

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:

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Hoofdstuk 2

Bestand Index

2.1 Bestandslijst

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

src/hmm.h	13
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src/hmm_lib.h	30
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src/run_hmm.h	34
src/sprintf_irc.c	??
src/sprintf_irc.h	??
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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]
```

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]
```

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_I_I

```
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]
```

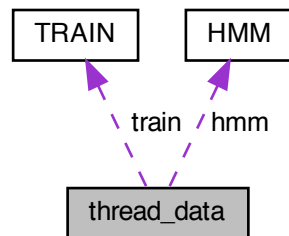
De documentatie voor deze struct is gegenereerd op grond van het volgende bestand:

- [src/hmm.h](#)

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

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]
```

3.3.1.4 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](#)

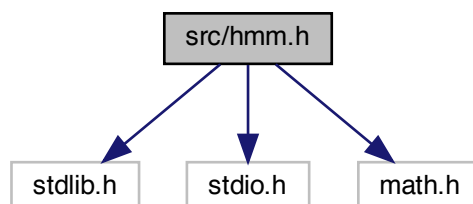
Hoofdstuk 4

Bestand Documentatie

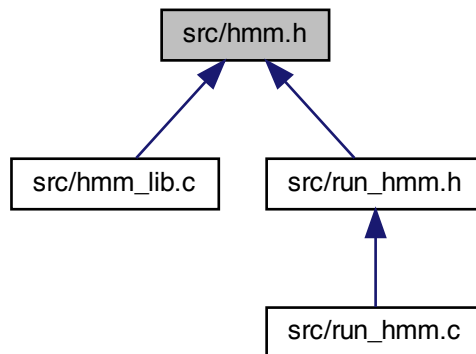
4.1 src/hmm.h Bestand Referentie

```
#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 I3_STATE 19
- #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_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

Funcities

- int `get_prob_from_cg` (HMM *hmm, TRAIN *train, char *O)
- void `get_train_from_file` (char *filename, HMM *hmm_ptr, char *mfilename, char *mfilename1, char *nfilename, char *sfilename, char *pfilename, char *s1filename, char *p1filename, char *dfilename, TRAIN *train_ptr)
- 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 `get_protein` (char *dna, char *protein, int strand, int whole_genome)
- void `get_rc_dna` (char *dna, char *dna1)

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
```

4.1.2.15 I5_STATE_1

```
#define I5_STATE_1 27
```

4.1.2.16 I6_STATE

```
#define I6_STATE 22
```

4.1.2.17 I6_STATE_1

```
#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

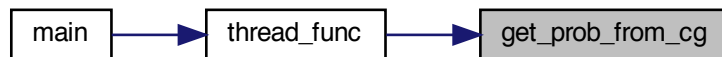
```
#define TR_RS 10
```

4.1.3 Documentatie van functies

4.1.3.1 get_prob_from_cg()

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

Hier is de caller graaf voor deze functie:



4.1.3.2 get_protein()

```
void get_protein (  
    char * dna,  
    char * protein,  
    int strand,  
    int whole_genome )
```

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:



4.1.3.3 get_rc_dna()

```
void get_rc_dna (
    char * dna,
    char * dna1 )
```

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



Hier is de caller graaf voor deze functie:



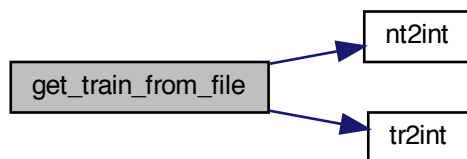
4.1.3.4 get_train_from_file()

```
void get_train_from_file (
    char * filename,
    HMM * hmm_ptr,
    char * mfilename,
    char * mfilename1,
    char * nfilename,
    char * sfilename,
    char * pfilename,
    char * slfilename,
    char * plfilename,
    char * dfilename,
    TRAIN * train_ptr )
```

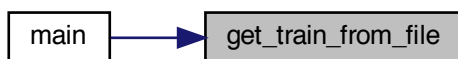
Reads files.

