

# Density Matrix Calculation

## Misc.

```
n = Length[Union[SET]] (* i.e. n = number of unique elements in the set *)  
H[α_, SET_] :=  $\frac{1}{1-\alpha} \text{Log}[\text{Sum}[(\text{prob}_i)^\alpha, \{i, 1, n\}]]$   
  
Count[{a, b, a, d, b, c, b}, b]  
Length[{a, b, a, d, b, c, b}]  
 $\frac{\text{Count}[\{a, b, a, d, b, c, b\}, b]}{\text{Length}[\{a, b, a, d, b, c, b\}]}$  (* Prob of getting b at random *)  
Length[Union[{a, b, a, d, b, c, b}]] (* i.e. = number of unique elements in the set *)  
3  
7  
 $\frac{3}{7}$   
4
```

## PrimePi Renyi Entropy Calc

## PCV1 Renyi Entropy Calc

### General Definitional Terms

```
(* M=PCV11; *)  
(* M={Join[PCV11[[1]],PCV12[[1]],PCV13[[1]],PCV14[[1]]]}; *)  
(* M={Table[PCV11[[1,i]],{i,1,64}]}; *)  
  
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;
```

## PCV Type 1 samples

Renyi Entropies of PCV1 data sets

Density matrix calc Using  $2 \times 2^{(n/2)}$  defn of  $W_0$  (not valid)

Density Matrix Calculation

Wrong  $\rho$ 's (as inner and outer products of  $W$ ,  $\text{Transpose}[W]$ )

Defn of  $W$  (RUN THIS before doing the below)

Misc. Code

---

## Pre/detailed

---

## 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;
LetterDNAtNum[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,"}]}],
    ],
    ",}" → ""]}

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

numgenesample = LetterDNAtNum[lettersample];
Flatten[numgenesample] [[3]]
2

lettersample = {ACGTAGTCAATT} // ToString;
LetterDNAtNum[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 = LetterDNAtNum[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 ×
GGCCAAAAAGAGCGGCACCTCGGAAGGCGAGTTATTTACCAAGCACTGGAGTAATATTGTAGATAAAAAAT ×
GCCTGTTGGATGCAAGAGAGGCCAACATTTTTTGAATTTTAAAGACGCGGTGCAATCAAGCAGATTTA ×
GGACCAATAAGCCTTAATTGGTTGAAGAACTTTCTTTAGAAGCTCCACCCTATAATTCTGAACCCACAG ×
AAGAATCTGGTTATAAAATCAGCTATGAACCAACCTATTTAAACACCACAAAGGAAACCTTATAATCA ×
GTTGGCTTCAACTCCAATAGTATTAGAGAGCCAATATACCAACAATCTCCTTTAAAGAATTAGATAAA ×
TACAGATTAGATTCAGGAAAGGATATTACCGATAGTAAACATAAAAGTTGTTGCACAAATGAAGCTAAAA ×
TGGATCGAGCAAAATGATGTTACCAGCCACCTCTAAATTCTTATCTTAGTGAAAGTCCTGTTCTACGATG ×
TACACATGTAACACCACAGAGAGAAAAAGTCGGTGGTATGTGGAAGCTTATTCATACACCGAAGCTTATG ×
AAGGGTCAGACACCAAAACGCATTTCTGAAAGCTAGGAGCTGAGGTGGATTCTGATATGTCTTGGTCAA ×
GTTCTTTAGCCACACCACCAACCCTTAGTTCTACTGTGCTAATAGTCAGAGATGAAGAAGTATCTGCAGC ×
TGTAATTTCTAATGACACTACTGTATTTTTTAAAGCTGTTTTTCTAACCATGATGAAAGCTGAAGAAA ×
AATGATCGGTTTATCCCTTGTGGTCCAGGCAAAGAAAACAAAATCAAAGGGAAGCTAAAAGTCAAAGTT ×
TGGGGAATTCATTTGGTAAAGTAAATAGCACCAAGACCATTTGTAAAGTCTACACCAATGTCCTAGA ×
GGATGAAGTACATGAAAAAGTTCTAGATGTTTCTGAAGAAGAAGATAGTTTTTCATTATGTGTTCTCTAAA ×
TATAAAACAAGAAATCTACAAAAATAAACTAGCAAACTAGGAAAAATATTTTAAATGAGACAAAAA ×
CCAGTGAATGTGAAGAAGCTAAAAAGCAAATGAAAGAAAATAAACATTCATTGGTATCTGAAATGGAACC ×
AAATGACAGTCATCCATTAGATTGGAATGTAACACATGAGAAGCCCTTTGGGAATGGAAGTACAAAAATC ×
TCCAAGGAAATTGTACTGTCTTCAGCCTCTGGATGTTCTGACCTAACCTCTCAAGTCTAAATGGAGCTC ×

```

AGATGGAGAAAAACACCTCTATTGCATACTTCTTATGACCAAAATAATTGAGAAAAAGACCTCATAATCAC ×  
 AGATAAAGAATGCACCAACTTCATTACTTTGGAAAATCTTGGCCACAGATTTCAAATGTACCAAAGTAT ×  
 TCAGAGAAGACGTTAAATGAGGAAATAGTAGTAAATAAGATAAACGAAGGGCAGTGTCTTGAATCTCATG ×  
 AAGATTCCGTTGTTTCGGTAAAGCAAGCAATATATGAACTACTTTAATAGCTTCTCCACTTCAGGGTAT ×  
 CAGAAAGTCTATATTAGGATAAGAGAATCACCTGAAGGGATGTCCAATGCAATGTTCTCAAATAATATG ×  
 ACTAATCCAACTTTAAAGAACCTGAAGCCTCTGAAAGTGGATTGGAAAAACATACTATTTGCTCTCAGA ×  
 AAGAGGATTCTTTATGTACAAGTTCAATTGATGATGGAAGCTGGCCAGCAACTATCAAACATACTTCTGT ×  
 AGCTTTGAAGAATTTAGGTTTAAATATCTAGTTTGAAAAAGAAAAACAAAAAGTTTATTTACGTTATAAAT ×  
 GATGAAACATCTAATCAAGGCCGTGAAAACACAGAAAGACCAAGAGTCAAGACTAATTAACCTTTTCGACCC ×  
 AATTTGAAGCAAATGCTTTTGAAGGACCCCTGACATTTACAAATGCTGATTGAGGTTTATTGCATTCTTC ×  
 TTCCATCAAAAAAAGTGTTCACAGAATGACTCAGAAAAACCAGCTTTGTCTTTAACCAGCTCTTTTGGG ×  
 ACAATTCTGAGAAAAGTTTCCAGTAATGGAGCCAGTTCTCCTAATAATAAAATAATATCTCAGGATCCTG ×  
 ATTATAAAGAAGCAAAAATTAATAAGAAAAAATTGGAGTCATTTATAACCACAGAACTGATTGTCTGTC ×  
 ATCCCTGCAGGAAAAACATTGGGAAGATGATGCAAAAAACAAAGAGTTTCAGATATAAAAGAAAAAGTC ×  
 TTGCCTACAGTAAGTCACCTCCTGTGCCACATTGAGAAGTGAAGGTAGTGATATTCACTTTTCAGTCTC ×  
 CAGAAAGCTTTTCATTGACTGTGATAATACCAGTCTGTTAACTCCTAGCTCTAGGGATTCTCCATCAAG ×  
 CCTAGTTGTGATGTCTAGAGGAAAAAGAAATCATATAAAATATCAGAGAACTAAAATGTAAGAATCATGAA ×  
 ACTGTTTTTGAATTAACCAAAAATATTCCCATGGAAGAAGTCAAGACATACATGTTTTAAATGCAGATT ×  
 CTAAAAATGCTAAACTGTTGTCAACTGAAAAACATATAACAGTAGCATCATCTTCAGTAAAGGTTTCAGTT ×  
 CAACCAAAATGCAATCTCACCACAATCCAAAAAGACCAAAAAGAACTACTTTAATTTCAAAAAATACT ×  
 GTTAATCCAACTCTGAAGAATTTTCCAGATGATGAAAATAATTTTGTCTTAAAGATAACTAATGAAA ×  
 GTAATACTCCTGTTTTAGGAAATACTAAGGAACTACATGATTCAAACCTCTGTTGTGTAAGAGATTCTGT ×  
 TCCTAAGAACTCTACCATGGTAGTATGTACAGACCTGGATGACAAACAAACAGCCAAAGTGTGATTATG ×  
 AAAGATTGTTATTCATCAAGCATAGATGATCTTACAGAAAGGAACAGAAGTACCATAAAGCAACAATAA ×  
 AAATGACTCTAGATCAAGATTCAAAATCAGACATTACCTCAGATATAGTTAGGAAATCAAATGGAAACAG ×  
 TGATTATATGGATAATTGGGCAAGACTGTCTGATCCAATTTCAAATCACAGTTTTGAAATGGCTTCAAA ×  
 ACAGCTTCTAATAAAGAGATAAACTCTCTGAAAACAACATTAGGAAAAGTAAAATGCTTTTCAAAGATA ×  
 TTGAGGAACATTATCCTACTAAGTCTAGCATGTCTTGAATTTGTAATACTTCATCATTAGAAAAGTCAAAA ×  
 GAAACCAAGCAAATCTCATGCACTTGATCCACAGTCAATTAATATCATATCTGGGTTTGTGCAGAATAGC ×  
 ACATATGTTTTCTGATAGTGAAAGTGGTCACACAGCTCCTCCAACTTTATCTTTAAAGCAAGATTTTGATT ×  
 CAAATCGTAATTTAACTCCTAGTCAAAAGGCAGAAATTACAGAACTTTCTACTATTTTGAAGAATCAGG ×  
 AAGCCAGTTTGAATTTACACAGTTTAGAAAACCAAGCCACATAATACAGAAAAATCCATTTGAAATGCCT ×  
 GAAAACCAAGCTGACTATCTTGAATAGCACTTCTAAGGAATGGAAAGATGATGATCTTCATCTCACAACTA ×  
 ATGCTCCATCTATCAGTCAGGTAGATAGCAAGAAATCTGAAGGTATAATTGGAGGTAAGCAGAAGTTTGC ×  
 TTGCTTGTCAAGAACCAGCTGTAAACAGAAGTGCTTCTGGCTATTCAACAGATAAAAATGAAGTGGAGTTT ×  
 AGAGGCTTTTATTCTGCTCGTGGCACAAAACCTGAATGTTGGTAGTGAAGCATTGCAAAAAGCTAAGAAAC ×  
 TGTTTCAGTGACCTTGAGAATATCAATGAGGAACTTCTGTAGAAGTAGATAGAAGTTTCTCCTCAAGCAA ×  
 ATACAATGATTCTGTCTCAATGATTGAGATAGAAGATTGTAATGATAAAAAATTTAAATGAGAAAAATAAT ×  
 AAATGCGCGCTAATACTACAAAATAATATTGAAATGACTACTGACATTTTTGTTGAAGAATATACTGAAA ×  
 GTTACAGGAGAAAATACAGAAAATGAAGGTAACCAATGTACTGACGCTGGTAGAAAATCTTGTAACCTCAGA ×  
 ATCTGATGGCAGTGATTCAAGTAAAAATGATACAGTTTATATTGATGAAGAAGAAAATGGCTTGCCCTGT ×  
 ATTGATCAGCACACATAGATCTGAAATTATTTAGCCAGTTTATGAAGGAGGGGAACACTCAAATTAAG ×  
 AAGGTTTGTGAGATTTAACTGTTTGAAGTTATGAAAGCTGAAGAAACATCTCATGTTACTATGTCAAA ×  
 TAAACAGCAGTTAACAGCTAATACGGGGCAAAACATAAAAGATTTTGACACTTTTTATTTATCCTTTTCAG ×  
 ACTGCAAGCAGAAAAAATAAAGGCTCTCCAAAGAGTCATTAATAAAGCTAGAAGTCTCCTTAATCAAA ×  
 AATGGACAGAAGAAGAATTAATAACTTTTCAGATTCCTTGAATTTCTGAATTACTTCTGGCATAGATAT ×  
 CAAGAAAACAGACATCTCAAATCATGAGGTAAATAGAAAATACTGAAAGAAAAGACAAAATAACGAAAGAA ×  
 AGTGACCTAATTGGTACTGAAAATATATTACTGATCTGCAGCAAGACCAGAAAGTAAAATAAAAAAGA ×  
 TCAAAGAATCTGCTGTGTTGGGTTTTATACAGCTAGTGGGAAAAAATAGAAATTACAAAGGAATCTTT ×  
 GGACAAAGTAAAAATCTTTTTGAAGAAAAAGAGCAAGATAATAGTGAAATCACTAATTTTAGCCATCGA ×  
 GGGGCAAAGATGTCCAAGGACAGAGAAGAATGTAAGATGGGCGTGAATTAGCTTGTGGGACAACTGAAA ×



TAACAACTACCCAGAGTATGAAGAACTCACAGTTCTCTAGAGAAGAAAAAATTGTTTCTAATGAGAT ×  
 TGCAGCCTTAAGACCCAGGCTCTTAAGTGATAATTTATACAAACAACTGAAAATCTTAAATATCAGAT ×  
 CATGCCTCTCAGAAAGTTGATGTACATGAAAAACAGAAAAAGCAAAAAAGCCTACAATGTATA ×  
 CAAATCAATCCACTTATTCTAGCCATTGAAAACCTACCTTTAACATTTTACACAGGACACGGAAGAAAAAT ×  
 TTCTGTGAGTGAGGCTTCACTATTTGAAGCAAAAAAATGGCTTAGAGAAGGAGAATGGGATGATCAATCA ×  
 GAAAGAATAAATGCTGCCAAGGTTAACTGCTTAAAGAATATCCTGATGATTACGTAGAAAAATCCTTCAT ×  
 GTGGAAATAGTTCAAATAGTGCCATAACTGAAAATGACAAAAATCATCTCTCTGAAAAACAAGGCTCAAC ×  
 TTATTTAAGTAATAGTACCATGTCTAACAGCTATTCATACCATCCTGGCTTTTGTCTATTCTAGTGAAGTG ×  
 TATAATAAATCAGAATATCTTTCAAGAAGTAAATTTGATAATTTGGTATTGAACCAGTAATAAAGAATA ×  
 TTAGAGAGAGAAAAACATTGGTTTTCTGAAATAATGTCCCCTGGAAGAGAAGCAGACACAGACCCACA ×  
 AAGTGTAATGAAGATATTTGTGTTGAGAACTTGCGACTAACTCTTCATGCAAAAATAAAAATACAGCC ×  
 ATTAAGTGGCCATATCTGACTCAAATAATTTAATACAATTCAAAAGTTGAATTCTGATTCAAATAATT ×  
 CTGTACCTGCATACAGTACAGTAAATAGTAAAGAGTCTTTGTTGCACACCAGACAAAAGTGACAGAGGG ×  
 GTTTACAGACAACTGCAGCATGGTAACTAAACAAAACACCAAGAGTAAATCAGACACTTGCCATGCAGAA ×  
 ATTGTGGCAGATTATCCTAAGGCACTGGATGATTGAGAGGCTATTTTTCTAACTCTCTGGGTGCTATAG ×  
 AATGTTACCTTCACATAAGGTTTTTGTGACATTCAAAGTGAACAACTTCACAACCTAACCAAGTAT ×  
 GTCTGGATTGGAGAAAGTTTCTGAAACACCACCTTGTGAGATTAATTCAAAAACTTCTGATAGATGTGAA ×  
 CTTCTAGGGGGAAGCTTCCCAAGTCAGTCTCTTACACAAATGCATGTGGGATTTTTAGCACAGCAAGTG ×  
 GAAAATCTGTACAAGTATCAGATGCTGCAATACAAAAGGCAAGAGAGGTGTTTTCTAAGCTAGAAGATAG ×  
 TGCCAAGCAACTCTTCTGAAGTATCACTTAAAGATAATGAAGAACATTGAGAAAAGTTCACAAATGAA ×  
 GAAAATACTGTGATATATACCTCCCAAATTTACTATCATCTGCTTTCTCTGGATTTAGGACAGCAAGTG ×  
 GGAACAAGTTCAGTTTCTGAAAGTGCTTATGCAAAGTTAAGGGAATGTTAGAAGAATTCAATCTGAT ×  
 CAGAACTGAAAGTTGTCTTCAGCATTCTACTTCTAGACAAGATGTATCAAAAATGCCTCCTCCCTCT ×  
 TGTATTGGTAAGAGAACCCAGAACACTCCAGAACTCCAAATTGGATAAAGCCTGCAATAAAGAATTTA ×  
 GATTATCAAGTAACTGTAAACATCAGAGTGGTCTTCAGAAAATCATCACTCTATTAAAGTTTCTCCATG ×  
 TCCCTCTCAATTGAAGCGAGACAAACCACAGTTGCTAGTCGGAAGCAAAGGATCACTTGTTGAGAACATT ×  
 CATCCTTTGGGAAAAGAACAAGCTTTACCTAAAAATATAAAAACAGAGATTGGGAAAGCTGAACTTTTTC ×  
 CTAATCTTCTGTGAAAACAAATATAGAATTTTGTCTACTTACTCCAAGGATCCAGAAAACTATTTTGA ×  
 AACAGAAACCGTAGAGATTGCCAAAGCTTTTATGGAAGATGGTGAGCTGACAGATTCCGAACTGCTAAGT ×  
 CATGCCAAACACTTTGTTTTTACATGCCAAAACACTAAGGAAATGGTTTTGTTAAATCAAGAATTGGAA ×  
 AAAGAAGAGGAGATGCACTTGTCTCAGTTGGAGAACCCCAATTAAGAAACTTGTAAATGAATTCGA ×  
 CAGGATAATAAAAAATCAAGAAACATCTTTAAAGCTTCAAAAAGCACTCCAGACGGCATCTAAAGAC ×  
 AGAAGCTTGTATGCATCATATTTCTTTAGAGCCAATTTCTGTGGACCTTTTCGCACAACTGAGGAAC ×  
 GGCAAGAAATACAGAAATCCAAATTTCACTGCACCTGGTCAAGAAATTTTGCCTAAATCTCATTTTTATGA ×  
 ACACCTGGCTTCAGAAAAATCTTCAAGTAATTTATCAGTTTCACGGCAACCATTTTGTATGGTTCCTGCC ×  
 ACAGGAAATGAAAAAAGGAGACACTTGATTGCTCCAGGCAAACCAAGTGAAGTCTTTGTCCCACCTTTTA ×  
 AAACTAAATCACAATTTTACAGAGATGAGCAGTGCATTAGCAAGAATACTAAATTGGAAAAAACAAACA ×  
 AAACTCCAAAGACATAGATGAACCTGGCTCTGGTGATAGTGAAGAAAAATATTAATGACAGTGGAATCCAT ×  
 CAGCTTAAGAAAAATAACTCCAATCAAGCAGCAACTATAATATTCACAAAGAATGAAAAAGAACCTTTAG ×  
 ATTTAATTACAAATCTTCAGAACGCCAGAGATATACAGGATATGCGGATTAAGAAAGAAACAAAGGCAGCA ×  
 TATTTTTCCACAGCCAGGTAGTCTGTATCTTGCAAAAACCTCCACTTTGCCTAGAATCTCTCTGAGAGAA ×  
 GCAGTAGAAGGCCGAGTCCCCTCTGCATGTTCTCATAAACAGCTCTATATGTATGGTGTTCCAAACATT ×  
 GTGTAATAAATAACAGCAAAAATGCAGAGTCTTTTCAGTTTCATGCTCAGGATTATTTTGGTAAGGAAGG ×  
 CCTATGGTCTGGAGAAGGAATACAATTGGCTGATGGTGGATGGCTCATACCCTCCAATGATGGAAAGATT ×  
 GGAAGAAGAAGATTTTATAGGGCTCTGTGTGACACCCAGGTGTGGATCCAAATTGTATTTCTAGAGTTT ×  
 GGGTATATAATCACTATAGATGGATTATATGGAATTTGGCAGCCATGGAATTTGCCTTTCTTAAGGAATT ×  
 TGCTAATAGGTGTCTAAGTCCAGAAAGAGTGCTTCTTCAACTAAAATACAGATATGATGTGGAAATTGAT ×  
 AAAAGCAGAAGATCAGCTATAAAGAAGATAATGGAAGGGATGACACAGCTGCAAAAACACTTGTCTCT ×  
 GTATTTCTGAAATCATTTCTGCAAGTGCAGATATATCTGAACTTCTAGTAGTAAACTAGTAGTGTGGG ×  
 TACCAAAAAAGTGGGCATTATTGAGCTCACAGATGGGTGGTATGCTATTAAGGCCAGTTAGACCCTCCC ×  
 CTCTTAGCTCTCGTAAAGAACGGGAGATTGACTGTGGGTGAGAAGATCACTATTCTGGAGCAGAACTGG ×

```

TAGGCTCTCCTGATGCCTGCACACCACTTGAAGCCCCAGAATCTCTTATGTTAAAGATTTCTGCTAACAG ×
TACTCGTCCTGCTTGGTATACCAAACCTGGATTCTCTCCTGATCCTAGACCTTTCCCTCTCCCTTG ×
TCATCACTTTTCAGTGATGGAGGAAATGTTGGTTGTGTTGATGTAGTTGTTCAAAGAGCATACCCAATAC ×
AGTGGATGGAGAGGACCCCATCTGGATTATGCATATTTGCAATGAAAGAGAGGAAGAAAAGGAAGCAAC ×
AAAAATATGCAGAAATCCAACAAAAGAACTAGAAGTTTTATTCAATAAAATTCAAGCAGAATTTGAAAAG ×
AATGATGAAAAATATAACAAAGCAGTGTATACCATCATGTGCATTAACAAGACAGCAGATCTGTGCTCTGC ×
AAGATGGTGCAGAGCTTTATGAAGCAGTGACAAATGCACCAGACCCAAGTGACCTGGAGGGTTATTTTAG ×
TGAAGAGCAGTTAAGAGCCTTGAATAATCACAGACAGATGTTGAATGATAAGAAGCAAGCACAGATCCAG ×
TTAGAATTCAAGAAGGCTATGGAATCTGCTGAGCAAGGAGAACAAATCTACCAAGGGATGTTACAAC TG ×
TGTGGAAGTTACGTATCATAAGCTACAGGAAAAAAGAAAAAGATTCAGTTACATTGAGTATCTGGCGTCC ×
ATCACCAGATTTATATTCCTGTTAATAGAAGGAAAGAGATACAGAATCTATCATCTTGCAGCATCACAA ×
TCTAAAAGTAAATCTGGAAGGCAACACACAGCTAACAGCAACAAAGAAAACTCAGTACCAGCAACTAC ×
CAGCATCAGATGAAATCCTATCCCAAGTTTATCAGCCAAGGGAACCCCTTTACTTCAACAAACTGTTGGA ×
TCCGGACTTCCAACCACCTTGTTCTGAGGTGGACCTAATAGGATTTGTAGTTTCTGTTGTGAAAAAATA ×
GGTCTTGCTCCTGTGGTCTATTTGTGAGATGAATGCCATAATTTATTGGCAATAAAGTTCTGGACTGATT ×
TTAATGAAGACATTATTAACCTTACACATTAATTGCTGCAAGCAACCTCCAGTGGCGACCAGAAGCCAA ×
ATCAGGAATTCCTACTTTATTTGCTGGAGATTTTTCCAGTTTTCTGCCAGTCCAAAGGAGGAGCATTTT ×
CAAGAGACATTCCACAAAATGAAAAATACTGTTGAGAATATTGGTATGTTTTACAATGATGCAGAAAACA ×
AACTTGTGCATATACTTAATGCAAATGATCCCAAGTTGTCCACCCGACTAAAGACTATGCTTCAGAGCC ×
ACACACAGCTCAAATAGTCCTTGGCATAGGAAATAAATTTCTGATGTCTTCTCCAATAATGAGATGAAT ×
TATCAGAGTCCTTTATCACTTTGTAAGCCAAAAGAGAAGTCTGTCCCATACTGGATCAACCCAAATGA ×
CTTCAAAGTCTTATTGTAAGAGGAGAAAGAGATGGATGACCCAAAACCTGCAAAAAGAGAAGAGCCTT ×
GGACTTTTTGAGTAGAGTGCCTTTACCTCCATCTGTGAGTCCCATTGTACATTTGTTTCTCCAGCTGCA ×
CAGAAGGCATTTAGCCACCACGGAGTTGCGGCACCAAAATATGAAACACTGATGAAGAAAGAGTTGAATT ×
CTCCACAGATGACTCCACGTAAATTTAATGACCTTTCCCTTTTGGAAAGTGATTCAATAGCAGACGAAGA ×
ACTCGCAATGATAAACACCCAAGCCCTTTTGTGGGTTACCAGGAGAACATCAACTTGTGTCTGTGAGT ×
GACTCTACCAGGACTGCTCCACGAGCTCAAAAGATTATCTTGGACTGAAAAGGCATTCTACTGCACCCG ×
GGGTGAGAGGACCCGAGAGCCCCAGGCCCTGCACCAGGAAGCGGGAGCCCCGTGTACAGAACACAAGTGA ×
TCTGAAAAGGACATCTCTGAGACTGCAGAGGCAACAAACACAAAAATGACAATGAATTGGTGACTGACTC ×
AACCTTTCCAATGTGTGGAACACAGCCTCAACCTGTATGTCAAGATGTGCATAATGAGACAAGAAAGA ×
CCACATCCCAAATCTCCTGTGTGCTTGTCTATCTTAGGAAACCTGGCCTATCTCTGTACTGGTCGGTGTA ×
CTTTATTTTCAGTTATGTGTCTGAAAATTGTGTATTTATTAGCTAATCAGGAAAAAAATCTCCTTTAAAC ×
TCTTATGATTGGATATGATCAAGTATATTTACAAAGTAAACACACTTTTTCTTTAAATTGTGTCCCTAA ×
TTAAATGAAAGTAGGTTTCAAAGTACTGTTATTTTGAAGTCTGTTAGTCTTTTGGTGACTTGGTTTTG ×
TTTTGTTTTTCGGAGGTAACCTACTATGAACCAAGTTTCTTAATAAACGTGTTGGTTCTCTTATAGTTG ×
TATCCTGATCAAAAGTCAGGAGGAGTAAGGAACAAACAGCAGTGCTCTCTCTGGACCAGTTCTTTAACCT ×
TACGTCAGCATAAGTGCAAGAAAAACAGAATCCTCAATGTGATTCTTTTATGATTCTAGTGTGATTGC ×
TGAATTATTTCAATTAATAAATTCAAATGCTTTTAAAAAAGAAAAAAGAAAAAAGAAAAAAGAAAAAAG // 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

### Process

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)";

```

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 *)

```

Human chromosome9 scaffold 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]]
Human chromosome9 scaffold genesample.txt

lengthofgeneitself = Length[Flatten[numgenesample]]
(*To make sure no base pairs are left out *)
59027

```

## 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 PCV1 samples

W for H1A

Run this for output and pdf

```

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[α_] :=  $\frac{1}{1 - \alpha} \text{Log}[2, \text{Sum}[(\text{set}[[i]])^\alpha, \{i, 1, n\}]] // N$ 
H0 = Log[n] // N; (* H0 = Hartley Entropy*)
H1 = -Sum[(set[[i]]) (Log[2, set[[i]]]), {i, 1, n}] // N;
(* H1 = Shannon Entropy*)
H2onward = Table[H[a], {a, 2, 20}] // N; (* H2 onward *)
RenyiEntropyofEigenvalues = Join[{H0}, {H1}, H2onward];

```

Human H1A gene

Null<sup>2</sup>

```

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],

```





```

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,
  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}};
    
```

```
M = H1Avec;
```

```

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;
    
```

## H1A Renyi Entropies

```

Length[M[[1]]] (*Should be 781 *)
Length[FilledVec[M]] (*Should be 1024 *)

781

1024
    
```

```
set = M[[1]]; (* Define the set here,
then the following functions are defined using this set *)
```

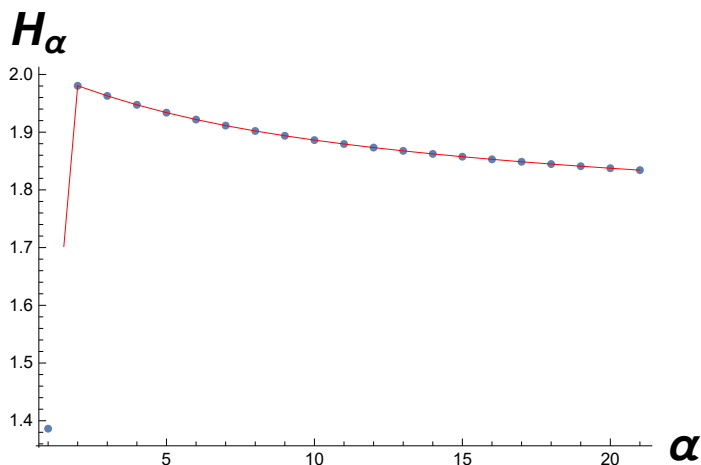
```
set = FilledVec[M]; (* Define the set here,
then the following functions are defined using this set *)
```

```
n = Length[Union[set]] (* i.e. n = number of unique elements in the set *)
prob[i_] :=  $\frac{\text{Count}[\text{set}, \text{Union}[\text{set}][[i]]]}{\text{Length}[\text{set}]}$  // N
 $H[\alpha_] := \frac{1}{1-\alpha} \text{Log}[2, \text{Sum}[(\text{prob}[i])^\alpha, \{i, 1, n\}]]$  // N
```

4

```
(* Gives a plot of the Renyi Entropies for  $\alpha \geq 0$ , RUN BLUE CELL FIRST *)
H0 = Log[n] // N; (*  $H_0$  = Hartley Entropy*)
H1 = -Sum[(prob[i]) (Log[2, prob[i]]), {i, 1, n}]; (*  $H_1$  = Shannon Entropy*)
H2onward = Table[H[a], {a, 2, 20}] // N; (*  $H_2$  onward *)
RenyiEntropyofH1ADData = Join[{H0}, {H1}, H2onward]
Show[
  ListPlot[RenyiEntropyofH1ADData, PlotRange -> All,
    AxesLabel -> {Style[" $\alpha$ ", Large, Bold], Style[" $H_\alpha$ ", Large, Bold]}],
  ListLinePlot[RenyiEntropyofH1ADData, PlotStyle -> {Red, Thin}]
]
```

```
{1.38629, 1.98039, 1.96288, 1.94741, 1.93382, 1.92189,
 1.91138, 1.90206, 1.89375, 1.88627, 1.87949, 1.8733, 1.86762, 1.86237,
 1.85749, 1.85296, 1.84872, 1.84475, 1.84102, 1.83752, 1.83423}
```



```
Union[set] (* prob[i] measures prob of obtaining Union[set][[i]] *)
```

```
prob[1]
```

```
prob[2]
```

```
prob[3]
```

```
prob[4]
```

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

```
0.297055
```

```
0.261204
```

```
0.256082
```

```
0.185659
```

```
Table[{"The", i - 1, "-th Renyi Entropy is =", RenyiEntropyofH1AData[[i]]}, {i, 1, 21}]
```

```
{ {The, 0, -th Renyi Entropy is =, 1.38629},
  {The, 1, -th Renyi Entropy is =, 1.98039}, {The, 2, -th Renyi Entropy is =, 1.96288},
  {The, 3, -th Renyi Entropy is =, 1.94741}, {The, 4, -th Renyi Entropy is =, 1.93382},
  {The, 5, -th Renyi Entropy is =, 1.92189}, {The, 6, -th Renyi Entropy is =, 1.91138},
  {The, 7, -th Renyi Entropy is =, 1.90206}, {The, 8, -th Renyi Entropy is =, 1.89375},
  {The, 9, -th Renyi Entropy is =, 1.88627}, {The, 10, -th Renyi Entropy is =, 1.87949},
  {The, 11, -th Renyi Entropy is =, 1.8733}, {The, 12, -th Renyi Entropy is =, 1.86762},
  {The, 13, -th Renyi Entropy is =, 1.86237}, {The, 14, -th Renyi Entropy is =, 1.85749},
  {The, 15, -th Renyi Entropy is =, 1.85296}, {The, 16, -th Renyi Entropy is =, 1.84872},
  {The, 17, -th Renyi Entropy is =, 1.84475}, {The, 18, -th Renyi Entropy is =, 1.84102},
  {The, 19, -th Renyi Entropy is =, 1.83752}, {The, 20, -th Renyi Entropy is =, 1.83423} }
```