FragScanTibo

0.1

Gegenereerd door Doxygen 1.8.18

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# Hoofdstuk 1

# **Klasse Index**

# 1.1 Klasse Lijst

Hieronder volgen de klassen, structs en unions met voor elk een korte beschrijving:

HMM									 														5
thread	_dat	a							 														8
TRAIN									 							 							10

2 Klasse Index

# Hoofdstuk 2

# **Bestand Index**

# 2.1 Bestandslijst

Hieronder volgt de lijst met alle bestanden, elk met een korte beschrijving:

src/hmm.h																	 					13
src/hmm_lib.c																	 					26
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4 Bestand Index

# Hoofdstuk 3

# Klassen Documentatie

## 3.1 HMM Struct Referentie

```
#include <hmm.h>
```

#### **Public Attributen**

- double pi [29]
- int N
- double tr [14]
- double e\_M\_1 [6][16][4]
- double e\_M [6][16][4]
- double tr\_R\_R [4][4]
- double tr\_I\_I [4][4]
- double tr\_M\_I [4][4]
- double tr\_S [61][64]
- double tr\_E [61][64]
- double tr\_S\_1 [61][64]
- double tr\_E\_1 [61][64]
- double S\_dist [6]
- double E\_dist [6]
- double S1\_dist [6]
- double E1\_dist [6]

#### 3.1.1 Documentatie van data members

## 3.1.1.1 E1\_dist

```
double E1_dist[6]
```

6 Klassen Documentatie

# 3.1.1.2 E\_dist double E\_dist[6] 3.1.1.3 e\_M double e\_M[6][16][4] 3.1.1.4 e\_M\_1 double e\_M\_1[6][16][4] 3.1.1.5 N int N 3.1.1.6 pi double pi[29] 3.1.1.7 S1\_dist double S1\_dist[6] 3.1.1.8 S\_dist double S\_dist[6]

3.1.1.9 tr

double tr[14]

#### Gegenereerd door Doxygen

3.1 HMM Struct Referentie 7

## 3.1.1.10 tr\_E

double tr\_E[61][64]

## 3.1.1.11 tr\_E\_1

double tr\_E\_1[61][64]

## 3.1.1.12 tr\_l\_l

double tr\_I\_I[4][4]

## 3.1.1.13 tr\_M\_I

double tr\_M\_I[4][4]

## 3.1.1.14 tr\_R\_R

double tr\_R\_R[4][4]

## 3.1.1.15 tr\_S

double tr\_S[61][64]

# 3.1.1.16 tr\_S\_1

double tr\_S\_1[61][64]

De documentatie voor deze struct is gegenereerd op grond van het volgende bestand:

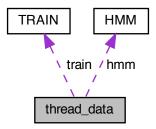
• src/hmm.h

8 Klassen Documentatie

# 3.2 thread\_data Struct Referentie

#include <run\_hmm.h>

Collaboratie diagram voor thread\_data:



#### **Public Attributen**

- FILE \* out
- FILE \* aa
- FILE \* dna
- char \* obs\_head
- char \* obs\_seq
- · int wholegenome
- int cg
- int format
- HMM \* hmm
- TRAIN \* train

## 3.2.1 Documentatie van data members

3.2.1.1 aa

FILE\* aa

3.2.1.2 cg

int cg

## 3.2.1.3 dna

FILE\* dna

#### 3.2.1.4 format

int format

## 3.2.1.5 hmm

HMM\* hmm

# 3.2.1.6 obs\_head

char\* obs\_head

#### 3.2.1.7 obs\_seq

char\* obs\_seq

## 3.2.1.8 out

FILE\* out

#### 3.2.1.9 train

TRAIN\* train

10 Klassen Documentatie

#### 3.2.1.10 wholegenome

```
int wholegenome
```

De documentatie voor deze struct is gegenereerd op grond van het volgende bestand:

• src/run hmm.h

## 3.3 TRAIN Struct Referentie

```
#include <hmm.h>
```

#### **Public Attributen**

- double trans [44][6][16][4]
- double rtrans [44][6][16][4]
- double noncoding [44][4][4]
- double start [44][61][64]
- double stop [44][61][64]
- double start1 [44][61][64]
- double stop1 [44][61][64]
- double S\_dist [44][6]
- double E\_dist [44][6]
- double S1\_dist [44][6]
- double E1\_dist [44][6]

#### 3.3.1 Documentatie van data members

#### 3.3.1.1 E1\_dist

```
double E1_dist[44][6]
```

#### 3.3.1.2 E\_dist

double E\_dist[44][6]

#### 3.3.1.3 noncoding

double noncoding[44][4][4]

3.3 TRAIN Struct Referentie

#### 3.3.1.4 rtrans

double rtrans[44][6][16][4]

## 3.3.1.5 S1\_dist

double S1\_dist[44][6]

## 3.3.1.6 S\_dist

double S\_dist[44][6]

#### 3.3.1.7 start

double start[44][61][64]

#### 3.3.1.8 start1

double start1[44][61][64]

#### 3.3.1.9 stop

double stop[44][61][64]

#### 3.3.1.10 stop1

double stop1[44][61][64]

## 3.3.1.11 trans

double trans[44][6][16][4]

De documentatie voor deze struct is gegenereerd op grond van het volgende bestand:

• src/hmm.h

12 Klassen Documentatie

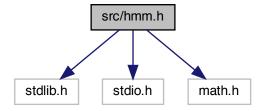
# Hoofdstuk 4

# **Bestand Documentatie**

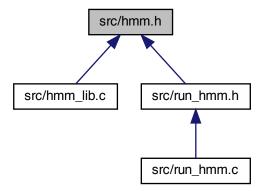
# 4.1 src/hmm.h Bestand Referentie

#include <stdlib.h>
#include <stdio.h>
#include <math.h>

 $Include\ afhankelijkheidsgraaf\ voor\ hmm.h:$ 



Deze graaf geeft aan welke bestanden direct of indirect afhankelijk zijn van dit bestand:



#### Klassen

- struct HMM
- struct TRAIN

#### **Macros**

- #define A 0
- #define C 1
- #define G 2
- #define T 3
- #define NUM STATE 29
- #define NOSTATE -1
- #define S\_STATE 0
- #define E\_STATE 1
- #define R\_STATE 2
- #define S\_STATE\_1 3
- #define E\_STATE\_1 4
- #define M1\_STATE 5
- #define M2\_STATE 6
- #define M3\_STATE 7
- #define M4\_STATE 8
- #define M5\_STATE 9#define M6\_STATE 10
- #define M1\_STATE\_1 11
- #define M2\_STATE\_1 12
- #define M3\_STATE\_1 13
- #define M4\_STATE\_1 14
- #define M5\_STATE\_1 15
- #define M6\_STATE\_1 16
- #define I1\_STATE 17
- #define I2\_STATE 18

#define I4\_STATE 20
#define I5\_STATE 21
#define I6\_STATE 22
#define I1\_STATE\_1 23
#define I2\_STATE\_1 24

• #define I3\_STATE 19

- #define I3\_STATE\_1 25
- #define I4\_STATE\_1 26
- #define I5\_STATE\_1 27
- #define I6\_STATE\_1 28
- #define TR MM 0
- #define TR MI 1
- #define TR\_MD 2
- #define TR II 3
- #define TR\_IM 4
- #define TR\_DD 5
- #define TR DM 6
- #define TR GE 7
- #define TR GG 8
- #define TR\_ER 9
- #define TR RS 10
- #define TR\_RR 11
- #define TR\_ES 12
- #define TR ES1 13

#### **Functies**

- int get\_prob\_from\_cg (HMM \*hmm, TRAIN \*train, char \*O)
- void get\_train\_from\_file (char \*filename, HMM \*hmm\_ptr, char \*mfilename, char \*mfilename, char \*mfilename, char \*sfilename, char \*sfilename,
- void viterbi (HMM \*hmm\_ptr, TRAIN \*train\_ptr, char \*O, FILE \*out\_filename, FILE \*log\_filename, FILE \*dna filename, char \*head, int metagene, int cg, int format)
- void free\_hmm (HMM \*hmm)
- void get\_protein (char \*dna, char \*protein, int strand, int whole\_genome)
- void get\_rc\_dna (char \*dna, char \*dna1)
- void get\_corrected\_dna (char \*dna, char \*dna\_f)

#### 4.1.1 Gedetailleerde Beschrijving

This is the header file for the hmm datastructure.

#### 4.1.2 Documentatie van macro's

#### 4.1.2.1 A

#define A 0

## 4.1.2.2 C

#define C 1

# 4.1.2.3 **E\_STATE**

#define E\_STATE 1

## 4.1.2.4 E\_STATE\_1

#define E\_STATE\_1 4

#### 4.1.2.5 G

#define G 2

## 4.1.2.6 I1\_STATE

#define I1\_STATE 17

## 4.1.2.7 I1\_STATE\_1

#define I1\_STATE\_1 23

# 4.1.2.8 I2\_STATE

#define I2\_STATE 18

## 4.1.2.9 I2\_STATE\_1

#define I2\_STATE\_1 24

## 4.1.2.10 I3\_STATE

#define I3\_STATE 19

## 4.1.2.11 I3\_STATE\_1

#define I3\_STATE\_1 25

## 4.1.2.12 I4\_STATE

#define I4\_STATE 20

## 4.1.2.13 I4\_STATE\_1

#define I4\_STATE\_1 26

## 4.1.2.14 I5\_STATE

#define I5\_STATE 21

## 

#define I5\_STATE\_1 27

## 4.1.2.16 I6\_STATE

#define I6\_STATE 22

## 

#define I6\_STATE\_1 28

## 4.1.2.18 M1\_STATE

#define M1\_STATE 5

## 4.1.2.19 M1\_STATE\_1

#define M1\_STATE\_1 11

## 4.1.2.20 M2\_STATE

#define M2\_STATE 6

## 4.1.2.21 M2\_STATE\_1

#define M2\_STATE\_1 12

## 4.1.2.22 M3\_STATE

#define M3\_STATE 7

## 4.1.2.23 M3\_STATE\_1

#define M3\_STATE\_1 13

# 4.1.2.24 M4\_STATE

#define M4\_STATE 8

## 4.1.2.25 M4\_STATE\_1

#define M4\_STATE\_1 14

## 4.1.2.26 M5\_STATE

#define M5\_STATE 9

## 4.1.2.27 M5\_STATE\_1

#define M5\_STATE\_1 15

## 4.1.2.28 M6\_STATE

#define M6\_STATE 10

#### 4.1.2.29 M6\_STATE\_1

#define M6\_STATE\_1 16

#### 4.1.2.30 NOSTATE

#define NOSTATE -1

## 4.1.2.31 NUM\_STATE

#define NUM\_STATE 29

Total number of states, mainly used in for loops.

## 4.1.2.32 R\_STATE

#define R\_STATE 2

#### 4.1.2.33 S\_STATE

#define S\_STATE 0

## 4.1.2.34 S\_STATE\_1

#define S\_STATE\_1 3

#### 4.1.2.35 T

#define T 3

## 4.1.2.36 TR\_DD

#define TR\_DD 5

## 4.1.2.37 TR\_DM

#define TR\_DM 6

## 4.1.2.38 TR\_ER

#define TR\_ER 9

## 4.1.2.39 TR\_ES

#define TR\_ES 12

## 4.1.2.40 TR\_ES1

#define TR\_ES1 13

## 4.1.2.41 TR\_GE

#define TR\_GE 7

## 4.1.2.42 TR\_GG

#define TR\_GG 8

## 4.1.2.43 TR\_II

#define TR\_II 3

## 4.1.2.44 TR\_IM

#define TR\_IM 4

# 4.1.2.45 TR\_MD

#define TR\_MD 2

#### 4.1.2.46 TR\_MI

#define TR\_MI 1

## 4.1.2.47 TR\_MM

#define TR\_MM 0

#### 4.1.2.48 TR\_RR

#define TR\_RR 11

## 4.1.2.49 TR\_RS

#define TR\_RS 10

#### 4.1.3 Documentatie van functies

#### 4.1.3.1 free\_hmm()

```
void free_hmm ( _{\rm HMM} * _{hmm} )
```

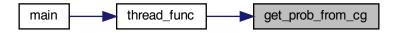
Hier is de call graaf voor deze functie:



#### 4.1.3.2 get\_corrected\_dna()

#### 4.1.3.3 get\_prob\_from\_cg()

```
int get_prob_from_cg (
    HMM * hmm,
    TRAIN * train,
    char * 0 )
```



#### 4.1.3.4 get\_protein()

Get a protein of dna if Whole\_genome equals to zero, then we want a short read and stop early. Hier is de caller graaf voor deze functie:



#### 4.1.3.5 get\_rc\_dna()

copies dna to dna1 in reverse. and Hier is de call graaf voor deze functie:



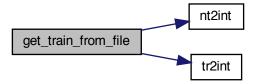


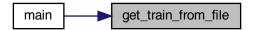
#### 4.1.3.6 get\_train\_from\_file()

Reads files.

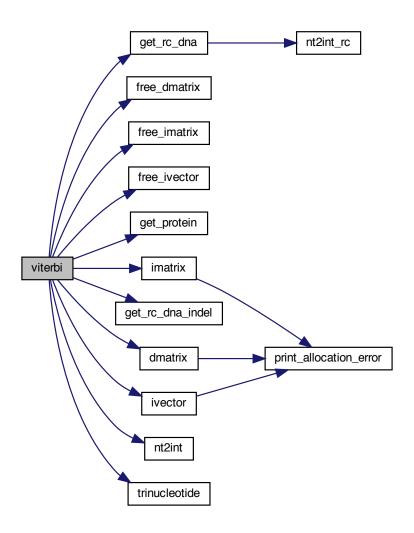
1. Reads trasition file and store in hmm datastructure

Hier is de call graaf voor deze functie:

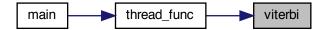




#### 4.1.3.7 viterbi()



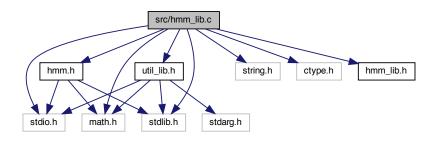
Hier is de caller graaf voor deze functie:



# 4.2 src/hmm lib.c Bestand Referentie

```
#include <stdio.h>
#include <math.h>
#include <string.h>
#include <stdlib.h>
#include <ctype.h>
#include "hmm.h"
#include "util_lib.h"
#include "hmm_lib.h"
```

Include afhankelijkheidsgraaf voor hmm\_lib.c:



#### **Functies**

- void viterbi (HMM \*hmm\_ptr, TRAIN \*train\_ptr, char \*O, FILE \*fp\_out, FILE \*fp\_aa, FILE \*fp\_dna, char \*head, int whole\_genome, int cg, int format)
- int get\_prob\_from\_cg (HMM \*hmm\_ptr, TRAIN \*train\_ptr, char \*O)
- void get\_train\_from\_file (char \*filename, HMM \*hmm\_ptr, char \*mfilename, char \*mfilename1, char \*nfilename, char \*sfilename, char \*sfilename, char \*sfilename, char \*sfilename, char \*sfilename, char \*filename, char \*sfilename, char \*sfilename,
- void free hmm (HMM \*hmm ptr)
- void dump memory (void \*p, int size)

#### 4.2.1 Documentatie van functies

#### 4.2.1.1 dump\_memory()

```
void dump_memory ( \label{eq:condition} \mbox{void} \, * \, p, \\ \mbox{int } size \; )
```

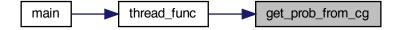
#### 4.2.1.2 free\_hmm()

```
void free_hmm ( {\tt HMM} \, * \, \mathit{hmm\_ptr} \, )
```

Hier is de call graaf voor deze functie:



#### 4.2.1.3 get\_prob\_from\_cg()

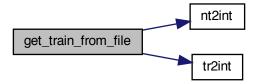


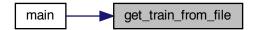
#### 4.2.1.4 get\_train\_from\_file()

Reads files.

1. Reads trasition file and store in hmm datastructure

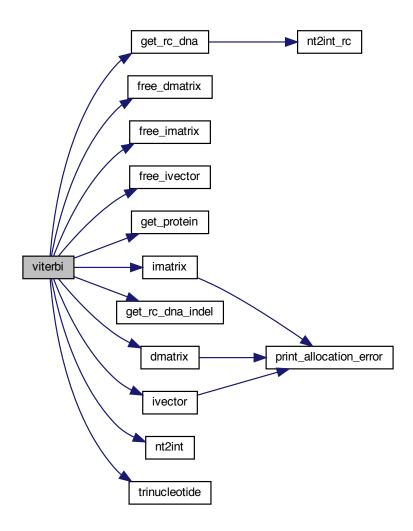
Hier is de call graaf voor deze functie:



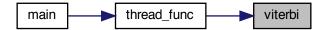


# 4.2.1.5 viterbi()

```
void viterbi (
    HMM * hmm_ptr,
    TRAIN * train_ptr,
    char * O,
    FILE * fp_out,
    FILE * fp_dna,
    char * head,
    int whole_genome,
    int cg,
    int format )
```

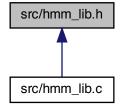


Hier is de caller graaf voor deze functie:



# 4.3 src/hmm\_lib.h Bestand Referentie

Deze graaf geeft aan welke bestanden direct of indirect afhankelijk zijn van dit bestand:



# **Functies**

- void dump\_memory (void \*p, int size)
- void <a href="mailto:get\_rc\_dna\_indel">get\_rc\_dna\_indel</a> (char dna\_f[300000], char dna\_f1[300000])

# 4.3.1 Documentatie van functies

# 4.3.1.1 dump\_memory()

```
void dump_memory ( \label{eq:condition} \mbox{void} \, * \, p, \\ \mbox{int } size \; )
```

# 4.3.1.2 get\_rc\_dna\_indel()

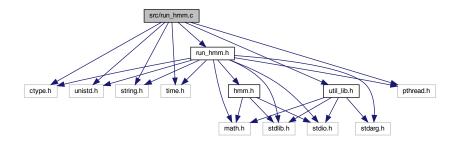
Hier is de caller graaf voor deze functie:



# 4.4 src/run hmm.c Bestand Referentie

```
#include <ctype.h>
#include <unistd.h>
#include <string.h>
#include <time.h>
#include "run_hmm.h"
#include "util_lib.h"
#include <pthread.h>
```

Include afhankelijkheidsgraaf voor run\_hmm.c:



#### **Macros**

- #define ADD LEN 1024
- #define STRINGLEN 4096

## **Functies**

- int main (int argc, char \*\*argv)
- void \* thread\_func (void \*threadarr)
- int appendSeq (char \*input, char \*\*seq, int input\_max)
- void print\_error (const char \*error\_message,...)
- void print\_file\_error (const char \*error\_message, char \*file)

# 4.4.1 Documentatie van macro's

# 4.4.1.1 ADD\_LEN

```
#define ADD_LEN 1024
```

# 4.4.1.2 STRINGLEN

```
#define STRINGLEN 4096
```

# 4.4.2 Documentatie van functies

# 4.4.2.1 appendSeq()

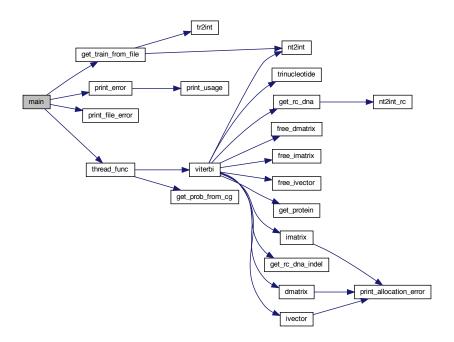
# 4.4.2.2 main()

```
int main (
          int argc,
          char ** argv )
```

# Entry point of program

- 1. Initialization of variables and datatypes
- 2. Check File acessiblity

Hier is de call graaf voor deze functie:



# 4.4.2.3 print\_error()

Error function:

- 1. Print error message
- 2. Call print\_usage() from util\_lib
- 3. EXIT program



Hier is de caller graaf voor deze functie:



# 4.4.2.4 print\_file\_error()

Error function:

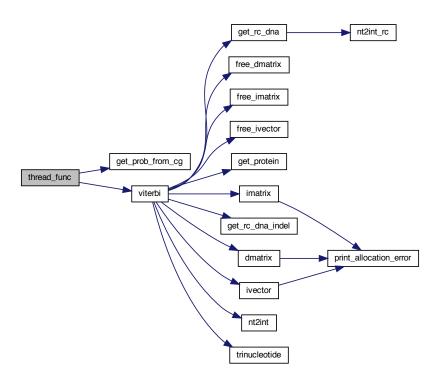
- 1. Print error message
- 2. EXIT program

Hier is de caller graaf voor deze functie:



# 4.4.2.5 thread\_func()

Hier is de call graaf voor deze functie:



Hier is de caller graaf voor deze functie:

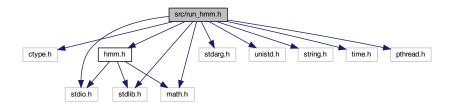


# 4.5 src/run\_hmm.h Bestand Referentie

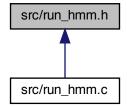
```
#include <ctype.h>
#include <stdio.h>
#include <stdlib.h>
#include <stdarg.h>
#include <math.h>
#include <unistd.h>
#include <string.h>
#include <time.h>
#include "hmm.h"
```

#include <pthread.h>

Include afhankelijkheidsgraaf voor run\_hmm.h:



Deze graaf geeft aan welke bestanden direct of indirect afhankelijk zijn van dit bestand:



# Klassen

• struct thread\_data

# **Typedefs**

• typedef struct thread\_data thread\_data

# **Functies**

- void \* thread func (void \*threadarr)
- void print\_error (const char \*error\_message,...)
- void print\_file\_error (const char \*error\_message, char \*file)

# 4.5.1 Documentatie van typedefs

# 4.5.1.1 thread\_data

```
typedef struct thread_data thread_data
```

# 4.5.2 Documentatie van functies

# 4.5.2.1 print\_error()

Error function:

- 1. Print error message
- 2. Call print\_usage() from util\_lib
- 3. EXIT program

Hier is de call graaf voor deze functie:





# 4.5.2.2 print\_file\_error()

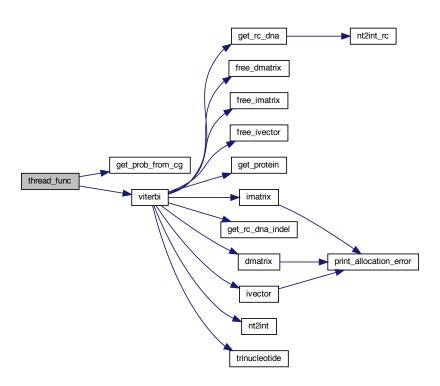
Error function:

- 1. Print error message
- 2. EXIT program

Hier is de caller graaf voor deze functie:



# 4.5.2.3 thread\_func()



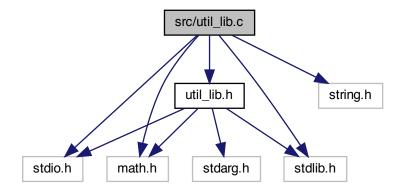
Hier is de caller graaf voor deze functie:



# 4.6 src/util\_lib.c Bestand Referentie

```
#include <stdio.h>
#include <math.h>
#include <stdlib.h>
#include <string.h>
#include "util_lib.h"
```

Include afhankelijkheidsgraaf voor util\_lib.c:



## **Macros**

• #define TR\_SIZE 14

## **Functies**

- double \*\* dmatrix (int num\_row, int num\_col)
- int \*\* imatrix (int num\_row, int num\_col)
- double \* dvector (int nh)
- int \* ivector (int nh)
- void free\_dvector (double \*v)
- void free\_ivector (int \*v)

```
void free_dmatrix (double **m, int num_row)
void free_imatrix (int **m, int num_row)
int tr2int (char *tr)
int nt2int (char nt)
int nt2int_rc (char nt)
int nt2int_rc_indel (char nt)
int trinucleotide (char a, char b, char c)
int trinucleotide_pep (char a, char b, char c)
void get_rc_dna (char *dna, char *dna1)
void get_rc_dna_indel (char *dna, char *dna1)
void get_protein (char *dna, char *protein, int strand, int whole_genome)
void print_usage ()
void print_allocation_error (const char *format,...)
```

#### Variabelen

```
    const char * tr_list [TR_SIZE] = { "MM","MI","MD","II","IM","DD","DM","GE","GG","ER","RS","RR","ES","ES1" }
    const char codon5 [5] = { 'A', 'C', 'G', 'T', 'N' }
    const char codon11 [11] = { 'A', 'C', 'G', 'T', 'N', 'a', 'c', 'g', 't', 'n', 'x' }
    const char codon_code [65]
    const char anti_codon_code [65]
```

#### 4.6.1 Documentatie van macro's

## 4.6.1.1 TR\_SIZE

```
#define TR_SIZE 14
```

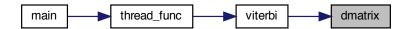
#### 4.6.2 Documentatie van functies

### 4.6.2.1 dmatrix()

Makes an matrix with datatype double. Elements are double pointers en matrix is a double double pointer (\*\*pointer). Exits when allocation fails. Hier is de call graaf voor deze functie:



Hier is de caller graaf voor deze functie:



# 4.6.2.2 dvector()

```
double* dvector (
          int nh )
```

Makes an vector (array) with datatype double. Elements are doubles en vector is a double pointer. Exits when allocation fails. Hier is de call graaf voor deze functie:



#### 4.6.2.3 free\_dmatrix()

Frees the memory allocation of an matrix with datatype double. Hier is de caller graaf voor deze functie:



# 4.6.2.4 free\_dvector()

```
void free_dvector ( \mbox{double } * \ v \ )
```

Frees the memory allocation of an vector with datatype double. Hier is de caller graaf voor deze functie:



### 4.6.2.5 free\_imatrix()

Frees the memory allocation of an matrix with datatype int. Hier is de caller graaf voor deze functie:



## 4.6.2.6 free\_ivector()

```
void free_ivector ( int \, * \, v \,)
```

Frees the memory allocation of an vector with datatype int. Hier is de caller graaf voor deze functie:



## 4.6.2.7 get\_protein()

Get a protein of dna if Whole\_genome equals to zero, then we want a short read and stop early. Hier is de call graaf voor deze functie:





#### 4.6.2.8 get\_rc\_dna()

copies dna to dna1 in reverse. and Hier is de call graaf voor deze functie:



Hier is de caller graaf voor deze functie:



# 4.6.2.9 get\_rc\_dna\_indel()

copies dna to dna1 in reverse. and Hier is de call graaf voor deze functie:



# 4.6.2.10 imatrix()

```
int** imatrix (
                int num_row,
                int num_col )
```

Makes an matrix with datatype int. Elements are int pointers en matrix is a double int pointer. Exits when allocation fails. Hier is de call graaf voor deze functie:



Hier is de caller graaf voor deze functie:



#### 4.6.2.11 ivector()

```
int* ivector ( \quad \quad \text{int } nh \ )
```

Makes an vector array) with datatype int. Elements are ints en vector is a int pointer. Exits when allocation fails. Hier is de call graaf voor deze functie:

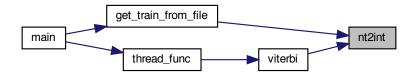


Hier is de caller graaf voor deze functie:



# 4.6.2.12 nt2int()

Hier is de caller graaf voor deze functie:



# 4.6.2.13 nt2int\_rc()



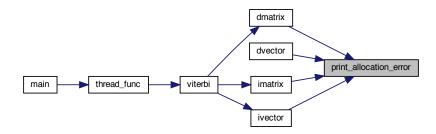
## 4.6.2.14 nt2int\_rc\_indel()

Hier is de caller graaf voor deze functie:



# 4.6.2.15 print\_allocation\_error()

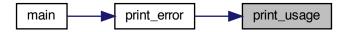
Custom error function to print allocation errors. Mostly called from matrix or vector functions. Hier is de caller graaf voor deze functie:



# 4.6.2.16 print\_usage()

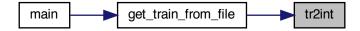
```
void print_usage ( )
```

Print how the program should be used. called mainly on help or error. Hier is de caller graaf voor deze functie:



# 4.6.2.17 tr2int()

Converts a given transition to int. Use for example as indexing. switch case not possible due the fact that strings are not constonant. Hier is de caller graaf voor deze functie:



## 4.6.2.18 trinucleotide()



## 4.6.2.19 trinucleotide\_pep()

```
int trinucleotide_pep (  \begin{array}{ccc} \text{char } a, \\ \text{char } b, \\ \text{char } c \end{array} )
```

Hier is de caller graaf voor deze functie:



# 4.6.3 Documentatie van variabelen

#### 4.6.3.1 anti\_codon\_code

const char anti\_codon\_code[65]

#### 4.6.3.2 codon11

```
const \ char \ codon11[11] \ = \ \{ \ 'A', \ 'C', \ 'G', \ 'T', \ 'N', \ 'a', \ 'c', \ 'g', \ 't', \ 'n', \ 'x' \ \}
```

### 4.6.3.3 codon5

```
const char codon5[5] = { 'A', 'C', 'G', 'T', 'N' }
```

#### 4.6.3.4 codon\_code

```
const char codon_code[65]
```

# Initiële waarde:

```
'A','A','A','A',
'G','G','G','G',
'V','V','V','V',
'*','Y','*','Y',
'S','S','S','S',
'*','C','W','C',
'L','F','L','F', 'X' }
```

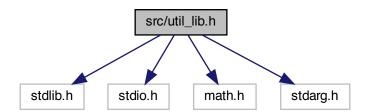
#### 4.6.3.5 tr\_list

```
\texttt{const char* tr\_list[TR\_SIZE]} \ = \ \{ \ "MM", "MI", "MD", "II", "IM", "DD", "DM", "GE", "GG", "ER", "RS", "ER", "ES", "E \leftrightarrow TABLE (TR\_SIZE) \} \\ + (TR\_SIZE) + (TR_SIZE) + (
S1" }
```

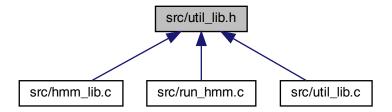
# 4.7 src/util\_lib.h Bestand Referentie

```
#include <stdlib.h>
#include <stdio.h>
#include <math.h>
#include <stdarg.h>
```

Include afhankelijkheidsgraaf voor util\_lib.h:



Deze graaf geeft aan welke bestanden direct of indirect afhankelijk zijn van dit bestand:



#### **Functies**

- double \*\* dmatrix (int num\_row, int num\_col)
- double \* dvector (int nh)
- int \*\* imatrix (int num\_row, int num\_col)
- int \* ivector (int nh)
- void free\_dvector (double \*v)
- void free dmatrix (double \*\*m, int num row)
- void free ivector (int \*v)
- void free imatrix (int \*\*m, int num row)
- int tr2int (char \*nt)
- int nt2int (char nt)
- int nt2int\_rc (char nt)
- int trinucleotide (char a, char b, char c)
- void get\_protein (char \*dna, char \*protein, int strand, int whole\_genome)
- void print\_usage ()
- void print allocation error (const char \*format,...)

#### 4.7.1 Documentatie van functies

#### 4.7.1.1 dmatrix()

Makes an matrix with datatype double. Elements are double pointers en matrix is a double double pointer (\*\*pointer). Exits when allocation fails. Hier is de call graaf voor deze functie:



Hier is de caller graaf voor deze functie:



# 4.7.1.2 dvector()

```
double* dvector ( \quad \text{int } nh \ )
```

Makes an vector (array) with datatype double. Elements are doubles en vector is a double pointer. Exits when allocation fails. Hier is de call graaf voor deze functie:



# 4.7.1.3 free\_dmatrix()

Frees the memory allocation of an matrix with datatype double. Hier is de caller graaf voor deze functie:



## 4.7.1.4 free\_dvector()

```
void free_dvector ( \mbox{double } * \ v \ )
```

Frees the memory allocation of an vector with datatype double. Hier is de caller graaf voor deze functie:



# 4.7.1.5 free\_imatrix()

Frees the memory allocation of an matrix with datatype int. Hier is de caller graaf voor deze functie:



#### 4.7.1.6 free\_ivector()

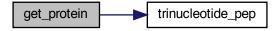
```
void free_ivector ( int \, * \, v \,)
```

Frees the memory allocation of an vector with datatype int. Hier is de caller graaf voor deze functie:



# 4.7.1.7 get\_protein()

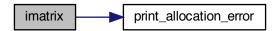
Get a protein of dna if Whole\_genome equals to zero, then we want a short read and stop early. Hier is de call graaf voor deze functie:

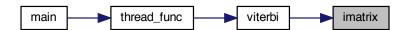


#### 4.7.1.8 imatrix()

```
int** imatrix (
                int num_row,
                int num_col )
```

Makes an matrix with datatype int. Elements are int pointers en matrix is a double int pointer. Exits when allocation fails. Hier is de call graaf voor deze functie:

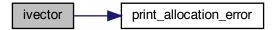




## 4.7.1.9 ivector()

```
int* ivector ( \quad \quad \text{int } nh \ )
```

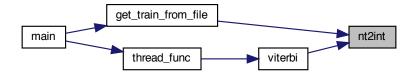
Makes an vector array) with datatype int. Elements are ints en vector is a int pointer. Exits when allocation fails. Hier is de call graaf voor deze functie:



Hier is de caller graaf voor deze functie:



# 4.7.1.10 nt2int()



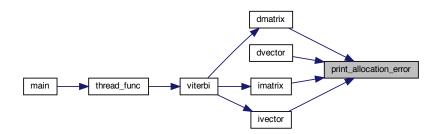
# 4.7.1.11 nt2int\_rc()

Hier is de caller graaf voor deze functie:



# 4.7.1.12 print\_allocation\_error()

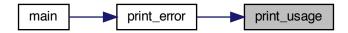
Custom error function to print allocation errors. Mostly called from matrix or vector functions. Hier is de caller graaf voor deze functie:



#### 4.7.1.13 print\_usage()

```
void print_usage ( )
```

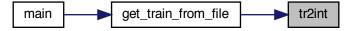
Print how the program should be used. called mainly on help or error. Hier is de caller graaf voor deze functie:



# 4.7.1.14 tr2int()

```
int tr2int ( {\tt char} \, * \, tr \,)
```

Converts a given transition to int. Use for example as indexing. switch case not possible due the fact that strings are not constonant. Hier is de caller graaf voor deze functie:



# 4.7.1.15 trinucleotide()