1. Reads transition file and store in hmm datastructure

Hier is de call graaf voor deze functie:



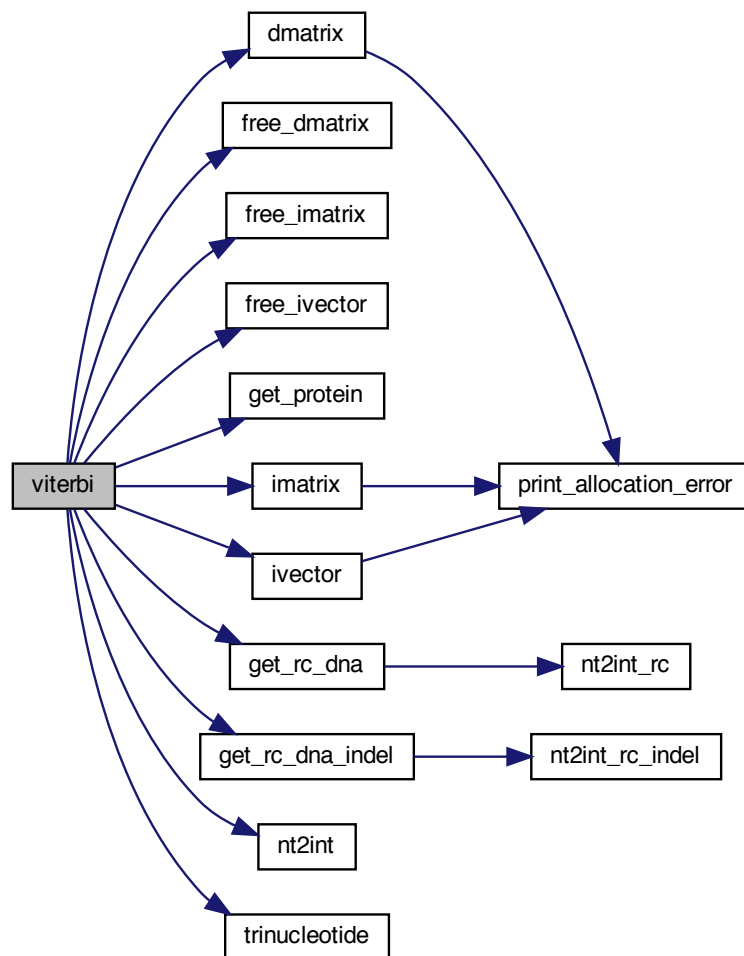
Hier is de caller graaf voor deze functie:



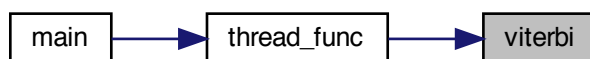
4.1.3.5 viterbi()

```
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 )
```


Hier is de call graaf voor deze functie:



Hier is de caller graaf voor deze functie:

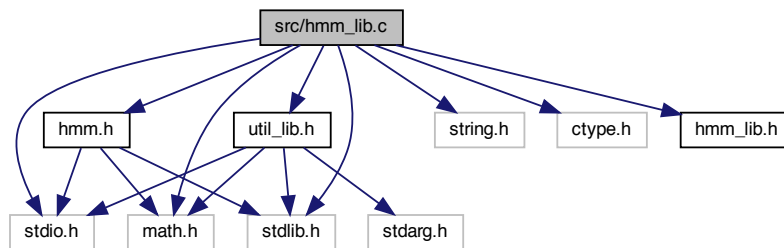


4.2 src/hmm_lib.c Bestand Referentie

```
#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 *pfilename, char *s1filename, char *p1filename, char *dfilename, `TRAIN` *train_ptr)

Variabelen

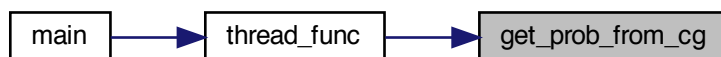
- const double `log53` = -0.634878
- const double `log16` = -1.832581
- const double `log30` = -1.203973
- const double `log25` = -1.386294
- const double `log95` = -0.051293
- const double `log54` = -0.616186
- const double `log83` = -0.186330
- const double `log07` = -2.659260

4.2.1 Documentatie van functies

4.2.1.1 get_prob_from_cg()

```
int get_prob_from_cg (  
    HMM * hmm_ptr,  
    TRAIN * train_ptr,  
    char * O )
```

Hier is de caller graaf voor deze functie:



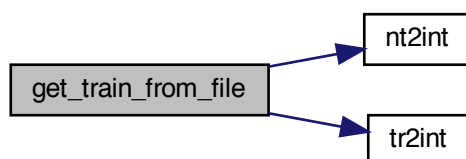
4.2.1.2 get_train_from_file()

```
void get_train_from_file (  
    char * filename,  
    HMM * hmm_ptr,  
    char * mfilename,  
    char * mfilename1,  
    char * nfilename,  
    char * sfilename,  
    char * pfilename,  
    char * slfilename,  
    char * plfilename,  
    char * dfilename,  
    TRAIN * train_ptr )
```

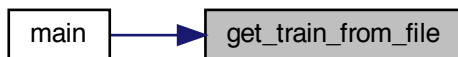
Reads files.

1. Reads transition file and store in hmm datastructure

Hier is de call graaf voor deze functie:



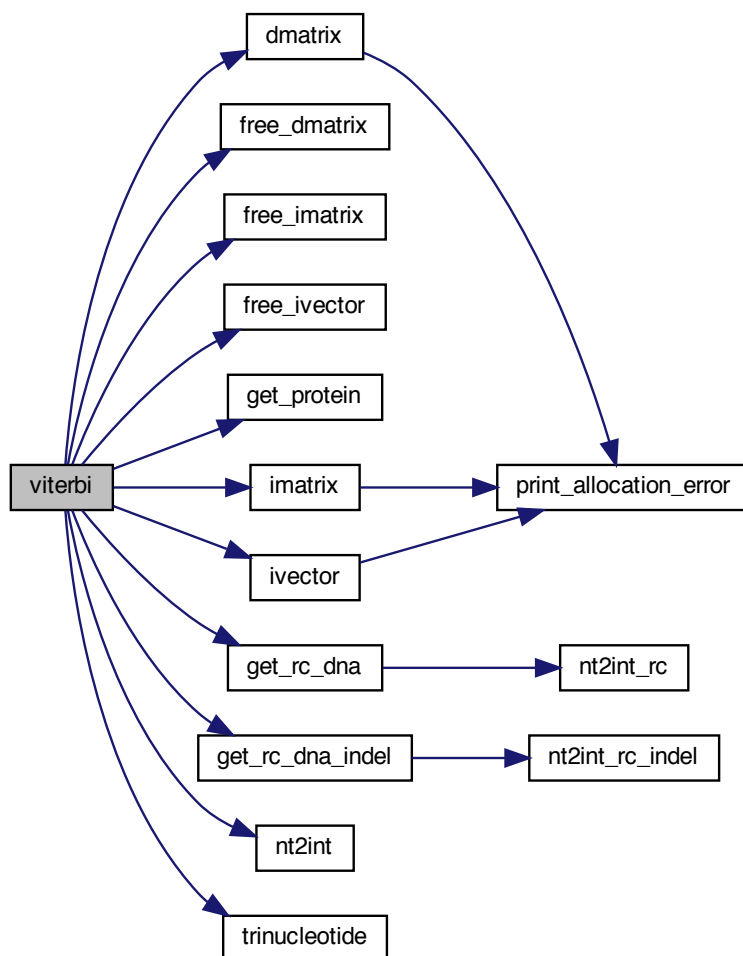
Hier is de caller graaf voor deze functie:



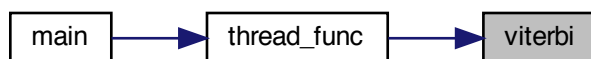
4.2.1.3 viterbi()

```
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 )
```

Hier is de call graaf voor deze functie:



Hier is de caller graaf voor deze functie:



4.2.2 Documentatie van variabelen

4.2.2.1 log07

```
const double log07 = -2.659260
```

4.2.2.2 log16

```
const double log16 = -1.832581
```

4.2.2.3 log25

```
const double log25 = -1.386294
```

4.2.2.4 log30

```
const double log30 = -1.203973
```

4.2.2.5 log53

```
const double log53 = -0.634878
```

4.2.2.6 log54

```
const double log54 = -0.616186
```

4.2.2.7 log83

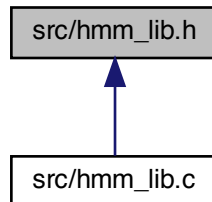
```
const double log83 = -0.186330
```

4.2.2.8 log95

```
const double log95 = -0.051293
```

4.3 src/hmm_lib.h Bestand Referentie

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



Functies

- void [get_rc_dna_indel](#) (char *dna_f, char *dna_f1)

4.3.1 Documentatie van functies

4.3.1.1 get_rc_dna_indel()

```
void get_rc_dna_indel (  
    char * dna,  
    char * dna1 )
```

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



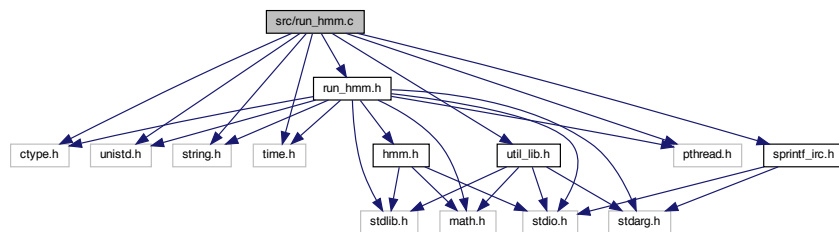
Hier is de caller graaf voor deze functie:



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 "sprintf_irc.h"
#include <pthread.h>
```

Include afhankelijkheidsgraaf voor run_hmm.c:



Macros

- #define `ADD_LEN` 1024
- #define `STRINGLEN` 4096+1
- #define `DELIMI` " "

Functies

- int `main` (int argc, char **argv)
- void * `thread_func` (void *threadarr)
- 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 DELIMI

```
#define DELIMI " "
```

4.4.1.3 STRINGLEN

```
#define STRINGLEN 4096+1
```

4.4.2 Documentatie van functies

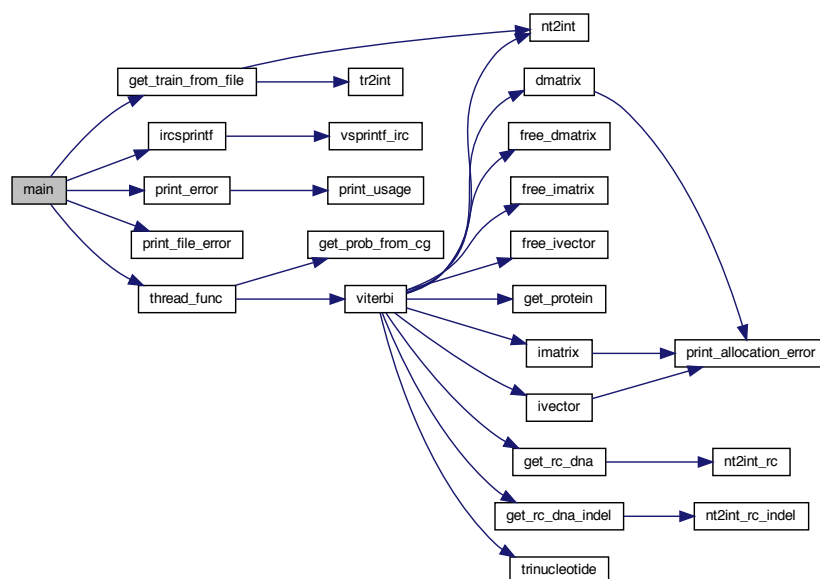
4.4.2.1 main()

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

Entry point of program

1. Initialization of variables and datatypes
2. Check File acessibility

Hier is de call graaf voor deze functie:



4.4.2.2 print_error()

```
void print_error (
    const char * error_message,
    ... )
```

Error function:

1. Print error message
2. Call [print_usage\(\)](#) from util_lib
3. EXIT program

Hier is de call graaf voor deze functie:



Hier is de caller graaf voor deze functie:



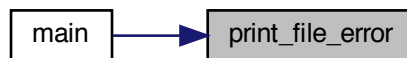
4.4.2.3 print_file_error()

```
void print_file_error (
    const char * error_message,
    char * file )
```

Error function:

1. Print error message
2. EXIT program

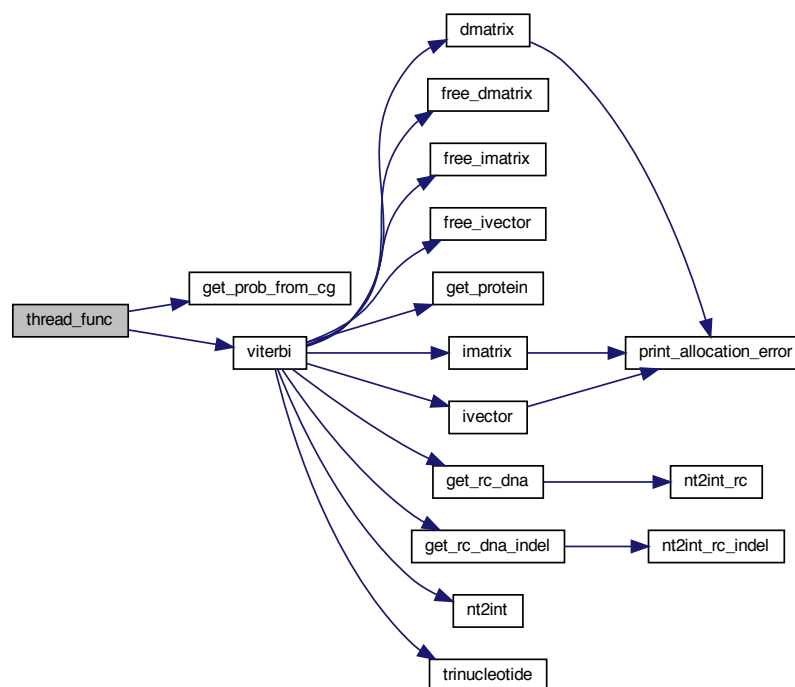
Hier is de caller graaf voor deze functie:



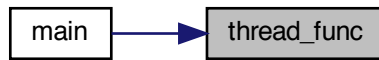
4.4.2.4 thread_func()

```
void* thread_func (  
    void * threadarr )
```

Function given to a thread during his creation. This thread will then execute this function. Hier is de call graaf voor deze functie:



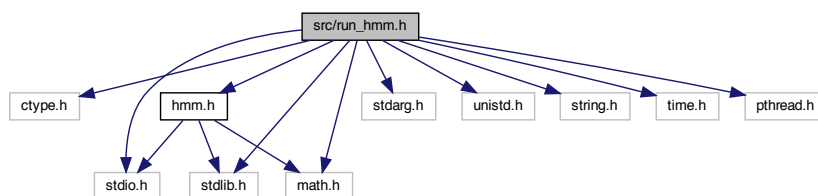
Hier is de caller graaf voor deze functie:



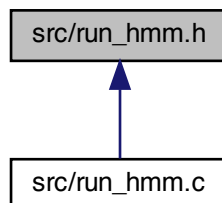
4.5 src/run_hmm.h Bestand Referentie

```
#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](#)

Funcities

- 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()

```
void print_error (  
    const char * error_message,  
    ... )
```

Error function:

1. Print error message
2. Call [print_usage\(\)](#) from util_lib
3. EXIT program

Hier is de call graaf voor deze functie:



Hier is de caller graaf voor deze functie:



4.5.2.2 `print_file_error()`

```
void print_file_error (
    const char * error_message,
    char * file )
```

Error function:

1. Print error message
2. EXIT program

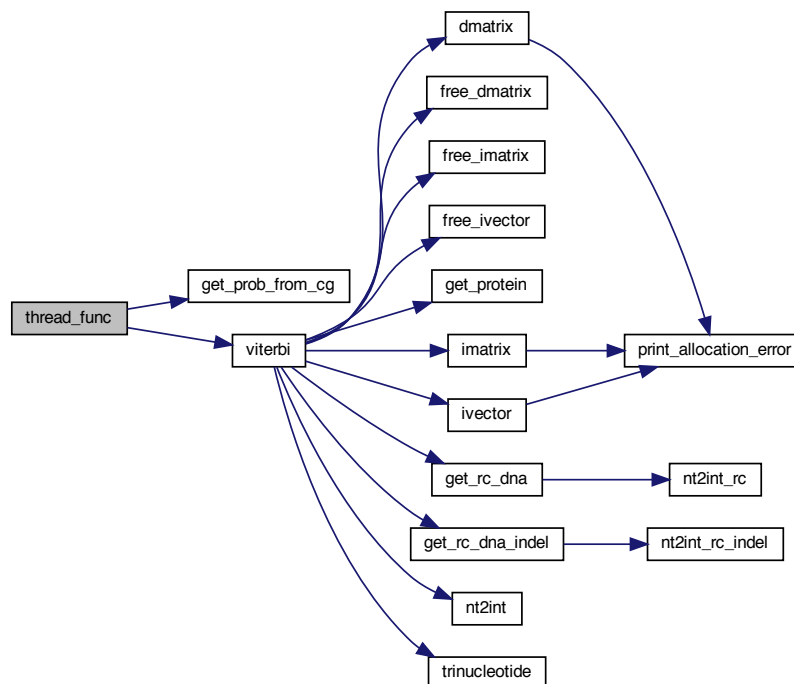
Hier is de caller graaf voor deze functie:



4.5.2.3 thread_func()

```
void* thread_func (
    void * threadarr )
```

Function given to a thread during his creation. This thread will then execute this function. Hier is de call graaf voor deze functie:



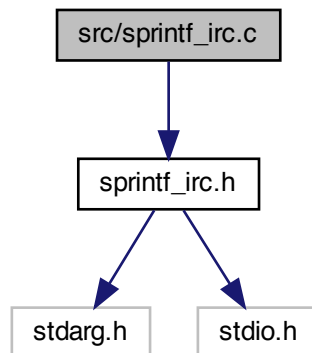
Hier is de caller graaf voor deze functie:



4.6 src/sprintf_irc.c Bestand Referentie

```
#include "sprintf_irc.h"
```

Include afhankelijkheidsgraaf voor `sprintf_irc.c`:



Functies

- int `vsprintf_irc` (char *str, const char *format, va_list args)
- int `ircsprintf` (char *str, const char *format,...)

4.6.1 Documentatie van functies

4.6.1.1 ircsprintf()

```
int ircsprintf (  
    char * str,  
    const char * format,  
    ... )
```

Hier is de call graaf voor deze functie:



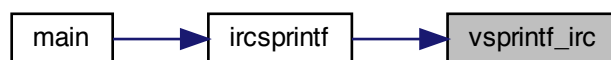
Hier is de caller graaf voor deze functie:



4.6.1.2 vsprintf_irc()

```
int vsprintf_irc (
    char * str,
    const char * format,
    va_list args )
```

Hier is de caller graaf voor deze functie:

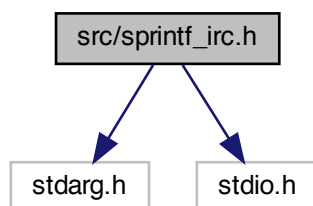


4.7 src/sprintf_irc.h Bestand Referentie

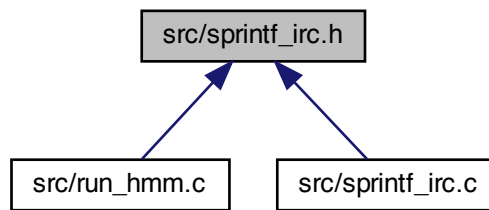
```
#include <stdarg.h>
```

```
#include <stdio.h>
```

Include afhankelijkheidsgraaf voor sprintf_irc.h:



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



Functies

- int [ircsprintf](#) (char *, const char *,...)

4.7.1 Documentatie van functies

4.7.1.1 ircsprintf()

```
int ircsprintf (  
    char * ,  
    const char * ,  
    ... )
```

Hier is de call graaf voor deze functie:



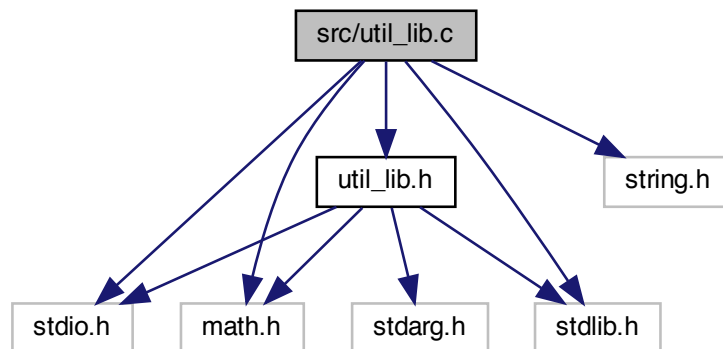
Hier is de caller graaf voor deze functie:



4.8 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`

Funcities

- `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

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

4.8.1 Documentatie van macro's

4.8.1.1 TR_SIZE

```
#define TR_SIZE 14
```

4.8.2 Documentatie van functies

4.8.2.1 dmatrix()

```
double** dmatrix (  
    int num_row,  
    int num_col )
```

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:



4.8.2.2 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:



4.8.2.3 free_dmatrix()

```
void free_dmatrix (
    double ** m,
    int num_row )
```

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



4.8.2.4 free_dvector()

```
void free_dvector (
    double * v )
```

Frees the memory allocation of an vector with datatype double.

4.8.2.5 free_imatrix()

```
void free_imatrix (
    int ** m,
    int num_row )
```

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



4.8.2.6 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:



4.8.2.7 get_protein()

```
void get_protein (
    char * dna,
    char * protein,
    int strand,
    int whole_genome )
```

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:



Hier is de caller graaf voor deze functie:



4.8.2.8 get_rc_dna()

```
void get_rc_dna (
    char * dna,
    char * dna1 )
```

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



Hier is de caller graaf voor deze functie:



4.8.2.9 get_rc_dna_indel()

```
void get_rc_dna_indel (
    char * dna,
    char * dna1 )
```

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



Hier is de caller graaf voor deze functie:



4.8.2.10 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:



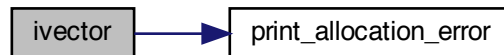
Hier is de caller graaf voor deze functie:



4.8.2.11 ivector()

```
int* ivector (
    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:



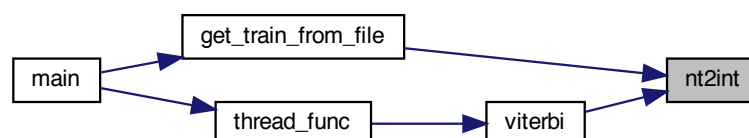
Hier is de caller graaf voor deze functie:



4.8.2.12 nt2int()

```
int nt2int (
    char nt )
```

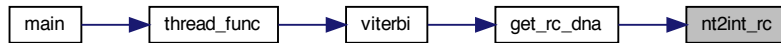
Hier is de caller graaf voor deze functie:



4.8.2.13 nt2int_rc()

```
int nt2int_rc (  
    char nt )
```

Hier is de caller graaf voor deze functie:



4.8.2.14 nt2int_rc_indel()

```
int nt2int_rc_indel (  
    char nt )
```

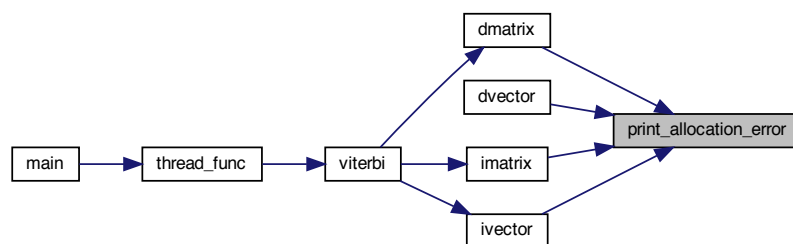
Hier is de caller graaf voor deze functie:



4.8.2.15 print_allocation_error()

```
void print_allocation_error (  
    const char * format,  
    ... )
```

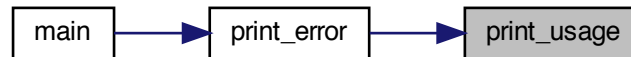
Custom error function to print allocation errors. Mostly called from matrix or vector functions. Hier is de caller graaf voor deze functie:



4.8.2.16 print_usage()

```
void print_usage ( )
```

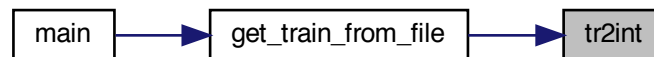
Print how the program should be used. called mainly on help or error. Hier is de caller graaf voor deze functie:



4.8.2.17 tr2int()

```
int tr2int (
    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:



4.8.2.18 trinucleotide()

```
int trinucleotide (
    char a,
    char b,
    char c )
```

Hier is de caller graaf voor deze functie:



4.8.2.19 trinucleotide_pep()

```
int trinucleotide_pep (
    char a,
    char b,
    char c )
```

Hier is de caller graaf voor deze functie:



4.8.3 Documentatie van variabelen

4.8.3.1 anti_codon_code

```
char anti_codon_code[65]
```

Initiële waarde:

```
= { 'F', 'V', 'L', 'I',
    'C', 'G', 'R', 'S',
    'S', 'A', 'P', 'T',
    'Y', 'D', 'H', 'N',
    'L', 'V', 'L', 'M',
    'W', 'G', 'R', 'R',
    'S', 'A', 'P', 'T',
    '*', 'E', 'Q', 'K',
    'F', 'V', 'L', 'I',
    'C', 'G', 'R', 'S',
    'S', 'A', 'P', 'T',
    'Y', 'D', 'H', 'N',
    'L', 'V', 'L', 'I',
    '*', 'G', 'R', 'R',
    'S', 'A', 'P', 'T',
    '*', 'E', 'Q', 'K', 'X' }
```

4.8.3.2 codon11

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

4.8.3.3 codon5

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

4.8.3.4 codon_code

```
char codon_code[65]
```

Initiële waarde:

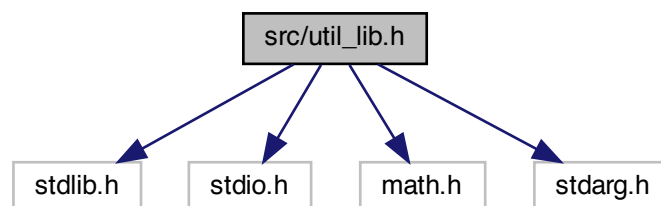
```
= { 'K', 'N', 'K', 'N',  
    'T', 'T', 'T', 'T',  
    'R', 'S', 'R', 'S',  
    'I', 'I', 'M', 'I',  
    'Q', 'H', 'Q', 'H',  
    'P', 'P', 'P', 'P',  
    'R', 'R', 'R', 'R',  
    'L', 'L', 'L', 'L',  
    'E', 'D', 'E', 'D',  
    'A', 'A', 'A', 'A',  
    'G', 'G', 'G', 'G',  
    'V', 'V', 'V', 'V',  
    '*', 'Y', '*', 'Y',  
    'S', 'S', 'S', 'S',  
    '*', 'C', 'W', 'C',  
    'L', 'F', 'L', 'F', 'X'  
}
```

4.8.3.5 tr_list

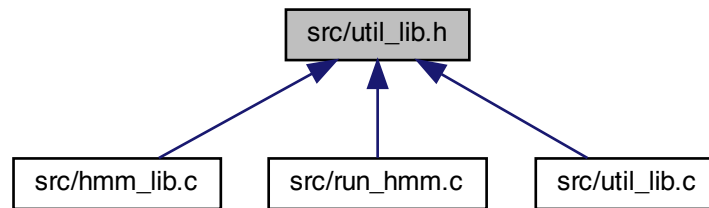
```
char* tr_list[TR_SIZE] = { "MM", "MI", "MD", "II", "IM", "DD", "DM", "GE", "GG", "ER", "RS", "RR", "ES", "E←  
S1" }
```

4.9 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,...)

4.9.1 Documentatie van functies

4.9.1.1 [dmatrix\(\)](#)

```
double** dmatrix (
    int num_row,
    int num_col )
```

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:



4.9.1.2 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:



4.9.1.3 free_dmatrix()

```
void free_dmatrix (
    double ** m,
    int num_row )
```

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



4.9.1.4 free_dvector()

```
void free_dvector (
    double * v )
```

Frees the memory allocation of an vector with datatype double.

4.9.1.5 free_imatrix()

```
void free_imatrix (
    int ** m,
    int num_row )
```

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



4.9.1.6 free_ivector()

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

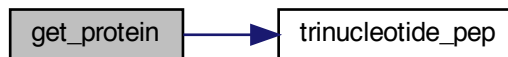
Frees the memory allocation of an vector with datatype int. Hier is de caller graaf voor deze functie:



4.9.1.7 get_protein()

```
void get_protein (
    char * dna,
    char * protein,
    int strand,
    int whole_genome )
```

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:



4.9.1.8 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:



Hier is de caller graaf voor deze functie:



4.9.1.9 ivector()

```
int* ivector (
    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:



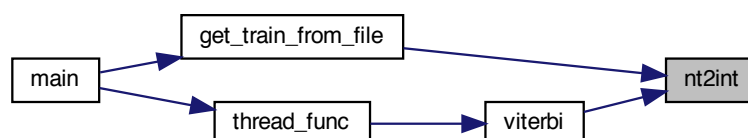
Hier is de caller graaf voor deze functie:



4.9.1.10 nt2int()

```
int nt2int (
    char nt )
```

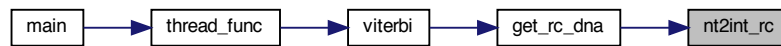
Hier is de caller graaf voor deze functie:



4.9.1.11 nt2int_rc()

```
int nt2int_rc (  
    char nt )
```

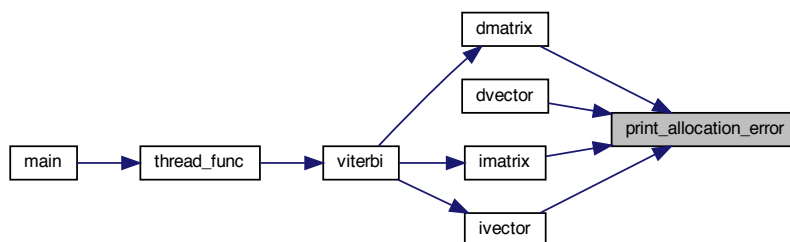
Hier is de caller graaf voor deze functie:



4.9.1.12 print_allocation_error()

```
void print_allocation_error (  
    const char * format,  
    ... )
```

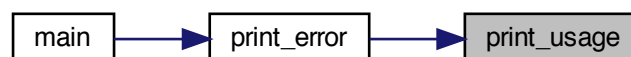
Custom error function to print allocation errors. Mostly called from matrix or vector functions. Hier is de caller graaf voor deze functie:



4.9.1.13 print_usage()

```
void print_usage ( )
```

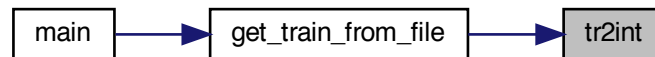
Print how the program should be used. called mainly on help or error. Hier is de caller graaf voor deze functie:



4.9.1.14 tr2int()

```
int tr2int (  
    char * tr )
```

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



4.9.1.15 trinucleotide()

```
int trinucleotide (  
    char a,  
    char b,  
    char c )
```

Hier is de caller graaf voor deze functie:

