EvaluationNotebook[], "rhoCalcOutput"];
SystemOpen["rhoEigenEntropies.pdf"]
(*DeleteFile[StringReplace["rhoEigenEntropies.pdf", "rho"→ Wgenesample]]
RenameFile["rhoEigenEntropies.pdf",
StringReplace["rhoEigenEntropies.pdf", "rho"→ Wgenesample]]
SystemOpen[StringReplace["rhoEigenEntropies.pdf", "rho"→ Wgenesample]]*)
, Background → Green];
nb = CreateDocument[];
Paste[nb, button]
NotebookEvaluate[nb];

```

Possible Issues

Be sure that W contains only integer values (e.g. 0, 1, 2, 3, 4).

Including floats (e.g. 0.0, 1.0, 2.0, 3.0, 4.0) may incorrectly make some of the eigenvalues negative or imaginary because of how mathematica handles floats differently than it handles integers.