**Modeling Notebook notes for TRACE – BLT Model**

This is not a TRACE document, but a TRACE notebook. General guidelines for TRACE are found on [Grimm et al 2014](http://dx.doi.org/10.1016/j.ecolmodel.2014.01.018) (structure on Table 1).

Interface gráfica do usuário, Diagrama

Descrição gerada automaticamente com confiança média

|  |  |
| --- | --- |
| **TRACE element/MN entry tag** | **MN keyword** |
| 1. Problem formulation | Model purpose; Research questions |
| 2. Model description | Model development; **Design decisions** |
| 3. Data evaluation | **Parameterization**; **Patterns** |
| 4. Conceptual model evaluation | **Conceptual design decisions** |
| 5. Implementation verification | **Debugging**  Software verification/Testing  Usability tools design |
| 6. Model output verification | Output verification/Goodness- of-fit **Calibration**; Tests on environmental drivers |
| 7. Model analysis and application | Sensitivity analysis; Uncertainty analysis Robustness analysis; Simulation experiment |
| 8. Model output corroboration | Output corroboration/**Validation** |

Implementation 1 refers to Step length + turning angles

implementation 2 refers to Resource visitation rules

implementation 3 refers to Territoriality

**Examples of topics to be included each day and a brief description:**

/purpose

To validate a simulation model of BLT dispersing seeds

/data

Check .xlsx

/verification

/calibration

/calibration/direct parametrization

Obtaining parameter values directly from the literature or experts

/calibration/inverse parametrization

Obtaining parameter values inversely by calibrating the model to observa- tions

/sensitivity analysis

/sensitivity analysis/submodels/energy

/sensitivity analysis/local

Varying one parameter at a time

/sensitivity analysis/global

Varying several or all parameters over their whole ranges

/alternatives

\_\_\_\_\_\_\_

\_\_\_\_\_\_\_

**2. Model description**

**February 04th 2022**

Discussion with Ronald about implementing turning angles and step lengths. We realized we need to check the empirical data a little bit further in order to implement a rule based on sl and ta in the model. I decided to check in [a published ABM](https://doi.org/10.1111/oik.07431) how it was done. Turning angles was implemented with von Mises distribution base on rvm-function from {CircStats} package, however I didn’t have a code for step length. Other two papers did similar implementations (Raghunathan et al 2020 and Gazagne et al 2020) with an HMM model but used also a gamma distribution to step lengths. With this I’ve checked the “Check if the impl. 1 needs model parameters” task. The answer is: Yes, it does. Gamma distribution for step lengths and von Mises distribution for turning angles. **For DPL and Path Twisting: don't know yet**.

After meeting with Ronald at 13:30 we decided to first have the nlrx workflow (to check results after running experiments) ready before making an implementation. Most importantly, we discussed more about step length/turning angles relation, I showed him the results from the ATBC and runned emptyspace analysis, explaining again the idea of taking time windows of resources and clumping measurements to make the home range emerge together with a DPL and Path Twisting based on a random point process specifying the tree aggregation. For last, I explained to him why I didn’t believe tamarins were going after resources and cited the mechanistic home range/central place forager hypothesis.

**February 09th 2022**

*Keywords: Model development; Design decisions*

From Feb 07th to now I’ve been troubleshooting Mayara’s code to match with the nlrx workflow. Two problems have risen:

1. Model output with nlrx to set workflow with R: even though it works with the Wolf-Sheep model, when I try with the BLT model, the px and py positions of the simulations don’t come out with **unnest\_simoutput()** function. Alternatively, [the nlrx vignette (see Step 3) explaining how to take manual output](https://docs.ropensci.org/nlrx/articles/manual-output.html) and relate to the nlrx object requires a reporter in NetLogo code, but when I insert it as a global, NetLogo gives me an error saying I should define it.
2. to-sleeping-trees procedure does not work when using the code for empirical sleeping trees. Thus I’ve created a Chooser **“sleeping-trees-scenario”** of whether simulated sleeping and resting trees are used or not (but only the code for simulated sleeping/resting trees work -> check **to-sleeping-trees** and **search-sleeping** procedure and sleeping-trees-here object is unused)

I also did the following:

* settled a **“tree-scenario”** Chooser for selecting which .shp with feeding trees was being used as input;
* settled an /runtime directory in model interface.

**February 15th 2022**

On the Feb 09th I started receiving this “Exception in thread "main" java.lang.IllegalStateException: unknown NetLogo version: NetLogo 6.2.0”. Ronald and I tried setting a stop condition because sometimes the model would run without stopping. Then I had to leave, and Ronald tried debugging the nlrx code. He was able to run the Mayara’s (the version he had) model, but mine was giving the following error together with same thread:

Exception in thread "main" Expected a literal value. at position 0 in

at org.nlogo.core.Fail$.exception(Fail.scala:27)

at org.nlogo.core.Fail$.exception(Fail.scala:25)

at org.nlogo.core.Fail$.exception(Fail.scala:23)

at org.nlogo.parse.LiteralParser.readLiteralPrefix(LiteralParser.scala:83)

at org.nlogo.parse.LiteralParser.getLiteralValue(LiteralParser.scala:33)

at org.nlogo.parse.CompilerUtilities$.$anonfun$readFromString$3(CompilerUtilities.scala:22)

at org.nlogo.parse.CompilerUtilities$.$anonfun$numberOrElse$1(CompilerUtilities.scala:37)

at scala.util.Either$RightProjection.getOrElse(Either.scala:665)

at org.nlogo.parse.CompilerUtilities$.numberOrElse(CompilerUtilities.scala:36)

at org.nlogo.parse.CompilerUtilities$.readFromString(CompilerUtilities.scala:22)

at org.nlogo.fileformat.LabLoader.$anonfun$readProtocolElement$4(LabLoader.scala:70)

at scala.collection.TraversableLike$WithFilter.$anonfun$map$2(TraversableLike.scala:874)

at scala.collection.Iterator.foreach(Iterator.scala:943)

at scala.collection.Iterator.foreach$(Iterator.scala:943)

at scala.collection.AbstractIterator.foreach(Iterator.scala:1431)

at scala.collection.IterableLike.foreach(IterableLike.scala:74)

at scala.collection.IterableLike.foreach$(IterableLike.scala:73)

at scala.collection.AbstractIterable.foreach(Iterable.scala:56)

at scala.collection.TraversableLike$WithFilter.map(TraversableLike.scala:873)

at org.nlogo.fileformat.LabLoader.readEnumeratedValueSetElement$1(LabLoader.scala:66)

at org.nlogo.fileformat.LabLoader.$anonfun$readProtocolElement$5(LabLoader.scala:77)

at scala.collection.immutable.List.flatMap(List.scala:366)

at org.nlogo.fileformat.LabLoader.valueSets$1(LabLoader.scala:74)

at org.nlogo.fileformat.LabLoader.readProtocolElement(LabLoader.scala:94)

at org.nlogo.fileformat.LabLoader.$anonfun$apply$1(LabLoader.scala:45)

at scala.collection.immutable.List.map(List.scala:293)

at org.nlogo.fileformat.LabLoader.apply(LabLoader.scala:45)

at org.nlogo.fileformat.LabLoader.apply(LabLoader.scala:30)

at org.nlogo.fileformat.LabFormat.load(NLogoLabFormat.scala:37)

at org.nlogo.fileformat.LabFormat.load$(NLogoLabFormat.scala:35)

at org.nlogo.fileformat.NLogoLabFormat.load(NLogoLabFormat.scala:40)

at org.nlogo.headless.BehaviorSpaceCoordinator$.$anonfun$selectProtocol$1(BehaviorSpaceCoordinator.scala:34)

at scala.Option.map(Option.scala:230)

at org.nlogo.headless.BehaviorSpaceCoordinator$.selectProtocol(BehaviorSpaceCoordinator.scala:32)

at org.nlogo.headless.Main$.runExperiment(Main.scala:23)

at org.nlogo.headless.Main$.$anonfun$main$1(Main.scala:12)

at org.nlogo.headless.Main$.$anonfun$main$1$adapted(Main.scala:12)

at scala.Option.foreach(Option.scala:407)

at org.nlogo.headless.Main$.main(Main.scala:12)

at org.nlogo.headless.Main.main(Main.scala)

**Error in util\_gather\_results(nl, outfile, seed, siminputrow) :**

**Output file is empty - simulation aborted due to a runtime error!**

**Make sure that parameter value definitions of the experiment are valid and the model code is running properly!**

So I went to the model (BLT\_model\_developm.nlogo) which I did the last implementations and realized that the model was breaking (running eternally) sometimes it was initiated. Then I checked the go, move-monkeys, run\_days and next\_day procedure. I did the following:

* go: nothing
* move-monkeys: moved in the “timestep > simulation-time” condition from the to go procedure and broke the multiple ifelses with the ‘if timestep < simulation-time’. I also took out from the setup the ‘if timestep = simulation-time’ condiditon (it was before the move-monkeys) and included it in the move-monkeys procedure
* run\_days: blocked the output code
* next\_day: added a stop condition: if day > no\_days [ stop ]
* commented out all the output procedures

The model was not breaking anymore but it still didn’t work out in the nlrx.

After one or two hours trying to debug it (BLT\_model\_developm-debug.nlogo), I assessed [this issue in nlrx](https://github.com/ropensci/nlrx/issues/27) and realized that the categorical variables should have a double quote like this:

constants = list( 'tree-scenario' = "\"trees\_all\_2\"",  
 'sleeping-trees-scenario' = "\"simulated\"" )

Now the problem is: I still can’t take pxcor and pycox coordinates. Maybe because I was doing one run with only a simple design? But how, if the tamarin run for days and 108 steps per day in different pxcor and pycor? The solution I found was setting up the number of runs in nlrx (and also in the model interface) to 1, and then specifying more runs in “repetitions” on the @experiment design

Milles et al 2020 model had an export-view function linked to an interrupter which allowed him to export the view of the run when it ended. I implemented it in a separate model (BLT\_model\_2022-developm\_issue-print-final-landscape.nlogo) but I was not able to see the tracks tamarins did. I let it like that for now.

Fiquei de fazer o cabeçalho dos arquivos output (que nem em Milles et al 2020) e de lembrar que diabos é aquele número no início de cada arquivo (o Ronald tinha descoberto já).

By the start of the night, I reassessed the move-monkeys code related to routine (the one I changed the ifelse condition in timestep < simulation time) and when I used the same code as Mayara I had a model break (infinite running). I could not understand what it was, but I think there lacks a stop condition.

I have also noticed that the tamarin was moving much less than I was used to see in the model, but it might be because I’m restricting the no\_days. -> Nope. It was because I had commented out the ‘let n random 100’ argument from the ‘search-sleeping-tree’ procedure. This is basically what moves tamarins.

**Left to fix/decide:**

* (**SOLVED**) Why the output of seeds and sleeping, resting and feeding trees text files contain information only for the last ran day even though the Mayara model works perfectly.

**February 16th 2022**

I have realized that Mayara’s model does not output seed and trees location by day. The problem was that I had copied the code ‘file-write day’ and ‘file-write timestep’ into the procedures ‘write-seeds’, ‘write-sleep’, ‘write-rest’ and ‘write-trees’.

Thus, I proceeded to add a header on each of those files and I also changed it to .csv

While running Mayara’s model I also realized that the simulations are faster when the locations of each day are written in a different file

I was succeed in extracting the pxcor and pycor from each agent by unnest\_simoutput() and proceeded with the spatial plot vignette. With this plot (one run of I don’t know how many days) it was possible to visualize that the rule of creating sleeping sites is not working properly (too many sleeping sites, a lot of them in the matrix):

Tela de jogo de vídeo game

Descrição gerada automaticamente com confiança média

**Left to fix/decide:**

* Check if using different files for each day of monkey locations increase velocity of the model run
* go/run\_days/next\_day procedures have two stop conditions. I think this is breaking the execution from nlrx
* include day in the nlrx simoutput
* gif animation and sequential plots not working: try to include an informative unique variable for each run (seed-number is not informative at all) -> patches explodindo
* Make the model run with the empirical sleeping sites

**March 04th 2022**

**TRACE Tag: 2. Model development**

**Keyword: Design decisions**

Me and Ronald had a meeting and a very nice discussion about:

1. **Theoretical landscapes of Tierry et al 2 022**

* We basically ignored this landscape generation because it is useful for the simulations but not for the implementations. Thus I scheduled to discuss it in other date;
* I wrote a ‘contact Vincenot’ task in Google Calendar to discuss about manually defining landscapes (as Vincenot itself and Tierry did).

Furthermore, my questions for discussion were:

- Não explica se as paisagens foram feitas "na mão" (apesar de parecerem ter sido feitas)

- Destaca que é modelado apenas o seed rain, sendo que seedling recruitment é outro processo: "seed dispersal patterns may not be accurate indicators of recruitment patterns (Schupp and Fuentes 1995)"

1. **The methods and model of memory + resource of Ranc et al 2022.** Ronald gave the following directions and suggestions:

* Play around with working memory (how fast they forget trees visited)
* Spatial memory is fixed at the moment
* RSF is not so interesting for the moment
* GA for the energy model to maximize DPL. ONLY AFTER GA RESULTS we can change model rules/implementationss. This allows for us to say that there's something in the structure
* IF there is no combination of energy variables that reproduces the expected patterns (DPL, etc), we can implement other rules
* BUT FOR THIS I should finish the R workflow, set all variables defined manually as slider
* Compare model diagrams (Mayara and Ronald)

Furthermore, my questions for discussion were:

- Usa Wasserstein metric para comparar diferentes distribuições de NSD, step lenght, turning angles e number of revisits, and associated times since last visit.

R: Ronald suggested me to use other patterns (such as step length, turning angles and number of revisits), but I didn’t comment about using the Wasserstein metric

- Is Fig 4 enough to say that simulations based only on resources are going to underestimate tendencies of directional movement?

R: I didn’t ask this one

- Revisitation is important -> they cite Bracis et al 2015 (othr publication of the author of recurse package) but don't use the package

R: Does not matter

- What would be equivalent to the deltaAIC approach they did?

R: I don’t remember what Ronald said about this

- Conspecific avoidance can't be discarded from our model

R: ok

- What this next paragraph helps with Parameterization 1 in terms of paramtrization/validation? "In this study, we characterised the biological drivers of fine-scale behavioural decisions through the fitting of a mechanistic movement model to empirical trajectories, and evaluated resulting predictions of space- use patterns. Although challenging, this approach is appealing because the space- use pattern itself is not fitted to data, but rather arises as an emergent property from the underlying movement process (Potts & Lewis, 2014)"

R: See the first topic of Wasserstein metric

- Where Resource Selection Functions/Habitat preference goes into the tamarin model? In your opinion, should it? Think about Suzano movements and also into Taquara group that uses a wet part of the home range which was not sampled

R: I showed him the cases where tamarin were using other parts of the home range but he said that RSF is not so interesting for the moment

**Decision for the BLT Model:**

* **GA for the energy model to maximize DPL. ONLY AFTER GA RESULTS we can change model rules/implementationss**. This allows for us to say that there's something in the structure
* **IF there is no combination of energy variables that reproduces the expected patterns (DPL, etc), we can implement other rules**
* **BUT FOR THIS I should finish the R workflow**
* **For this, set all variables defined manually as chooser. This also includes the timevisit variables** (which represents working memory, or how fast they forget trees visited)
* **Compare model diagrams (Mayara and Ronald)**

For this, I created a new model version called “**BLT\_model\_2022-developm\_ga.nlogo**“ from the model “BLT\_model\_2022-developm.nlogo”

**Left to fix/decide:**

* I’ve got an error with tree\_current/tree\_pot\_list for the first time:

Interface gráfica do usuário, Texto, Aplicativo

Descrição gerada automaticamente

* As far as I understood the workflow through the implementation process will be: 1) Implement rule -> 2) run GA for parameters > 3) See if the values reach the desired values (DPL for example)
* Remove sleeping sites being created out of nowhere
* Check Monthly trees .shp
* Print activity in the output file (activity bugdet)
* Check .shp from github not working in Ronald’s computer

**March 09th 2022**

**(continuing from March 04th)**

**Files:** BLT\_model\_2022-developm\_ga.nlogo

I reached Ronald for a briefing meeting and I decided to do this for the day:

1) model diagram

2) get .shp of sleeping, feeding and resting sites separetely

2) remove simulation trees

3) Do First R workflow to verification

4) Check memory procedures (forget\_trees, forget\_seeds procedure)

When possible - See why commenting "action" in in the model makes

And then the GA in the next day

He would take a look in the optimization within

1. **Model diagram**

Extraordinary fail to understand how to put loops in the diagram.

[Done on 17/03/22 but I still had some questions]

1. **get .shp of sleeping, feeding and resting sites separately**

[Done one 16/03/22]

1. **remove simulation trees**
2. **Do First R workflow to verification**

[Done on 11/03/22 (but still the kernel analysis were not good)]

1. **Check memory procedures (forget\_trees, forget\_seeds procedure)**

[specific implementation on Implementation 2 designed by Ronald (check Model\_development/memory folder)]

**Left to fix/decide:**

* [DONE; Ronald say it is wrong to renew the energy] energy when the day starts: ‘set energy start-energy’ commented out -> tamarins accumulate a lot of energy

**March 10th 2022**

Context: BLT Model Meeting 1 with Laurence, Eckhard and Ronald is going to happen tomorrow. I started doing the R workflow for analysis of pattern validation (on “Model\_analysis/Workflow”) to present the contrasting results of runs with input\_forget low and high values. The results of this comparison are in “Model\_development/BLT\_Model\_Meeting1/00\_Input\_forget.pdf”

**March 11th 2022 – Day of BLT Meeting 1 (w/ Eckhard and Laurence)**

I showed Ronald how my code version (Mayara’s) was different from the one he was running. He realized that there was completely different stuff (for example, not using travel\_speed to move, etc). THEFEFORE THE DAY THAT I WAS LISTENED ABOUT THE FACT THAT MAYARA DID SOME REALLY WEIRD STUFF). Thus, I asked for his code (the model he was using) to implement my modifications into. He didn’t send me. We had a very nice discussion and **that’s what we are doing next week (14/03):**

**1) Troubleshoot code + flowchart** (to identify where else Mayara changed the code and if the code we were running makes sense)

**2) Distances to next feeding tree** (step length output (see “00\_Input\_forget.pdf”) is too small.

**3) GA/recurse package** -> Ronald quer explorar os padrões de step lenght e turning angles em escalas maiores de tempo por algum motivo. Disse que podemos fazer tanto o GA quanto analisar métricas de revisitação com o pacote ao mesmo tempo. Ele também concordou e meu tentar contatar o Ranc (autor da publicação de memória)

**Let to fix/decide:**

* Seminar dates
* [Done on 16/03/22] month shapefiles with sleeping, feeding and resting trees in different files
* Phenology values (does it stay?)

**March 14th 2022**

* Finished doing model diagram with Ronald but didn’t finish including the layers in the diagram to check the status and action each step
* Identified that the step when “night” is set does not change energy (thus it is wrong, it shouldn’t count as one timestep)
* Included a column “behavior” that matches the activity budget

**Let to fix/decide:**

* [Kind of done on 15/03/22 but still get tree target errors some times] clean run\_days/go
* clean set status/target etc (colocar tudo dentro dos procedures e tirar do loop)
* [?] colocar no workflow quando status, action (e se der tempo, tree\_target/tree\_current)
* checar se defecation ocorre somente antes do timestep 84 (depois de 3 pm)
* ifelse random ( 2 \* 5) > action time -> conferir quantas vezes eles podem executar a mesma atividade seguida e colocar como slider (APENAS FORAGING OU RESTING)
* Desimplementar "species-time"
* Include “idle” behavior
* Esse erro em to-feeding-tree

Interface gráfica do usuário, Texto, Aplicativo

Descrição gerada automaticamente

**March 15-17th 2022**

* Button go does not work if I click Run Days or Next Day before
* [Kind of done on 15/03/22 but still get tree target errors some times] Tried cleaning the go/next\_day procedures and it was a pain in the ass. I settled a ‘step’ procedure only for debug porposes but could not get rid of the next\_day/run\_days structure
* Continued having the run-time error from the previous day (but only sometimes)
* Error in sleeping procedure (target 0) when running ‘go’ after Run Days or Next Day -> set heading towards tree\_target -> I think I have solved by setting the tree\_target correctly in each procedure
* [solved -> check “Data/Guarei-Trees/Tree-points-unique.R”] Bug: sleeping-tree 1112 e 1113 são a mesma -> checar no shp -> acontece quando a árvore se repete nas observaçõe (linhas) do dataset que é transformado em .shp

Interface gráfica do usuário, Aplicativo

Descrição gerada automaticamente

* I did distinct shape files for the model. One with all trees (for all months and for each month, which is equal as what Mayara did) and separate shapes for all months or each month with sleeping, feeding or resting trees.
* Added sleeping-trees? interrupter in the interface (to don’t have to make distinct shapefiles for each type of tree)
* After selecting shape files, as there were few resources each month, the tamarins were dying and the model would not stop running the run\_days, so I did a stop condition that for some reason does not work:

if not any? monkeys [ stop ]

* I realized that for debugging purposes I should print the DPL in the end of the day

**Meeting with Ronald 17/03 afternoon:**

- [Done on 17/03, check “Data/Trees-Guarei/Tree-points-unique.R”) do shapefiles with all sleeping sites

- Path Dependence in Operational Research (pathways of modelling)

- [Fixed on 17/03, but the status is kind of wrong and we should look at it again] wrong behavior when going to resting tree

**Left to fix/decide:**

* Gráfico, Gráfico de dispersão

  Descrição gerada automaticamenteAfter I specified a distinct input for all sleeping trees (all-slp-trees? chooser), monkeys started to die earlier than expected. It looks like it is because of the sleeping sites because they keep changing sleeping-site in target:
* [18/03] Troubleshooting this problem with Ronald: the problem was something related to the position of the set tree\_target in the code, but I don’t remember where exactly
* REVIEW search-resting-defined (input de árvores de Descanso não deveria ser necessário)

**18/03/22**

[18/3] For debugging:

> let agentset [self] of sleeping-trees print agentset

We realized that an error was happening: the tamarins would forget all the potential trees thus the list would become empty, giving the error of 0 agentset. I did a graph ploting the length of the list and realized the problem was the *‘set tree\_pot\_list remove-item ( position [who] of tree\_current tree\_pot\_list ) tree\_pot\_list* of the *to-feeding-tree* procedure. If this is commented out, the tree\_pot\_list increases (surprisingly)

In the image below, basically, the forget\_trees procedure is commented out, but the memory procedures inside the move-monkeys procedure are not. The only thing that is running that influences the memory is the move-monkeys code with the tree\_ate\_list and tree\_pot\_list being changed inside the move-monkeys. Similar code inside feeding etc procedures are commented out.

* tree\_pot\_list increases more than the number of available trees
* **Interface gráfica do usuário, Aplicativo

  Descrição gerada automaticamente**tree\_mem\_list does not increase too much

Then that’s what happens:

Interface gráfica do usuário, Texto, Aplicativo

Descrição gerada automaticamente

Interface gráfica do usuário, Texto, Aplicativo

Descrição gerada automaticamenteThe problem was in this procedure, probably because it is being called 3 times in the same move\_monkeys loop.

Interface gráfica do usuário

Descrição gerada automaticamenteNow when it is taken out, that’s the result:

Interface gráfica do usuário, Texto, Aplicativo

Descrição gerada automaticamenteBut if I keep running, eventually I get this:

So, everything seems to be solved, but we came back to getting the error I was having with Ronald before.

**Left to fix/decide:**

1. Now, the memory increases a lot, not getting any asymptote
2. The tamarin does not move from a specific patch of trees -> probably related to the distance to the next feeding tree. This might be solved by the patch based implementation planned by Ronald

**March 21st 2022**

* Now the model puts 0 values in the tree\_mem\_list, resulting in a list like: [0 0 0 0 0 0 0]. Furthermore, this increases much slower than the version of 2022-03-11d
* Had to go back at the code and check where the problem was. Basically:

1. I had removed the set action-time and set status from inside the move monkey to inside the called procedure but positioned it wrong.
2. The procedure in yellow above didn’t work if I put it in the *forget-tree* procedures
3. I could not identify why the tree\_mem\_list was not getting bigger in the graph (pen-plot) -> it was too close to the red line, the seed\_mem\_list (they have the exact value)
4. For some reason, the tree\_pot\_list was getting bigger after a certain time even though the tamarin was barely moving [image below]

Gráfico

Descrição gerada automaticamente com confiança média

**March 22nd**

Conversei com o Ronald e expus os problmas dos dias anteriores. Concordamos em:

1) Fazer o flowchart focando no set status/tree\_target (como eu estava fazendo antes)

2) Primeiro iremos mexer no componente de memória (forget\_trees, diminuição e aumento das litas), depois o critério de seleção de árvores (que provavelmente vai entrar em step lengths)

* Changed input\_forget to step\_forget (makes more sense)
* step\_forget has to be dependent on the number of trees available, otherwise the pot\_list gets empty
* the pot\_list diminishes and receives the consumed trees back, but for some reason it keeps getting smaller

**6. Model output verification**

**18/03/22**

Preparing for the Genetic analysis next week, I realized we need the following to proceed:

1. Model not breaking
2. Prepare the nrlx setup again
3. Multi core run

**5. Implementation verification [?]**

**Jan/Feb (não coloquei a data)**

/extensions/gis

Got the code of nlrx running for Mayara’s model. Problem: It is generating one file for each day, thus it dumps everything in github

Files and code: [path](file:///D:\Data\Documentos\github\BLT_IBM-Model\Model_development)

Script: 00\_start-nlrx.R

Left to fix/decide:

* If I’m keeping model output as external files

/extensions/gis

**Feb 2nd 2022**

Debugged shapefiles of the three areas (Guareí, Suzano and Taquara) to initialize with a chooser + if condition.

Files and code: [path](file:///D:\Data\Documentos\github\BLT_IBM-Model\Model_development\gis-extension)

Included a procedure to check if turtles are inside the shapefile: ‘check-agent-in-fragment’

Tried looking how to scale the different shapes (scale parameter in gis extension). Didn’t find anything.

Left to fix/decide:

* Better .shp files to NetLogo (for Suzano and Taquara, too big)

**6. Model output verification/calibration**

**Day 1 – January 14th**

Six steps for making a good calibration (Railsback & Grimm 2012 Chap 20):

1. Identify a few good parameters

Important, uncertain and independent parameters. To know convincingly which parameters are important, we need sensitivity analysis. It is prudent to reconsider model calibration after this sensitivity analysis. A parameter that is highly uncertain but has little effect on results should not be used for calibration.

Decision for the BLT model: activity budget (e.g. time spent travelling?)

1. Choose Categorical vs. Best-Fit Calibration

Categorial gives a range of values (e.g. mean number of agents between 120-150), while Best-Fit Calibration gives single values (e.g. mean number of agents = 127).

Decision for the BLT model: Categorical seems the best option

1. Decide Whether and How to Use Time-Series Calibration

“If our model's purpose includes representing how results change over time (e.g., how long does it take the system to recover from some perturbation? How is the system affected by changes in its environment over time?), then it usually does make sense to use time-series calibration. But some ABMs (e.g., the woodhoopoe model of section 19.4.3) are intended to explain long-term average conditions, so they intentionally do not contain all the processes that cause the real system to change over time and use no input data to represent how the agents’ environment changes over time. In such cases, time-series calibration may not be useful or necessary.”

Decision for the BLT model: As I’ll only be focusing on the mean and sd numbers at the end of the runs, I believe I don’t need Time-Series Calibration

1. Identify Calibration Criteria

* Calibrate the model against all interested patterns
* Of apples and oranges: the observations need to match the same time and space scale
* There are kinds and kinds of variation: Calibrating variability measures (CV, sd) should be done carefully
* Often our data is inaccurate and we need to know if it is inaccurate by 10, 20 or 50%. Using uncertain data is OK and unavoidable: when the calibration patterns are more uncertain, we don't worry so much about matching them exactly, and we recognize that calibrated parameter values are less certain. But we need to have at least some idea how accurate or certain the observations are, so we know how much information they really contain
* We must specify how we will compare the observed patterns to model results. Of particular concern is how to calibrate several different kinds of model results at once: if we want to calibrate a model to reproduce the number, size, and wealth of agents, how do we decide between a set of parameters that reproduces number and wealth well but not size, and a parameter set that reproduces size and wealth well but not the number of agents?
* At the end of this step of defining calibration criteria, we should have a specific algorithm for quantifying how well a set of model results reproduces the selected observations

Decision for the BLT Model:

* Calibrate against SDD, DPL, Home range and Activity Budget
* (In order of importance) prioritize SDD, DPL, Home range and Activity Budget
* Algorithm: the mean (min and max too?) SDD, the Home range and Activity Budget evaluated after 30 running days and the DPL of each of these days compared to data of Guareí (PEMD and Suzano to be independent observations serving as validation?)

1. Design and Conduct Simulation Experiments

This calibration experiment executes the model many times, using combinations of values over the feasible range of all parameters. The results of this experiment will tell us what ranges of parameter values produce results that meet the calibration criteria. Steps:

* Select values for the non-calibration parameters and input data (if any) that represent the conditions (the same time period, environment, etc.) under which calibration patterns were observed.
* Define *parameter space* (It is usually good to include values that bound the range of feasibility) (see Figure 20.3)
* If the model is stochastic, analyze means seems right, but plot model results against parameter values [? p. 357]

Decision for the BLT model:

* Select all non-calibration parameters an input data from Guareí (using values that bound the range of feasibility, e.g. energy-from-fruit 0)

1. Analyze Calibration Experiment Results

* If your ABM does meet all the calibration criteria for one or more combinations of parameter values, then you can in fact complete this step and move on to the kinds of analysis we address in part IV
* What should you do if you cannot meet all the calibration criteria at once?

1. Screen for errors (code, submodels, etc)
2. If there isn’t any error, your model is too simple or too simple in the wrong ways, to reproduce the observed patterns you chose as calibration criteria. You could consider going back to the theory development stage and seeing if you can reproduce the observed patterns you chose as calibration criteria. You could consider going back to the theory development stage and seeing if you can improve the traits for agent behavior, and you could consider adding processes or behaviors that were left out the first time. But keep in mind that there are costs to adding complexity to your model, especially if it is not very clear what change needs to be made.
3. [Alternatively] It is very common for good modelers to keep their model simple instead of adding stuff to it until the model can reproduce all the calibration criteria. Keep in mind the overfitting issue: it can be risky to try too hard to make the model reproduce a limited set of observations. It may make sense to revise your calibration criteria so they are not as restrictive. If you choose not to revise the model to make it fit more of the calibration criteria, simply document your calibration results and the extent to which the model does not meet some criteria under your “best” parameter values, and your decision not to revise the model. Then, when you use the model to solve problems, keep in mind which results are less certain as indicated by the calibration experiment.

Decision for the BLT Model: I can’t make any decision right now, but the comment 3) seems very important. I’ll probably proceed if I find a set of values that meet the first 2 variables (SDD, DPL), and then Home Range size (not core home range as it is mostly not stable throughout time)

Left to fix/decide:

**March 21st, 2022**

**MN Keyword: 2. Model description**

Although scent marking behavior was already taken out from the model, this process was not documented. Reading Heymann ([2021](https://www.tandfonline.com/doi/full/10.1080/03949370.2021.2015454)), which cites Franklin et al. 2007 study on *Leontopithecus rosalia* assigning a function of marking to reducing predation at sleeping sites, and considering that we don’t have enough data to perform the *quasi experiment* scenario which fragmentation produces (n=4 sites), I think this is enough for us to disregard scent marking as a process that influences movement and seed dispersal.