
SAMPLING FROM THE BAYESIAN POSTER OF AN AGENT-BASED MODEL GIVEN PARTIAL OBSERVATIONS

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ABSTRACT

abstract

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1 Introduction

- Problem
 - Agent-based models are often used to explain how and why systems behave in the way that they do; i.e. ‘generative social science’ (Epstein, 1999).
 - With empirical models, we need to be able to find the most likely model configurations that might have generated the observations we have from the real system. In other words we need to find the model posterior.
- Dan: I'd like to discuss the difference in calibration v.s. finding specific trajectories
- (With simple models there might be a formal solution but this is rarely (if ever) the case with ABMs).
 - Typically this is done by running the model forward using different parameter values (calibration).
 - But this is very computationally expensive, even with modern, efficient calibration algorithms (Thiele, Kurth, & Grimm, 2014) as many attempts to find the ‘correct’ parameter configurations will fail, wasting resource that could have been better spent running models that may have generated realistic observations.
 - MCMC methods approach this problem by XXXX, but XXXX
- Solution
 - We propose a new MCMC method that works directly on ABMs.
 - Benefits
 - Computational efficiency - but why? Generates more plausible solutions?
 - In addition can find the most likely trajectories? (which may be interesting in their own right)
 - Experiments
 - We test the new method

what's the word??

Notes

- (Kennedy & O'Hagan, 2001) makes a good argument about the weaknesses of ‘traditional’ calibration

2 Related Research

NM to start on background section

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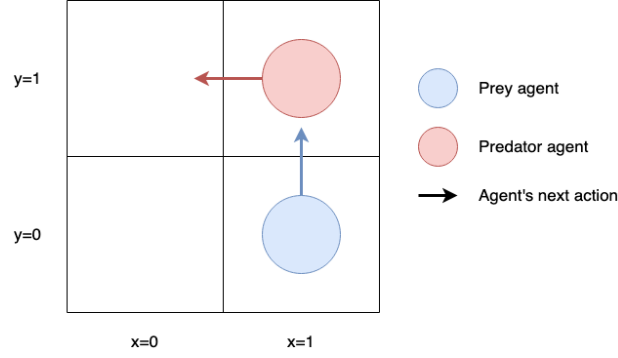


Figure 1: An simple predator-prey example.

3 ABMCMC

Suppose we have a timestepping ABM where agents have a finite number of possible internal states and a finite number of ways of acting on their world. Given this, we can define an ABM as:

- A domain of agent actions $\mathcal{A} = \{a_0 \dots a_n\}$
- A domain of agent states $\mathcal{S} = \{\psi_0 \dots \psi_m\}$
- An *agent timestep*, which is a computer program, $\pi(\psi, \Psi)$, where $\psi \in \mathcal{S}$ is the agent's state and $\Psi \in \mathbb{Z}^m$ is a (sparse) vector whose i^{th} element is the number of agents in state ψ_i at the start of the timestep. The program returns an action $a \in \mathcal{A}$ which defines the behaviour of the agent in a timestep. The program may make calls to a random number generator that returns a value $0 \leq r < 1$ with uniform probability, so its returned value may be non-deterministic and we write $P(\pi(\psi, \Psi) = a)$ to be the probability that the program will return action a , given inputs ψ and Ψ .
- An *action function*, $F : \mathcal{S} \times \mathcal{A} \rightarrow \mathbb{Z}^m$, which defines the effect of an action on the world, where $F(\psi, a)$ gives a (sparse) vector, Φ , whose i^{th} element gives the number of agents in state i that result from an agent in state ψ performing act a (this includes the final state of the acting agent).

As a very simple demonstration, we'll take a spatial predator-prey model where agents are either predator or prey moving about a 2-dimensional grid. In this case, the domain of actions are $\mathbb{A} = \{a_0 = \text{move-left}, a_1 = \text{move-right}, a_2 = \text{move-up}, a_3 = \text{move-down}, a_4 = \text{give-birth}, a_5 = \text{die}\}$, and the domain of agent states, \mathcal{S} , can be thought of as the class

```
class Agent {
    Boolean isaPredator
    Integer xPositiion
    Integer yPositiion
}
```

To simplify our methodological explanation further, we will use a worked example of a 2×2 predator-prey model, with a single predator and a single prey. See Figure 1.

It will be useful in the following to define an ordering of agent states. The ordering we choose isn't important but we choose to place agent a in the $(a.isaPredator * G^2 + a.yPositiion * G + a.xPositiion)^{th}$ position, where G is the size of the grid.

The timestep function for a predator/prey agent is shown in algorithm 1 (where we assume the grid has periodic boundary conditions so we don't need to worry about its edges). The action function encodes the results of the various actions. So, for example, the zeroth state, ψ_0 , corresponds to a prey ($isaPredator = 0$) at position ($yPositiion = 0, xPositiion = 0$), i.e. $\psi_0 = [0, 0, 0]$. If this agent moves right (i.e. performs action a_1) then the result is a prey at position (1,0) (i.e. in state $\psi_1 = [0, 1, 0]$) so $F(0, 1) = \langle 0, 1, 0 \dots \rangle$ etc.

The following illustrates all possible states in our simple 2×2 model, as per Figure 1, where ψ_2 represents the state of the prey and ψ_7 represents the state of the predator:

Algorithm 1 Timestep of a predator/prey agent

```

function TIMESTEP(agent, otherAgents)
  if agent.isaPredator then
    if RANDOM( ) <  $\beta_0$  and number of prey on neighbouring gridsquares of otherAgents > 0 then
      return give-birth ▷ more likely to reproduce when food is available
    end if
    if RANDOM( ) <  $\gamma_0$  then
      return die
    end if
  else
    if RANDOM( ) <  $\beta_1$  then
      return give-birth
    end if
    if RANDOM( ) <  $\gamma_1$  then
      return die
    end if
    if RANDOM( ) <  $\delta$  and number of predators on this or neighbouring gridsquares of otherAgents > 0 then
      return die ▷ been eaten by predator
    end if
  end if
  return move-left, move-right, move-up or move-down based on a call to RANDOM( )
end function
    
```

$\psi_0 = [0, 0, 0]$	
$\psi_1 = [1, 0, 0]$	
$\psi_2 = [0, 1, 0]$	state of the prey
$\psi_3 = [1, 1, 0]$	
$\psi_4 = [0, 0, 1]$	
$\psi_5 = [1, 0, 1]$	
$\psi_6 = [0, 1, 1]$	
$\psi_7 = [1, 1, 1]$	state of the predator

Let a model timestep consist of a matrix E whose elements $e_{\psi a}$ are the number of agents in state ψ that perform act a in this timestep. The matrix sparse; in our simple case all cells are empty except those representing the states and actions of the predator prey agents; (ψ_2, a_2) and (ψ_7, a_0) respectively:

$$E = \begin{matrix} & \begin{matrix} a_0 & a_1 & a_2 & a_3 & a_4 & a_5 \end{matrix} \\ \begin{matrix} \psi_0 \\ \psi_1 \\ \psi_2 \\ \psi_3 \\ \psi_4 \\ \psi_5 \\ \psi_6 \\ \psi_7 \end{matrix} & \begin{bmatrix} 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 1 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 \\ 1 & 0 & 0 & 0 & 0 & 0 \end{bmatrix} \end{matrix}$$

Now let a model trajectory be a tensor, T , whose elements, $T_{\psi a}^t$, are the number of agents in state ψ that perform act a in timestep t . I.e. in the first timestep, $t = 0$, then $T_{\psi a}^0 = E$, and in subsequent timesteps the trajectory consists of multiple E matrices, one for each timestep.

A trajectory must satisfy a number of constraints in order to be a possible trajectory of an ABM. Since the elements of a trajectory are counts, they must all be non-negative integers, we'll call this the *non-negative integer constraint*

$$\forall t, \psi, a : T_{\psi a}^t \geq 0, T_{\psi a}^t \in \mathbb{Z} \quad (1)$$

A trajectory, $T_{\psi a}^t$, must also be *continuous* which means that the number of agents in each state at the end of timestep $t - 1$ must be the number of agents in each state at the beginning of timestep t . This can be expressed in terms of the

continuity constraints:

$$\forall t \in 1 \dots n : \forall \phi : \sum_{\psi, a} F(\psi, a)_{\phi} T_{\psi a}^{t-1} - \sum_a T_{\phi a}^t = 0 \quad (2)$$

where the trajectory runs from $t = 0$ to $t = n$. If a trajectory satisfies constraints 1 and 2 we say that it is *valid*.

If we let Ψ_{ψ}^t be the number of agents in state ψ at the beginning of timestep t then the prior probability of a trajectory is

$$P(T) = \begin{cases} P(\Psi_*^0 = \sum_a T_{*a}^0) \prod_{\psi, t} P(T_{\psi*}^t | \Psi_*^t = \sum_a T_{*a}^t) & \text{if } T \text{ is valid} \\ 0 & \text{otherwise} \end{cases}$$

where $P(\Psi_*^0)$ is our prior beliefs about the state at time $t = 0$, and we use $*$ as an index when we want to emphasise that the index remains unbound.

For example, in the predator/prey model, we may know the probability of finding a predator or prey at each gridpoint at $t = 0$, may not know the exact positions of each agent so the prior would be a product of Poisson distributions

$$P(\Psi_*^0) = \prod_{\psi} \frac{\lambda_{\psi}^{\Psi_{\psi}^0} e^{-\lambda_{\psi}}}{\Psi_{\psi}^0!}.$$

However, we know the probability that a single agent in a given state will perform an action given the state of the other agents, this is defined by the agent timestep function, so the joint probability that Ψ_{ψ}^t agents will perform actions $T_{\psi*}^t$ is the multinomial distribution

$$P(T_{\psi*}^t | \Psi_{\psi}^t) = \Psi_{\psi}^t! \prod_a \frac{P(\pi(\psi, \Psi_{\psi}^t) = a)^{T_{\psi a}^t}}{T_{\psi a}^t!}.$$

So the prior probability of a trajectory is

$$P(T) = \begin{cases} P(\Psi_*^0 = \sum_a T_{*a}^0) \prod_{t, \psi} (\sum_a T_{\psi a}^t)! \prod_a \frac{P(\pi(\psi, \sum_b T_{\psi b}^t) = a)^{T_{\psi a}^t}}{T_{\psi a}^t!} & \text{if } T_{\psi a}^t \text{ is valid} \\ 0 & \text{otherwise} \end{cases}$$

Suppose we also have a set of noisy, aggregate observations, Ω , that have a likelihood function $P(\Omega | T_{\psi a}^t)$. In our running example, let's suppose we periodically note down when and where we see footprints of predator and prey. Our aim is to generate a number of samples from the posterior distribution

$$P(T_{\psi a}^t | \Omega) \propto P(\Omega | T_{\psi a}^t) P(T_{\psi a}^t)$$

We assume that observations are independent given the trajectory and that each individual observation in Ω can be simulated by a computer program $\omega(T_{\psi a}^t)$ which may make calls to a random number generator. The probability, $P(\omega(T_{\psi a}^t) = v)$, that the program returns a value v defines the likelihood of observing value v upon making observation ω on trajectory $T_{\psi a}^t$. i.e. $\omega(T_{\psi a}^t)$ is just a simulation of the observation, which is usually quite easy to express as a computer program and to compute.

In the case of the predator prey model, let's suppose that at the end of every day we go out to a fixed set of inspection sites and check for footprints. Suppose that if predators were present in the grid-square containing an inspection site on that day, there's a 50% chance we'll see a footprint, and if prey were present, there's a 25% chance. The observation function for a site is shown in algorithm 2

Given this

$$P(\Omega | T_{\psi a}^t) = \prod_{(\omega, v) \in \Omega} P(\omega(T_{\psi a}^t) = v)$$

Also, for notational convenience, and without loss of generality, we express our prior beliefs in terms of a set of observations, Ω_0 , at $t = 0$, so that $P_0(\sum_a T_{\psi a}^0) \propto \prod_{(\omega, v) \in \Omega_0} P(\omega(T_{\psi a}^0) = v)$ (this also allows any priors that aren't at $t = 0$). The posterior can now be written as

$$P(T_{\psi a}^t | \Omega) \propto \begin{cases} \prod_{(\omega, v) \in \Omega} P(\omega(T_{\psi a}^t) = v) \prod_{t, \psi} (\sum_a T_{\psi a}^t)! \prod_a \frac{P(\pi(\psi, \sum_b T_{\psi b}^t) = a)^{T_{\psi a}^t}}{T_{\psi a}^t!} & \text{if } T_{\psi a}^t \text{ is valid} \\ 0 & \text{otherwise} \end{cases} \quad (3)$$

where Ω now includes any priors.

In many practical applications, sampling from this posterior is difficult because it has zero probability for the vast majority of trajectories (i.e. most trajectories are invalid, contain an impossible action or are refuted by the observations). This is true of our predator/prey model, for example. Even though we can generate valid trajectories that fit the prior by

Algorithm 2 Function for observing predator/prey footprints at a site

```

function OBSERVEFOOTPRINTS(T)                                ▷ T is the ABM trajectory
    t ← time of observation
    x ← x-position of observation
    y ← y-position of observation
    G ← size of grid
    ψ ← G2 + Gy + x                                           ▷ state of predator in this gridsquare
    φ ← Gy + x                                                  ▷ state of prey in this gridsquare
    Fpred ← FALSE                                              ▷ did we observe predator footprints?
    Fprey ← FALSE                                              ▷ did we observe prey footprints?
    if ∑a Tψat > 0 and RANDOM( ) < 0.5 then
        Fpred ← TRUE
    end if
    if ∑a Tφat > 0 and RANDOM( ) < 0.25 then
        Fprey ← TRUE
    end if
    return Fpred, Fprey
end function
    
```

simply performing a forward execution of the model from an initial state drawn from the prior, if the trajectory has no predator or prey at a time and place that we observed footprints, then the observations refute the trajectory and the probability falls to zero. It doesn't take many observations until the probability of randomly choosing a trajectory that fits the observations becomes very small indeed. So simple techniques such as rejection sampling are not practical.

In this paper we'll use the Metropolis-Hastings algorithm to generate samples. However, this method requires a proposal function which randomly generates the next sample given the current one. However, given the current sample, we still have the problem of finding a trajectory with non-zero probability. For example if we generate a new trajectory by perturbing one or more elements of the last sample at random (a common strategy with Metropolis-Hastings), it's very unlikely that we'll end up with a trajectory that is valid, contains only possible actions and satisfies the observations. So, the proposed next sample would almost certainly be rejected and we'd probably end up stuck on the first sample until we grew old.

4 Approximating the support of the posterior

To solve this problem we'd like to constrain our proposal function to the support of the posterior, $\text{supp}(P(T_{\psi a}^t | \Omega))$ (i.e. the set of trajectories that have non-zero probability).

If we let

$$\pi'(s, \xi, T_{\psi a}^t) = \pi(\xi, \sum_b T_{\phi b}^s)$$

Then from equation 3

$$\text{supp}(P(\cdot | \Omega)) = \bigcap_{(\omega, v) \in \Omega} \text{supp}(P(\omega(\cdot) = v)) \cap \bigcap_{t, \psi, a} (\text{supp}(P(\pi'(t, \psi, \cdot) = a)) \cup \{T_{\xi b}^s : T_{\psi a}^t = 0\}) \cap \{T_{\psi a}^t \mid T_{\psi a}^t \text{ is valid}\} \quad (4)$$

i.e. in order for the posterior to be non-zero, all the observation likelihoods must be non-zero, the probability of each agent's action at every timestep must be non-zero and the trajectory must be valid (note that we've dropped the exponent in the agent action term as it doesn't affect the support).

The first two terms in equation 4 consists of the supports of computer programs whose inputs are ABM trajectories and whose outputs are given, i.e. the set of trajectories that, when passed to a computer program, would produce a given output.

Calculating the support of a computer program for a given output is, in full generality, NP-complete² but it is possible to use a technique known as *abstract interpretation* (Cousot & Cousot, 1977) to efficiently calculate a superset of the support. So, given a computer program ρ , we can calculate a set $\mathcal{P}(\rho, v)$ such that

$$\text{supp}(P(\rho(\cdot) = y)) \subset \mathcal{P}(\rho, v)$$

Tools to perform abstract interpretation already exist (e.g. PAGAI (Henry, Monniaux, & Moy, 2012)) and are used widely in applications such as the verification of safety critical systems (Blanchet et al., 2003) and in practice $\mathcal{P}(\rho, v)$

²Consider, for example, a program that accepts an assignment of variables to truth values, and returns true if that assignment satisfies a Boolean formula. Deciding whether the support of this program, given that it returns true, is empty or not is equivalent to solving the Boolean satisfiability problem, which is known to be NP-complete (Cook, 1971)

is often reasonably tight (i.e. most members of $\mathcal{P}(\rho, v)$ are in $\text{supp}(P(\rho(\cdot) = y))$). For our application we choose to express $\mathcal{P}(\rho, v)$ in terms of a set of linear inequalities on ρ 's inputs, this corresponds to the abstract domain of convex polyhedra (Cousot & Halbwachs, 1978) (Becchi & Zaffanella, 2018). Calls to the random number generator can be dealt with in the abstract domain by generating a new variable, r , that satisfies $0 \leq r < 1$ for each call to $\text{Random}()$. These can either be left in as ‘‘auxiliary’’ variables in the same way as slack variables, or removed as soon as the variable goes out of scope by finding the convex hull of the projection into a lower dimensional space (this can be done using the double description method (Motzkin, Raiffa, Thompson, & Thrall, 1953)).

If we replace the supports in equation 4 with their equivalent supersets we end up with a superset of the posterior

$$\text{supp}(P(\cdot | \Omega)) \subset \bigcap_{(\omega, v) \in \Omega} \mathcal{P}(\omega, v) \cap \bigcap_{t, \psi, a} (\mathcal{P}(\pi'(t, \psi, \cdot), a) \cup \{T_{\xi b}^s : T_{\psi a}^t = 0\}) \cap \{T_{\psi a}^t \mid T_{\psi a}^t \text{ is valid}\} \quad (5)$$

Constraining our proposal function to members of the superset in equation 5 instead of the true support won't affect the stationary distribution of the Markov Chain. If the proposal function happens to return a trajectory that isn't in $\text{supp}(P(\rho(\cdot) = y))$ then it will just be rejected. This is fine as long as we generate acceptable proposals at a reasonable rate.

The final term in equation 5, which requires the trajectories in the support to be valid, consists of the continuity constraints of equation 2, which are also linear constraints so can be treated along with the other constraints, and the non-negative integer constraints of equation 1 which we leave as separate constraints.

The union term in equation 5 can also be transformed to a set of linear constraints, but the way we do this depends on the type of ABM we're dealing with, so we defer the explanation until sections 5.1 and ...

Without loss of generality, we express inequalities as equalities by introducing slack variables so that

$$\sum_i c_i x_i \leq y$$

is equivalent to

$$\begin{aligned} \sum_i c_i x_i + \lambda &= y \\ \text{subject to } \lambda &\geq 0 \end{aligned}$$

All the constraints can now be gathered into a standard matrix form

$$\begin{aligned} AX &= B \\ \text{subject to } x_i &\geq 0 \\ x_i &\in \mathbb{Z} \end{aligned} \quad (6)$$

where the vector X contains the elements of the trajectory tensor $T_{\psi a}^t$ and the slack variables λ .

5 From convex polyhedron to Markov process

Given a set of inequalities that contain the support of the posterior, we next need to define a stochastic proposal function $f : \mathbb{S} \rightarrow \mathbb{S}$ on a set of Markov states, \mathbb{S} , where each state, S , is associated with a trajectory $T(S)$. The proposal function should have the following properties:

- There should be a set of transitions with non-zero probability between any two Markov states. This ensures proper mixing as time tends to infinity.
- At the stationary point, the probability of being in a state associated with trajectory T should be our target posterior probability $P(T \mid \Omega)$.
- For any transition from state $S_a \rightarrow S_b$ the probability of transitioning from $S_b \rightarrow S_a$ should be non-zero. This allows us to attain detailed balance by using the Metropolis Hastings algorithm.
- The proposal function should be computationally efficient.

5.1 A proposal function for a Fermionic ABM

We define a Fermionic ABM to be one where no two agents can share the same state at the same time. i.e. in addition to the validity constraints of equations 1 and 2 the trajectory of a Fermionic ABM also satisfies the *Fermionic constraint*

$$\begin{aligned} \forall t, \psi : \sum_a T_{\psi a}^t &\leq 1 \\ \forall \phi : \sum_{\psi, a} F(\psi, a)_\phi T_{\psi a}^n &\leq 1 \end{aligned} \quad (7)$$

in which case we call it a *Fermionic trajectory* [this can be weakened to $T_{\psi a}^t \leq 1$].

It is clear that every element of a Fermionic trajectory must be either 0 or 1. This means we can easily convert the union term in equation 5

$$\mathcal{P}(\pi'(t, \psi, \cdot), a) \cup \{T_{\xi b}^s : T_{\psi a}^t = 0\}$$

into a set of linear constraints. Expanding \mathcal{P} we have

$$\left(\bigcap_j \sum_i a_{ij} x_i \leq b_j \right) \cup \{X : x_k = 0\} = \bigcap_j \left(\sum_i a_{ij} x_i \leq b_j \cup \{X : x_k = 0\} \right)$$

where we've projected the elements of the trajectory $T_{\psi a}^t$ into a vector X . But

$$\sum_i a_{ij} x_i \leq b_j \cup \{X : x_k = 0\} = \sum_i a_{ij} x_i + \left(\sum_i \frac{|a_{ij}| + a_{ij}}{2} - b_j \right) x_k \leq \sum_i \frac{|a_{ij}| + a_{ij}}{2}$$

since all x_i are also either 0 or 1 so $\sum_i a_{ij} x_i$ must be less than $\frac{|a_{ij}| + a_{ij}}{2}$ and the constraint is satisfied if x_k is 0, whereas we have the original constraint if x_k is 1.

So, taking our predator prey agent of algorithm 1 as an example, we can see that the move, reproduce and die actions can be returned irrespective of the state of other agents³, so for these actions the union is also unconditionally true. However, the reproduction actions of a predator are conditional on there being some prey on neighbouring gridsquares. So, suppose we have a predator in state ψ at time t , the support of it's timestep function given that it returns a reproduction action is

$$\sum_{i \in N_{\psi}^t} -x_i \leq -1$$

where N_{ψ}^t is the set of indices that correspond to elements of $T_{\phi a}^t$ where ϕ is a prey on a gridsquare neighbouring ψ at time t . So, taking the union with $\{X : x_k = 0\}$, the final constraint for a predator in state ψ at time t is

$$\sum_{i \in N_{\psi}^t} -x_i + x_k \leq 0$$

Theorem 1. *All trajectories that satisfy equations 1 and 7 are at extreme points. Where an extreme point is one that can't be expressed as a linear combination of two other points that satisfy the constraints.*

Proof. Inequality 7 can only be satisfied by non-negative integers if, for any given t and ψ either all $T_{\psi a}^t$ are zero or one element of $T_{\psi a}^t$ is one and the rest zero. In the first case the solution is touching all $|a|$ of the $T_{\psi a}^t \geq 0$ constraints, in the second case the solution is touching $|a| - 1$ of the $T_{\psi a}^t \geq 0$ constraints and one $\sum_a T_{\psi a}^t \leq 1$ constraint to give a total of $|a|$.

[Conversely, we can show that all extreme points are integer points, so we have the polytope of integer solutions]

[In the case of the action Fermionic constraint, this can be seen immediately since equations 1 and the action Fermionic constraint describe a unit hypercube, so all integer solutions are on vertices.]

□

This seemingly abstract fact has a profound practical consequence for the efficiency of our proposal function. Given a system of m linear constraints on n variables expressed in the form $AX = B$, let a *pivot state* be a partition of the variables into m *basic variables* and $n - m$ *non-basic variables* so we can express the constraints in the form

$$A_b X_b + A_n X_n = B \quad (8)$$

where X_b and X_n are the basic and non-basic variables respectively, A_b is the matrix formed from the columns in A that correspond to the basic variables and A_n is the matrix formed from the columns of A that correspond to the non-basic variables. If we now constrain all non-basic variables to zero then we're left with $A_b X_b = B$ but since A_b is an $m \times m$ square matrix then $X_b = A_b^{-1} B$ gives a unique solution, as long as A_b is invertable. If A_b is invertable and X_b has no negative entries then we say that the pivot state is valid. It can be shown that all valid pivot states of equations 6 correspond to extreme points (Dantzig, Orden, Wolfe, et al., 1955).

This idea leads to a computationally very efficient way of perturbing trajectories. We begin by representing the constraints and the current trajectory in the form of equation 8. If we multiply any row of A_b by some constant, c , then the truth of equation 8 can be maintained by multiplying the same row of A_n and of B by c as well. Similarly if we add one row of A_b to another row, the equations can be maintained by performing the same row operation on A_n and B .

³the Fermionic constraint is dealt with separately, so we don't need to worry about that here

If the pivot state is valid, then A_b is invertable so we can use Gaussian elimination, by performing a sequence of row additions and multiplications, to transform A_b into a form such that each row and column is zero on all elements except one, which is 1. We'll call this the standard form. Once in this form, if we assume $X_n = 0$ the solution for X_b can be read off directly from B .

Given a valid pivot state in standard form, a perturbed valid solution can be generated by choosing a column, C , of A_n , with at least one positive-valued row, and choosing from among those rows the one, C_i , that minimises B_i/C_i . We then make the variable corresponding to column C a basic variable and make the basic variable corresponding to row i of A_b non-basic by swapping column C with the column on row i of A_b that has value 1. A_b is now not in standard form as column C will, in general, have more than one non-zero value so we perform Gaussian elimination (row additions and multiplications) to return it to standard form. This perturbation is known as a *pivot* operation and is the same method as employed in the Simplex algorithm (Dantzig et al., 1955) (Vanderbei, 2020). This is exactly what we need for our proposal function since it efficiently generates a new valid trajectory from a current one.

5.1.1 Fractional solutions and feasibility pumps

Although pivoting ensures solutions are positive, there may exist extreme points that don't fall on the grid of integer solutions and so do not satisfy the integer constraint. However, if we posit the existence of a *null trajectory* and associate all fractional solutions with the null trajectory, then simply ignore null trajectory samples, then we end up with samples from the posterior, as required (i.e. the Markov chain can pass through fractional states, but no sample is generated when doing so). However, during the sample process we don't want to spend too much computational effort moving between fractional states without generating any usable samples. There are two ways of achieving this, firstly we should make transitions between fractional states as efficient as possible, and secondly we should ensure that the sum of the probabilities of all fractional states is a small fraction of the sum of probabilities of all states (but not so small as to endanger proper mixing of the Markov process).

In order to achieve this we define the probability of a fractional state, X , to be

$$P_f = \frac{\overline{P}_i \beta \pi_1(X) e^{-k\|X - [X]\|}}{\overline{\pi}_1} \quad (9)$$

where \overline{P}_i is the average probability of the integer states, $\beta = \frac{n_i}{n_f}$ is the ratio of integer states to fractional states $\pi_1(X)$ is the number of non-degenerate pivots from state X , k is a tunable parameter, $[X]$ is the vector that results from rounding each element of X to the nearest integer and $\|\cdot\|$ is the L1-norm and $\overline{\pi}_1$ is the average number of non-degenerate pivots in fractional states.

When calculating the probability of acceptance between fractional states X and Y

$$\alpha = \frac{e^{-k\|Y - [Y]\|}}{e^{-k\|X - [X]\|}}$$

This can be calculated without doing a pivot by just calculating B so we only need to pivot if the proposal is accepted, making transitions between fractional states much more efficient than between integer states.

The proportion of samples that are null samples (i.e. samples coming from fractional states) is expected to be simply the normalised sum of probabilities of all fractional states. If we assume the correlation between the number of non-degenerate pivots and the fractionality is small then the normalised total fractional probability is approximated by

$$P_F \approx \frac{\overline{e^{-k\|X - [X]\|}}}{1 + \overline{e^{-k\|X - [X]\|}}}$$

where the overbar on the exponential denotes the average over all fractional states.

Since we don't know \overline{P}_i , β , $\overline{\pi}_1$ or $\overline{e^{-k\|X - [X]\|}}$ during the sampling process, in practice we can approximate $\overline{\pi}_1$ with its expectation value calculated from the samples so far.

The sum of probabilities of Integer points can be estimated from the expectation value of $\frac{1}{P(X)}$

Also from the samples so far we can build an approximation of the distribution of $\|X - [X]\| e^{-k\|X - [X]\|}$ by kernel density approximation from the samples from fractional states. If we divide this by $e^{-k\|X - [X]\|}$ we get the probability distribution over $\|X - [X]\|$ which allows us to calculate P_F and to set k to ensure that P_F is of a desirable value (say 0.01). We can then compare this against the proportion of integer to fractional samples and adjust our estimate of $\overline{P}_i \beta$ accordingly. Initially we set $\beta = 1$ and approximate $\overline{P}_i \beta$ with the square root of the expectation value of P calculated from the samples so far (since the expectation of P is the mean square of P).

approximate them by taking the averages over the states we have visited so far. Since we start on a valid sample, this is always well defined. On the assumption that the number of fractional states is of the same order of magnitude (or less) than the number of integer states, and since transition between fractional states is computationally cheaper than between integer states, then we expect to spend only a small fraction of computational effort on fractional states.

[Alternatively, transitions are always between integer states, but with fixed probability choose to make a “hyperspace” proposal: a new integer state that is generated by a *feasibility pump* (Fischetti, Glover, & Lodi, 2005) - but how do we calculate probability of transition?? Use row ordering as pseudo-random flip direction (e.g. flip first 40 rows) but does this guarantee a return transition?. Non-random perturbation during flip? Or very reduced space. Or, look at the objective: All integer states that have V as L1 closest valid state must have a fully positive objective (costly for basis variables). Generate flip by reverse feasibility pump? - Starting in state X , choose an integer state X' that has X as L1 closest valid point (how to choose with known probability?) - pump in reverse (away from X' -objective) until the next pivot would reach a state C such that $[C] \neq X'$. (need to check whether we go through a state that is the closest valid point to any integer point - likely every point is the closest to a number of integer points and choosing one is as hard as choosing a vertex from a convex polyhedron!) - now choose any pivot that changes the rounding (we know there is at least one) - perform the pivot and then pump forward to an integer solution.] [Alternatively, the original plan works if it is true that there is always a pivot that reduces the current state’s distance to its rounding...however, it’s easy to construct an example where this isn’t true (just in 2D)] [Finally, could just start many chains! (but then don’t know relative weights of each chain)]

5.1.2 Dealing with degeneracy

The use of pivoting is complicated somewhat when the solutions are degenerate. A degenerate solution is one that can be generated by more than one pivot state. This can occur when a pivot state has basic variables with value zero (in which case we say the pivot state is degenerate and call the zero variables *degenerate variables* of the pivot state). Pivoting on a degenerate variable does not change the solution since during a pivot the outgoing basic variable becomes constrained to zero; but if the basic-variable’s value was zero to start with, constraining it to zero has no effect on the solution (for a thorough theoretical development of degeneracy see (Zörnig, 1993)). This is problematic because we need to assign a probability to each Markov state but under degeneracy a pivot state can’t be given the probability of its trajectory because this would result in degenerate trajectories being “counted” multiple times, once for each degenerate pivot state of the trajectory. To maintain the correct distribution, the probabilities of all the degenerate pivot states of a given trajectory should sum to the probability of that trajectory.

To solve this problem in a computationally efficient way we introduce the concept of an *ordered pivot state* which consists of a partition of the variables into basic and non-basic variables as before, but in addition there is now an ordering of the basic variables. Since we represent A_b as a matrix in our standard form representation, it is easy to represent this ordering in the ordering of the rows of A_b . As before, the validity of equation 8 can be maintained under permutations of the ordering of the rows of A_b by performing the same row permutations on A_n and B . We now associate a Markov state with each ordered pivot state, assigning it a probability according to algorithm 3:

Algorithm 3 Algorithm to calculate probability of an ordered pivot state

```

function PROBABILITY( $A_b, A_n, B$ ) ▷ Standard representation of an ordered pivot state
     $P_T \leftarrow$  the probability of the trajectory of the current solution
     $P_d \leftarrow 1.0$ 
     $C \leftarrow \emptyset$ 
     $r \leftarrow 0$ 
    for  $i = |B| - 1$  down to 0 do
        if  $B[i] == 0$  then
             $C \leftarrow C \cup$  the indices of the non-zero entries in row  $i$  of  $A_n$ 
             $P_d \leftarrow P_d / (|C| + r + 1)$ 
             $r \leftarrow r + 1$ 
        end if
    end for
    return  $\frac{P_d P_T (|B| - \sigma)!}{|B|!}$ 
end function
    
```

The proposal function for ordered pivot states is shown in algorithm 4

We now show that this assignment of probability to ordered pivot states has the property that the sum of the probabilities of ordered pivot states associated with a trajectory is the probability of that trajectory.

First some notation: let $\mathbb{E}(T)$ be the set of all ordered pivot states with solution T , let $T(S)$ be the trajectory associated with pivot state S , let $D(S)$ be the tuple of degenerate variables in ordered pivot state S , let $D_{\leq n}(S)$ be the n -tuple consisting of the first n elements of $D(S)$ and let

$$C(\mathbb{S} | V) = \left\| \left\{ V^+ : S \in \mathbb{S}, V^+ = D_{\leq |V|+1}(S), V_{\leq |V|}^+ = V \right\} \right\| \quad (10)$$

i.e. among the members of \mathbb{S} that have the degenerate variable prefix V , $C(\mathbb{S} | V)$ counts how many distinct degenerate variable prefixes of length $|V| + 1$ there are.

Algorithm 4 Proposal function for ordered pivot states

```

function PROPOSAL( $A_b, A_n, B$ )                                ▷ Standard representation of the current ordered pivot state
     $A'_b, A'_n, B' \leftarrow A_b, A_n, B$ 
     $\alpha, \beta \leftarrow$  constant parameters
     $p_0 \leftarrow$  the set of degenerate pivots of  $(A_b, A_n, B)$  (i.e. the ones that do not change the solution)
     $p_1 \leftarrow$  the set of non-degenerate pivots of  $(A_b, A_n, B)$  (i.e. the ones that change the solution)
     $P_{f/b} \leftarrow 1$                                           ▷ the ratio of probabilities of making this transition forwards/backwards
    if BERNOULLI( $\alpha$ ) then
         $r \leftarrow$  choose member of  $p_1$  with uniform probability
        perform pivot  $r$  on  $(A'_b, A'_n, B')$ 
         $p'_1 \leftarrow$  number of non-degenerate pivots of  $(A'_b, A'_n, B')$ 
         $P_{f/b} \leftarrow \frac{|p'_1|}{|p_1|}$ 
    else if BERNOULLI( $\beta$ ) then
         $r \leftarrow$  choose member of  $p_0$  with uniform probability
        perform pivot  $r$  on  $(A'_b, A'_n, B')$ 
         $p'_0 \leftarrow$  number of degenerate pivots of  $(A'_b, A'_n, B')$ 
         $P_{f/b} \leftarrow \frac{|p'_0|}{|p_0|}$ 
    else
        choose two rows  $a$  and  $b$  with uniform probability
        swap rows  $a$  and  $b$  of  $(A'_b, A'_n, B')$ 
    end if
    return  $(A'_b, A'_n, B', P_{f/b})$ 
end function
    
```

Theorem 2. If we assign the probability

$$P(S) = \frac{(m - \sigma_T(S))!}{m!} \frac{P_T(S)}{\prod_{q=0}^{\sigma_T(S)-1} C(\mathbb{E}(T(S)) \mid D_{\leq q}(S))} \quad (11)$$

to pivot state S , where m is the number of constraints and σ_T is the degeneracy of trajectory T (i.e. m minus the number of non-zero variables in T) then

$$\sum_{S \in \mathbb{E}(T)} P(S) = P_T$$

Proof. Expanding the sum over degenerate states

$$\sum_{S \in \mathbb{E}(T)} P(S) = P_T \frac{(m - \sigma_T)!}{m!} \sum_{S \in \mathbb{E}(T)} \frac{1}{\prod_{q=0}^{\sigma_T-1} C(\mathbb{E}(T) \mid D_{\leq q}(S))}$$

Notice first that all pivot states with the same σ_T -tuple of degenerate variables, $D(S)$, have the same probability. There are exactly $\frac{m!}{(m - \sigma_T)!}$ ordered pivot states with a given $D(S)$, corresponding to the different placings of the $m - \sigma_T$ non-degenerate basic variables among the m possible positions. So, if we let

$$\mathbb{D}(T) = \{D(S) : S \in \mathbb{E}(T)\}$$

then

$$\sum_{S \in \mathbb{E}(T)} P(S) = P_T \sum_{D \in \mathbb{D}(T)} \frac{1}{\prod_{q=0}^{\sigma_T-1} C(\mathbb{E}(T) \mid D_{\leq q})}$$

Since $C(\mathbb{S} \mid V)$ depends only on the order of the degenerate variables in the members of \mathbb{S}

$$C(\mathbb{E}(T) \mid V) = C(\mathbb{D}(T) \mid V)$$

so

$$\sum_{S \in \mathbb{E}(T)} P(S) = P_T \sum_{D \in \mathbb{D}(T)} \frac{1}{\prod_{q=0}^{\sigma_T-1} C(\mathbb{D}(T) \mid D_{\leq q})} \quad (12)$$

if we separate the terms in the sum into groups that share the same prefix apart from the last variable then we get

$$\sum_{S \in \mathbb{E}(T)} P(S) = P_T \sum_{V: D \in \mathbb{D}(T), V = D_{\leq (\sigma_T-1)}} \left(\frac{1}{\prod_{q=0}^{\sigma_T-2} C(\mathbb{D}(T|V_{\leq q})} \sum_{V+: V+ \in \mathbb{D}(T), V = V_{\leq (\sigma_T-1)}^+} \frac{1}{C(\mathbb{D}(T|V))} \right)$$

So

$$\sum_{S \in \mathbb{E}(T)} P(S) = P_T \sum_{V: D \in \mathbb{D}(T), V = D_{\leq (\sigma_T - 1)}} \frac{1}{\prod_{q=0}^{\sigma_T - 2} C(\mathbb{D}(T|V_{\leq q}))} \frac{\left\| \left\{ V^+ : V^+ \in \mathbb{D}(T), V = V^+_{\leq (\sigma_T - 1)} \right\} \right\|}{C(\mathbb{D}(T|V_{\leq (\sigma_T - 1)}))}$$

but the final fraction here is 1 by equation 10, so

$$\sum_{S \in \mathbb{E}(T)} P(S) = P_T \sum_{V: D \in \mathbb{D}(T), V = D_{\leq (\sigma_T - 1)}} \frac{1}{\prod_{q=0}^{\sigma_T - 2} C(\mathbb{D}(T|V_{\leq q}))}$$

we can now use the same argument again to reduce the upper bound of q iteratively until we show that the sum on the right hand equals one, so

$$\sum_{S \in \mathbb{E}(T)} P(S) = P_T$$

□

Theorem 3. *Algorithm 3 assigns a probability*

$$P(S) = \frac{(m - \sigma_{T(S)})!}{m!} \frac{P_{T(S)}}{\prod_{q=0}^{\sigma_{T(S)} - 1} C(\mathbb{E}(T(S)) \mid D_{\leq q}(S))}$$

to pivot state S

Proof. By inspection of the algorithm, it can be seen that the algorithm assigns probability

$$P(S) = \frac{(m - \sigma_{T(S)})!}{m!} \frac{P_{T(S)}}{\prod_{r=0}^{\sigma_{T(S)} - 1} (\|C'(S, r)\| + r + 1)}$$

where

$$\begin{aligned} C'(S, r) &= C'(S, r - 1) \cup N(S, r) \\ C'(S, -1) &= \emptyset \end{aligned}$$

and $N(S, r)$ is the set of indices of the non-zero entries in the r -from-last degenerate row of A_n . So, $C'(S, r)$ is the set of columns that have at least one non-zero entry on a degenerate row on or below the r -from-last.

However, $C(\mathbb{E}(T(S)) \mid D_{\leq q}(S))$ is, by definition, the number of different degenerate variables that can take the $(q + 1)^{th}$ place in the list of degenerate variables, given that the first q are given by $D_{\leq q}(S)$.

Now, given an ordered pivot state, S in standard form; if there exists a column, j , with a non-zero entry in a degenerate row, i , of A_n or A_b below the q^{th} degenerate row, then we can generate a valid member of $\mathbb{E}(T(S))$ with j the $(q + 1)^{th}$ variable by pivoting on column j and row i (if not already pivoted in), then swapping row i with whichever row makes it the $(q + 1)^{th}$ degenerate variable. Conversely, if there exists a member of $\mathbb{E}(T(S))$ with j as the $(q + 1)^{th}$ variable, it must be possible to pivot on the j^{th} column without pivoting out any of the first q degenerate variables or the non-degenerate variables. The only way this is possible is if there is either a non-zero entry in the j^{th} column of A_n or A_b below the q^{th} degenerate row. So, the set of degenerate variables that can take the $(q + 1)^{th}$ place is the set of columns that have non-zero entries in degenerate rows below the q^{th} in A_n or A_b . From the form of A_b we know immediately that there are $r + 1$ columns with non-zero degenerate entries on or below the r -to-last degenerate row. So

$$C(\mathbb{E}(T(S)) \mid D_{\leq \sigma_{T(S)} - 1 - r}(S)) = \|C'(S, r)\| + r + 1$$

□

5.1.3 Metropolis-Hastings on ordered pivot states

All this can now be assembled into a sampling algorithm based on Metropolis-Hastings. Starting with the constraints in the form of equation 6, recursively choose a non-zero element on an un-pivoted row and pivot on that element, until we reach an ordered pivot state. In general, B will contain negative elements which means that the solution will not conform to the non-negative constraint. In order to find an initial positive solution let $I^- = \{i : B_i < 0\}$ be the set of rows of B that are negative and pivot on the column, j , that minimises $\sum_{i \in I^-} A_{ij}$. Repeat until a positive solution is found (or no column has a negative sum, in which case no positive solution exists).

From the initial valid pivot state, S , generate a proposal pivot state S' and a transition probability ratio $\frac{P(S \rightarrow S')}{P(S' \rightarrow S)}$ using algorithm 4 and accept with probability

$$\alpha = \frac{P(S')}{P(S)} \frac{P(S' \rightarrow S)}{P(S \rightarrow S')}$$

where the probability of a pivot state is given by algorithm 3.

Given this, it's clear that all the necessary properties of the proposal function are fulfilled.

6 Experimental Results

Outline the predator-prey experiments and the results

7 Conclusions

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