

GenAlg

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Contents

1	Definitions	2
1.1	Selection	2
1.2	Reproduction	2
1.3	Mutation	2
2	Interface	3
3	Code	9
3.1	genalg.c	9
3.2	genalg-inline.c	24
4	Makefile	32
5	Unit tests	33
6	Unit tests output	43

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 $\frac{1}{nbElite} \sum_{i=1}^{nbElite} \frac{\|\vec{adn}(elite_i) - \vec{adn}(elite_0)\|}{\|\vec{bound}_{max} - \vec{bound}_{min}\| * Age(elite_i)}$ 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
    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(int id, int lengthAdnF,
    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
VecFloat* GAAdnAdnF(GenAlgAdn* that);

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

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

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

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

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

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

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

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

```

```

inline
#endif
bool GAAdnIsNew(GenAlgAdn* 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
    int _lengthAdnF;
    // Length of adn for integer value
    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;
} 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(int nbEntities, int nbElites, int lengthAdnF,
    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(GenAlg* that);

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

// Return the nb of elites of the GenAlg 'that'
#if BUILDMODE != 0

```

```

inline
#endif
int GAGetNbElites(GenAlg* that);

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

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

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

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

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

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

// Get the bounds for the 'iGene'-th gene of adn for integer values
#if BUILDMODE != 0
inline
#endif
VecShort2D* GABoundsAdnInt(GenAlg* that, 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
#endif

```

```

void GASetBoundsAdnFloat(GenAlg* that, int iGene, VecFloat2D* 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* that, int iGene, VecShort2D* bounds);

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

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

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

// Get the level of diversity of curent 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
float GAGetDiversity(GenAlg* that);

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

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

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

// ===== 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(int id, int lengthAdnF,
    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 GAAAdnInit(GenAlgAdn* that, GenAlg* ga) {
#ifdef 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 GAAAdnPrintln(GenAlgAdn* that, FILE* 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", GAAAdnGetId(that), GAAAdnGetAge(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(GenAlg* that, int* 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* that, int* parents, 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* that, int* parents, int iChild);

// Reset the GenAlg 'that'
// Randomize all the gene except those of the first adn
void GAKTEvent(GenAlg* that);

// ===== 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(int nbEntities, int nbElites, int lengthAdnF,
    int lengthAdnI) {
    // Allocate memory
    GenAlg* that = PBErrMalloc(GenAlgErr, sizeof(GenAlg));
    // Set the properties
    that->_adns = GSetCreate();
    that->_curEpoch = 0;
    that->_lengthAdnF = lengthAdnF;
    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->_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 GSetNbEntities(GenAlg* that, int nb) {
    #if BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PErrCatch(GenAlgErr);
    }
    if (nb <= 1) {
        GenAlgErr->_type = PErrTypeInvalidArg;
        sprintf(GenAlgErr->_msg, "'nb' is invalid (%d>1)", nb);
        PErrCatch(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)
        GSetNbElites(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* that, 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* 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* 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);
    do {
        // Get the adn
        GenAlgAdn* adn = GSetIterGet(&iter);
        // 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
void GASep(GenAlg* that) {
#ifdef BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PErrCatch(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 < GENALG_DIVERSITYTHRESHOLD) {
        // Break the diversity by applying 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(GenAlg* that, int* parents) {
#ifdef BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PErrCatch(GenAlgErr);
    }
    if (parents == NULL) {
        GenAlgErr->_type = PErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'parents' is null");
        PErrCatch(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* that, int* parents, int iChild) {
#if 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 probabilitly
        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 probabilitly
        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* that, int* parents, 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
            VecFloat2D* 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
        VecShort2D* 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(GenAlg* that, FILE* 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));
}

```

```

// Get the level of diversity of curent 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
float GAGetDiversity(GenAlg* that) {
#ifdef BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
#endif
    // Declare a variable to memorize the result
    float diversity = 0.0;
    // Declare a variable for calculation
    int nb = 1;
    // If there are adn for floating point values
    if (GAGetLengthAdnFloat(that) > 0) {
        // Declare a vector to memorize the ranges in 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
        float normRange = VecNorm(range);
        // For each elite entity except the first one
        for (int iEnt = 1; iEnt < GAGetNbElites(that); ++iEnt) {
            // Get the difference in adn with the first entity
            VecFloat* diff = VecGetOp(GAAdnAdnF(GAAdn(that, iEnt)), 1.0,
                GAAdnAdnF(GAAdn(that, 0)), -1.0);
            // Calculate the diversity
            diversity += VecNorm(diff) /
                (normRange * (float)GAAdnGetAge(GAAdn(that, iEnt)));
            // Free memory
            VecFree(&diff);
        }
        // Calculate the diversity
        nb += GAGetNbElites(that);
        // Free memory
        VecFree(&range);
    }
    // If there are adn for floating point values
    if (GAGetLengthAdnInt(that) > 0) {
        // Declare a vector to memorize the ranges in gene values
        VecFloat* range = VecFloatCreate(GAGetLengthAdnInt(that));
        // Calculate the ranges in gene values
        for (int iGene = GAGetLengthAdnInt(that); iGene--;)
            VecSet(range, iGene,
                (float)(VecGet(GABoundsAdnInt(that, iGene), 1) -
                VecGet(GABoundsAdnInt(that, iGene), 0)));
        // Calculate the norm of the range
        float normRange = VecNorm(range);
        // For each elite entity except the first one
        for (int iEnt = 1; iEnt < GAGetNbElites(that); ++iEnt) {
            // Get the difference in adn with the first entity
            VecShort* diff = VecGetOp(GAAdnAdnI(GAAdn(that, iEnt)), 1,
                GAAdnAdnI(GAAdn(that, 0)), -1);
            VecFloat* diffF = VecShortToFloat(diff);
            // Calculate the diversity
            diversity += VecNorm(diffF) /
                (normRange * (float)GAAdnGetAge(GAAdn(that, iEnt)));
        }
    }
}

```

```

        // Free memory
        VecFree(&diffF);
        VecFree(&diff);
    }
    // Calculate the diversity
    nb += GAGetNbElites(that);
    // Free memory
    VecFree(&range);
}
// Calculate the diversity
diversity /= (float)nb;
// Return the result
return diversity;
}

// Function which return the JSON encoding of 'that'
JSONNode* GAAdnEncodeAsJSON(GenAlgAdn* that, 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(GenAlg* that) {
#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 diversity threshold

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    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) {
        GSetElem* setElem = GSetGetElem(GAAdns(that), iEnt);
        GenAlgAdn* ent = (GenAlgAdn*)(setElem->_data);
        JSONArrayStructAdd(&setAdn,
            GAAdnEncodeAsJSON(ent, setElem->_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, JSONNode* json) {
    #if 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 (JSONGetNbValue(prop) != lengthAdnF)
            return false;
        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 (JSONGetNbValue(prop) != lengthAdnI)
            return false;
        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, JSONNode* json) {

```

```

#if 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
        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");

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    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);
        }
    }
    // 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);
        GSetElem* setElem = GSetGetElem(GAAdns(*that), iEnt);
        if (!GAAdnDecodeAsJSON((GenAlgAdn**)&(setElem->_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* stream) {
    #if 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)) {

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```

        return false;
    }
    // Decode the data from the JSON
    if (!GADecodeAsJSON(that, json)) {
        return false;
    }
    // Free the memory used by the JSON
    JSONFree(&jjson);
    // 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(GenAlg* that, FILE* stream, 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(&jjson);
    // 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'
#ifdef BUILDMODE != 0
inline
#endif
VecFloat* GAAdnAdnF(GenAlgAdn* that) {
#ifdef BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }

```



```

    }
#endif
    return that->_adnF;
}

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

```

```

        PBErCatch(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(GenAlgAdn* that, int iGene) {
    if (BUILDMODE == 0
        if (that == NULL) {
            GenAlgErr->_type = PBErTypeNullPointer;
            sprintf(GenAlgErr->_msg, "'that' is null");
            PBErCatch(GenAlgErr);
        }
    )
    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* that, int iGene, float gene) {
    if (BUILDMODE == 0
        if (that == NULL) {
            GenAlgErr->_type = PBErTypeNullPointer;
            sprintf(GenAlgErr->_msg, "'that' is null");
            PBErCatch(GenAlgErr);
        }
    )
    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* that, int iGene, float delta) {
    if (BUILDMODE == 0
        if (that == NULL) {
            GenAlgErr->_type = PBErTypeNullPointer;
            sprintf(GenAlgErr->_msg, "'that' is null");
            PBErCatch(GenAlgErr);
        }
    )
    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* that, int iGene, 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(GenAlgAdn* 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(GenAlgAdn* 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(GenAlgAdn* that) {
#if BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PErrCatch(GenAlgErr);
    }
#endif
    return (that->_age == 1);
}

// ----- GenAlg

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

// Return the GSet of the GenAlg 'that'
#if BUILDMODE != 0
inline

```

```

#endif
GSet* GAAdns(GenAlg* that) {
#if BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PErrCatch(GenAlgErr);
    }
#endif
    return that->_adns;
}

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

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

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

// Get the length of adn for floating point value
#if BUILDMODE != 0
inline
#endif
int GAGetLengthAdnFloat(GenAlg* 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(GenAlg* 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* that, int iGene, VecFloat2D* 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");
        PBErrCatch(GenAlgErr);
    }
    if (iGene < 0 || iGene >= that->_lengthAdnF) {
        GenAlgErr->_type = PBErrTypeInvalidArg;
        sprintf(GenAlgErr->_msg, "'iGene' is invalid (0<=%d<%d)",
            iGene, that->_lengthAdnF);
        PBErrCatch(GenAlgErr);
    }
#endif
    VecCopy(that->_boundsF + iGene, 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* that, int iGene, VecShort2D* bounds) {
#if BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
    }
#endif
}

```

```

        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);
}

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

// Get the bounds for the 'iGene'-th gene of adn for integer values
#if BUILDMODE != 0
inline
#endif
VecShort2D* GABoundsAdnInt(GenAlg* that, int iGene) {
    if (BUILDMODE == 0
        if (that == NULL) {
            GenAlgErr->_type = PBErTypeNullPointer;
            sprintf(GenAlgErr->_msg, "'that' 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);
        }
    )
    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(GenAlg* that, int iRank) {
#if BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PErrCatch(GenAlgErr);
    }
    if (iRank < 0 || iRank >= GAGetNbAdns(that)) {
        GenAlgErr->_type = PErrTypeInvalidArg;
        sprintf(GenAlgErr->_msg, "'iRank' is invalid (0<=%d<%d)",
            iRank, GAGetNbAdns(that));
        PErrCatch(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* that, GenAlgAdn* adn, float val) {
#if BUILDMODE == 0
    if (that == NULL) {
        GenAlgErr->_type = PErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PErrCatch(GenAlgErr);
    }
    if (adn == NULL) {
        GenAlgErr->_type = PErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'adn' is null");
        PErrCatch(GenAlgErr);
    }
#endif
    GSetGetFirstElem(GAAdns(that), adn)->_sortVal = val;
}

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

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

```

```

    if (that == NULL) {
        GenAlgErr->_type = PBErrTypeNullPointer;
        sprintf(GenAlgErr->_msg, "'that' is null");
        PBErrCatch(GenAlgErr);
    }
#endif
    that->_diversityThreshold = div;
}

```

4 Makefile

```

#directory
PBERRDIR=../PBErr
PBATHDIR=../PBMath
GSETDIR=../GSet
BCURVEDIR=../BCurve
SHAPOIDDIR=../Shapoid
GTREEDIR=../GTree
PBJSONDIR=../PBJson

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

include $(PBERRDIR)/Makefile.inc

INCPATH=-I./ -I$(PBERRDIR)/ -I$(GSETDIR)/ -I$(PBATHDIR)/ -I$(ELORANKDIR)/ -I$(BCURVEDIR)/ -I$(SHAPOIDDIR)/ -I$(PBJSONDIR)/
BUILDOPTIONS=$(BUILDPARAM) $(INCPATH)

# compiler
COMPILER=gcc

#rules
all : main getSCurve

main: main.o pberr.o gset.o pbmath.o genalg.o genalg.o pbjson.o gtree.o Makefile
$(COMPILER) main.o pberr.o gset.o pbjson.o gtree.o pbmath.o genalg.o $(LINKOPTIONS) -o main

main.o : main.c $(PBERRDIR)/pberr.h $(GSETDIR)/gset.h $(ELORANKDIR)/ genalg.h genalg-inline.c Makefile
$(COMPILER) $(BUILDOPTIONS) -c main.c

getSCurve: getSCurve.o pberr.o gset.o pbmath.o genalg.o genalg.o bcurve.o shapoid.o pbjson.o gtree.o Makefile
$(COMPILER) getSCurve.o pberr.o gset.o pbjson.o gtree.o pbmath.o genalg.o bcurve.o shapoid.o $(LINKOPTIONS) -o getSCurve

pbjson.o : $(PBJSONDIR)/pbjson.c $(PBJSONDIR)/pbjson-inline.c $(PBJSONDIR)/pbjson.h Makefile
$(COMPILER) $(BUILDOPTIONS) -c $(PBJSONDIR)/pbjson.c

gtree.o : $(GTREEDIR)/gtree.c $(GTREEDIR)/gtree.h $(GTREEDIR)/gtree-inline.c Makefile $(GSETDIR)/gset-inline.c $(GSETDIR)/gset.h
$(COMPILER) $(BUILDOPTIONS) -c $(GTREEDIR)/gtree.c

getSCurve.o : getSCurve.c $(PBERRDIR)/pberr.h $(GSETDIR)/gset.h $(ELORANKDIR)/ $(BCURVEDIR)/bcurve.h genalg.h genalg-inline.c
$(COMPILER) $(BUILDOPTIONS) -c getSCurve.c

genalg.o : genalg.c genalg.h genalg-inline.c Makefile
$(COMPILER) $(BUILDOPTIONS) -c genalg.c

```



```

shapoid.o : $(SHAPOIDDIR)/shapoid.c $(SHAPOIDDIR)/shapoid.h $(SHAPOIDDIR)/shapoid-inline.c Makefile
$(COMPILER) $(BUILDOPTIONS) -c $(SHAPOIDDIR)/shapoid.c

bcurve.o : $(BCURVEDIR)/bcurve.c $(BCURVEDIR)/bcurve.h $(BCURVEDIR)/bcurve-inline.c Makefile
$(COMPILER) $(BUILDOPTIONS) -c $(BCURVEDIR)/bcurve.c

pberr.o : $(PBERRDIR)/pberr.c $(PBERRDIR)/pberr.h Makefile
$(COMPILER) $(BUILDOPTIONS) -c $(PBERRDIR)/pberr.c

pbmath.o : $(PBATHDIR)/pbmath.c $(PBATHDIR)/pbmath-inline.c $(PBATHDIR)/pbmath.h Makefile $(PBERRDIR)/pberr.h
$(COMPILER) $(BUILDOPTIONS) -c $(PBATHDIR)/pbmath.c

gset.o : $(GSETDIR)/gset.c $(GSETDIR)/gset-inline.c $(GSETDIR)/gset.h Makefile $(PBERRDIR)/pberr.h
$(COMPILER) $(BUILDOPTIONS) -c $(GSETDIR)/gset.c

clean :
rm -rf *.o main

valgrind :
valgrind -v --track-origins=yes --leak-check=full --gen-suppressions=yes --show-leak-kinds=all ./main

unitTest :
main > unitTest.txt; diff unitTest.txt unitTestRef.txt

```

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");
    }
}

```

```

    PBErriCatch(GenAlgErr);
}
if (GAAdnIsNew(ent) != true) {
    GenAlgErr->_type = PBErriTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GAAdnIsNew failed");
    PBErriCatch(GenAlgErr);
}
ent->_age = 2;
if (GAAdnIsNew(ent) != false) {
    GenAlgErr->_type = PBErriTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GAAdnIsNew failed");
    PBErriCatch(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);
    VecCopy(GABoundsAdnFloat(ga, 0), &boundsF);
    VecCopy(GABoundsAdnFloat(ga, 1), &boundsF);
    VecCopy(GABoundsAdnInt(ga, 0), &boundsI);
    VecCopy(GABoundsAdnInt(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 = PBErriTypeUnitTestFailed;
        sprintf(GenAlgErr->_msg, "GAAdnInit failed");
        PBErriCatch(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");
    }
}

```

```

    PBErCatch(GenAlgErr);
}
if (GAGetLengthAdnInt(ga) != lengthAdnI) {
    GenAlgErr->_type = PBErTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GAGetLengthAdnInt failed");
    PBErCatch(GenAlgErr);
}
if (GABoundsAdnFloat(ga, 1) != ga->_boundsF + 1) {
    GenAlgErr->_type = PBErTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GABoundsAdnFloat failed");
    PBErCatch(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 = PBErTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GASetBoundsAdnFloat failed");
    PBErCatch(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 = PBErTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GASetBoundsAdnInt failed");
    PBErCatch(GenAlgErr);
}
if (GABoundsAdnInt(ga, 1) != ga->_boundsI + 1) {
    GenAlgErr->_type = PBErTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GABoundsAdnInt failed");
    PBErCatch(GenAlgErr);
}
GASetAdnValue(ga, GAAdn(ga, 0), 1.0);
if (ISEQUALF(ga->_adns->_tail->_sortVal, 1.0) == false) {
    GenAlgErr->_type = PBErTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GASetAdnValue failed");
    PBErCatch(GenAlgErr);
}
if (ISEQUALF(GAGetDiversityThreshold(ga),
ga->_diversityThreshold) == false) {
    GenAlgErr->_type = PBErTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GAGetDiversityThreshold failed");
    PBErCatch(GenAlgErr);
}
GASetDiversityThreshold(ga, 0.5);
if (ISEQUALF(GAGetDiversityThreshold(ga), 0.5) == false) {
    GenAlgErr->_type = PBErTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GASetDiversityThreshold failed");
    PBErCatch(GenAlgErr);
}
GenAlgFree(&ga);
printf("UnitTestGenAlgGetSet OK\n");
}

void UnitTestGenAlgInit() {
    srand(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);
VecCopy(GABoundsAdnFloat(ga, 0), &boundsF);
VecCopy(GABoundsAdnFloat(ga, 1), &boundsF);
VecCopy(GABoundsAdnInt(ga, 0), &boundsI);
VecCopy(GABoundsAdnInt(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);
    VecCopy(GABoundsAdnFloat(ga, 0), &boundsF);
    VecCopy(GABoundsAdnFloat(ga, 1), &boundsF);
    VecCopy(GABoundsAdnInt(ga, 0), &boundsI);
    VecCopy(GABoundsAdnInt(ga, 1), &boundsI);
    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);
    VecCopy(GABoundsAdnFloat(ga, 0), &boundsF);
    VecCopy(GABoundsAdnFloat(ga, 1), &boundsF);
    VecCopy(GABoundsAdnInt(ga, 0), &boundsI);
    VecCopy(GABoundsAdnInt(ga, 1), &boundsI);
    GASetNbElites(ga, 2);
    GASetNbEntities(ga, 3);
    GAInit(ga);
    if (ISEQUALF(GAGetDiversity(ga), 0.182041) == 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(GAAdnGetDiversity(ga), 0.0) == false) {
    GenAlgErr->_type = PBErrTypeUnitTestFailed;
    sprintf(GenAlgErr->_msg, "GAAdnGetDiversity failed");
    PBErrCatch(GenAlgErr);
}
GenAlgFree(&ga);
printf("UnitTestGenAlgGetDiversity OK\n");
}

void UnitTestGenAlgStep() {
    srand(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);
    VecCopy(GABoundsAdnFloat(ga, 0), &boundsF);
    VecCopy(GABoundsAdnFloat(ga, 1), &boundsF);
    VecCopy(GABoundsAdnInt(ga, 0), &boundsI);
    VecCopy(GABoundsAdnInt(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);
    GASetAdnValue(ga, child, 3.0 - (float)i);
    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, "GAStep failed");
        PBErrCatch(GenAlgErr);
    }
    VecCopy(GAAdn(ga, 1)->_adnF, GAAdn(ga, 0)->_adnF);
    VecCopy(GAAdn(ga, 1)->_adnI, GAAdn(ga, 0)->_adnI);
    GASetAdnValue(ga, child, 3.0 - (float)i);
    printf("After StepEpoch with interbreeding:\n");
    GAPrintln(ga, stdout);
    if (ga->_nextId != 6 || GAAdnGetId(child) != 5 ||
        GAAdnGetAge(child) != 1 ||
        ISEQUALF(GAAdnGetGeneF(child, 0), 0.289982) == false ||
        ISEQUALF(GAAdnGetGeneF(child, 1), -0.910199) == false ||
        ISEQUALF(GAAdnGetDeltaGeneF(child, 0), 0.081678) == false ||
        ISEQUALF(GAAdnGetDeltaGeneF(child, 1), 0.0) == false ||
        GAAdnGetGeneI(child, 0) != 9 ||

```

```

    GAAdnGetGeneI(child, 1) != 8 ||
    GAAdn(ga, 2) != child ||
    GAAdnGetAge(GAAdn(ga, 0)) != 2 ||
    GAAdnGetAge(GAAdn(ga, 1)) != 1 ||
    GAAdnGetId(GAAdn(ga, 0)) != 0 ||
    GAAdnGetId(GAAdn(ga, 1)) != 4) {
        GenAlgErr->_type = PBErrTypeUnitTestFailed;
        sprintf(GenAlgErr->_msg, "GASstep failed");
        PBErrCatch(GenAlgErr);
    }
    GenAlgFree(&ga);
    printf("UnitTestGenAlgStep 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);
    VecCopy(GABoundsAdnFloat(ga, 0), &boundsF);
    VecCopy(GABoundsAdnFloat(ga, 1), &boundsF);
    VecCopy(GABoundsAdnInt(ga, 0), &boundsI);
    VecCopy(GABoundsAdnInt(ga, 1), &boundsI);
    GASetDiversityThreshold(ga, 0.02);
    GAInit(ga);
    GASstep(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 ||
        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(VecFloat* adnF, 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() {
    srand(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--;) {
        VecCopy(GABoundsAdnFloat(ga, i), &boundsF);
        VecCopy(GABoundsAdnInt(ga, i), &boundsI);
    }
}

```

```

    GAInit(ga);
    //float best = 1.0;
    //int step = 0;
    /*float ev = evaluate(GABestAdnF(ga), GABestAdnI(ga));
    printf("%d %f %f\n", GAGetCurEpoch(ga), ev, GAGetDiversity(ga));*/
    do {
        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);
    } while (GAGetCurEpoch(ga) < 20000 ||
        evaluate(GABestAdnF(ga), GABestAdnI(ga)) < PBMath_EPSILON);
    //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 > PBMath_EPSILON) {
        best = ev;
        printf("%d %f ", GAGetCurEpoch(ga), best);
        VecFloatPrint(GABestAdnF(ga), stdout, 6);
        printf(" ");
        VecPrint(GABestAdnI(ga), stdout);
        printf("\n");
    }*/
    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();
    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>
After StepEpoch with interbreeding:
epoch:2
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:4 age:1
  adnF:<0.700961,0.779526>
  deltaAdnF:<0.000000,0.000000>
  adnI:<2,4>
#2 value:1.000000 id:5 age:1
  adnF:<0.289982,-0.910199>
  deltaAdnF:<0.081678,0.000000>
  adnI:<9,8>
UnitTestGenAlgStep OK
UnitTestGenAlgLoadSave OK
target:  $-0.5*x^3 + 0.314*x^2 - 0.7777*x + 0.1$ 
approx:
id:983022 age:7634
  adnF:<-0.772051,0.313140,-0.502480,0.101328>
  deltaAdnF:<-0.006612,-0.000041,0.085838,0.741578>
  adnI:<1,2,3,0>
error: 0.002613
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 0.522828 <-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 0.224411 <-0.505146,-0.496834,0.808183,-0.689687> <1,2,2,3>
6 0.150845 <-0.408103,-0.345099,0.456368,-0.569714> <3,1,2,1>
8 0.123301 <-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 0.073451 <-0.549889,-0.646563,0.432221,-0.076964> <3,1,2,2>
59 0.073127 <-0.504843,0.334721,-0.700384,-0.077186> <3,2,1,1>
153 0.071830 <-0.389386,0.342553,-0.856339,-0.077186> <3,2,1,3>
198 0.064972 <0.353165,-0.442254,-0.461739,-0.429105> <2,1,3,1>
214 0.062897 <0.110719,0.342553,-0.461739,-0.885047> <0,2,3,1>
319 0.054397 <0.110719,0.342553,-0.461739,-0.830428> <0,2,3,1>
442 0.041706 <0.334213,-0.528332,-0.749878,0.075020> <2,3,1,0>

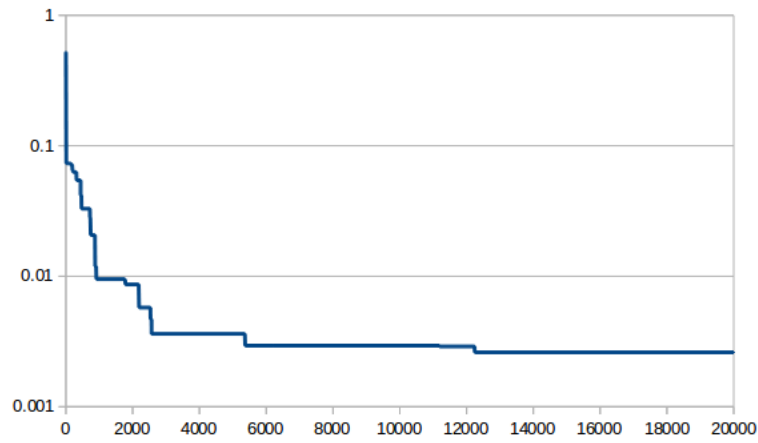
```

```

469 0.032929 <-0.334213,-0.490436,-0.770205,0.075020> <2,3,1,0>
721 0.028211 <-0.820933,-0.490436,0.315181,0.075020> <1,3,2,0>
743 0.020676 <-0.334213,-0.503733,-0.770205,0.075020> <2,3,1,0>
878 0.011919 <-0.770117,0.313181,-0.507948,0.095455> <1,2,3,0>
909 0.010774 <-0.770117,0.313181,-0.507948,0.103277> <1,2,3,0>
914 0.009503 <-0.765439,0.313181,-0.507948,0.095455> <1,2,3,0>
1773 0.009382 <-0.767748,0.313181,-0.507948,0.097955> <1,2,3,0>
1783 0.008650 <-0.765439,0.313181,-0.507948,0.097955> <1,2,3,0>
2192 0.005756 <-0.765439,0.313181,-0.503778,0.097955> <1,2,3,0>
2534 0.004692 <-0.772051,0.313181,-0.503778,0.097955> <1,2,3,0>
2565 0.003620 <-0.772051,0.313181,-0.502827,0.097955> <1,2,3,0>
5369 0.002942 <-0.772051,0.313181,-0.502827,0.101665> <1,2,3,0>
11186 0.002902 <-0.772051,0.313140,-0.502827,0.101328> <1,2,3,0>
12245 0.002613 <-0.772051,0.313140,-0.502480,0.101328> <1,2,3,0>

```

eval() of best genes over epoch:



inbreeding over epoch:

