

# GenAlg

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## Introduction

GenAlg is a C library providing structures and functions implementing a Genetic Algorithm.

The genes are memorized as a VecFloat and/or VecShort. The user can defined a range of possible values for each gene. The user can define the size of the pool of entities and the size of the breeding pool. Selection, reproduction and mutation are designed to efficiently explore all the possible gene combination, and avoid local optimum. It is also possible to save and load

the GenAlg.

It uses the PBErr, PBMath and GSet libraries.

## 1 Definitions

A genetic algorithm has 3 steps. In a pool of entities it discards a given number of entities based on their ranking (given by a mean external to the algorithm). Then it replaces each of the discarded entity by a new one created from two selected entities from the non discarded one. The newly created entity's properties are a mix of these two selected entities, plus a certain amount of random modification. The detail of the implementation in GenAlg of these 3 steps (selection, reproduction and mutation) are given below.

### 1.1 Selection

The non discarded entities are called 'elite' in GenAlg. The size of the pool of elite is configurable by the user. The selection of two elite entities is simply a random selection in the pool of elites. Selection of the same elite twice is allowed.

### 1.2 Reproduction

The reproduction step copies the genes of the elite entity into the new entity. Each gene has a probability of 50% to be chosen in one or the other elite.

### 1.3 Mutation

The mutation occurs as follow. First we calculate the probability of mutation for every gene as follow:  $P = \frac{rank}{nbEntity}$  where rank is the rank of the discarded entity in the pool of entities, and nbEntity is the number of entities in the pool. A gene affected by a mutation according to this probability is modified as follow. The amplitude of the mutation is equal to  $1 - \frac{1}{\sqrt{age+1}}$  where age is the age of the oldest elite entity used during the reproduction step for the entity. Then the new value of the gene is equals

to  $gene + range * amp * (rnd + delta)$  where  $gene$  is the current value of the gene,  $range$  is equal to  $max_{gene} - min_{gene}$  (the difference of the maximum allowed value for this gene and its minimum value),  $amp$  is the amplitude calculated above,  $rnd$  is a random value between -0.5 and 0.5, and  $delta$  is the mutation that has been applied to this gene in the corresponding elite entity. Genes' value is kept in bounds by bouncing it on the bounds when necessary ( $gene = 2 * bound - gene$ )

To counteract inbreeding (the algorithm getting stuck into a local minimum), we also apply mutation to all the entities except the best one when the diversity level of the elite pool fall below a threshold (set to 0.01 by default). The diversity level is calculated as follow  $\frac{1}{nbElite} \sum_{i=1}^{nbElite} \frac{\|\vec{adn}(elite_i) - \vec{adn}(elite_0)\|}{\|\vec{bound}_{max} - \vec{bound}_{min}\|}$  where  $nbElite$  is the number of elite entities,  $\vec{adn}(elite_i)$  is the genes vector of the  $i$ -th elite entity, and  $\vec{bound}_{max}$  and  $\vec{bound}_{min}$  are the vector of maximum and minimum values of the genes.

Some explanation:  $delta$  bias the mutation toward the direction that improved the result at previous step; in the pool of discarded entities high ranked ones tend to have few mutations and low ranked ones tend to have more mutation, this tends to cover any possibilities of evolution; entities newly entered in the elite pool tends to produce new entities near to them (in term of distance in the genes space), while older ones tend to produce more diverse new entities, thus the exploration of solution space occurs from the vicinity of newly better solutions toward larger areas; from the previous point, a good entity tends to create a lot of similar entity, which may lead to an elite pool saturated with very similar entities (inbreeding) from which the algorithm can't escape, this is prevented by the forced mutation of elites when the inbreeding level gets too high.

## 2 Interface

```
// ===== GENALG.H =====

#ifndef GENALG_H
#define GENALG_H

// ===== Include =====

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

```

#include <stdbool.h>
#include "pberr.h"
#include "pbmath.h"
#include "gset.h"

// ===== Define =====

#define GENALG_NBENTITIES 100
#define GENALG_NBELITES 20
#define GENALG_DIVERSITYTHRESHOLD 0.01

// ----- GenAlgAdn

// ===== Data structure =====

typedef struct GenAlg GenAlg;

typedef struct GenAlgAdn {
    // ID
    unsigned long int _id;
    // Age
    unsigned long int _age;
    // Adn for floating point value
    VecFloat* _adnF;
    // Delta Adn during mutation for floating point value
    VecFloat* _deltaAdnF;
    // Adn for integer point value
    VecShort* _adnI;
} GenAlgAdn;

// ===== Functions declaration =====

// Create a new GenAlgAdn with ID 'id', 'lengthAdnF' and 'lengthAdnI'
// 'lengthAdnF' and 'lengthAdnI' must be greater than or equal to 0
GenAlgAdn* GenAlgAdnCreate(const int id, const int lengthAdnF,
    const int lengthAdnI);

// Free memory used by the GenAlgAdn 'that'
void GenAlgAdnFree(GenAlgAdn** that);

// Return the adn for floating point values of the GenAlgAdn 'that'
#if BUILDMODE != 0
inline
#endif
const VecFloat* GAAdnAdnF(const GenAlgAdn* const that);

// Return the delta of adn for floating point values of the
// GenAlgAdn 'that'
#if BUILDMODE != 0
inline
#endif
const VecFloat* GAAdnDeltaAdnF(const GenAlgAdn* const that);

// Return the adn for integer values of the GenAlgAdn 'that'
#if BUILDMODE != 0
inline
#endif
const VecShort* GAAdnAdnI(const GenAlgAdn* const that);

// Initialise randomly the genes of the GenAlgAdn 'that' of the
// GenAlg 'ga'
void GAAdnInit(const GenAlgAdn* const that, const GenAlg* ga);

```

```

// Get the 'iGene'-th gene of the adn for floating point values of the
// GenAlgAdn 'that'
#if BUILDMODE != 0
inline
#endif
float GAAdnGetGeneF(const GenAlgAdn* const that, const int iGene);

// Get the delta of the 'iGene'-th gene of the adn for floating point
// values of the GenAlgAdn 'that'
#if BUILDMODE != 0
inline
#endif
float GAAdnGetDeltaGeneF(const GenAlgAdn* const that, const int iGene);

// Get the 'iGene'-th gene of the adn for int values of the
// GenAlgAdn 'that'
#if BUILDMODE != 0
inline
#endif
int GAAdnGetGeneI(const GenAlgAdn* const that, const int iGene);

// Set the 'iGene'-th gene of the adn for floating point values of the
// GenAlgAdn 'that' to 'gene'
#if BUILDMODE != 0
inline
#endif
void GAAdnSetGeneF(GenAlgAdn* const that, const int iGene,
    const float gene);

// Set the delta of the 'iGene'-th gene of the adn for floating point
// values of the GenAlgAdn 'that' to 'delta'
#if BUILDMODE != 0
inline
#endif
void GAAdnSetDeltaGeneF(GenAlgAdn* const that, const int iGene,
    const float delta);

// Set the 'iGene'-th gene of the adn for int values of the
// GenAlgAdn 'that' to 'gene'
#if BUILDMODE != 0
inline
#endif
void GAAdnSetGeneI(GenAlgAdn* const that, const int iGene,
    const short gene);

// Get the id of the GenAlgAdn 'that'
#if BUILDMODE != 0
inline
#endif
unsigned long int GAAdnGetId(const GenAlgAdn* const that);

// Get the age of the GenAlgAdn 'that'
#if BUILDMODE != 0
inline
#endif
unsigned long int GAAdnGetAge(const GenAlgAdn* const that);

// Print the information about the GenAlgAdn 'that' on the
// stream 'stream'
void GAAdnPrintln(const GenAlgAdn* const that, FILE* const stream);

```

```

// Return true if the GenAlgAdn 'that' is new, i.e. is age equals 1
// Return false
#if BUILDMODE != 0
inline
#endif
bool GAAdnIsNew(const GenAlgAdn* const that);

// ----- GenAlg

// ===== Define =====

#define GABestAdnF(that) GAAdnAdnF(GAAdn(that, 0))
#define GABestAdnI(that) GAAdnAdnI(GAAdn(that, 0))

// ===== Data structure =====

typedef struct GenAlg {
    // GSet of GenAlgAdn, sortval == score so the head of the set is the
    // worst adn and the tail of the set is the best
    GSet* _adns;
    // Current epoch
    unsigned long int _curEpoch;
    // Nb elite entities in population
    int _nbElites;
    // Id of the next new GenAlgAdn
    unsigned long int _nextId;
    // Length of adn for floating point value
    const int _lengthAdnF;
    // Length of adn for integer value
    const int _lengthAdnI;
    // Bounds (min, max) for floating point values adn
    VecFloat2D* _boundsF;
    // Bounds (min, max) for integer values adn
    VecShort2D* _boundsI;
    // Diversity threshold for KTEvent
    float _diversityThreshold;
    // Norm of the range value for adns (optimization for diversity
    // calculation)
    float _normRangeFloat;
    float _normRangeInt;
} GenAlg;

// ===== Functions declaration =====

// Create a new GenAlg with 'nbEntities', 'nbElites', 'lengthAdnF'
// and 'lengthAdnI'
// 'nbEntities' must greater than 2
// 'nbElites' must greater than 1
// 'lengthAdnF' and 'lengthAdnI' must be greater than or equal to 0
GenAlg* GenAlgCreate(const int nbEntities, const int nbElites,
    const int lengthAdnF, const int lengthAdnI);

// Free memory used by the GenAlg 'that'
void GenAlgFree(GenAlg** that);

// Return the GSet of the GenAlg 'that'
#if BUILDMODE != 0
inline
#endif
GSet* GAAdns(const GenAlg* const that);

// Return the nb of entities of the GenAlg 'that'

```

```

#if BUILDMODE != 0
inline
#endif
int GAGetNbAdns(const GenAlg* const that);

// Return the nb of elites of the GenAlg 'that'
#if BUILDMODE != 0
inline
#endif
int GAGetNbElites(const GenAlg* const that);

// Return the diversity threshold of the GenAlg 'that'
#if BUILDMODE != 0
inline
#endif
float GAGetDiversityThreshold(const GenAlg* const that);

// Set the diversity threshold of the GenAlg 'that' to 'div'
#if BUILDMODE != 0
inline
#endif
void GASetDiversityThreshold(GenAlg* const that, const float div);

// Return the current epoch of the GenAlg 'that'
#if BUILDMODE != 0
inline
#endif
unsigned long int GAGetCurEpoch(const GenAlg* const that);

// Set the nb of entities of the GenAlg 'that' to 'nb'
// 'nb' must be greater than 1, if 'nb' is lower than the current nb
// of elite the number of elite is set to 'nb' - 1
void GASetNbEntities(GenAlg* const that, const int nb);

// Set the nb of elites of the GenAlg 'that' to 'nb'
// 'nb' must be greater than 0, if 'nb' is greater or equal to the
// current nb of entities the number of entities is set to 'nb' + 1
void GASetNbElites(GenAlg* const that, const int nb);

// Get the length of adn for floating point value
#if BUILDMODE != 0
inline
#endif
int GAGetLengthAdnFloat(const GenAlg* const that);

// Get the length of adn for integer value
#if BUILDMODE != 0
inline
#endif
int GAGetLengthAdnInt(const GenAlg* const that);

// Get the bounds for the 'iGene'-th gene of adn for floating point
// values
#if BUILDMODE != 0
inline
#endif
const VecFloat2D* GABoundsAdnFloat(const GenAlg* const that,
    const int iGene);

// Get the bounds for the 'iGene'-th gene of adn for integer values
#if BUILDMODE != 0
inline

```

```

#endif
const VecShort2D* GABoundsAdnInt(const GenAlg* const that,
    const int iGene);

// Set the bounds for the 'iGene'-th gene of adn for floating point
// values to a copy of 'bounds'
#if BUILDMODE != 0
inline
#endif
void GASetBoundsAdnFloat(GenAlg* const that, const int iGene,
    const VecFloat2D* const bounds);

// Set the bounds for the 'iGene'-th gene of adn for integer values
// to a copy of 'bounds'
#if BUILDMODE != 0
inline
#endif
void GASetBoundsAdnInt(GenAlg* const that, const int iGene,
    const VecShort2D* bounds);

// Get the GenAlgAdn of the GenAlg 'that' currently at rank 'iRank'
#if BUILDMODE != 0
inline
#endif
GenAlgAdn* GAAdn(const GenAlg* const that, const int iRank);

// Init the GenAlg 'that'
// Must be called after the bounds have been set
// The random generator must have been initialised before calling this
// function
void GAInit(GenAlg* const that);

// Step an epoch for the GenAlg 'that' with the current ranking of
// GenAlgAdn
void GAStep(GenAlg* const that);

// Step an epoch for the GenAlg 'that' with the current ranking of
// GenAlgAdn, only considering the first 'nbGeneF' genes of float adn
// and the first 'nbGeneI' of int adn
void GASetSubset(GenAlg* const that, const int nbGeneF,
    const int nbGeneI);

// Print the information about the GenAlg 'that' on the stream 'stream'
void GAPrintLn(const GenAlg* const that, FILE* const stream);

// Get the average diversity of current entities of the GenAlg 'that'
// The return value is in [0.0, 1.0]
// 0.0 means all the elite entities have exactly the same adns
// 1.0 means all the elite entities except the first one have adns
// as different compare to the first one's adn as possible given the
// range of adn values
float GAGetDiversity(const GenAlg* const that);

// Function which return the JSON encoding of 'that'
JSONNode* GAEncodeAsJSON(const GenAlg* const that);

// Function which decode from JSON encoding 'json' to 'that'
bool GADecodeAsJSON(GenAlg** that, const JSONNode* const json);

// Load the GenAlg 'that' from the stream 'stream'
// If the GenAlg is already allocated, it is freed before loading
// Return true in case of success, else false

```



```

bool GALoad(GenAlg** that, FILE* const stream);

// Save the GenAlg 'that' to the stream 'stream'
// If 'compact' equals true it saves in compact form, else it saves in
// readable form
// Return true in case of success, else false
bool GASave(const GenAlg* const that, FILE* const stream,
            const bool compact);

// Set the value of the GenAlgAdn 'adn' of the GenAlg 'that' to 'val'
#if BUILDMODE != 0
inline
#endif
void GASetAdnValue(GenAlg* const that, const GenAlgAdn* const adn,
                  const float val);

// Update the norm of the range value for adans of the GenAlg 'that'
void GAUpdateNormRange(GenAlg* const that);

// ===== Polymorphism =====

// ===== Inliner =====

#if BUILDMODE != 0
#include "genalg-inline.c"
#endif

#endif

```

## 3 Code

### 3.1 genalg.c

```

// ===== GENALG.C =====

// ===== Include =====

#include "genalg.h"
#if BUILDMODE == 0
#include "genalg-inline.c"
#endif

// ----- GenAlgAdn

// ===== Functions declaration =====

// ===== Functions implementation =====

// Create a new GenAlgAdn with ID 'id', 'lengthAdnF' and 'lengthAdnI'
// 'lengthAdnF' and 'lengthAdnI' must be greater than or equal to 0
GenAlgAdn* GenAlgAdnCreate(const int id, const int lengthAdnF,
                          const int lengthAdnI) {
#if BUILDMODE == 0
    if (lengthAdnF < 0) {
        GenAlgErr->_type = PBErrTypeInvalidArg;
        sprintf(GenAlgErr->_msg, "'lengthAdnF' is invalid (%d>=0)",

```

```

        lengthAdnF);
    PBErrCatch(GenAlgErr);
}
if (lengthAdnI < 0) {
    GenAlgErr->_type = PBErrTypeInvalidArg;
    sprintf(GenAlgErr->_msg, "'lengthAdnI' is invalid (%d>=0)",
        lengthAdnI);
    PBErrCatch(GenAlgErr);
}
#endif
// Allocate memory
GenAlgAdn* that = PBErrMalloc(GenAlgErr, sizeof(GenAlgAdn));
// Set the properties
that->_age = 1;
that->_id = id;
if (lengthAdnF > 0) {
    that->_adnF = VecFloatCreate(lengthAdnF);
    that->_deltaAdnF = VecFloatCreate(lengthAdnF);
} else {
    that->_adnF = NULL;
    that->_deltaAdnF = NULL;
}
if (lengthAdnI > 0)
    that->_adnI = VecShortCreate(lengthAdnI);
else
    that->_adnI = NULL;
// Return the new GenAlgAdn
return that;
}

// Free memory used by the GenAlgAdn 'that'
void GenAlgAdnFree(GenAlgAdn** that) {
    // Check the argument
    if (that == NULL || *that == NULL) return;
    // Free memory
    if ((*that)->_adnF != NULL)
        VecFree(&((*that)->_adnF));
    if ((*that)->_deltaAdnF != NULL)
        VecFree(&((*that)->_deltaAdnF));
    if ((*that)->_adnI != NULL)
        VecFree(&((*that)->_adnI));
    free(*that);
    // Set the pointer to null
    *that = NULL;
}

// Initialise randomly the genes of the GenAlgAdn 'that' of the
// GenAlg 'ga'
void GAAdnInit(const GenAlgAdn* const that, const GenAlg* const ga) {
    #if BUILDMODE == 0
        if (that == NULL) {
            GenAlgErr->_type = PBErrTypeNullPointer;
            sprintf(GenAlgErr->_msg, "'that' is null");
            PBErrCatch(GenAlgErr);
        }
    #endif
    // For each floating point value gene
    for (int iGene = GAGetLengthAdnFloat(ga); iGene--;) {
        float min = VecGet(GABoundsAdnFloat(ga, iGene), 0);
        float max = VecGet(GABoundsAdnFloat(ga, iGene), 1);
        float val = min + (max - min) * rnd();
        VecSet(that->_adnF, iGene, val);
    }
}

```

```

    }
    // For each integer value gene
    for (int iGene = GAGetLengthAdnInt(ga); iGene--;) {
        short min = VecGet(GABoundsAdnInt(ga, iGene), 0);
        short max = VecGet(GABoundsAdnInt(ga, iGene), 1);
        short val = (short)round((float)min + (float)(max - min) * rnd());
        VecSet(that->_adnI, iGene, val);
    }
}

// Print the information about the GenAlgAdn 'that' on the
// stream 'stream'
void GAAdnPrintln(const GenAlgAdn* const that, FILE* const stream) {
#ifdef BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
    if (stream == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'stream' is null");
        PBErrCatch(GenAlgErr);
    }
#endif
    fprintf(stream, "id:%lu age:%lu", GAAdnGetId(that), GAAdnGetAge(that));
    fprintf(stream, "\n");
    fprintf(stream, "  adnF:");
    VecFloatPrint(GAAdnAdnF(that), stream, 6);
    fprintf(stream, "\n");
    fprintf(stream, "  deltaAdnF:");
    VecFloatPrint(GAAdnDeltaAdnF(that), stream, 6);
    fprintf(stream, "\n");
    fprintf(stream, "  adnI:");
    VecPrint(GAAdnAdnI(that), stream);
    fprintf(stream, "\n");
}

// ----- GenAlg

// ===== Functions declaration =====

// Select the rank of two parents for the SRM algorithm
// Return the ranks in 'parents', with parents[0] <= parents[1]
void GASelectParents(const GenAlg* const that, int* const parents);

// Set the genes of the entity at rank 'iChild' as a 50/50 mix of the
// genes of entities at ranks 'parents[0]' and 'parents[1]'
void GAReproduction(GenAlg* const that, const int* const parents,
    const int iChild);

// Mute the genes of the entity at rank 'iChild'
// The probability of mutation for one gene is equal to
// 'rankChild'/'that'->_nbEntities
// The amplitude of the mutation
// is equal to
// (max-min).(gauss(0.0, 1.0)+deltaAdn).ln('parents[0]'.age)
void GAMute(GenAlg* const that, const int* const parents,
    const int iChild);

// Reset the GenAlg 'that'
// Randomize all the gene except those of the first adn

```

```

void GAKTEvent(GenAlg* const that);

// Get the diversity value of 'adnA' against 'adnB'
// The diversity is equal to
float GAAdnGetDiversity(const GenAlgAdn* const adnA,
    const GenAlgAdn* const adnB, const GenAlg* const ga);

// ===== Functions implementation =====

// Create a new GenAlg with 'nbEntities', 'nbElites', 'lengthAdnF'
// and 'lengthAdnI'
// 'nbEntities' must greater than 2
// 'nbElites' must greater than 1
// 'lengthAdnF' and 'lengthAdnI' must be greater than or equal to 0
GenAlg* GenAlgCreate(const int nbEntities, const int nbElites,
    const int lengthAdnF, const int lengthAdnI) {
    // Allocate memory
    GenAlg* that = PBErrMalloc(GenAlgErr, sizeof(GenAlg));
    // Set the properties
    that->_adns = GSetCreate();
    that->_curEpoch = 0;
    *(int*)&(that->_lengthAdnF) = lengthAdnF;
    *(int*)&(that->_lengthAdnI) = lengthAdnI;
    if (lengthAdnF > 0) {
        that->_boundsF =
            PBErrMalloc(GenAlgErr, sizeof(VecFloat2D) * lengthAdnF);
        for (int iGene = lengthAdnF; iGene--;)
            that->_boundsF[iGene] = VecFloatCreateStatic2D();
    } else
        that->_boundsF = NULL;
    if (lengthAdnI > 0) {
        that->_boundsI =
            PBErrMalloc(GenAlgErr, sizeof(VecShort2D) * lengthAdnI);
        for (int iGene = lengthAdnI; iGene--;)
            that->_boundsI[iGene] = VecShortCreateStatic2D();
    } else
        that->_boundsI = NULL;
    that->_normRangeFloat = 1.0;
    that->_normRangeInt = 1.0;
    that->_nbElites = 0;
    that->_nextId = 0;
    that->_diversityThreshold = GENALG_DIVERSITYTHRESHOLD;
    GSetNbEntities(that, nbEntities);
    GSetNbElites(that, nbElites);
    // Return the new GenAlg
    return that;
}

// Free memory used by the GenAlg 'that'
void GenAlgFree(GenAlg** that) {
    // Check the argument
    if (that == NULL || *that == NULL) return;
    // Free memory
    GSetIterForward iter = GSetIterForwardCreateStatic(GAAdns(*that));
    do {
        GenAlgAdn* gaEnt = GSetIterGet(&iter);
        GenAlgAdnFree(&gaEnt);
    } while (GSetIterStep(&iter));
    GSetFree(&((*that)->_adns));
    if ((*that)->_boundsF != NULL)
        free((*that)->_boundsF);
    if ((*that)->_boundsI != NULL)

```

```

    free((*that)->_boundsI);
    free(*that);
    // Set the pointer to null
    *that = NULL;
}

// Set the nb of entities of the GenAlg 'that' to 'nb'
// 'nb' must be greater than 1, if 'nb' is lower than the current nb
// of elite the number of elite is set to 'nb' - 1
void GASetNbEntities(GenAlg* const that, const int nb) {
#ifdef BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
    if (nb <= 1) {
        GenAlgErr->_type = PBErrTypeInvalidArg;
        sprintf(GenAlgErr->_msg, "'nb' is invalid (%d>1)", nb);
        PBErrCatch(GenAlgErr);
    }
#endif
    while (GSetNbElem(GAAdns(that)) > nb) {
        GenAlgAdn* gaEnt = GSetPop(GAAdns(that));
        GenAlgAdnFree(&gaEnt);
    }
    while (GSetNbElem(GAAdns(that)) < nb) {
        GenAlgAdn* ent = GenAlgAdnCreate(that->_nextId++,
            GAGetLengthAdnFloat(that), GAGetLengthAdnInt(that));
        GSetPush(GAAdns(that), ent);
    }
    if (GAGetNbElites(that) >= nb)
        GASetNbElites(that, nb - 1);
}

// Set the nb of elites of the GenAlg 'that' to 'nb'
// 'nb' must be greater than 0, if 'nb' is greater or equal to the
// current nb of entities the number of entities is set to 'nb' + 1
void GASetNbElites(GenAlg* const that, const int nb) {
#ifdef BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
    if (nb <= 1) {
        GenAlgErr->_type = PBErrTypeInvalidArg;
        sprintf(GenAlgErr->_msg, "'nb' is invalid (%d>1)", nb);
        PBErrCatch(GenAlgErr);
    }
#endif
    if (GAGetNbAdns(that) <= nb)
        GASetNbEntities(that, nb + 1);
    that->_nbElites = nb;
}

// Init the GenAlg 'that'
// Must be called after the bounds have been set
// The random generator must have been initialised before calling this
// function
void GAINit(GenAlg* const that) {
#ifdef BUILDMODE == 0

```

```

    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
#endif
    // For each adn
    GSetIterForward iter = GSetIterForwardCreateStatic(GAAdns(that));
    do {
        // Get the adn
        GenAlgAdn* adn = GSetIterGet(&iter);
        // Initialise randomly the genes of the adn
        GAAdnInit(adn, that);
    } while (GSetIterStep(&iter));
}

// Reset the GenAlg 'that'
// Randomize all the gene except those of the best adn
void GAKTEvent(GenAlg* const that) {
#ifdef BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
#endif
    // For each adn except the best one
    GSetIterBackward iter = GSetIterBackwardCreateStatic(GAAdns(that));
    GSetIterStep(&iter);
    // We suppose here there is at least 2 adns in the pool
    do {
        // Get the adn
        GenAlgAdn* adn = GSetIterGet(&iter);
        // Get the diversity of this adn against the first one
        float diversity = GAAdnGetDiversity(adn, GAAdn(that, 0), that);
        // If the diversity is under the threshold
        if (diversity < GAGetDiversityThreshold(that)) {
            // Initialise randomly the genes of the adn
            GAAdnInit(adn, that);
            // Reset the age of the child
            adn->_age = 1;
            // Set the id of the child
            adn->_id = (that->_nextId)++;
        }
    } while (GSetIterStep(&iter));
}

// Step an epoch for the GenAlg 'that' with the current ranking of
// GenAlgAdn, only considering the first 'nbGeneF' genes of float adn
// and the first 'nbGeneI' of int adn
void GASTepSubset(GenAlg* const that, const int nbGeneF,
    const int nbGeneI) {
#ifdef BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
    if (nbGeneF <= 0 || nbGeneF > GAGetLengthAdnFloat(that)) {
        GenAlgErr->_type = PBErrTypeInvalidArg;
        sprintf(GenAlgErr->_msg, "'nbGeneF' is invalid (0<=%d<=%d)",
            nbGeneF, GAGetLengthAdnFloat(that));
    }

```

```

    PBErCatch(GenAlgErr);
}
if (nbGeneI <= 0 || nbGeneI > GAGetLengthAdnInt(that)) {
    GenAlgErr->_type = PBErTypeInvalidArg;
    sprintf(GenAlgErr->_msg, "'nbGeneI' is invalid (0<=%d<=%d)",
        nbGeneI, GAGetLengthAdnInt(that));
    PBErCatch(GenAlgErr);
}
#endif
// Declare variables to memorize the real size of float and int adn
int realNbGeneF = GAGetLengthAdnFloat(that);
int realNbGeneI = GAGetLengthAdnInt(that);
// Trick the length of adns
*(int*)&(that->_lengthAdnF) = nbGeneF;
*(int*)&(that->_lengthAdnI) = nbGeneI;
for (int iAdn = GAGetNbAdns(that); iAdn--;) {
    GenAlgAdn* adn = GAAdn(that, iAdn);
    adn->_adnF->_dim = nbGeneF;
    adn->_deltaAdnF->_dim = nbGeneF;
    adn->_adnI->_dim = nbGeneI;
}
// Step the GenAlg
GAStep(that);
// Reset the true length of adns
*(int*)&(that->_lengthAdnF) = realNbGeneF;
*(int*)&(that->_lengthAdnI) = realNbGeneI;
for (int iAdn = GAGetNbAdns(that); iAdn--;) {
    GenAlgAdn* adn = GAAdn(that, iAdn);
    adn->_adnF->_dim = realNbGeneF;
    adn->_deltaAdnF->_dim = realNbGeneF;
    adn->_adnI->_dim = realNbGeneI;
}
}

// Step an epoch for the GenAlg 'that' with the current ranking of
// GenAlgAdn
void GAStep(GenAlg* const that) {
#if BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErCatch(GenAlgErr);
    }
#endif
    // Selection, Reproduction, Mutation
    // Ensure the set of adns is sorted
    GSetSort(GAAdns(that));
    // Declare a variable to memorize the parents
    int parents[2];
    // Get the diversity level
    float diversity = GAGetDiversity(that);
    // If the diversity level is too low
    if (diversity < GAGetDiversityThreshold(that)) {
        // Break the diversity by applying a KT event (in memory of
        // chickens' grand pa and grand ma)
        GAKTEvent(that);
    }
    // Else, the diversity level is ok
    } else {
        // For each adn which is an elite
        for (int iAdn = 0; iAdn < GAGetNbElites(that); ++iAdn) {
            // Increment age

```

```

        (GAAdn(that, iAdn)->_age)++;
    }
    // For each adn which is not an elite
    for (int iAdn = GAGetNbElites(that); iAdn < GAGetNbAdns(that);
        ++iAdn) {
        // Select two parents for this adn
        GASelectParents(that, parents);
        // Set the genes of the adn as a 50/50 mix of parents' genes
        GAReproduction(that, parents, iAdn);
        // Mute the genes of the adn
        GAMute(that, parents, iAdn);
    }
}
// Increment the number of epochs
++(that->_curEpoch);
}

// Select the rank of two parents for the SRM algorithm
// Return the ranks in 'parents', with parents[0] <= parents[1]
void GASelectParents(const GenAlg* const that, int* const parents) {
#ifdef BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
    if (parents == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'parents' is null");
        PBErrCatch(GenAlgErr);
    }
#endif
    // Declare a variable to memorize the parents' rank
    int p[2];
    for (int i = 2; i--;)
        // p[i] below may be equal to the rank of the highest non elite
        // adn, but it's not a problem so leave it and let's call that
        // the Hawking radiation of this function in memory of this great
        // man.
        p[i] = (int)floor(rnd() * (float)GAGetNbElites(that));
    // Memorize the sorted parents' rank
    if (p[0] < p[1]) {
        parents[0] = p[0];
        parents[1] = p[1];
    } else {
        parents[0] = p[1];
        parents[1] = p[0];
    }
}

// Set the genes of the adn at rank 'iChild' as a 50/50 mix of the
// genes of adns at ranks 'parents[0]' and 'parents[1]'
void GAReproduction(GenAlg* const that, const int* const parents,
    const int iChild) {
#ifdef BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
    if (parents == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
    }

```



```

    sprintf(GenAlgErr->_msg, "'parents' is null");
    PBErrCatch(GenAlgErr);
}
if (iChild < 0 || iChild >= GAGetNbAdns(that)) {
    GenAlgErr->_type = PBErrTypeInvalidArg;
    sprintf(GenAlgErr->_msg, "'child' is invalid (0<=%d<=%d)",
        iChild, GAGetNbAdns(that));
    PBErrCatch(GenAlgErr);
}
#endif
// Get the parents and child
GenAlgAdn* parentA = GAAdn(that, parents[0]);
GenAlgAdn* parentB = GAAdn(that, parents[1]);
GenAlgAdn* child = GAAdn(that, iChild);
// For each gene of the adn for floating point value
for (int iGene = GAGetLengthAdnFloat(that); iGene--;) {
    // Get the gene from one parent or the other with equal probability
    if (rnd() < 0.5) {
        VecSet(child->_adnF, iGene, VecGet(parentA->_adnF, iGene));
        VecSet(child->_deltaAdnF, iGene,
            VecGet(parentA->_deltaAdnF, iGene));
    } else {
        VecSet(child->_adnF, iGene, VecGet(parentB->_adnF, iGene));
        VecSet(child->_deltaAdnF, iGene,
            VecGet(parentB->_deltaAdnF, iGene));
    }
}
// For each gene of the adn for int value
for (int iGene = GAGetLengthAdnInt(that); iGene--;) {
    // Get the gene from one parent or the other with equal probability
    if (rnd() < 0.5)
        VecSet(child->_adnI, iGene, VecGet(parentA->_adnI, iGene));
    else
        VecSet(child->_adnI, iGene, VecGet(parentB->_adnI, iGene));
}
// Reset the age of the child
child->_age = 1;
// Set the id of the child
child->_id = (that->_nextId)++;
}

// Mute the genes of the entity at rank 'iChild'
// The probability of mutation for one gene is equal to
// 'rankChild'/'that'->_nbEntities
// The amplitude of the mutation
// is equal to (max-min).(gauss(0.0, 1.0)+deltaAdn).ln('parents[0]'.age)
void GAMute(GenAlg* const that, const int* const parents,
    const int iChild) {
#ifdef BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
    if (parents == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'parents' is null");
        PBErrCatch(GenAlgErr);
    }
    if (iChild < 0 || iChild >= GAGetNbAdns(that)) {
        GenAlgErr->_type = PBErrTypeInvalidArg;
        sprintf(GenAlgErr->_msg, "'child' is invalid (0<=%d<=%d)",

```

```

        iChild, GAGetNbAdns(that));
    PBErrCatch(GenAlgErr);
}
#endif
// Get the first parent and child
GenAlgAdn* parentA = GAAdn(that, parents[0]);
GenAlgAdn* child = GAAdn(that, iChild);
// Get the proba amplitude of mutation
float probMute = ((float)iChild) / ((float)GAGetNbAdns(that));
float amp = 1.0 - 1.0 / sqrt((float)(parentA->_age + 1));
// For each gene of the adn for floating point value
for (int iGene = GAGetLengthAdnFloat(that); iGene--;) {
    // If this gene mutes
    if (rnd() < probMute) {
        // Get the bounds
        const VecFloat2D* const bounds = GABoundsAdnFloat(that, iGene);
        // Declare a variable to memorize the previous value of the gene
        float prevVal = GAAdnGetGeneF(child, iGene);
        // Apply the mutation
        GAAdnSetGeneF(child, iGene, GAAdnGetGeneF(child, iGene) +
            (VecGet(bounds, 1) - VecGet(bounds, 0)) * amp *
            (rnd() - 0.5 + GAAdnGetDeltaGeneF(child, iGene)));
        // Keep the gene value in bounds
        while (GAAdnGetGeneF(child, iGene) < VecGet(bounds, 0) ||
            GAAdnGetGeneF(child, iGene) > VecGet(bounds, 1)) {
            if (GAAdnGetGeneF(child, iGene) > VecGet(bounds, 1))
                GAAdnSetGeneF(child, iGene,
                    2.0 * VecGet(bounds, 1) - GAAdnGetGeneF(child, iGene));
            else if (GAAdnGetGeneF(child, iGene) < VecGet(bounds, 0))
                GAAdnSetGeneF(child, iGene,
                    2.0 * VecGet(bounds, 0) - GAAdnGetGeneF(child, iGene));
        }
        // Update the deltaAdn
        GAAdnSetDeltaGeneF(child, iGene,
            GAAdnGetGeneF(child, iGene) - prevVal);
    }
}
// For each gene of the adn for int value
for (int iGene = GAGetLengthAdnInt(that); iGene--;) {
    // If this gene mutes
    if (rnd() < probMute) {
        // Get the bounds
        const VecShort2D* const boundsI = GABoundsAdnInt(that, iGene);
        VecFloat2D bounds = VecShortToFloat2D(boundsI);
        // Apply the mutation (as it is int value, ensure the amplitude
        // is big enough to have an effect
        float ampI = MIN(2.0,
            (float)(VecGet(&bounds, 1) - VecGet(&bounds, 0)) * amp);
        GAAdnSetGeneI(child, iGene, GAAdnGetGeneI(child, iGene) +
            (short)round(ampI * (rnd() - 0.5)));
        // Keep the gene value in bounds
        while (GAAdnGetGeneI(child, iGene) < VecGet(&bounds, 0) ||
            GAAdnGetGeneI(child, iGene) > VecGet(&bounds, 1)) {
            if (GAAdnGetGeneI(child, iGene) > VecGet(&bounds, 1))
                GAAdnSetGeneI(child, iGene,
                    2 * VecGet(&bounds, 1) - GAAdnGetGeneI(child, iGene));
            else if (GAAdnGetGeneI(child, iGene) < VecGet(&bounds, 0))
                GAAdnSetGeneI(child, iGene,
                    2 * VecGet(&bounds, 0) - GAAdnGetGeneI(child, iGene));
        }
    }
}
}

```

```

}

// Print the information about the GenAlg 'that' on the stream 'stream'
void GAPrintln(const GenAlg* const that, FILE* const stream) {
#ifdef BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
    if (stream == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'stream' is null");
        PBErrCatch(GenAlgErr);
    }
#endif
    fprintf(stream, "epoch:%lu\n", GAGetCurEpoch(that));
    fprintf(stream, "%d entities, %d elites\n", GAGetNbAdns(that),
        GAGetNbElites(that));
    GSetIterBackward iter = GSetIterBackwardCreateStatic(GAAdns(that));
    int iEnt = 0;
    do {
        GenAlgAdn* ent = GSetIterGet(&iter);
        fprintf(stream, "%d value:%f ", iEnt,
            GSetIterGetElem(&iter)->_sortVal);
        if (iEnt < GAGetNbElites(that))
            fprintf(stream, "elite ");
        GAAdnPrintln(ent, stream);
        ++iEnt;
    } while (GSetIterStep(&iter));
}

// Update the norm of the range value for adans of the GenAlg 'that'
void GAUpdateNormRange(GenAlg* const that) {
#ifdef BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
#endif
    // If there are float adn
    if (GAGetLengthAdnFloat(that) > 0) {
        // Declare a vector to memorize the ranges in float gene values
        VecFloat* range = VecFloatCreate(GAGetLengthAdnFloat(that));
        // Calculate the ranges in gene values
        for (int iGene = GAGetLengthAdnFloat(that); iGene--;)
            VecSet(range, iGene,
                VecGet(GABoundsAdnFloat(that, iGene), 1) -
                VecGet(GABoundsAdnFloat(that, iGene), 0));
        // Calculate the norm of the range
        that->_normRangeFloat = VecNorm(range);
        // Free memory
        VecFree(&range);
    }

    // If there are int adn
    if (GAGetLengthAdnInt(that) > 0) {
        // Declare a vector to memorize the ranges in int gene values
        VecFloat* range = VecFloatCreate(GAGetLengthAdnInt(that));
        // Calculate the ranges in gene values
        for (int iGene = GAGetLengthAdnInt(that); iGene--;)

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        VecSet(range, iGene,
            VecGet(GABoundsAdnInt(that, iGene), 1) -
            VecGet(GABoundsAdnInt(that, iGene), 0));
    // Calculate the norm of the range
    that->_normRangeInt = VecNorm(range);
    // Free memory
    VecFree(&range);
}
}

// Get the diversity value of 'adnA' against 'adnB'
// The diversity is equal to
float GAAdnGetDiversity(const GenAlgAdn* const adnA,
    const GenAlgAdn* const adnB, const GenAlg* const ga) {
#ifdef BUILDMODE == 0
    if (adnA == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'adnA' is null");
        PBErrCatch(GenAlgErr);
    }
    if (adnB == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'adnB' is null");
        PBErrCatch(GenAlgErr);
    }
#endif
    // Declare a variable to memorize the result
    float diversity = 0.0;
    // If there are adn for floating point values
    if (GAAdnAdnF(adnA) != NULL && GAAdnAdnF(adnB) != NULL) {
        // Get the difference in adn with the first entity
        VecFloat* diff =
            VecGetOp(GAAdnAdnF(adnA), 1.0, GAAdnAdnF(adnB), -1.0);
        // Calculate the diversity
        diversity += VecNorm(diff) / ga->_normRangeFloat;
        // Free memory
        VecFree(&diff);
    }
    // If there are adn for int values
    if (GAAdnAdnI(adnA) != NULL && GAAdnAdnI(adnB) != NULL) {
        // Get the difference in adn with the first entity
        VecShort* diffI =
            VecGetOp(GAAdnAdnI(adnA), 1, GAAdnAdnI(adnB), -1);
        VecFloat* diff = VecShortToFloat(diffI);
        // Calculate the diversity
        diversity += VecNorm(diff) / ga->_normRangeInt;
        // Free memory
        VecFree(&diffI);
        VecFree(&diff);
    }
    // Correct diversity if there was both float and int adns
    if (GAAdnAdnF(adnA) != NULL && GAAdnAdnF(adnB) != NULL &&
        GAAdnAdnI(adnA) != NULL && GAAdnAdnI(adnB) != NULL)
        diversity /= 2.0;
    // Return the result
    return diversity;
}

// Get the average diversity of current entities of the GenAlg 'that'
// The return value is in [0.0, 1.0]
// 0.0 means all the elite entities have exactly the same adns

```

```

// 1.0 means all the elite entities except the first one have adns
// as different compare to the first one's adn as possible given the
// range of adn values
float GAGetDiversity(const GenAlg* const that) {
#ifdef BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
#endif
    // Declare a variable for calculation of the average of diversities
    float sumDiversity = 0.0;
    // For each elite entity except the first one
    for (int iEnt = 1; iEnt < GAGetNbElites(that); ++iEnt) {
        // Sum the diversity of this entity against the first one
        // for float values
        sumDiversity +=
            GAAdnGetDiversity(GAAdn(that, 0), GAAdn(that, iEnt), that);
    }
    // Calculate the average diversity
    float diversity = sumDiversity / (float)(GAGetNbElites(that) - 1);
    // Return the result
    return diversity;
}

// Function which return the JSON encoding of 'that'
JSONNode* GAAdnEncodeAsJSON(const GenAlgAdn* const that,
    const float elo) {
#ifdef BUILDMODE == 0
    if (that == NULL) {
        PBMathErr->_type = PBErrTypeNullPointer;
        sprintf(PBMathErr->_msg, "'that' is null");
        PBErrCatch(PBMathErr);
    }
#endif
    // Create the JSON structure
    JSONNode* json = JSONCreate();
    // Declare a buffer to convert value into string
    char val[100];
    // Encode the id
    sprintf(val, "%lu", that->_id);
    JSONAddProp(json, "_id", val);
    // Encode the age
    sprintf(val, "%lu", that->_age);
    JSONAddProp(json, "_age", val);
    // Encode the elo
    sprintf(val, "%f", elo);
    JSONAddProp(json, "_elo", val);
    // Encode the genes
    if (that->_adnF != NULL) {
        JSONAddProp(json, "_adnF", VecEncodeAsJSON(that->_adnF));
        JSONAddProp(json, "_deltaAdnF", VecEncodeAsJSON(that->_deltaAdnF));
    }
    if (that->_adnI != NULL)
        JSONAddProp(json, "_adnI", VecEncodeAsJSON(that->_adnI));
    // Return the created JSON
    return json;
}

// Function which return the JSON encoding of 'that'
JSONNode* GAEncodeAsJSON(const GenAlg* const that) {

```

```

#if BUILDMODE == 0
    if (that == NULL) {
        PBMathErr->_type = PBErrTypeNullPointer;
        sprintf(PBMathErr->_msg, "'that' is null");
        PBErrCatch(PBMathErr);
    }
#endif
    // Create the JSON structure
    JSONNode* json = JSONCreate();
    // Declare a buffer to convert value into string
    char val[100];
    // Encode the diversity threshold
    sprintf(val, "%f", GAGetDiversityThreshold(that));
    JSONAddProp(json, "_diversityThreshold", val);
    // Encode the nb adns
    sprintf(val, "%d", GAGetNbAdns(that));
    JSONAddProp(json, "_nbAdns", val);
    // Encode the nb elites
    sprintf(val, "%d", GAGetNbElites(that));
    JSONAddProp(json, "_nbElites", val);
    // Encode the length adn float
    sprintf(val, "%d", GAGetLengthAdnFloat(that));
    JSONAddProp(json, "_lengthAdnF", val);
    // Encode the length adn int
    sprintf(val, "%d", GAGetLengthAdnInt(that));
    JSONAddProp(json, "_lengthAdnI", val);
    // Encode the epoch
    sprintf(val, "%lu", GAGetCurEpoch(that));
    JSONAddProp(json, "_curEpoch", val);
    // Encode the next id
    sprintf(val, "%lu", that->nextId);
    JSONAddProp(json, "_nextId", val);
    // Encode the bounds
    JSONArrayStruct setBoundFloat = JSONArrayStructCreateStatic();
    if (GAGetLengthAdnFloat(that) > 0) {
        for (int iBound = 0; iBound < GAGetLengthAdnFloat(that); ++iBound)
            JSONArrayStructAdd(&setBoundFloat,
                VecEncodeAsJSON((VecFloat*)GABoundsAdnFloat(that, iBound)));
        JSONAddProp(json, "_boundFloat", &setBoundFloat);
    }
    JSONArrayStruct setBoundInt = JSONArrayStructCreateStatic();
    if (GAGetLengthAdnInt(that) > 0) {
        for (int iBound = 0; iBound < GAGetLengthAdnInt(that); ++iBound)
            JSONArrayStructAdd(&setBoundInt,
                VecEncodeAsJSON((VecShort*)GABoundsAdnInt(that, iBound)));
        JSONAddProp(json, "_boundInt", &setBoundInt);
    }
    // Save the adns
    JSONArrayStruct setAdn = JSONArrayStructCreateStatic();
    for (int iEnt = 0; iEnt < GAGetNbAdns(that); ++iEnt) {
        GenAlgAdn* ent = GSetElemData(GSetElement(GAAdns(that), iEnt));
        float sortVal = GSetElemGetSortVal(GSetElement(GAAdns(that), iEnt));
        JSONArrayStructAdd(&setAdn, GAAdnEncodeAsJSON(ent, sortVal));
    }
    JSONAddProp(json, "_adns", &setAdn);
    // Free memory
    JSONArrayStructFlush(&setBoundFloat);
    JSONArrayStructFlush(&setBoundInt);
    JSONArrayStructFlush(&setAdn);
    // Return the created JSON
    return json;
}

```

```

// Function which decode from JSON encoding 'json' to 'that'
bool GAAdnDecodeAsJSON(GenAlgAdn** that, const JSONNode* const json) {
#ifdef BUILDMODE == 0
    if (that == NULL) {
        PBMathErr->_type = PBErrTypeNullPointer;
        sprintf(PBMathErr->_msg, "'that' is null");
        PBErrCatch(PBMathErr);
    }
    if (json == NULL) {
        PBMathErr->_type = PBErrTypeNullPointer;
        sprintf(PBMathErr->_msg, "'json' is null");
        PBErrCatch(PBMathErr);
    }
#endif
    // If 'that' is already allocated
    if (*that != NULL)
        // Free memory
        GenAlgAdnFree(that);
    // Get the id from the JSON
    JSONNode* prop = JSONProperty(json, "_id");
    if (prop == NULL) {
        return false;
    }
    int id = strtoul(JSONLabel(JSONValue(prop, 0)), NULL, 10);
    // Get the lengthAdnF from the JSON
    int lengthAdnF = 0;
    prop = JSONProperty(json, "_adnF");
    if (prop != NULL) {
        JSONNode* subprop = JSONProperty(prop, "_dim");
        lengthAdnF = atoi(JSONLabel(JSONValue(subprop, 0)));
    }
    // Get the lengthAdnI from the JSON
    int lengthAdnI = 0;
    prop = JSONProperty(json, "_adnI");
    if (prop != NULL) {
        JSONNode* subprop = JSONProperty(prop, "_dim");
        lengthAdnI = atoi(JSONLabel(JSONValue(subprop, 0)));
    }
    // Allocate memory
    *that = GenAlgAdnCreate(id, lengthAdnF, lengthAdnI);
    // Get the age from the JSON
    prop = JSONProperty(json, "_age");
    if (prop == NULL) {
        return false;
    }
    (*that)->_age = strtoul(JSONLabel(JSONValue(prop, 0)), NULL, 10);
    // Get the adnF from the JSON
    prop = JSONProperty(json, "_adnF");
    if (prop != NULL) {
        if (!VecDecodeAsJSON(&((*that)->_adnF), prop)) {
            return false;
        }
        prop = JSONProperty(json, "_deltaAdnF");
        if (prop == NULL) {
            return false;
        }
        if (!VecDecodeAsJSON(&((*that)->_deltaAdnF), prop)) {
            return false;
        }
    }
    // Get the adnI from the JSON

```

```

    prop = JSONProperty(json, "_adnI");
    if (prop != NULL)
        if (!VecDecodeAsJSON(&((*that)->_adnI), prop)) {
            return false;
        }
    // Return the success code
    return true;
}

// Function which decode from JSON encoding 'json' to 'that'
bool GADecodeAsJSON(GenAlg** that, const JSONNode* const json) {
#ifdef BUILDMODE == 0
    if (that == NULL) {
        PBMathErr->_type = PErrTypeNullPointer;
        sprintf(PBMathErr->_msg, "'that' is null");
        PBErreCatch(PBMathErr);
    }
    if (json == NULL) {
        PBMathErr->_type = PErrTypeNullPointer;
        sprintf(PBMathErr->_msg, "'json' is null");
        PBErreCatch(PBMathErr);
    }
#endif
    // If 'that' is already allocated
    if (*that != NULL)
        // Free memory
        GenAlgFree(that);
    // Decode the nb adns
    JSONNode* prop = JSONProperty(json, "_nbAdns");
    if (prop == NULL) {
        return false;
    }
    int nbAdns = atoi(JSONLabel(JSONValue(prop, 0)));
    // Decode the nb elites
    prop = JSONProperty(json, "_nbElites");
    if (prop == NULL) {
        return false;
    }
    int nbElites = atoi(JSONLabel(JSONValue(prop, 0)));
    // Decode the length adn float
    prop = JSONProperty(json, "_lengthAdnF");
    if (prop == NULL) {
        return false;
    }
    int lengthAdnF = atoi(JSONLabel(JSONValue(prop, 0)));
    // Decode the length adn int
    prop = JSONProperty(json, "_lengthAdnI");
    if (prop == NULL) {
        return false;
    }
    int lengthAdnI = atoi(JSONLabel(JSONValue(prop, 0)));
    // Allocate memory
    *that = GenAlgCreate(nbAdns, nbElites, lengthAdnF, lengthAdnI);
    // Decode the diversity threshold
    prop = JSONProperty(json, "_diversityThreshold");
    if (prop == NULL) {
        return false;
    }
    (*that)->_diversityThreshold = atof(JSONLabel(JSONValue(prop, 0)));
    // Decode the epoch
    prop = JSONProperty(json, "_curEpoch");
    if (prop == NULL) {

```



```

    return false;
}
(*that)->_curEpoch =
    strtoul(JSONLabel(JSONValue(prop, 0)), NULL, 10);
// Decode the next id
prop = JSONProperty(json, "_nextId");
if (prop == NULL) {
    return false;
}
(*that)->_nextId = strtoul(JSONLabel(JSONValue(prop, 0)), NULL, 10);
// Decode the bounds
prop = JSONProperty(json, "_boundFloat");
if (prop != NULL) {
    if (JSONGetNbValue(prop) != GAGetLengthAdnFloat(*that))
        return false;
    for (int iBound = 0; iBound < GAGetLengthAdnFloat(*that); ++iBound) {
        JSONNode* val = JSONValue(prop, iBound);
        VecFloat2D* b = NULL;
        if (!VecDecodeAsJSON((VecFloat**) &b, val)) {
            return false;
        }
        GASetBoundsAdnFloat(*that, iBound, b);
        VecFree((VecFloat**) &b);
    }
}
prop = JSONProperty(json, "_boundInt");
if (prop != NULL) {
    if (JSONGetNbValue(prop) != GAGetLengthAdnInt(*that))
        return false;
    for (int iBound = 0; iBound < GAGetLengthAdnInt(*that); ++iBound) {
        JSONNode* val = JSONValue(prop, iBound);
        VecShort2D* b = NULL;
        if (!VecDecodeAsJSON((VecShort**) &b, val)) {
            return false;
        }
        GASetBoundsAdnInt(*that, iBound, b);
        VecFree((VecShort**) &b);
    }
}
// Upadte the norm of the range values
GAUpdateNormRange(*that);
// Decode the adns
prop = JSONProperty(json, "_adns");
if (prop == NULL) {
    return false;
}
if (JSONGetNbValue(prop) != GAGetNbAdns(*that))
    return false;
for (int iEnt = 0; iEnt < GAGetNbAdns(*that); ++iEnt) {
    JSONNode* val = JSONValue(prop, iEnt);
    GenAlgAdn* data = GSetElemData(GSetElement(GAAdns(*that), iEnt));
    if (!GAAdnDecodeAsJSON(&data, val)) {
        return false;
    }
}
// Return the success code
return true;
}

// Load the GenAlg 'that' from the stream 'stream'
// If the GenAlg is already allocated, it is freed before loading
// Return true in case of success, else false

```

```

bool GALoad(GenAlg** that, FILE* const stream) {
#ifdef BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
    if (stream == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'stream' is null");
        PBErrCatch(GenAlgErr);
    }
#endif
    // Declare a json to load the encoded data
    JSONNode* json = JSONCreate();
    // Load the whole encoded data
    if (!JSONLoad(json, stream)) {
        return false;
    }
    // Decode the data from the JSON
    if (!GADecodeAsJSON(that, json)) {
        return false;
    }
    // Free the memory used by the JSON
    JSONFree(&json);
    // Return the success code
    return true;
}

// Save the GenAlg 'that' to the stream 'stream'
// If 'compact' equals true it saves in compact form, else it saves in
// readable form
// Return true in case of success, else false
bool GASave(const GenAlg* const that, FILE* const stream,
    const bool compact) {
#ifdef BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
    if (stream == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'stream' is null");
        PBErrCatch(GenAlgErr);
    }
#endif
    // Get the JSON encoding
    JSONNode* json = GAEncodeAsJSON(that);
    // Save the JSON
    if (!JSONSave(json, stream, compact)) {
        return false;
    }
    // Free memory
    JSONFree(&json);
    // Return success code
    return true;
}

```

## 3.2 genalg-inline.c

```
// ===== GENALG-INLINE.C =====

// ----- GenAlgAdn

// ===== Functions implementation =====

// Return the adn for floating point values of the GenAlgAdn 'that'
#if BUILDMODE != 0
inline
#endif
const VecFloat* GAAdnAdnF(const GenAlgAdn* const that) {
#if BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PErrCatch(GenAlgErr);
    }
#endif
    return that->_adnF;
}

// Return the delta of adn for floating point values of the
// GenAlgAdn 'that'
#if BUILDMODE != 0
inline
#endif
const VecFloat* GAAdnDeltaAdnF(const GenAlgAdn* const that) {
#if BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PErrCatch(GenAlgErr);
    }
#endif
    return that->_deltaAdnF;
}

// Return the adn for integer values of the GenAlgAdn 'that'
#if BUILDMODE != 0
inline
#endif
const VecShort* GAAdnAdnI(const GenAlgAdn* const that) {
#if BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PErrCatch(GenAlgErr);
    }
#endif
    return that->_adnI;
}

// Get the 'iGene'-th gene of the adn for floating point values of the
// GenAlgAdn 'that'
#if BUILDMODE != 0
inline
#endif
float GAAdnGetGeneF(const GenAlgAdn* const that, const int iGene) {
#if BUILDMODE == 0
    if (that == NULL) {
```

```

        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
#endif
    return VecGet(that->_adnF, iGene);
}

// Get the delta of the 'iGene'-th gene of the adn for floating point
// values of the GenAlgAdn 'that'
#if BUILDMODE != 0
inline
#endif
float GAAdnGetDeltaGeneF(const GenAlgAdn* const that, const int iGene) {
#if BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
#endif
    return VecGet(that->_deltaAdnF, iGene);
}

// Get the 'iGene'-th gene of the adn for int values of the
// GenAlgAdn 'that'
#if BUILDMODE != 0
inline
#endif
int GAAdnGetGeneI(const GenAlgAdn* const that, const int iGene) {
#if BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
#endif
    return VecGet(that->_adnI, iGene);
}

// Set the 'iGene'-th gene of the adn for floating point values of the
// GenAlgAdn 'that' to 'gene'
#if BUILDMODE != 0
inline
#endif
void GAAdnSetGeneF(GenAlgAdn* const that, const int iGene,
    const float gene) {
#if BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
#endif
    VecSet(that->_adnF, iGene, gene);
}

// Set the delta of the 'iGene'-th gene of the adn for floating point
// values of the GenAlgAdn 'that' to 'delta'
#if BUILDMODE != 0
inline
#endif

```

```

void GAAdnSetDeltaGeneF(GenAlgAdn* const that, const int iGene,
    const float delta) {
    #if BUILDMODE == 0
        if (that == NULL) {
            GenAlgErr->_type = PErrTypeNullPointer;
            sprintf(GenAlgErr->_msg, "'that' is null");
            PErrCatch(GenAlgErr);
        }
    #endif
    VecSet(that->_deltaAdnF, iGene, delta);
}

// Set the 'iGene'-th gene of the adn for int values of the
// GenAlgAdn 'that'to 'gene'
#if BUILDMODE != 0
    inline
#endif
void GAAdnSetGeneI(GenAlgAdn* const that, const int iGene,
    const short gene) {
    #if BUILDMODE == 0
        if (that == NULL) {
            GenAlgErr->_type = PErrTypeNullPointer;
            sprintf(GenAlgErr->_msg, "'that' is null");
            PErrCatch(GenAlgErr);
        }
    #endif
    VecSet(that->_adnI, iGene, gene);
}

// Get the id of the GenAlgAdn 'that'
#if BUILDMODE != 0
    inline
#endif
unsigned long int GAAdnGetId(const GenAlgAdn* const that) {
    #if BUILDMODE == 0
        if (that == NULL) {
            GenAlgErr->_type = PErrTypeNullPointer;
            sprintf(GenAlgErr->_msg, "'that' is null");
            PErrCatch(GenAlgErr);
        }
    #endif
    return that->_id;
}

// Get the age of the GenAlgAdn 'that'
#if BUILDMODE != 0
    inline
#endif
unsigned long int GAAdnGetAge(const GenAlgAdn* const that) {
    #if BUILDMODE == 0
        if (that == NULL) {
            GenAlgErr->_type = PErrTypeNullPointer;
            sprintf(GenAlgErr->_msg, "'that' is null");
            PErrCatch(GenAlgErr);
        }
    #endif
    return that->_age;
}

// Return true if the GenAlgAdn 'that' is new, i.e. is age equals 1
// Return false
#if BUILDMODE != 0

```

```

inline
#endif
bool GAAdnIsNew(const GenAlgAdn* const that) {
#if BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
#endif
    return (that->_age == 1);
}

// ----- GenAlg

// ===== Functions implementation =====

// Return the GSet of the GenAlg 'that'
#if BUILDMODE != 0
inline
#endif
GSet* GAAdns(const GenAlg* const that) {
#if BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
#endif
    return that->_adns;
}

// Return the nb of entities of the GenAlg 'that'
#if BUILDMODE != 0
inline
#endif
int GAGetNbAdns(const GenAlg* const that) {
#if BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
#endif
    return GSetNbElem(that->_adns);
}

// Return the nb of elites of the GenAlg 'that'
#if BUILDMODE != 0
inline
#endif
int GAGetNbElites(const GenAlg* const that) {
#if BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
#endif
    return that->_nbElites;
}

```

```

// Return the current epoch of the GenAlg 'that'
#if BUILDMODE != 0
inline
#endif
unsigned long int GAGetCurEpoch(const GenAlg* const that) {
#if BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
#endif
    return that->_curEpoch;
}

// Get the length of adn for floating point value
#if BUILDMODE != 0
inline
#endif
int GAGetLengthAdnFloat(const GenAlg* const that) {
#if BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
#endif
    return that->_lengthAdnF;
}

// Get the length of adn for integer value
#if BUILDMODE != 0
inline
#endif
int GAGetLengthAdnInt(const GenAlg* const that) {
#if BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
#endif
    return that->_lengthAdnI;
}

// Set the bounds for the 'iGene'-th gene of adn for floating point
// values to a copy of 'bounds'
#if BUILDMODE != 0
inline
#endif
void GASetBoundsAdnFloat(GenAlg* const that, const int iGene,
    const VecFloat2D* const bounds) {
#if BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
    if (bounds == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'bounds' is null");
    }
#endif
}

```

```

        PBErCatch(GenAlgErr);
    }
    if (iGene < 0 || iGene >= that->_lengthAdnF) {
        GenAlgErr->_type = PBErTypeInvalidArg;
        sprintf(GenAlgErr->_msg, "'iGene' is invalid (0<=%d<%d)",
            iGene, that->_lengthAdnF);
        PBErCatch(GenAlgErr);
    }
#endif
    VecCopy(that->_boundsF + iGene, bounds);
    GAUpdateNormRange(that);
}

// Set the bounds for the 'iGene'-th gene of adn for integer values
// to a copy of 'bounds'
#if BUILDMODE != 0
inline
#endif
void GASetBoundsAdnInt(GenAlg* const that, const int iGene,
    const VecShort2D* const bounds) {
    #if BUILDMODE == 0
        if (that == NULL) {
            GenAlgErr->_type = PBErTypeNullPointer;
            sprintf(GenAlgErr->_msg, "'that' is null");
            PBErCatch(GenAlgErr);
        }
        if (bounds == NULL) {
            GenAlgErr->_type = PBErTypeNullPointer;
            sprintf(GenAlgErr->_msg, "'bounds' is null");
            PBErCatch(GenAlgErr);
        }
        if (iGene < 0 || iGene >= that->_lengthAdnI) {
            GenAlgErr->_type = PBErTypeInvalidArg;
            sprintf(GenAlgErr->_msg, "'iGene' is invalid (0<=%d<%d)",
                iGene, that->_lengthAdnI);
            PBErCatch(GenAlgErr);
        }
    #endif
    VecCopy(that->_boundsI + iGene, bounds);
    GAUpdateNormRange(that);
}

// Get the bounds for the 'iGene'-th gene of adn for floating point
// values
#if BUILDMODE != 0
inline
#endif
const VecFloat2D* GABoundsAdnFloat(const GenAlg* const that,
    const int iGene) {
    #if BUILDMODE == 0
        if (that == NULL) {
            GenAlgErr->_type = PBErTypeNullPointer;
            sprintf(GenAlgErr->_msg, "'that' is null");
            PBErCatch(GenAlgErr);
        }
        if (iGene < 0 || iGene >= that->_lengthAdnF) {
            GenAlgErr->_type = PBErTypeInvalidArg;
            sprintf(GenAlgErr->_msg, "'iGene' is invalid (0<=%d<%d)",
                iGene, that->_lengthAdnF);
            PBErCatch(GenAlgErr);
        }
    #endif
}

```



```

    return that->_boundsF + iGene;
}

// Get the bounds for the 'iGene'-th gene of adn for integer values
#if BUILDMODE != 0
inline
#endif
const VecShort2D* GABoundsAdnInt(const GenAlg* const that,
    const int iGene) {
#if BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
    if (iGene < 0 || iGene >= that->_lengthAdnI) {
        GenAlgErr->_type = PBErrTypeInvalidArg;
        sprintf(GenAlgErr->_msg, "'iGene' is invalid (0<=%d<%d)",
            iGene, that->_lengthAdnI);
        PBErrCatch(GenAlgErr);
    }
#endif
    return that->_boundsI + iGene;
}

// Get the GenAlgAdn of the GenAlg 'that' currently at rank 'iRank'
// (0 is the best adn)
#if BUILDMODE != 0
inline
#endif
GenAlgAdn* GAAdn(const GenAlg* const that, const int iRank) {
#if BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
    if (iRank < 0 || iRank >= GAGetNbAdns(that)) {
        GenAlgErr->_type = PBErrTypeInvalidArg;
        sprintf(GenAlgErr->_msg, "'iRank' is invalid (0<=%d<%d)",
            iRank, GAGetNbAdns(that));
        PBErrCatch(GenAlgErr);
    }
#endif
    return (GenAlgAdn*)GSetGet(that->_adns,
        GSetNbElem(that->_adns) - iRank - 1);
}

// Set the value of the GenAlgAdn 'adn' of the GenAlg 'that' to 'val'
#if BUILDMODE != 0
inline
#endif
void GASetAdnValue(GenAlg* const that, const GenAlgAdn* const adn,
    const float val) {
#if BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
    if (adn == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
    }
#endif
}

```

```

        sprintf(GenAlgErr->_msg, "'adn' is null");
        PBErrCatch(GenAlgErr);
    }
#endif
    GSetElemSetSortVal((GSetElem*)GSetFirstElem(GAAdns(that), adn), val);
}

// Return the diversity threshold of the GenAlg 'that'
#if BUILDMODE != 0
inline
#endif
float GAGetDiversityThreshold(const GenAlg* const that) {
    if (BUILDMODE == 0
        if (that == NULL) {
            GenAlgErr->_type = PBErrTypeNullPointer;
            sprintf(GenAlgErr->_msg, "'that' is null");
            PBErrCatch(GenAlgErr);
        }
    )
    return that->_diversityThreshold;
}

// Set the diversity threshold of the GenAlg 'that' to 'div'
#if BUILDMODE != 0
inline
#endif
void GASetDiversityThreshold(GenAlg* const that, const float div) {
    if (BUILDMODE == 0
        if (that == NULL) {
            GenAlgErr->_type = PBErrTypeNullPointer;
            sprintf(GenAlgErr->_msg, "'that' is null");
            PBErrCatch(GenAlgErr);
        }
    )
    that->_diversityThreshold = div;
}

```

## 4 Makefile

```

# Build mode
# 0: development (max safety, no optimisation)
# 1: release (min safety, optimisation)
# 2: fast and furious (no safety, optimisation)
BUILD_MODE?=1

all: main

# Makefile definitions
MAKEFILE_INC=../PBMake/Makefile.inc
include $(MAKEFILE_INC)

# Rules to make the executable
repo=genalg
$(repo)_EXENAME: \
$(repo)_EXENAME.o \
$(repo)_EXE_DEP \
$(repo)_DEP
$(COMPILER) 'echo "$(repo)_EXE_DEP" "$(repo)_EXENAME.o" | tr ' ' '\n' | sort -u' $(LINK_ARG) $(repo)_LINK_ARG

```

```

$$($(repo)_EXENAME).o: \
$$($(repo)_DIR)/$$($(repo)_EXENAME).c \
$$($(repo)_INC_H_EXE) \
$$($(repo)_EXE_DEP)
$(COMPILER) $(BUILD_ARG) $$($(repo)_BUILD_ARG) 'echo "$$(repo)_INC_DIR" | tr ' ' '\n' | sort -u' -c $$($(repo)_DIR)/

```

## 5 Unit tests

```

#include <stdlib.h>
#include <stdio.h>
#include <time.h>
#include <string.h>
#include <time.h>
#include <unistd.h>
#include <sys/time.h>
#include "genalg.h"

#define RANDOMSEED 2

void UnitTestGenAlgAdnCreateFree() {
    unsigned long int id = 1;
    int lengthAdnF = 2;
    int lengthAdnI = 3;
    GenAlgAdn* ent = GenAlgAdnCreate(id, lengthAdnF, lengthAdnI);
    if (ent->_age != 1 ||
        ent->_id != id ||
        VecGetDim(ent->_adnF) != lengthAdnF ||
        VecGetDim(ent->_deltaAdnF) != lengthAdnF ||
        VecGetDim(ent->_adnI) != lengthAdnI) {
        GenAlgErr->_type = PBErrTypeUnitTestFailed;
        sprintf(GenAlgErr->_msg, "GenAlgAdnCreate failed");
        PBErrCatch(GenAlgErr);
    }
    GenAlgAdnFree(&ent);
    if (ent != NULL) {
        GenAlgErr->_type = PBErrTypeUnitTestFailed;
        sprintf(GenAlgErr->_msg, "GenAlgAdnFree failed");
        PBErrCatch(GenAlgErr);
    }
    printf("UnitTestGenAlgAdnCreateFree OK\n");
}

void UnitTestGenAlgAdnGetSet() {
    unsigned long int id = 1;
    int lengthAdnF = 2;
    int lengthAdnI = 3;
    GenAlgAdn* ent = GenAlgAdnCreate(id, lengthAdnF, lengthAdnI);
    if (GAAdnAdnF(ent) != ent->_adnF) {
        GenAlgErr->_type = PBErrTypeUnitTestFailed;
        sprintf(GenAlgErr->_msg, "GAAdnAdnF failed");
        PBErrCatch(GenAlgErr);
    }
    if (GAAdnDeltaAdnF(ent) != ent->_deltaAdnF) {
        GenAlgErr->_type = PBErrTypeUnitTestFailed;
        sprintf(GenAlgErr->_msg, "GAAdnDeltaAdnF failed");
        PBErrCatch(GenAlgErr);
    }
}

```

```

if (GAAdnAdnI(ent) != ent->_adnI) {
    GenAlgErr->_type = PBErrTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GAAdnAdnI failed");
    PBErrCatch(GenAlgErr);
}
GAAdnSetGeneF(ent, 0, 1.0);
if (ISEQUALF(VecGet(ent->_adnF, 0), 1.0) == false) {
    GenAlgErr->_type = PBErrTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GAAdnSetGeneF failed");
    PBErrCatch(GenAlgErr);
}
if (ISEQUALF(GAAdnGetGeneF(ent, 0), 1.0) == false) {
    GenAlgErr->_type = PBErrTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GAAdnGetGeneF failed");
    PBErrCatch(GenAlgErr);
}
GAAdnSetDeltaGeneF(ent, 0, 2.0);
if (ISEQUALF(VecGet(ent->_deltaAdnF, 0), 2.0) == false) {
    GenAlgErr->_type = PBErrTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GAAdnSetDeltaGeneF failed");
    PBErrCatch(GenAlgErr);
}
if (ISEQUALF(GAAdnGetDeltaGeneF(ent, 0), 2.0) == false) {
    GenAlgErr->_type = PBErrTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GAAdnGetDeltaGeneF failed");
    PBErrCatch(GenAlgErr);
}
GAAdnSetGeneI(ent, 0, 3);
if (VecGet(ent->_adnI, 0) != 3) {
    GenAlgErr->_type = PBErrTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GAAdnSetGeneI failed");
    PBErrCatch(GenAlgErr);
}
if (GAAdnGetGeneI(ent, 0) != 3) {
    GenAlgErr->_type = PBErrTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GAAdnGetGeneI failed");
    PBErrCatch(GenAlgErr);
}
if (GAAdnGetAge(ent) != 1) {
    GenAlgErr->_type = PBErrTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GAAdnGetAge failed");
    PBErrCatch(GenAlgErr);
}
if (GAAdnGetId(ent) != id) {
    GenAlgErr->_type = PBErrTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GAAdnGetId failed");
    PBErrCatch(GenAlgErr);
}
if (GAAdnIsNew(ent) != true) {
    GenAlgErr->_type = PBErrTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GAAdnIsNew failed");
    PBErrCatch(GenAlgErr);
}
ent->_age = 2;
if (GAAdnIsNew(ent) != false) {
    GenAlgErr->_type = PBErrTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GAAdnIsNew failed");
    PBErrCatch(GenAlgErr);
}
GenAlgAdnFree(&ent);
printf("UnitTestGenAlgAdnGetSet OK\n");
}

```

```

void UnitTestGenAlgAdnInit() {
    srandom(5);
    unsigned long int id = 1;
    int lengthAdnF = 2;
    int lengthAdnI = 2;
    GenAlgAdn* ent = GenAlgAdnCreate(id, lengthAdnF, lengthAdnI);
    GenAlg* ga = GenAlgCreate(GENALG_NBENTITIES, GENALG_NBELITES,
        lengthAdnF, lengthAdnI);
    VecFloat2D boundsF = VecFloatCreateStatic2D();
    VecShort2D boundsI = VecShortCreateStatic2D();
    VecSet(&boundsF, 0, -1.0); VecSet(&boundsF, 1, 1.0);
    VecSet(&boundsI, 0, 1); VecSet(&boundsI, 1, 10);
    GASetBoundsAdnFloat(ga, 0, &boundsF);
    GASetBoundsAdnFloat(ga, 1, &boundsF);
    GASetBoundsAdnInt(ga, 0, &boundsI);
    GASetBoundsAdnInt(ga, 1, &boundsI);
    GAAdnInit(ent, ga);
    if (ISEQUALF(VecGet(ent->_adnF, 0), -0.907064) == false ||
        ISEQUALF(VecGet(ent->_adnF, 1), -0.450509) == false ||
        VecGet(ent->_adnI, 0) != 2 ||
        VecGet(ent->_adnI, 1) != 10) {
        GenAlgErr->_type = PBErrTypeUnitTestFailed;
        sprintf(GenAlgErr->_msg, "GAAdnInit failed");
        PBErrCatch(GenAlgErr);
    }
    GenAlgFree(&ga);
    GenAlgAdnFree(&ent);
    printf("UnitTestGenAlgAdnInit OK\n");
}

void UnitTestGenAlgAdn() {
    UnitTestGenAlgAdnCreateFree();
    UnitTestGenAlgAdnGetSet();
    UnitTestGenAlgAdnInit();
    printf("UnitTestGenAlgAdn OK\n");
}

void UnitTestGenAlgCreateFree() {
    int lengthAdnF = 2;
    int lengthAdnI = 3;
    GenAlg* ga = GenAlgCreate(GENALG_NBENTITIES, GENALG_NBELITES,
        lengthAdnF, lengthAdnI);
    if (ga->_curEpoch != 0 ||
        ga->_nextId != GENALG_NBENTITIES ||
        ga->_nbElites != GENALG_NBELITES ||
        ga->_lengthAdnF != lengthAdnF ||
        ga->_lengthAdnI != lengthAdnI ||
        ISEQUALF(ga->_diversityThreshold,
            GENALG_DIVERSITYTHRESHOLD) == false ||
        GSetNbElem(GAAdns(ga)) != GENALG_NBENTITIES) {
        GenAlgErr->_type = PBErrTypeUnitTestFailed;
        sprintf(GenAlgErr->_msg, "GenAlgCreate failed");
        PBErrCatch(GenAlgErr);
    }
    GenAlgFree(&ga);
    if (ga != NULL) {
        GenAlgErr->_type = PBErrTypeUnitTestFailed;
        sprintf(GenAlgErr->_msg, "GenAlgFree failed");
        PBErrCatch(GenAlgErr);
    }
    printf("UnitTestGenAlgCreateFree OK\n");
}

```

```

}

void UnitTestGenAlgGetSet() {
    int lengthAdnF = 2;
    int lengthAdnI = 3;
    GenAlg* ga = GenAlgCreate(GENALG_NBENTITIES, GENALG_NBELITES,
        lengthAdnF, lengthAdnI);
    if (GAAdns(ga) != ga->_adns) {
        GenAlgErr->_type = PBErrTypeUnitTestFailed;
        sprintf(GenAlgErr->_msg, "GAElorank failed");
        PBErrCatch(GenAlgErr);
    }
    if (GAGetNbAdns(ga) != GENALG_NBENTITIES) {
        GenAlgErr->_type = PBErrTypeUnitTestFailed;
        sprintf(GenAlgErr->_msg, "GAGetNbAdns failed");
        PBErrCatch(GenAlgErr);
    }
    if (GAGetNbElites(ga) != GENALG_NBELITES) {
        GenAlgErr->_type = PBErrTypeUnitTestFailed;
        sprintf(GenAlgErr->_msg, "GAGetNbElites failed");
        PBErrCatch(GenAlgErr);
    }
    if (GAGetCurEpoch(ga) != 0) {
        GenAlgErr->_type = PBErrTypeUnitTestFailed;
        sprintf(GenAlgErr->_msg, "GAGetCurEpoch failed");
        PBErrCatch(GenAlgErr);
    }
    GASetNbEntities(ga, 10);
    if (GAGetNbAdns(ga) != 10 ||
        GAGetNbElites(ga) != 9 ||
        GSetNbElem(GAAdns(ga)) != 10) {
        GenAlgErr->_type = PBErrTypeUnitTestFailed;
        sprintf(GenAlgErr->_msg, "GASetNbEntities failed");
        PBErrCatch(GenAlgErr);
    }
    GASetNbElites(ga, 20);
    if (GAGetNbAdns(ga) != 21 ||
        GAGetNbElites(ga) != 20 ||
        GSetNbElem(GAAdns(ga)) != 21) {
        GenAlgErr->_type = PBErrTypeUnitTestFailed;
        sprintf(GenAlgErr->_msg, "GASetNbElites failed");
        PBErrCatch(GenAlgErr);
    }
    if (GAGetLengthAdnFloat(ga) != lengthAdnF) {
        GenAlgErr->_type = PBErrTypeUnitTestFailed;
        sprintf(GenAlgErr->_msg, "GAGetLengthAdnFloat failed");
        PBErrCatch(GenAlgErr);
    }
    if (GAGetLengthAdnInt(ga) != lengthAdnI) {
        GenAlgErr->_type = PBErrTypeUnitTestFailed;
        sprintf(GenAlgErr->_msg, "GAGetLengthAdnInt failed");
        PBErrCatch(GenAlgErr);
    }
    if (GABoundsAdnFloat(ga, 1) != ga->_boundsF + 1) {
        GenAlgErr->_type = PBErrTypeUnitTestFailed;
        sprintf(GenAlgErr->_msg, "GABoundsAdnFloat failed");
        PBErrCatch(GenAlgErr);
    }
    VecFloat2D boundsF = VecFloatCreateStatic2D();
    VecSet(&boundsF, 0, -1.0); VecSet(&boundsF, 1, 1.0);
    GASetBoundsAdnFloat(ga, 1, &boundsF);
    if (VecIsEqual(GABoundsAdnFloat(ga, 1), &boundsF) == false) {

```

```

    GenAlgErr->_type = PBErrTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GASetBoundsAdnFloat failed");
    PBErrCatch(GenAlgErr);
}
VecShort2D boundsS = VecShortCreateStatic2D();
VecSet(&boundsS, 0, -1); VecSet(&boundsS, 1, 1);
GASetBoundsAdnInt(ga, 1, &boundsS);
if (VecIsEqual(GABoundsAdnInt(ga, 1), &boundsS) == false) {
    GenAlgErr->_type = PBErrTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GASetBoundsAdnInt failed");
    PBErrCatch(GenAlgErr);
}
if (GABoundsAdnInt(ga, 1) != ga->_boundsI + 1) {
    GenAlgErr->_type = PBErrTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GABoundsAdnInt failed");
    PBErrCatch(GenAlgErr);
}
GASetAdnValue(ga, GAAdn(ga, 0), 1.0);
if (ISEQUALF(ga->_adns->_tail->_sortVal, 1.0) == false) {
    GenAlgErr->_type = PBErrTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GASetAdnValue failed");
    PBErrCatch(GenAlgErr);
}
if (ISEQUALF(GAGetDiversityThreshold(ga),
    ga->_diversityThreshold) == false) {
    GenAlgErr->_type = PBErrTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GAGetDiversityThreshold failed");
    PBErrCatch(GenAlgErr);
}
GASetDiversityThreshold(ga, 0.5);
if (ISEQUALF(GAGetDiversityThreshold(ga), 0.5) == false) {
    GenAlgErr->_type = PBErrTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GASetDiversityThreshold failed");
    PBErrCatch(GenAlgErr);
}
GenAlgFree(&ga);
printf("UnitTestGenAlgGetSet OK\n");
}

void UnitTestGenAlgInit() {
    srandom(5);
    int lengthAdnF = 2;
    int lengthAdnI = 2;
    GenAlg* ga = GenAlgCreate(GENALG_NBENTITIES, GENALG_NBELITES,
        lengthAdnF, lengthAdnI);
    VecFloat2D boundsF = VecFloatCreateStatic2D();
    VecShort2D boundsI = VecShortCreateStatic2D();
    VecSet(&boundsF, 0, -1.0); VecSet(&boundsF, 1, 1.0);
    VecSet(&boundsI, 0, 1); VecSet(&boundsI, 1, 10);
    GASetBoundsAdnFloat(ga, 0, &boundsF);
    GASetBoundsAdnFloat(ga, 1, &boundsF);
    GASetBoundsAdnInt(ga, 0, &boundsI);
    GASetBoundsAdnInt(ga, 1, &boundsI);
    GAINit(ga);
    GenAlgAdn* ent = (GenAlgAdn*)(GAAdns(ga)->_head->_data);
    if (ISEQUALF(VecGet(ent->_adnF, 0), -0.907064) == false ||
        ISEQUALF(VecGet(ent->_adnF, 1), -0.450509) == false ||
        VecGet(ent->_adnI, 0) != 2 ||
        VecGet(ent->_adnI, 1) != 10) {
        GenAlgErr->_type = PBErrTypeUnitTestFailed;
        sprintf(GenAlgErr->_msg, "GAINit failed");
        PBErrCatch(GenAlgErr);
    }
}

```

```

    }
    GenAlgFree(&ga);
    printf("UnitTestGenAlgInit OK\n");
}

void UnitTestGenAlgPrint() {
    srandom(5);
    int lengthAdnF = 2;
    int lengthAdnI = 2;
    GenAlg* ga = GenAlgCreate(3, 2, lengthAdnF, lengthAdnI);
    VecFloat2D boundsF = VecFloatCreateStatic2D();
    VecShort2D boundsI = VecShortCreateStatic2D();
    VecSet(&boundsF, 0, -1.0); VecSet(&boundsF, 1, 1.0);
    VecSet(&boundsI, 0, 1); VecSet(&boundsI, 1, 10);
    GASetBoundsAdnFloat(ga, 0, &boundsF);
    GASetBoundsAdnFloat(ga, 1, &boundsF);
    GASetBoundsAdnInt(ga, 0, &boundsI);
    GASetBoundsAdnInt(ga, 1, &boundsI);
    GAINit(ga);
    GAINit(ga);
    GAPrintln(ga, stdout);
    GenAlgFree(&ga);
    printf("UnitTestGenAlgInit OK\n");
}

void UnitTestGenAlgGetDiversity() {
    srandom(5);
    int lengthAdnF = 2;
    int lengthAdnI = 2;
    GenAlg* ga = GenAlgCreate(GENALG_NBENTITIES, GENALG_NBELITES,
        lengthAdnF, lengthAdnI);
    VecFloat2D boundsF = VecFloatCreateStatic2D();
    VecShort2D boundsI = VecShortCreateStatic2D();
    VecSet(&boundsF, 0, -1.0); VecSet(&boundsF, 1, 1.0);
    VecSet(&boundsI, 0, 1); VecSet(&boundsI, 1, 10);
    GASetBoundsAdnFloat(ga, 0, &boundsF);
    GASetBoundsAdnFloat(ga, 1, &boundsF);
    GASetBoundsAdnInt(ga, 0, &boundsI);
    GASetBoundsAdnInt(ga, 1, &boundsI);
    GASetNbElites(ga, 2);
    GASetNbEntities(ga, 3);
    GAINit(ga);
    if (ISEQUALF(GAGetDiversity(ga), 0.455102) == false) {
        GenAlgErr->_type = PBErrTypeUnitTestFailed;
        sprintf(GenAlgErr->_msg, "GAGetDiversity failed");
        PBErrCatch(GenAlgErr);
    }
    VecCopy(GAAdn(ga, 1)->_adnF, GAAdn(ga, 0)->_adnF);
    VecCopy(GAAdn(ga, 1)->_adnI, GAAdn(ga, 0)->_adnI);
    if (ISEQUALF(GAGetDiversity(ga), 0.0) == false) {
        GenAlgErr->_type = PBErrTypeUnitTestFailed;
        sprintf(GenAlgErr->_msg, "GAGetDiversity failed");
        PBErrCatch(GenAlgErr);
    }
    GenAlgFree(&ga);
    printf("UnitTestGenAlgGetDiversity OK\n");
}

void UnitTestGenAlgStep() {
    srandom(2);
    int lengthAdnF = 2;
    int lengthAdnI = 2;
    GenAlg* ga = GenAlgCreate(3, 2, lengthAdnF, lengthAdnI);

```



```

VecFloat2D boundsF = VecFloatCreateStatic2D();
VecShort2D boundsI = VecShortCreateStatic2D();
VecSet(&boundsF, 0, -1.0); VecSet(&boundsF, 1, 1.0);
VecSet(&boundsI, 0, 1); VecSet(&boundsI, 1, 10);
GASetBoundsAdnFloat(ga, 0, &boundsF);
GASetBoundsAdnFloat(ga, 1, &boundsF);
GASetBoundsAdnInt(ga, 0, &boundsI);
GASetBoundsAdnInt(ga, 1, &boundsI);
GAInit(ga);
for (int i = 3; i--;)
    GASetAdnValue(ga, GAAdn(ga, i), 3.0 - (float)i);
printf("Before Step:\n");
GAPrintln(ga, stdout);
GenAlgAdn* child = GAAdn(ga, 2);
GASetStep(ga);
printf("After Step:\n");
GAPrintln(ga, stdout);
if (ga->_nextId != 4 || GAAdnGetId(child) != 3 ||
    GAAdnGetAge(child) != 1 ||
    ISEQUALF(GAAdnGetGeneF(child, 0), 0.367611) == false ||
    ISEQUALF(GAAdnGetGeneF(child, 1), 0.174965) == false ||
    ISEQUALF(GAAdnGetDeltaGeneF(child, 0), 0.081678) == false ||
    ISEQUALF(GAAdnGetDeltaGeneF(child, 1), 0.0) == false ||
    GAAdnGetGeneI(child, 0) != 4 ||
    GAAdnGetGeneI(child, 1) != 9 ||
    GAAdn(ga, 2) != child ||
    GAAdnGetAge(GAAdn(ga, 0)) != 2 ||
    GAAdnGetAge(GAAdn(ga, 1)) != 2 ||
    GAAdnGetId(GAAdn(ga, 0)) != 0 ||
    GAAdnGetId(GAAdn(ga, 1)) != 1) {
    GenAlgErr->_type = PBErrTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GASetStep failed");
    PBErrCatch(GenAlgErr);
}
GenAlgFree(&ga);
printf("UnitTestGenAlgStep OK\n");
}

void UnitTestGenAlgStepSubset() {
    srandom(2);
    int lengthAdnF = 2;
    int lengthAdnI = 2;
    GenAlg* ga = GenAlgCreate(3, 2, lengthAdnF, lengthAdnI);
    VecFloat2D boundsF = VecFloatCreateStatic2D();
    VecShort2D boundsI = VecShortCreateStatic2D();
    VecSet(&boundsF, 0, -1.0); VecSet(&boundsF, 1, 1.0);
    VecSet(&boundsI, 0, 1); VecSet(&boundsI, 1, 10);
    GASetBoundsAdnFloat(ga, 0, &boundsF);
    GASetBoundsAdnFloat(ga, 1, &boundsF);
    GASetBoundsAdnInt(ga, 0, &boundsI);
    GASetBoundsAdnInt(ga, 1, &boundsI);
    GAInit(ga);
    for (int i = 3; i--;)
        GASetAdnValue(ga, GAAdn(ga, i), 3.0 - (float)i);
    printf("Before StepSubset:\n");
    GAAPrintln(ga, stdout);
    GenAlgAdn* child = GAAdn(ga, 2);
    GASetStepSubset(ga, 1, 1);
    printf("After StepSubset:\n");
    GAAPrintln(ga, stdout);
    if (ga->_nextId != 4 || GAAdnGetId(child) != 3 ||
        GAAdnGetAge(child) != 1 ||

```

```

ISEQUALF(GAAdnGetGeneF(child, 0), 0.698748) == false ||
ISEQUALF(GAAdnGetGeneF(child, 1), 0.401953) == false ||
ISEQUALF(GAAdnGetDeltaGeneF(child, 0), 0.412815) == false ||
ISEQUALF(GAAdnGetDeltaGeneF(child, 1), 0.0) == false ||
GAAdnGetGeneI(child, 0) != 4 ||
GAAdnGetGeneI(child, 1) != 2 ||
GAAdn(ga, 2) != child ||
GAAdnGetAge(GAAdn(ga, 0)) != 2 ||
GAAdnGetAge(GAAdn(ga, 1)) != 2 ||
GAAdnGetId(GAAdn(ga, 0)) != 0 ||
GAAdnGetId(GAAdn(ga, 1)) != 1) {
    GenAlgErr->_type = PBErrTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GAStepSubset failed");
    PBErrCatch(GenAlgErr);
}
GenAlgFree(&ga);
printf("UnitTestGenAlgStepSubset OK\n");
}

void UnitTestGenAlgLoadSave() {
    srand(5);
    int lengthAdnF = 2;
    int lengthAdnI = 2;
    GenAlg* ga = GenAlgCreate(3, 2, lengthAdnF, lengthAdnI);
    VecFloat2D boundsF = VecFloatCreateStatic2D();
    VecShort2D boundsI = VecShortCreateStatic2D();
    VecSet(&boundsF, 0, -1.0); VecSet(&boundsF, 1, 1.0);
    VecSet(&boundsI, 0, 1); VecSet(&boundsI, 1, 10);
    GASetBoundsAdnFloat(ga, 0, &boundsF);
    GASetBoundsAdnFloat(ga, 1, &boundsF);
    GASetBoundsAdnInt(ga, 0, &boundsI);
    GASetBoundsAdnInt(ga, 1, &boundsI);
    GASetDiversityThreshold(ga, 0.02);
    GAINit(ga);
    GASet(ga);
    GSet* rank = GSetCreate();
    for (int i = 3; i--;)
        GSetAddSort(rank, GAAdn(ga, i), 3.0 - (float)i);
    FILE* stream = fopen("./UnitTestGenAlgLoadSave.txt", "w");
    if (GASave(ga, stream, false) == false) {
        GenAlgErr->_type = PBErrTypeUnitTestFailed;
        sprintf(GenAlgErr->_msg, "GASave failed");
        PBErrCatch(GenAlgErr);
    }
    fclose(stream);
    stream = fopen("./UnitTestGenAlgLoadSave.txt", "r");
    GenAlg* gaLoad = NULL;
    if (GALoad(&gaLoad, stream) == false) {
        GenAlgErr->_type = PBErrTypeUnitTestFailed;
        sprintf(GenAlgErr->_msg, "GALoad failed");
        PBErrCatch(GenAlgErr);
    }
    fclose(stream);
    if (ga->_nextId != gaLoad->_nextId ||
        ga->_curEpoch != gaLoad->_curEpoch ||
        ga->_nbElites != gaLoad->_nbElites ||
        !ISEQUALF(ga->_diversityThreshold, gaLoad->_diversityThreshold) ||
        ga->_lengthAdnF != gaLoad->_lengthAdnF ||
        ga->_lengthAdnI != gaLoad->_lengthAdnI ||
        VecIsEqual(ga->_boundsF, gaLoad->_boundsF) == false ||
        VecIsEqual(ga->_boundsF + 1, gaLoad->_boundsF + 1) == false ||
        VecIsEqual(ga->_boundsI, gaLoad->_boundsI) == false ||

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```

VecIsEqual(ga->_boundsI + 1, gaLoad->_boundsI + 1) == false ||
GAAdnGetId(GAAdn(ga, 0)) != GAAdnGetId(GAAdn(gaLoad, 0)) ||
GAAdnGetId(GAAdn(ga, 1)) != GAAdnGetId(GAAdn(gaLoad, 1)) ||
GAAdnGetId(GAAdn(ga, 2)) != GAAdnGetId(GAAdn(gaLoad, 2)) ||
GAAdnGetAge(GAAdn(ga, 0)) != GAAdnGetAge(GAAdn(gaLoad, 0)) ||
GAAdnGetAge(GAAdn(ga, 1)) != GAAdnGetAge(GAAdn(gaLoad, 1)) ||
GAAdnGetAge(GAAdn(ga, 2)) != GAAdnGetAge(GAAdn(gaLoad, 2)) ||
VecIsEqual(GAAdn(ga, 0)->_adnF,
GAAdn(gaLoad, 0)->_adnF) == false ||
VecIsEqual(GAAdn(ga, 0)->_deltaAdnF,
GAAdn(gaLoad, 0)->_deltaAdnF) == false ||
VecIsEqual(GAAdn(ga, 0)->_adnI,
GAAdn(gaLoad, 0)->_adnI) == false ||
VecIsEqual(GAAdn(ga, 1)->_adnF,
GAAdn(gaLoad, 1)->_adnF) == false ||
VecIsEqual(GAAdn(ga, 1)->_deltaAdnF,
GAAdn(gaLoad, 1)->_deltaAdnF) == false ||
VecIsEqual(GAAdn(ga, 1)->_adnI,
GAAdn(gaLoad, 1)->_adnI) == false ||
VecIsEqual(GAAdn(ga, 2)->_adnF,
GAAdn(gaLoad, 2)->_adnF) == false ||
VecIsEqual(GAAdn(ga, 2)->_deltaAdnF,
GAAdn(gaLoad, 2)->_deltaAdnF) == false ||
VecIsEqual(GAAdn(ga, 2)->_adnI,
GAAdn(gaLoad, 2)->_adnI) == false) {
GenAlgErr->_type = PBErrTypeUnitTestFailed;
sprintf(GenAlgErr->_msg, "UnitTestGenAlgLoadSave failed");
PBErrCatch(GenAlgErr);
}
GSetFree(&rank);
GenAlgFree(&ga);
GenAlgFree(&gaLoad);
printf("UnitTestGenAlgLoadSave OK\n");
}

float ftarget(float x) {
return -0.5 * fastpow(x, 3) + 0.314 * fastpow(x, 2) - 0.7777 * x + 0.1;
}

float evaluate(const VecFloat* adnF, const VecShort* adnI) {
float delta = 0.02;
int nb = (int)round(4.0 / delta);
float res = 0.0;
float x = -2.0;
for (int i = 0; i < nb; ++i, x += delta) {
float y = 0.0;
for (int j = 4; j--;)
y += VecGet(adnF, j) * fastpow(x, VecGet(adnI, j));
res += fabs(ftarget(x) - y);
}
return res / (float)nb;
}

void UnitTestGenAlgTest() {
srandom(5);
int lengthAdnF = 4;
int lengthAdnI = lengthAdnF;
GenAlg* ga = GenAlgCreate(GENALG_NBENTITIES, GENALG_NBELITES,
lengthAdnF, lengthAdnI);
VecFloat2D boundsF = VecFloatCreateStatic2D();
VecShort2D boundsI = VecShortCreateStatic2D();
VecSet(&boundsF, 0, -1.0); VecSet(&boundsF, 1, 1.0);

```

```

VecSet(&boundsI, 0, 0); VecSet(&boundsI, 1, 4);
for (int i = lengthAdnF; i--;) {
    GASetBoundsAdnFloat(ga, i, &boundsF);
    GASetBoundsAdnInt(ga, i, &boundsI);
}
GAInit(ga);
//GASetDiversityThreshold(ga, 0.0);
//float best = 1.0;
//int step = 0;
do {
//float ev = evaluate(GABestAdnF(ga), GABestAdnI(ga));
//printf("%lu %f %f\n", GAGetCurEpoch(ga), ev, GAGetDiversity(ga));
    for (int iEnt = GAGetNbAdns(ga); iEnt--;)
        if (GAAdnIsNew(GAAdn(ga, iEnt)))
            GASetAdnValue(ga, GAAdn(ga, iEnt),
                -1.0 * evaluate(GAAdnAdnF(GAAdn(ga, iEnt)),
                    GAAdnAdnI(GAAdn(ga, iEnt))));
    GAStep(ga);
//float ev = evaluate(GABestAdnF(ga), GABestAdnI(ga));
//if (step == 10){
//    printf("%d %f %f\n", GAGetCurEpoch(ga), ev, GAGetDiversity(ga));
//    step = 0;
//} else step++;
/*if (best - ev > PBMMATH_EPSILON) {
    best = ev;
    printf("%lu %f ", GAGetCurEpoch(ga), best);
    VecFloatPrint(GABestAdnF(ga), stdout, 6);
    printf(" ");
    VecPrint(GABestAdnI(ga), stdout);
    printf("\n");
}*/
} while (GAGetCurEpoch(ga) < 20000 ||
    evaluate(GABestAdnF(ga), GABestAdnI(ga)) < PBMMATH_EPSILON);
printf("target: -0.5*x^3 + 0.314*x^2 - 0.7777*x + 0.1\n");
printf("approx: \n");
GAAdnPrintln(GAAdn(ga, 0), stdout);
printf("error: %f\n", evaluate(GABestAdnF(ga), GABestAdnI(ga)));
GenAlgFree(&ga);
printf("UnitTestGenAlgTest OK\n");
}

void UnitTestGenAlg() {
    UnitTestGenAlgCreateFree();
    UnitTestGenAlgGetSet();
    UnitTestGenAlgInit();
    UnitTestGenAlgPrint();
    UnitTestGenAlgGetDiversity();
    UnitTestGenAlgStep();
    UnitTestGenAlgStepSubset();
    UnitTestGenAlgLoadSave();
    UnitTestGenAlgTest();
    printf("UnitTestGenAlg OK\n");
}

void UnitTestAll() {
    UnitTestGenAlgAdn();
    UnitTestGenAlg();
    printf("UnitTestAll OK\n");
}

int main() {
    UnitTestAll();
}

```

```

    // Return success code
    return 0;
}

```

## 6 Unit tests output

```

UnitTestGenAlgAdnCreateFree OK
UnitTestGenAlgAdnGetSet OK
UnitTestGenAlgAdnInit OK
UnitTestGenAlgAdn OK
UnitTestGenAlgCreateFree OK
UnitTestGenAlgGetSet OK
UnitTestGenAlgInit OK
epoch:0
3 entities, 2 elites
#0 value:0.000000 elite id:0 age:1
  adnF:<0.788004,-0.003504>
  deltaAdnF:<0.000000,0.000000>
  adnI:<3,1>
#1 value:0.000000 elite id:1 age:1
  adnF:<-0.840711,-0.704622>
  deltaAdnF:<0.000000,0.000000>
  adnI:<5,4>
#2 value:0.000000 id:2 age:1
  adnF:<-0.907064,-0.450509>
  deltaAdnF:<0.000000,0.000000>
  adnI:<2,10>
UnitTestGenAlgInit OK
UnitTestGenAlgGetDiversity OK
Before Step:
epoch:0
3 entities, 2 elites
#0 value:3.000000 elite id:0 age:1
  adnF:<0.285933,0.174965>
  deltaAdnF:<0.000000,0.000000>
  adnI:<4,10>
#1 value:2.000000 elite id:1 age:1
  adnF:<-0.156076,-0.303386>
  deltaAdnF:<0.000000,0.000000>
  adnI:<2,7>
#2 value:1.000000 id:2 age:1
  adnF:<0.619353,0.401953>
  deltaAdnF:<0.000000,0.000000>
  adnI:<2,2>
After Step:
epoch:1
3 entities, 2 elites
#0 value:3.000000 elite id:0 age:2
  adnF:<0.285933,0.174965>
  deltaAdnF:<0.000000,0.000000>
  adnI:<4,10>
#1 value:2.000000 elite id:1 age:2
  adnF:<-0.156076,-0.303386>
  deltaAdnF:<0.000000,0.000000>
  adnI:<2,7>
#2 value:1.000000 id:3 age:1
  adnF:<0.367611,0.174965>
  deltaAdnF:<0.081678,0.000000>

```

```

    adnI:<4,9>
UnitTestGenAlgStep OK
Before StepSubset:
epoch:0
3 entities, 2 elites
#0 value:3.000000 elite id:0 age:1
  adnF:<0.285933,0.174965>
  deltaAdnF:<0.000000,0.000000>
  adnI:<4,10>
#1 value:2.000000 elite id:1 age:1
  adnF:<-0.156076,-0.303386>
  deltaAdnF:<0.000000,0.000000>
  adnI:<2,7>
#2 value:1.000000 id:2 age:1
  adnF:<0.619353,0.401953>
  deltaAdnF:<0.000000,0.000000>
  adnI:<2,2>
After StepSubset:
epoch:1
3 entities, 2 elites
#0 value:3.000000 elite id:0 age:2
  adnF:<0.285933,0.174965>
  deltaAdnF:<0.000000,0.000000>
  adnI:<4,10>
#1 value:2.000000 elite id:1 age:2
  adnF:<-0.156076,-0.303386>
  deltaAdnF:<0.000000,0.000000>
  adnI:<2,7>
#2 value:1.000000 id:3 age:1
  adnF:<0.698748,0.401953>
  deltaAdnF:<0.412815,0.000000>
  adnI:<4,2>
UnitTestGenAlgStepSubset OK
UnitTestGenAlgLoadSave OK
target:  $-0.5*x^3 + 0.314*x^2 - 0.7777*x + 0.1$ 
approx:
id:1448376 age:138
  adnF:<-0.501354,-0.774919,0.314189,0.099895>
  deltaAdnF:<0.000099,0.000207,-0.000430,0.000671>
  adnI:<3,1,2,0>
error: 0.001367
UnitTestGenAlgTest OK
UnitTestGenAlg OK
UnitTestAll OK

```

UnitTestGenAlgLoadSave.txt:

```

{
  "_diversityThreshold":"0.020000",
  "_nbAdns":"3",
  "_nbElites":"2",
  "_lengthAdnF":"2",
  "_lengthAdnI":"2",
  "_curEpoch":"1",
  "_nextId":"4",
  "_boundFloat":[
    {
      "_dim":"2",
      "_val":["-1.000000","1.000000"]
    },
    {

```

```

        "_dim": "2",
        "_val": ["-1.000000", "1.000000"]
    }
],
"_boundInt": [
    {
        "_dim": "2",
        "_val": ["1", "10"]
    },
    {
        "_dim": "2",
        "_val": ["1", "10"]
    }
],
"_adns": [
    {
        "_id": "3",
        "_age": "1",
        "_elo": "0.000000",
        "_adnF": {
            "_dim": "2",
            "_val": ["0.755265", "-0.209552"]
        },
        "_deltaAdnF": {
            "_dim": "2",
            "_val": ["-0.032739", "-0.206048"]
        },
        "_adnI": {
            "_dim": "2",
            "_val": ["4", "1"]
        }
    },
    {
        "_id": "1",
        "_age": "2",
        "_elo": "0.000000",
        "_adnF": {
            "_dim": "2",
            "_val": ["-0.840711", "-0.704622"]
        },
        "_deltaAdnF": {
            "_dim": "2",
            "_val": ["0.000000", "0.000000"]
        },
        "_adnI": {
            "_dim": "2",
            "_val": ["5", "4"]
        }
    },
    {
        "_id": "0",
        "_age": "2",
        "_elo": "0.000000",
        "_adnF": {
            "_dim": "2",
            "_val": ["0.788004", "-0.003504"]
        },
        "_deltaAdnF": {
            "_dim": "2",
            "_val": ["0.000000", "0.000000"]
        },
        "_adnI": {

```

```

    "_dim": "2",
    "_val": ["3", "1"]
  }
}
]
}

```

UnitTestGenAlgTest.txt:

```

1 0.522830 <-0.959931,0.745928,-0.259332,0.037688> <3,2,4,3>
3 0.272741 <-0.804893,-0.743738,0.349810,0.186053> <3,1,3,2>
5 0.224412 <-0.505146,-0.496834,0.808183,-0.689687> <1,2,2,3>
6 0.150846 <-0.408103,-0.345099,0.456368,-0.569714> <3,1,2,1>
8 0.123302 <-0.590015,-0.541932,0.303166,0.005749> <3,1,2,2>
9 0.095673 <-0.547299,-0.602595,0.303166,0.037688> <3,1,2,2>
11 0.073450 <-0.549889,-0.646563,0.432221,-0.076964> <3,1,2,2>
34 0.073053 <-0.549889,-0.668342,0.432221,-0.076964> <3,1,2,2>
50 0.072697 <-0.549889,-0.668342,0.432221,-0.064067> <3,1,2,2>
64 0.070609 <-0.547299,-0.668342,0.432221,-0.076964> <3,1,2,2>
87 0.070529 <-0.547299,-0.660726,0.432221,-0.076964> <3,1,2,2>
171 0.070492 <-0.547299,-0.660726,0.786521,-0.430374> <3,1,2,2>
184 0.060799 <-0.512247,-0.758888,0.432221,-0.076964> <3,1,2,2>
256 0.060471 <-0.512247,-0.758888,0.430695,-0.071411> <3,1,2,2>
269 0.060416 <-0.512247,-0.758888,0.430695,-0.072517> <3,1,2,2>
335 0.060040 <-0.487596,-0.820049,0.430695,-0.077548> <3,1,2,2>
370 0.059923 <-0.512247,-0.742234,0.430695,-0.072517> <3,1,2,2>
411 0.059110 <-0.512247,-0.742234,0.430695,-0.077548> <3,1,2,2>
418 0.058697 <-0.487596,-0.806099,0.430695,-0.077548> <3,1,2,2>
452 0.058641 <-0.487596,-0.727993,0.354763,-0.077548> <3,1,2,1>
628 0.058576 <-0.487596,-0.717782,0.354763,-0.089644> <3,1,2,1>
637 0.052264 <-0.487596,-0.806099,0.354763,0.012227> <3,1,2,0>
675 0.051898 <-0.487596,-0.806099,0.347155,0.012227> <3,1,2,0>
678 0.046308 <-0.487596,-0.806099,0.354763,0.028041> <3,1,2,0>
725 0.045874 <-0.487596,-0.800538,0.354763,0.028041> <3,1,2,0>
736 0.042052 <-0.487596,-0.806099,0.354763,0.057678> <3,1,2,0>
738 0.041833 <-0.487596,-0.800538,0.354763,0.045991> <3,1,2,0>
744 0.041471 <-0.487596,-0.792179,0.354763,0.045991> <3,1,2,0>
756 0.027894 <-0.487596,-0.775547,0.307387,0.104752> <3,1,2,0>
773 0.024908 <-0.487596,-0.824285,0.307387,0.123487> <3,1,2,0>
780 0.020777 <-0.487596,-0.811369,0.329454,0.074439> <3,1,2,0>
817 0.019937 <-0.484129,-0.811369,0.329454,0.081048> <3,1,2,0>
1070 0.017202 <-0.502090,-0.765516,0.329454,0.081048> <3,1,2,0>
1132 0.016983 <-0.502698,-0.765516,0.329454,0.081048> <3,1,2,0>
1197 0.016886 <-0.502698,-0.781629,0.329454,0.081048> <3,1,2,0>
1203 0.016112 <-0.502698,-0.769398,0.329454,0.081048> <3,1,2,0>
1243 0.015848 <-0.502698,-0.771443,0.329454,0.081048> <3,1,2,0>
2871 0.015816 <-0.503136,-0.771443,0.329454,0.081048> <3,1,2,0>
5008 0.015671 <-0.503136,-0.771512,0.329454,0.084652> <3,1,2,0>
5902 0.013948 <-0.493362,-0.796571,0.319417,0.106413> <3,1,2,0>
6212 0.010161 <-0.493362,-0.796571,0.315858,0.106413> <3,1,2,0>
6904 0.010081 <-0.493362,-0.796571,0.315858,0.106253> <3,1,2,0>
6984 0.007980 <-0.493362,-0.796571,0.315858,0.097262> <3,1,2,0>
7227 0.006951 <-0.495266,-0.793222,0.315858,0.097262> <3,1,2,0>
7252 0.006330 <-0.495266,-0.792201,0.315858,0.097262> <3,1,2,0>
7934 0.006205 <-0.495266,-0.792201,0.314732,0.097262> <3,1,2,0>
8113 0.006120 <-0.495266,-0.792201,0.314732,0.098331> <3,1,2,0>
8148 0.004850 <-0.495266,-0.786328,0.314732,0.098331> <3,1,2,0>
8932 0.004649 <-0.504552,-0.768060,0.314619,0.098331> <3,1,2,0>
9938 0.004629 <-0.504552,-0.768060,0.314619,0.098553> <3,1,2,0>
10750 0.004602 <-0.504552,-0.768060,0.314619,0.099105> <3,1,2,0>
15488 0.004586 <-0.504552,-0.768514,0.314619,0.099141> <3,1,2,0>

```

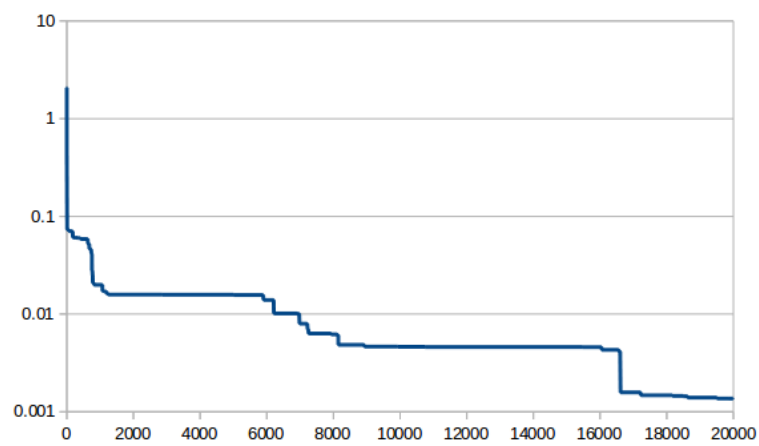


```

16054 0.004302 <-0.504237,-0.768514,0.314619,0.099141> <3,1,2,0>
16563 0.004162 <-0.501354,-0.770848,0.314619,0.099141> <3,1,2,0>
16613 0.001580 <-0.501354,0.099481,0.314619,-0.774215> <3,0,2,1>
17222 0.001476 <-0.501354,-0.774599,0.314619,0.099141> <3,1,2,0>
18170 0.001457 <-0.501354,-0.774599,0.314189,0.099141> <3,1,2,0>
18497 0.001441 <-0.501354,-0.774599,0.314189,0.099224> <3,1,2,0>
18632 0.001396 <-0.501354,-0.774599,0.314189,0.099895> <3,1,2,0>
19511 0.001369 <-0.501354,-0.775126,0.314189,0.099895> <3,1,2,0>

```

eval() of best genes over epoch:



inbreeding over epoch:

