

Framsticks scripting

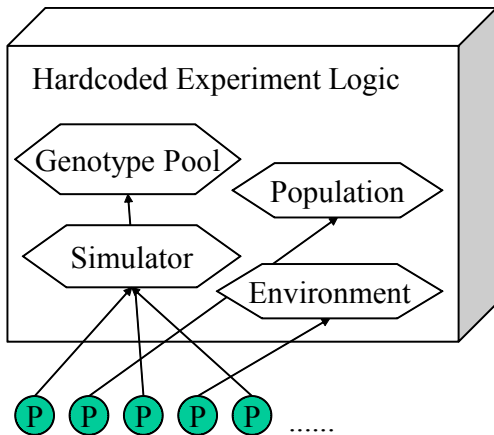
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www.framsticks.com

Why use script language?

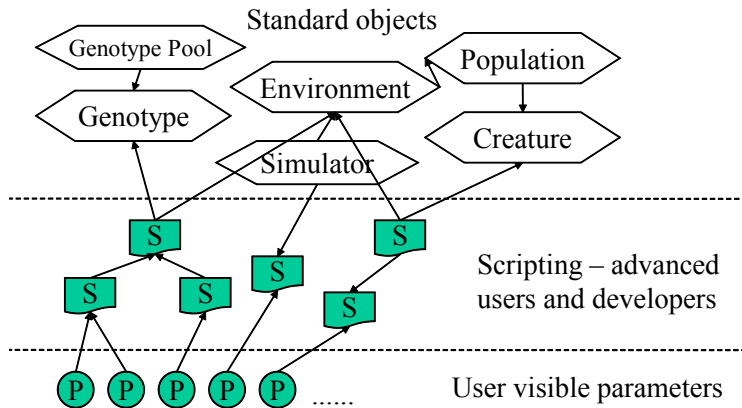
- In GUI: to automate common/repeating tasks (“macros”)
- Algorithmical approach is more straightforward for some purposes
- To simplify internal dependencies
- To give more control to advanced users
- An advanced user can define new experiments, evaluation methods, neuron types, creature interaction, visualization
- Regular users do not see the involved complexity

Popular Alife Software



User-visible parameters

Framsticks Architecture

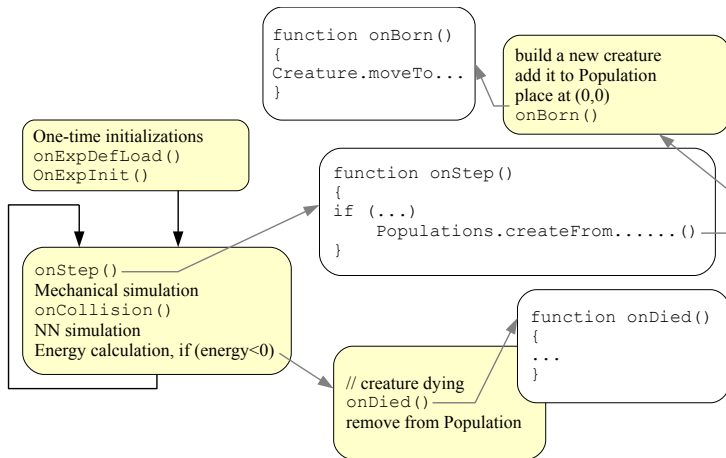


FramScript Language

- JavaScript-like
- 'C' syntax and semantics, untyped variables, functions, can reference Framsticks objects
- Sample code:

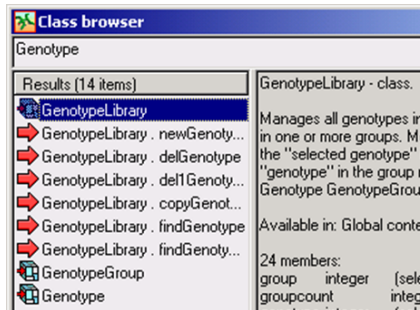
```
function test ( angle , mult )  
{  
  var s=mult*Math.sin ( angle );  
  return s+ExpProperties.uservalue ;  
}
```

Event-based architecture: main loop and script handlers



Object Model

- Core Simulator objects
(Genotype, Creature, World, ...)
- User Interface objects (CLI, GLDisplay, Chart, ...)
- General Purpose objects
(Math, Collections, ...)



Class reference available in the
Framsticks application:
Help/Class Browser

The screenshot shows the Framsticks software interface with several components labeled:

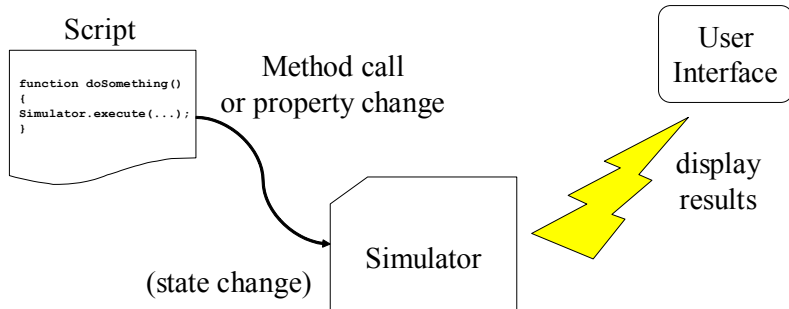
- Simulator**: Points to the top menu bar and toolbar.
- GLDisplay**: Points to the central 3D visualization window.
- World**: Points to the 3D environment within the GLDisplay window.
- Model**: Points to the 'Body & Brain' window showing a hierarchical tree of the selected entity.
- Genotype** and **Geno**: Point to the 'Genotypes' list in the 'Groups' panel.
- GenePools** and **GenePool**: Point to the 'Gene Pools' list in the 'Groups' panel.
- Populations**, **Population**, and **Population**: Point to the 'Populations' list in the 'Groups' panel.
- Creature**: Points to the 'Creatures' list in the 'Groups' panel.

The interface includes a top menu bar (File, Simulation, Genotypes, Individuals, Interface, Window, Help), a toolbar with icons for file operations and simulation control, and several panels for managing groups, displaying the 3D world, and viewing detailed information about selected entities.

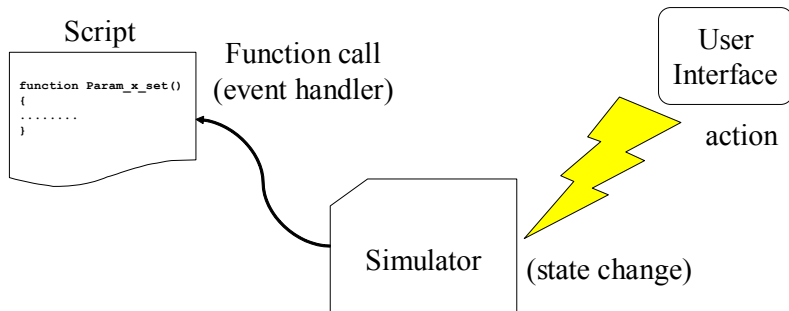
Name	Genotype	Body joints	Brain size	Instan...	Life span	Final fitness
Caterpillar	mXtBkIISSEFFXt	8	6	0	0	0 / ...

Name	Genotype	Energy	Performa...	NN enabl...
Caterpillar	mXtBkIISSEFFXt	106415	On	1
Food	//0m.Vstyle=foodp.	200	Off	1

Script/Simulator/User



User/Simulator/Script

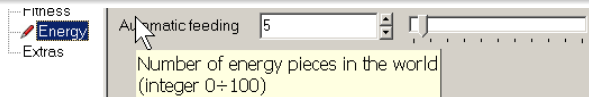


User-visible properties

- Definition:

```
property:  
id:feed  
name:Automatic feeding  
type:d 0 100 5  
group:Energy  
help:Number of energy pieces in the world
```

- Presentation:



- Script access:

```
ObjectName.feed=12;  
function ObjectName_set_feed() // 'feed' changed  
{  
    // handle modification of the 'feed' variable  
}
```

Property types

- **“d” – integer value**

- “d min max” – integer with constraints

- “d 0 1” – false/true switch (checkbox)



- “d min max ~name1~name2~name3...” – integer presented as named values (combo)

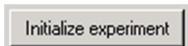


- **“f” – floating point value**

- “f min max” – floating point with constraints

Property types

- **“s” – text string**
 - “s 1 maxlength” – multiline text
 - “s ~value1~value2~value3...” – text with suggested default values (combo)
- **“p” – procedure call** (action), a button in GUI
- **“o”/“x”** – object property/unknown type property (not useful in the interface)



Editors

These syntax highlighting editors will make script writing easier:

- Emacs, www.gnu.org/software/emacs
- jEdit, www.jedit.org
- Crimson Editor, www.crimsoneditor.com
- (your favorite editor with syntax highlighting)

...but the recommended tool is Framclipse.

Framclipse: a plugin for Eclipse

- Fully integrated with the Eclipse platform
- Easy installation – "Install New Software" Eclipse feature

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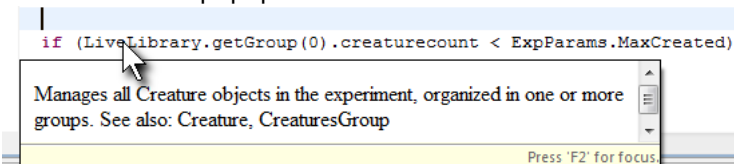
```
//comment|  
function onExpLoad()  
{  
    Simulator.message("expdef: onExpLoad not implemented", 3);  
}
```


Framclipse: a plugin for Eclipse

- Fully integrated with the Eclipse platform
- Easy installation – "Install New Software" Eclipse feature
- Syntax highlighting

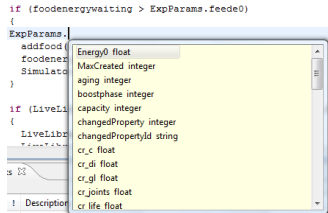
```
//comment  
function onExpLoad()  
{  
    Simulator.message("expdef: onExpLoad not implemented", 3);  
}
```

- Documentation popups



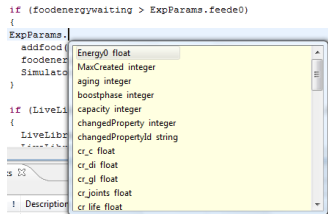
Framclipse: more features

- Content assistance

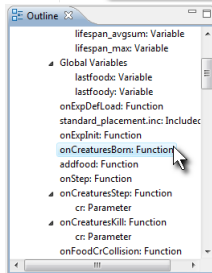


Framclipse: more features

- Content assistance



- Interactive outline
- And much more.



```
function onCreaturesBorn()
{
  Creature.idlelen = ExpParams.e_meta;
  Creature.energ0 = ExpParams.Energy0*0.8+Math.rndUni(0,E
  Creature.energy = Creature.energ0;

  Creature.user1 = null; //doesn't know where food is
  Creature.user2 = null;
  Creature.user3 = Math.rndUni(0,Math.twopi); //initial ar
  Creature.rotate(0,0,Creature.user3);
}

function addfood()
```

Script applications in Framsticks

- Neuron definitions
- Experiment definitions
- Advanced 3D visualization
- "Framsticks Theater" presentations
- General purpose scripting (macros)

Neuron definitions

- Neuron: a signal processing unit
- Direct control of the neuron output
- Can use any user defined functions, read from neuron inputs, preserve state in the neuron itself
- Public properties can be used to influence the neuron behavior. Genetic operators will accept and operate on those properties

Neuron definitions code

- Initialization function ("constructor") – set the internal state if needed

```
function init()  
{ NeuroProperties.somestate=0; }
```

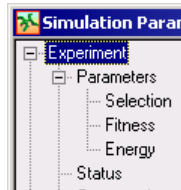
- Working function – calculate the new state

```
function go()  
{ Neuro.state=...something...; }
```

- Properties (accessible as "NeuroProperties")
 - Public properties (with flags=0) – initialized based on the genotype data, and available for genetic operators
 - Private properties (with flags=32) – only for a neuron script, for internal purposes

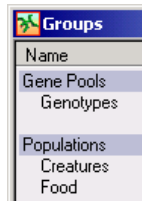
Experiment definition

- Controls the way objects are created in the world and how they are evaluated
- Can create user-visible parameters (ExpProperties) and experiment state variables (ExpState)
- Responsible for saving/restoring experiment state



Experiment definition: Groups

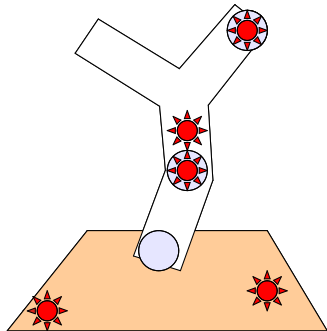
- Interaction with User Interface
- Script access:
 - GenePools
 - GenePool
 - Genotype
 - Populations
 - Population
 - Creature



Communication: Signal objects

Why use signals:

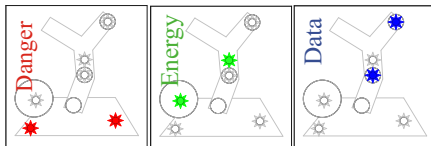
- Optimized for common cases
- Unified programming interface for stationary,  neuron-attached,  and creature-attached signals 
- Visualization in GUI and Theater



Communication: Signal objects

Signal properties:

- Channel

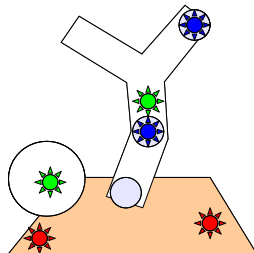


- Power



- Flavor

- Value (any object)



Communication: Signal objects

Receiving aggregated power

- Low level, "biological" senses
- Easy, one-line custom neurons:

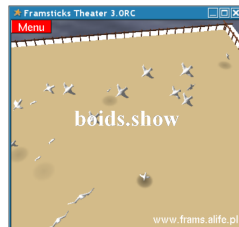
```
Neuro.state=Neuro.signals.  
receiveFilter("channel",  
max_range, flavor, filter);
```



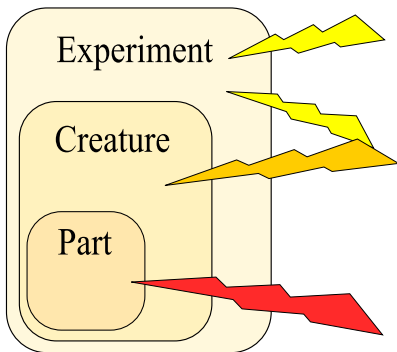
Receiving objects

- Can pass any kind of data
- More sophisticated processing:

```
var n=Creature.signals.  
receiveSet("flock",range);  
for(i=0;i<n.size;i++)  
doSomething(n[i].value);
```



Experiment definition: Events



onExpDefLoad(), onExplnit(),
onExpLoad(), onExpSave(),
onStep()

onBorn(), onDied(), onStep(),
onUpdate(), onCrCollision()

onCollision()

Experiment definition: Events

- `onExpDefLoad()`

The experiment definition is loaded.

Usual actions:

- Setup genotypes and creatures groups
- Initialize experiment parameters and state

- `onExpInit()`

New experiment started.

- Clear all state information

Experiment definition: Events

- `onBorn()`
A new creature was born (as a result of script or user action).
Usual handling:
 - Adjust placement
 - Initialize creature's properties
- `onGroupNameBorn(creature_arg)`
Group-specific handler for `onBorn()`

Experiment definition: Events

- `onDied()`
The creature has died. Usual handling:
 - Update performance data about its genotype.
- `onGroupNameDied(creature_arg)`
Group-specific handler for `onDied()`

Note: in older version of Framsticks, use `onKill()` instead of `onDied()`.

Experiment definition: Events

- `onStep()`
Called before each simulation step, once
- `onGroupNameStep(creature_arg)`
Called for each creature in a group
- `onUpdate()`
Called periodically as requested by the "performance sampling period" parameter. Creature's performance fields (distance, velocity) have been updated.
- `onGroupNameUpdate(creature_arg)`
Group-specific handler for `onUpdate()`

Experiment definition: Events

- onGroupNameCollision()
Called for each creature part touching another creature.
Usual handling:
 - Energy transfer (see "standard.expdef")
 - Update performance
- onGroupNameCrCollision()
Similar, for creature-level collisions

Experiment definition: Events

- onExpSave()

Should save experiment state using the "File" object.

Sample usage:

```
File.writeComment("my parameters");  
File.writeObject(sim_params.*);
```

- onExpLoad()

Restore the experiment state from the file created in the "ExpSave" event. The "Loader" object can parse and load multiple objects.

```
Loader.addClass(sim_params.*);  
Loader.run();
```

Experiment definition: Events

UserEvent: An action users can launch from the context menu, eg. "Place food" and "Drop food" found in most standard experiments (defined in **standard_events.inc**)

- `queryEventNames()`
Return the supported event names (vector of text strings)
- `onUserEvent(type,point,vector)`
Handle the event. Point and vector are 3-element vectors describing the world location where the user action occurred.

Experiment definition: Events

```
function queryEventNames()  
{  
    return ["Place food", "Drop food"];  
}  
  
function onUserEvent(type, point, vector)  
{  
    var p = WorldMap.intersect(point, vector);  
    if (p)  
    {  
        addfood();  
        var z = p[2];  
        if (type == 1) z += 10;  
        Creature.moveAbs(p[0] - Creature.size_x / 2,  
                          p[1] - Creature.size_y / 2, z);  
    }  
}
```

Experiment definition: Low-level mouse interaction

- `onMouseClicked(options, xy, point, vector)`
- `onMouseMove(options, xy, point, vector)`
- `onMouseUnclick(options, xy, point, vector)`

Arguments:

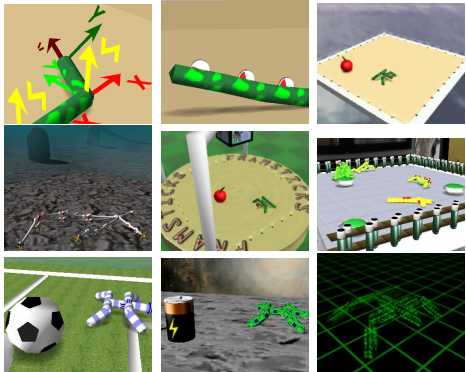
- **options** – mouse button and double click identification
- **xy** – 2D coordinates (`xy[0]` and `xy[1]`)
- **point, vector** – 3D "line of click" defined by point and vector

Advanced 3D visualization

- OpenGL – 3D rendering engine
- PLIB/SSG – Scene graph
- Script:
 - Create or load geometry and textures for all objects (environment, creatures)
 - Update object properties (positions)

Advanced 3D visualization: style definition

- Style: predefined visualization functions with associated graphics files



Examples: XYZ, pressure, standard, spooksticks, arena, laboratory, football, space, matrix

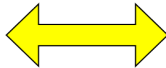
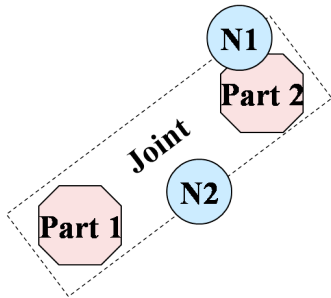
Advanced 3D visualization: objects

- **GeomBuilder**
 - Create/modify scene graph(SSG nodes manipulation)
 - Special support for VisualModel
- **VertexBuilder**
 - Create any shape (low level: points, polygons)
- **Loader**
 - Load 3D object files (SSG)
- **Material**
 - Create/modify object attributes – colors, textures (SSG)

Advanced 3D visualization: objects

- **VisualModel**

- Mapping: creature element $\leftrightarrow$ scene node



- ***Creature root node***
 - *Part1*
 - *N1*
 - *Part2*
 - *Joint*
 - *N2*

Advanced 3D visualization: script functions

- `onLoad()`
- `stylename_model_build()` – called once, before the visual object is created
- `world_build()` – called when the world parameters change
- `modelviewer_build_empty()`
- `modelviewer_build()`

Advanced 3D visualization: script functions

- `stylename_elementname_build()` – create a 3D object associated with the current element
- `stylename_elementname_update()` – update the 3D object according to the current properties
- **stylename**: “default”, unless set to another name in the model (eg. “food”, “manipulator”)
- **elementname**: “part”, “joint”, “neuro” unless set to another name in the model element

Advanced 3D visualization: script sample #1

Listing 1: Style definition

```
function default_flag_build()  
{...}
```

Genotype

```
...  
p:3,1.72,6,Vstyle=flag  
...
```

Result



Advanced 3D visualization: script sample #2

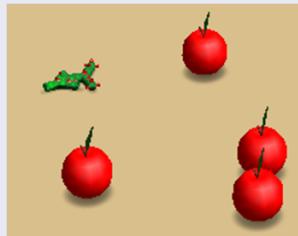
Listing 2: Style definition

```
function food_part_build()  
{...}
```

Genotype

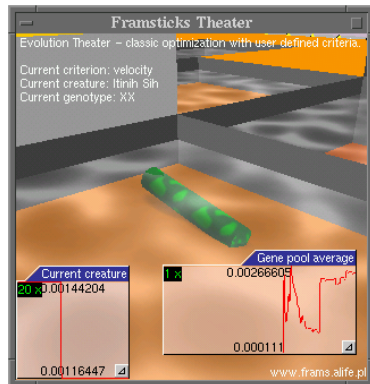
```
//0  
m:Vstyle=food  
p:
```

Result



Framsticks Theater presentations

- Setup the simulator for a predefined experiment
- Provides a simple, attractive graphical user interface
- Can define user-visible parameters
- Can create and use additional GUI objects



Framsticks Theater presentations

- UI capabilities:
 - **ShowProperties** object (user-visible parameters and actions presented in the menu)
 - **GLDisplay.banner="text"** (simple text display)
 - **GLDisplay.createWindow(style, title, client)**
 - **GLDisplay.newCreatureCharts()**
 - **GLDisplay.newSimStatsCharts()**



Framsticks Theater presentations

- Events:
 - onLoad(), onResize(), onShowStep(), onSimStep(), onSelectionChange()
- Objects:
 - GLDisplay, Camera, Window, DynaChart

General purpose scripting

- Used for repeating tasks and testing
- Can access rarely used features without a user interface
- Command line simulator: all user actions are actually a script
- GUI version: user scripts can be accessed from the "Parameters" window

Let's create a neuron that occasionally generates noise.

The neuron should occasionally generate random output, otherwise it should work like the regular “N” neuron.

Let's create a neuron that occasionally generates noise.

The neuron should occasionally generate random output, otherwise it should work like the regular “N” neuron.

General neuron information (inside the “class:” section)

```
class:
name:Nn
longname:Noisy neuron
prefinputs:-1
prefoutput:1
code:~
..... the script code will be here....
~
```

One public parameter – “Error rate” (probability of response perturbation)

```
property:
id:e
name>Error rate
type:f 0.0 0.1
```

Let's create a neuron that occasionally generates noise.

The neuron should occasionally generate random output, otherwise it should work like the regular “N” neuron.

General neuron information (inside the “class:” section)

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~
```

One public parameter – “Error rate” (probability of response perturbation)

```
property:
id:e
name:Error rate
type:f 0.0 0.1
```

Listing 5: Work function

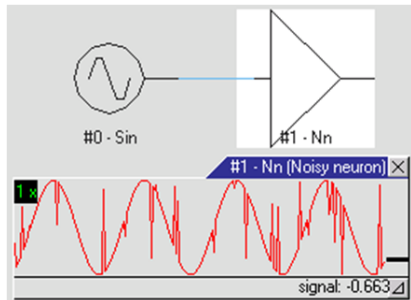
```
function go()
{
    var s=Neuro.weightedInputSum;
    if (Math.rnd01 < NeuroProperties.e
        s=(Math.rnd01*2)-1.0;
    Neuro.state=s;
}
```

Complete source and demonstration

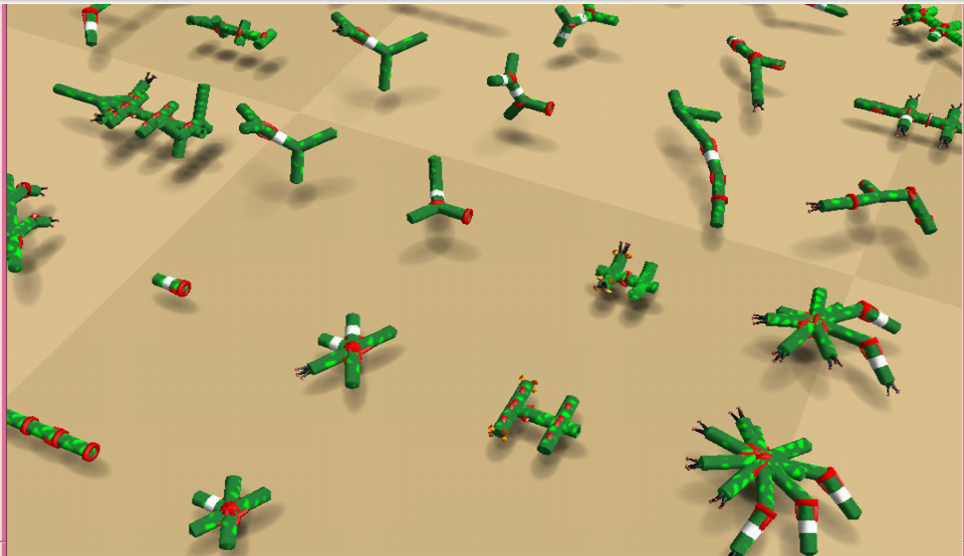
The “noisy.neuro” file

```
class:
name:Nn
longname:Noisy neuron
prefinputs:-1
prefoutput:1
code:~
function go()
{
  var s=Neuro.weightedInputSum;
  if (Math.rnd01 < NeuroProperties.e)
    s=(Math.rnd01*2)-1.0;
  Neuro.state=s;
}
~

property:
id:e
name>Error rate
type:f 0.0 0.1
```



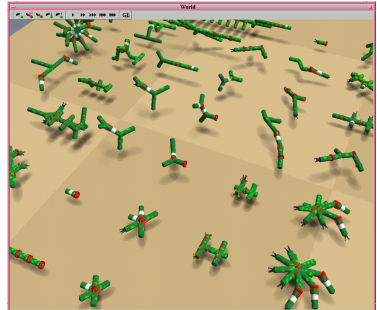
Let's place all creatures in a grid (e.g. for a presentation)



Script header

General script information (inside the “script:” section)

```
script:  
name:Gallery  
help:Create a grid of creatures from the first genotype list  
code:~  
..... the script will be here....  
~
```



Script source

```
var gridx = 6.0, gridy = 6.0; // grid spacing
var n = GenePools[0].size;
if (n==0) {
    Simulator.print("This script needs some genotypes "+
        "in the first group."); return;
}
var side = 0+(Math.sqrt(n)+0.999);
Simulator.print("Have "+n+" genotype(s), it will be "+
    side+" by "+side+" grid");

var i,j,g=0,x,y,z;
for(i=0;i<side;i++)
    for(j=0;j<side;j++)
    {
        if (g >= GenePools[0].size) return;
        Populations[0].createFromGeno(GenePools[0][g].geno);
        x=gridx*i; y=gridy*j;
        z=WorldMap.getHeight(x,y);
        Creature.moveAbs(x-Creature.size_x/2,y-Creature.size_y/2,z);
        g++;
    }
```


Let's create an interesting evolutionary experiment

- Creatures gain energy by eating (touching) food
- Each creature with a sufficient amount of energy produces an offspring, which may be a clone or a mutant
- New food pieces are created in the world and placed randomly

General experiment information (inside the "expdef:" section)

```
expdef:
name:Asexual reproduction in the world
info:~
Each creature with a sufficient amount of energy
produces an offspring, which is then put close to its parent.
Food is created at a constant rate and placed randomly.
~
code:~
...this will contain the script...
~
```

Experiment parameters

These parameters will be public, available in GUI and for other components

```
property:  
id:p_mutate  
name:Mutation probability  
type:f 0 100
```

```
property:  
id:reprEnergy  
name:Reproduction energy  
type:f 0 10000  
group:Energy  
help:Creature energy required to produce an offspring
```

```
property:  
id:e_meta  
name:Idle metabolism  
type:f 0 1  
group:Energy  
help:Each creature consumes this amount of energy in one time step
```

...

Key functions

```
global foodenergywaiting;

function onExpDefLoad()
{
    // define genotype and creature groups
    GenePools.clear();
    GenePool.name="Unused";
    Populations.clear();
    Population.name="Creatures";
    Population.nnsim=1;
    Population.enableperf=1;
    Population.colmask=4;
    Populations.addGroup("Food");
    Population.colmask=5;
    Population.nnsim=0;
    Population.enableperf=0;

    // initialize experiment parameters
    ExpProperties.p.mutate=0.5;
    ExpProperties.reprEnergy=1001;
    ExpProperties.e.meta=0.1;
    ExpProperties.feedrate=0.2;
    ExpProperties.feede0=200;
    foodenergywaiting=0.0;
}
```

```
function onExpInit()
{
    Populations.clearGroup(0);
    Populations.clearGroup(1);
    foodenergywaiting=0.0;
    var cr=Populations.createFromString(
        "... (initial genotype) ...");
    cr.name="Initial creature";
    place_randomly(cr);
}

function onStep()
{
    foodenergywaiting += ExpProperties.feedrate;

    if (foodenergywaiting > ExpProperties.feede0)
    {
        addFood();
        foodenergywaiting=0.0;
    }

    if (Populations[0].size==0)
    {
        Simulator.print("no more creatures");
        Simulator.stop();
    }
}
```

Creature management

```
function onCreaturesBorn()  
{  
  Creature.idleEn=ExpProperties.e_metal;  
  Creature.energy0=1000;  
  Creature.energy=Creature.energy0;  
}  
  
function onFoodCollision()  
{  
  var e=Collision.Part2.ing;  
  
  Collision.Creature1.energy_m=  
    Collision.Creature1.energy_m+e;  
  
  Collision.Creature2.energy_p=  
    Collision.Creature2.energy_p+e;  
}
```

```
function onCreaturesStep(cr)  
{  
  if (cr.energy>=ExpProperties.reprEnergy)  
  {  
    var newcreature;  
    if (Math.rnd01 > ExpProperties.p_mutate) // exact co  
    {  
      GenePools.getFromCreatureObject(cr);  
      newcreature=Populations[0].createFromGenotype();  
    }  
    else  
    { // mutation  
      GenePools.getFromCreatureObject(cr);  
      GenePools.mutate();  
      newcreature=Populations[0].createFromGenotype();  
    }  
    if (newcreature!=null)  
    {  
      newcreature.energy0=cr.energy/2;  
      newcreature.energy=newcreature.energy0;  
      newcreature.moveAbs(  
        cr.pos_x+(cr.size_x-newcreature.size_x)/2,  
        cr.pos_y+(cr.size_y-newcreature.size_y)/2,  
        cr.pos_z+(cr.size_z-newcreature.size_z)/2);  
      cr.energy=cr.energy/2;  
    }  
  }  
}
```