

Private and Population-Level Parameters for Hybridization Simulations

All Populations used in the simulations usually have these parameters:

BirthDeath: b0, d0, theta

Fertility: min age, max age, interbirth

OldAge: max age

WeightedMove: move probability

NPPCapacity: NPP->CC ramp function, water bonus, coastal bonus

Generally, all these parameters are population-level parameters; for the Hybridization-Simulations some of them are changed to individual parameters.

There are three main features used in the current implementations for the Hybridization-Simulations:

Corrected Birth Probability

Every agent has its hybridization value, and a saddle function defined by a population-level parameter **HybMinProb** is used to calculate a correction factor for the birth-probability from the parents' hybridization:

$$p_{\text{corr}} = a + s * (h_{\text{Mother}} - 0.5) * (h_{\text{Father}} - 0.5)$$

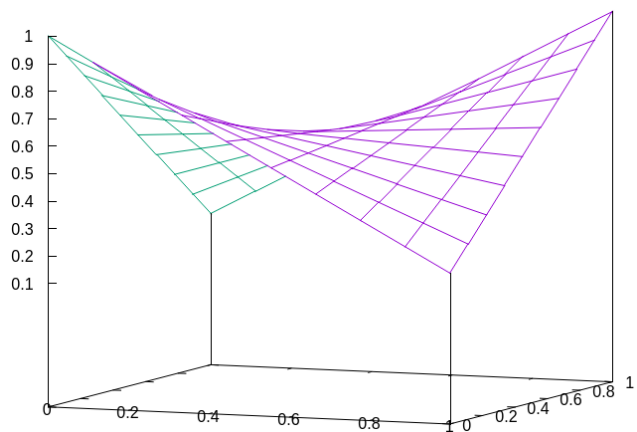
where

$$a = 2 * (1 - \text{HybMinProb})$$

$$s = 1 - (1 - \text{HybMinProb})/2$$

$$p_{\text{corr}}(0,0) = p_{\text{corr}}(1,1) = 1$$

$$p_{\text{corr}}(0,1) = p_{\text{corr}}(1,0) = \text{HybMinProb}$$



This factor p_{corr} is multiplied to the birth probability, which is in turn used to determine whether a birth takes place or not. The offspring's hybridization is currently calculated as the average of the parents' hybridization values, but could be changed to a recombination-dependent calculation.

Private Parameters

The parameters for **BirthDeath**, **Fertility**, **OldAge** and **WeightedMove** are private, i.e. each agent has a complete set of these values which are used for the various computations. The parents' parameter values can be combined arbitrarily to produce the child's parameter value. For example, calculating weighted averages based on the hybridization, or use the parameter values of one parent only, etc.

Dual NPP Parameters

There is a set of NPP population-level parameters describing "pure" sapiens and one set describing "pure" neanderthals. To determine the carrying capacity of a cell, the calculation is done once with the sapiens parameters and once with the neanderthal parameters. The actual carrying capacity the child sees is the weighted average of the two values.

Currently there are implementations for 3 different models for the Hybridization simulations, which use some or all of the features described above: “BirthProb Only”, “Partially Private”, “NPP-Mix”.

The following table gives an overview of which features are used by these implementations

	Corrected Birth Probability	Private Parameters	Dual NPP Parameters
“BirthProb Only”	Yes	No	No
“Partially Private”	Yes	Yes	No
“NPP-Mix”	Yes	Yes	Yes

I think for a system as discussed via skype we could use the “NPP-mix” implementation, where the private parameters are inherited from one parent (instead of mixing them).