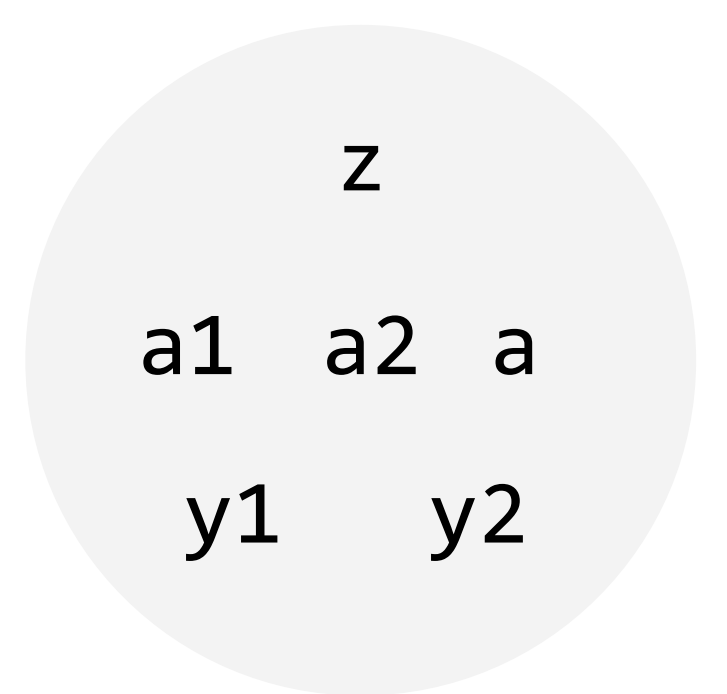


1. Address and traces in Gen

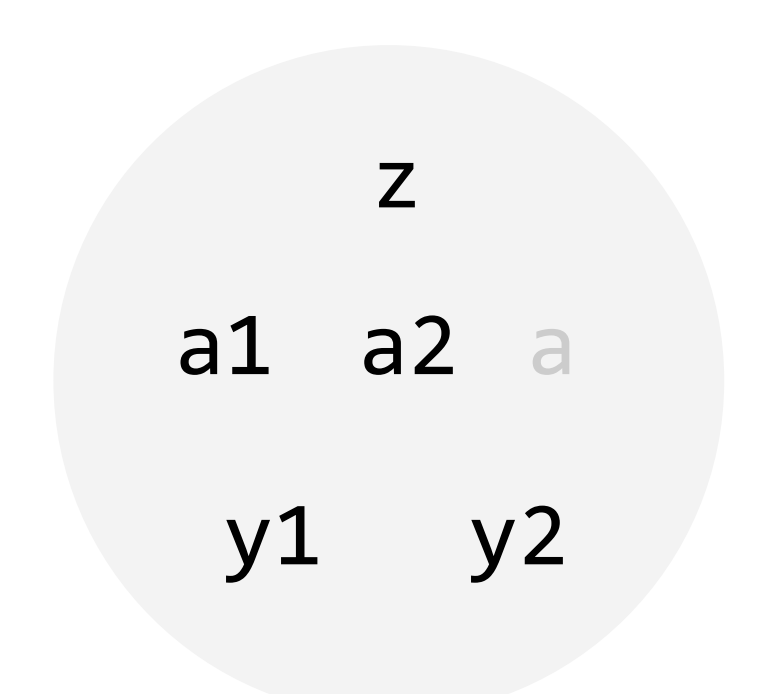
A probabilistic program

```
@gen function foo()
  if ({:z} ~ bernoulli(0.5))
    a1 = ({:a1} ~ gamma(1, 1))
    a2 = ({:a2} ~ gamma(1, 1))
  else
    a = ({:a} ~ gamma(1, 1))
    (a1, a2) = (a, a)
  end
  {:y1} ~ normal(a1, 0.1)
  {:y2} ~ normal(a2, 0.1)
end
```

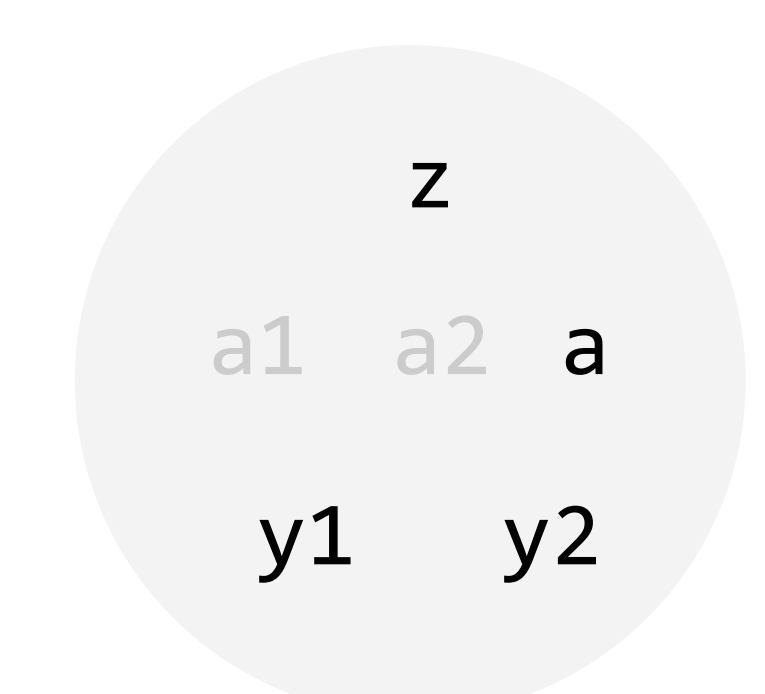
All random choice addresses



Addresses sampled if z = true

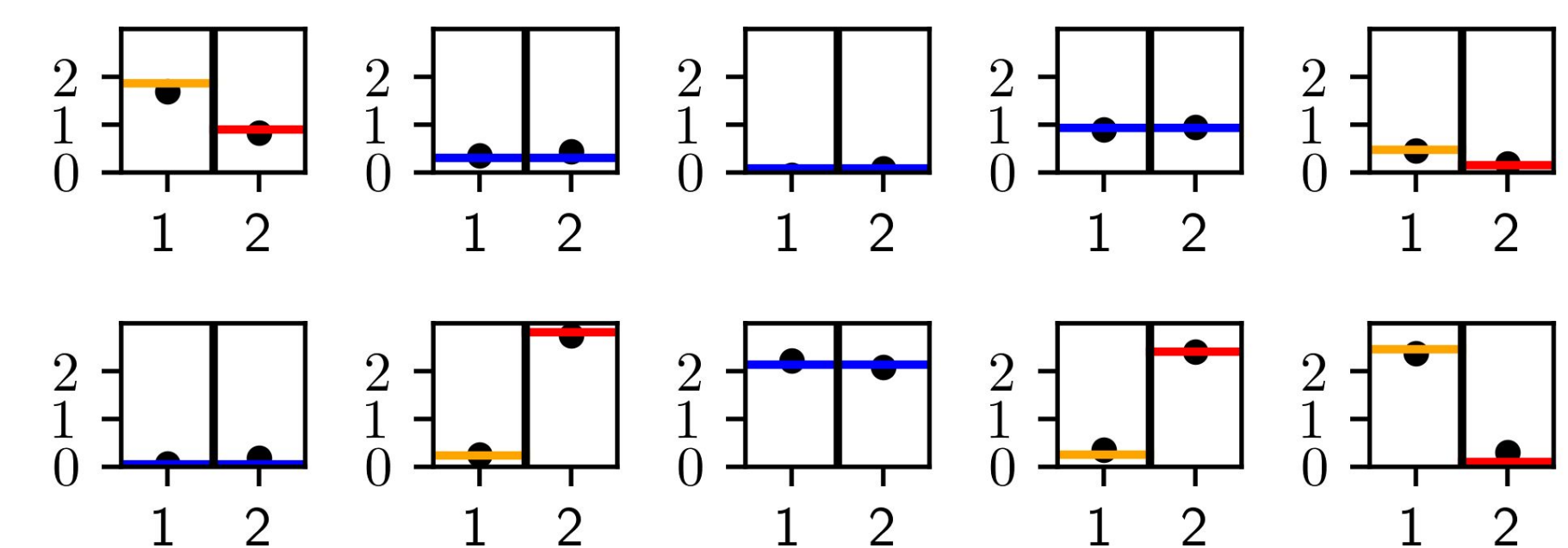


Addresses sampled if z = false



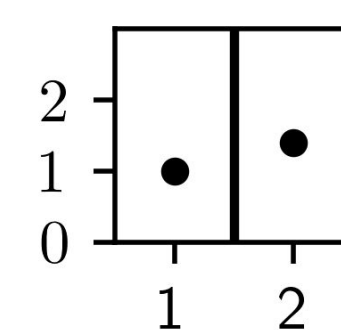
Some traces of the probabilistic program

z	a1	a2	y1	y2
true	1.87	0.89	1.69	0.82
false	0.32		0.36	0.44
false	0.09		-0.03	0.08
false	0.93		0.90	0.96
true	0.48	0.15	0.47	0.19
false	0.04		0.05	0.18
true	0.22	2.81	0.24	2.73
false	2.13		2.20	2.07
true	0.25	2.40	0.34	2.39
true	2.45	0.09	2.36	0.29



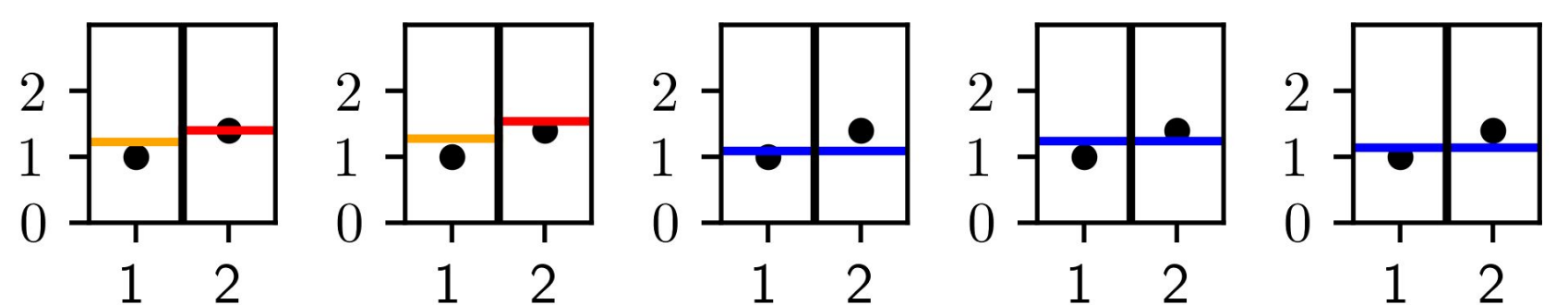
Observed data

(y1 = 1.0, y2 = 1.4)



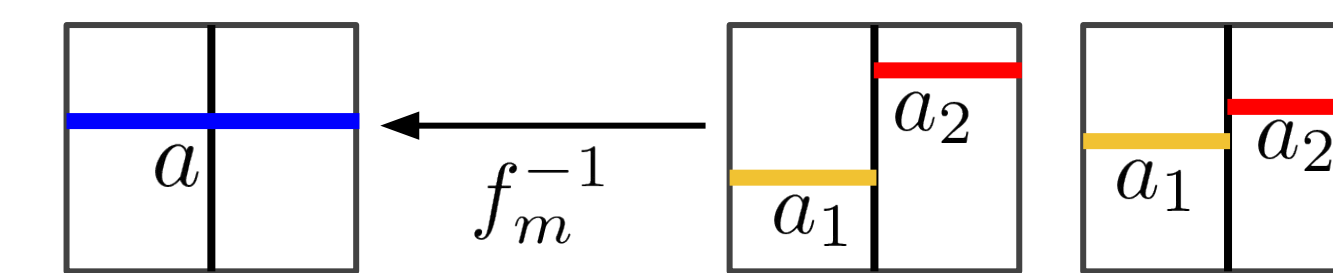
Posterior samples (traces)

z	a1	a2	y1	y2
true	1.22	1.40	1.0	1.4
true	1.27	1.53	1.0	1.4
false	1.09		1.0	1.4
false	1.23		1.0	1.4
false	1.13		1.0	1.4

Cusumano-Towner, Marco. Gen: A High-Level Programming Platform for Probabilistic Inference. PhD thesis. MIT 2020. [Link to PDF](#)

2. Involution MCMC and custom reversible jump MCMC in Gen

A custom kernel for switching branches efficiently



$$f_m^{-1}(a_1, a_2) = \left(\sqrt{a_1 a_2}, \frac{a_1}{a_1 + a_2} \right)$$

Geometric mean Degree of freedom

Auxiliary probabilistic program

```
@gen function q(trace)
  if !trace[z]
    # switch from one to two segments
    {:dof} ~ uniform_continuous(0, 1)
  end
end
```

Trace transform program (involution)

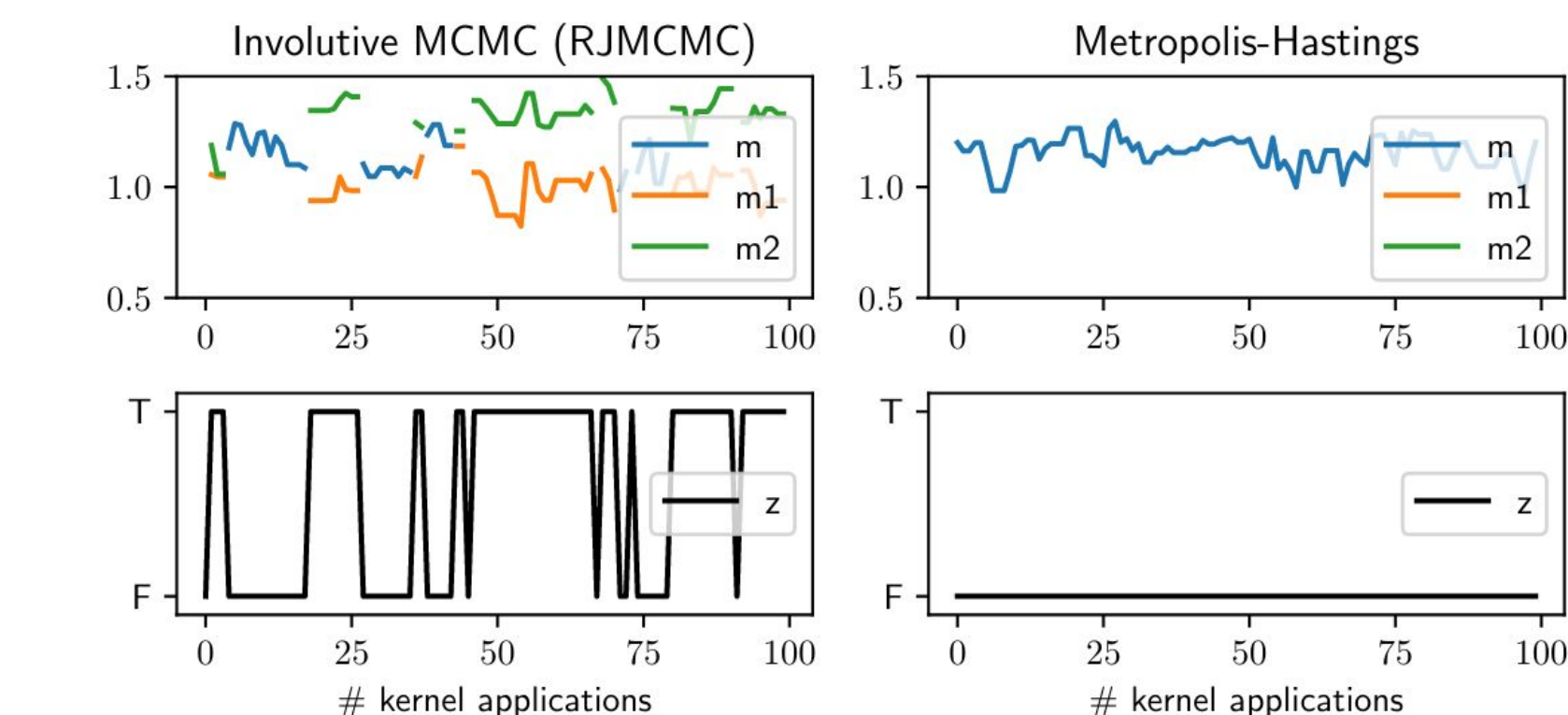
```
@transform h (model_in, aux_in) to (model_out, aux_out)
  if @read(model_in[z], :disc)
    # currently two segments, switch to one
    @write(model_out[z], false)
    m1 = @read(model_in[m1], :cont)
    m2 = @read(model_in[m2], :cont)
    (m, dof) = merge_means(m1, m2)
    @write(model_out[m], m, :cont)
    @write(aux_out[:dof], dof, :cont)
  else
    # currently one segment, switch to two
    @write(model_out[z], true, :disc)
    m = @read(model_in[:m], :cont)
    dof = @read(aux_in[:dof], :cont)
    (m1, m2) = split_mean(m, dof)
    @write(model_out[:m1], m1, :cont)
    @write(model_out[:m2], m2, :cont)
  end
end
```

Cusumano-Towner, Marco, Alexander K. Lew, and Vikash K. Mansinghka.

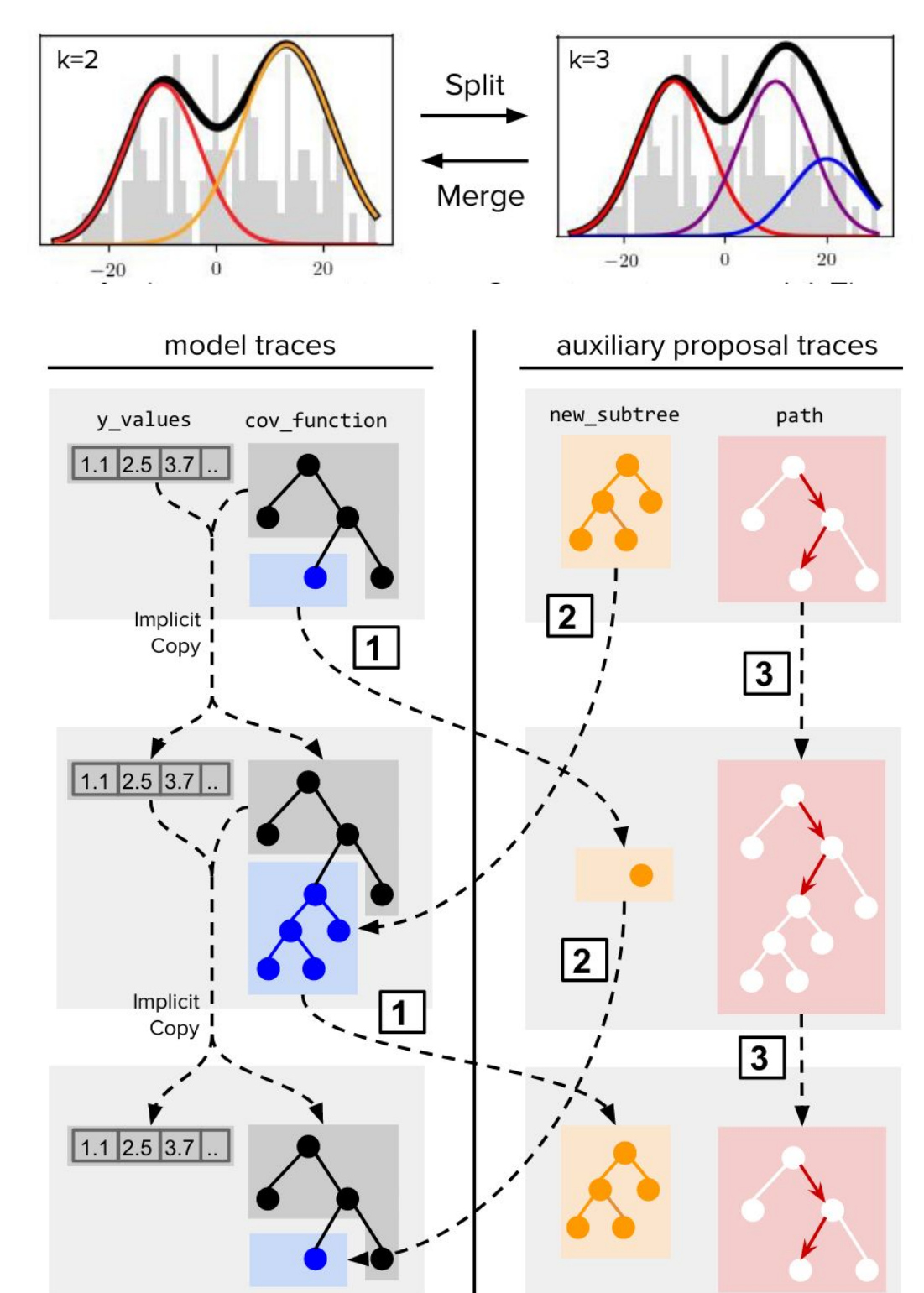
"Automating Involution MCMC using Probabilistic and Differentiable Programming."

arXiv preprint arXiv:2007.09871 (2020). [Link to PDF](#)

Custom kernel is more efficient

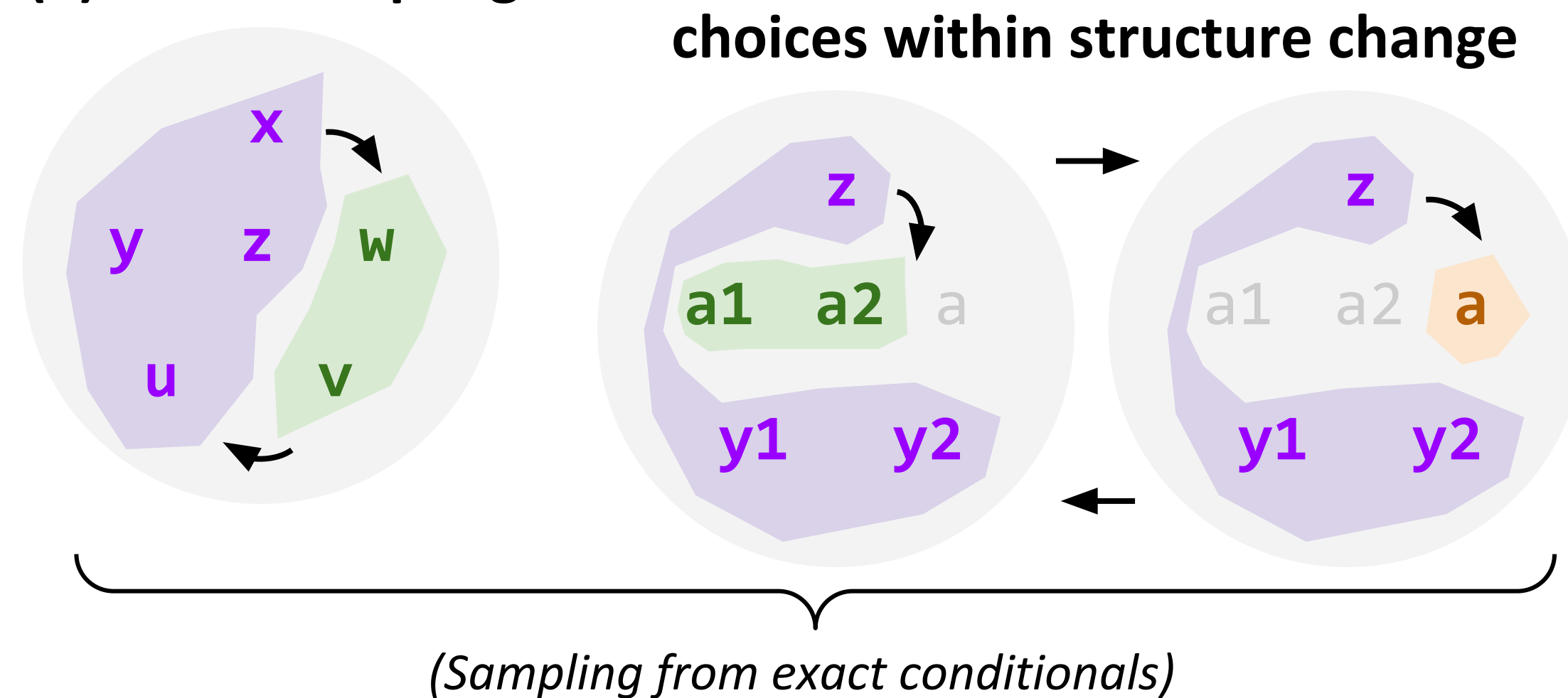


Other examples

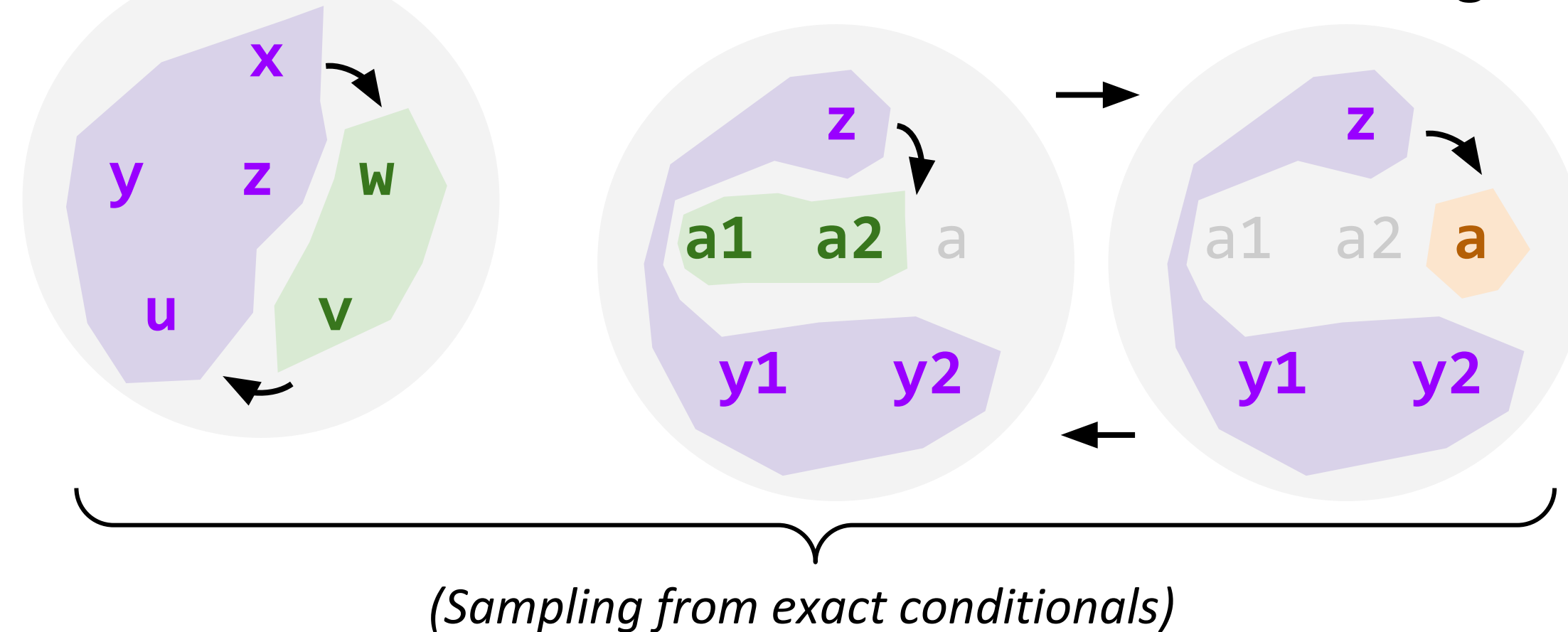


3. Use approx. samplers in the auxiliary probabilistic program to infer values of new addresses

(a) Gibbs sampling

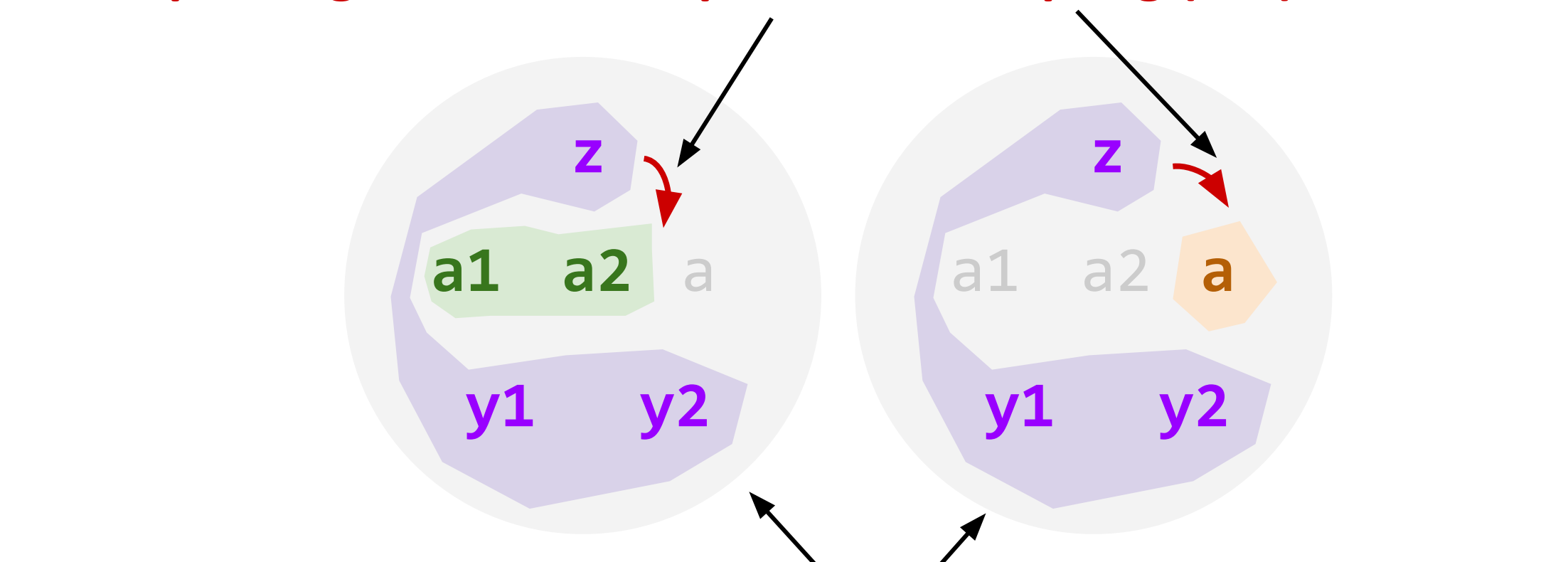


(b) Conditionally sample new choices within structure change



(c) This work: Approximately conditionally sample new (+ optionally, some old) choices within a structure change

Sampler e.g. annealed importance sampling (AIS), SMC, etc.

**"Subproblem" = Any previously nonexistent addresses, and any addressed in the (optional) "shared subproblem"**

Algorithm 1 Subproblem pseudomarginal reversible jump (SPRJ) MCMC kernel template

```
procedure SPRJ-KERNEL(z, θz)
  m ~ q(·; z)
  u', θ'(Iz ∪ Iz') ∪ Sm ~ q(·; m, z', θ(Iz ∪ Iz') ∪ Sm, y)
  θ'Iz' ← (θ(Iz ∪ Iz') ∪ Sm, θ'(Iz ∪ Iz') ∪ Sm)
  p̂(y | θ(Iz ∪ Iz') ∪ Sm; z') ← MLEST(m, z, z', u', θ'Iz')
  u ~ r(·; m, z, θ(Iz ∪ Iz') ∪ Sm, y, θ(Iz ∪ Iz') ∪ Sm)
  p̂(y | θ(Iz ∪ Iz') ∪ Sm; z) ← MLEST(m, z', z, u, θIz)
  a ← q(m; z) p(z') p̂(y | θ(Iz ∪ Iz') ∪ Sm; z')
  q(m; z) p(z) p̂(y | θ(Iz ∪ Iz') ∪ Sm; z)
  return ACCEPT-REJECT(a, (z, θIz), (z', θ'Iz'))
end procedure
procedure MLEST(m, z, z', u', θ'Iz')
  p(θ'Iz'; z') p(y; z', θ'Iz') r(u'; m, z', θ'(Iz ∪ Iz') ∪ Sm, y, θ'(Iz ∪ Iz') ∪ Sm)
  return q(u', θ'(Iz ∪ Iz') ∪ Sm; m, z', θ(Iz ∪ Iz') ∪ Sm, y)
end procedure
procedure ACCEPT-REJECT(a, x, x')
  w ~ Uniform(0, 1)
  if w ≤ a then return x' else return x
end procedure
```

Algorithm 2 SPRJ kernel using annealed importance sampling for subproblem inference (AIS-SPRJ)

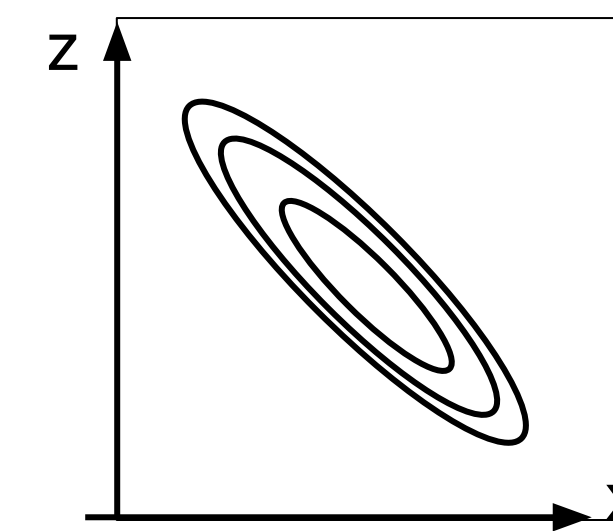
```
procedure SPRJ-KERNEL-AIS(z, θz)
  m ~ q(·; z)
  (θ(Iz ∪ Iz') ∪ Sm, p̂(y | θ(Iz ∪ Iz') ∪ Sm; z')) ~ AIS(m, z', θ(Iz ∪ Iz') ∪ Sm, y)
  p̂(y | θ(Iz ∪ Iz') ∪ Sm; z) ~ R-AIS(m, z, θ(Iz ∪ Iz') ∪ Sm, y, θ(Iz ∪ Iz') ∪ Sm)
  a ← q(m; z) p(z') p̂(y | θ(Iz ∪ Iz') ∪ Sm; z')
  q(m; z) p(z) p̂(y | θ(Iz ∪ Iz') ∪ Sm; z)
  return ACCEPT-REJECT(a, (z, θIz), (z', θ'Iz'))
end procedure
```

4. Approximates standard Gibbs sampling when there is no structure change

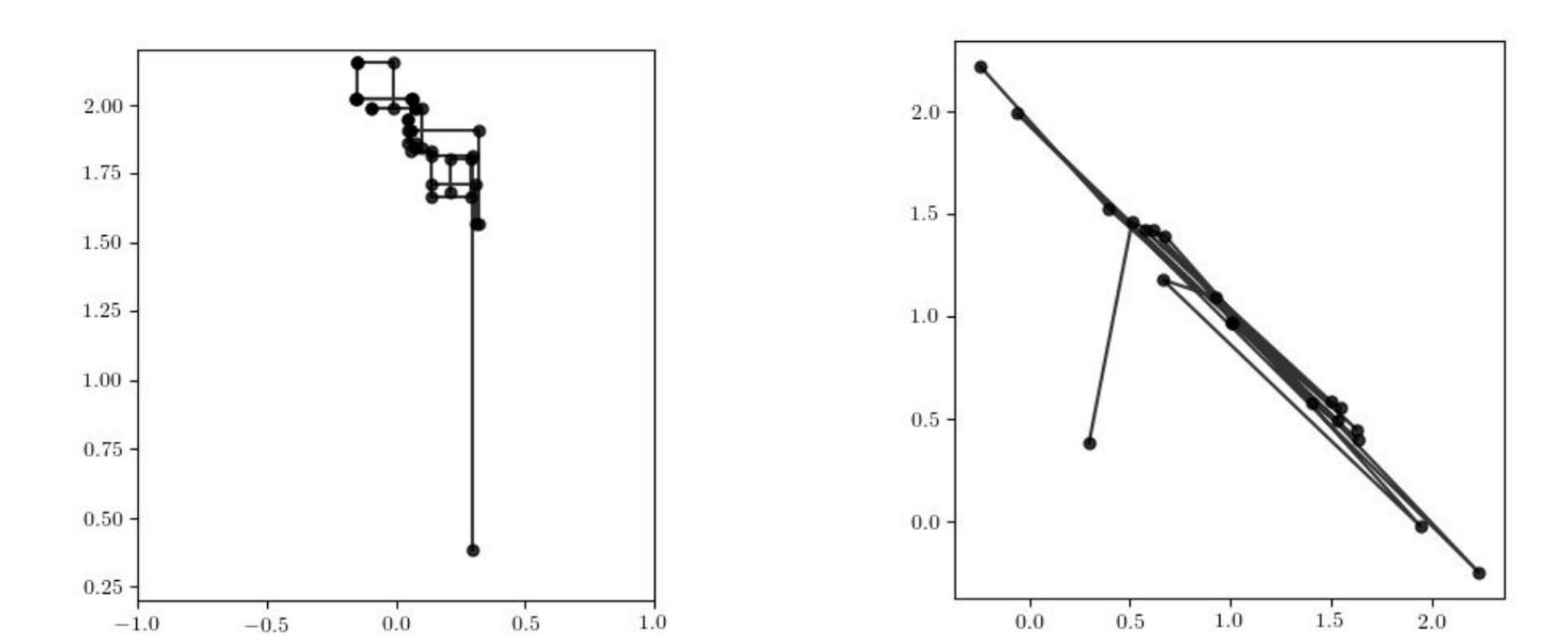
A model with fixed structure

```
@gen function model(noise::Float64)
  x = @trace(normal(0, 1), :x)
  z = @trace(normal(0, 1), :z)
  y = @trace(normal(x + z, noise), :y)
end
```

Conditional distribution on x, z given y



Iterates from two different AIS-SPRJ kernels

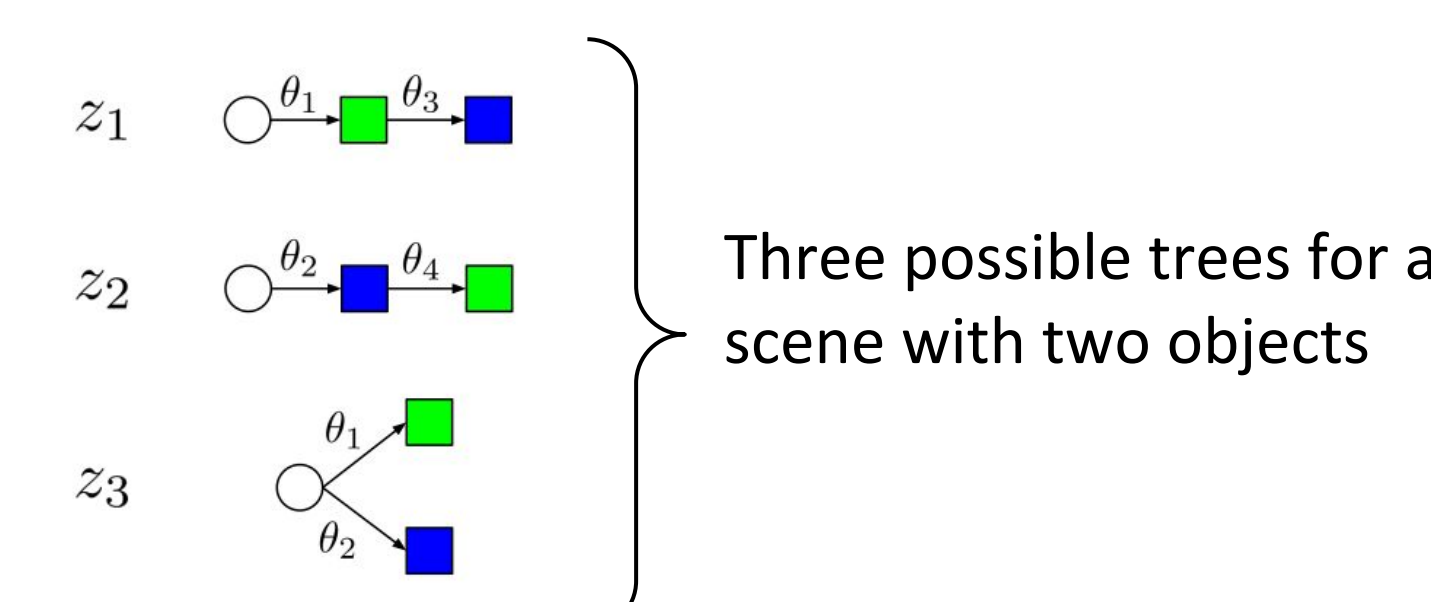


Cycle of two AIS-SPRJ kernels with subproblems {x} and {z}

An AIS-SPRJ kernel with subproblem {x,z}

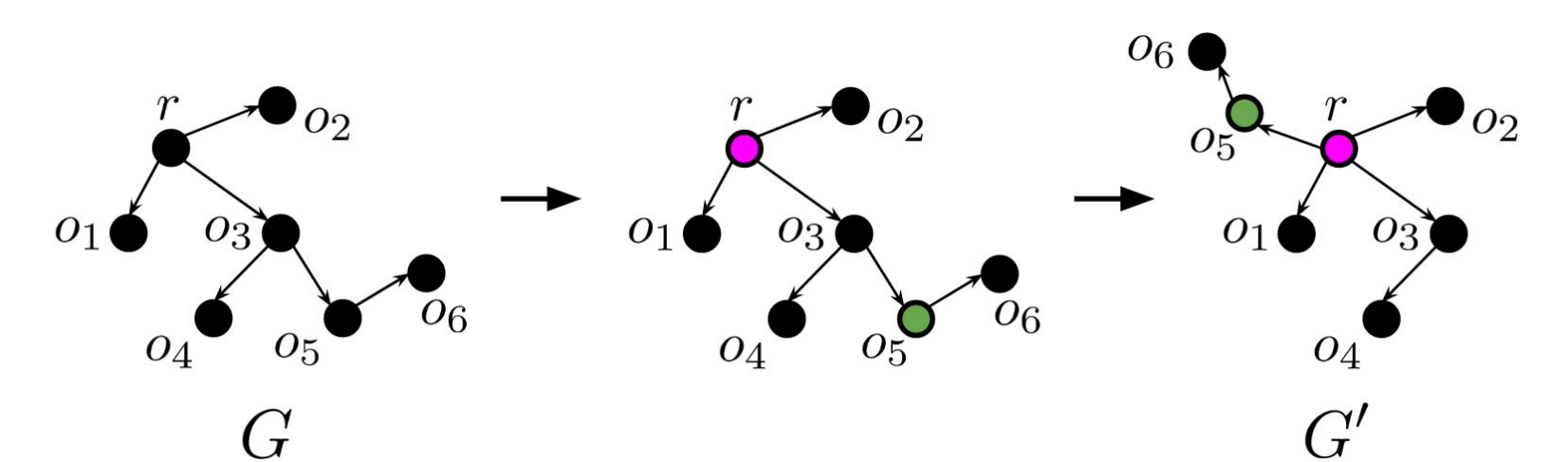
5. Application: Inferring geometric scene graph structure (tree-based pose parametrization)

Tree of coordinate frames in a scene

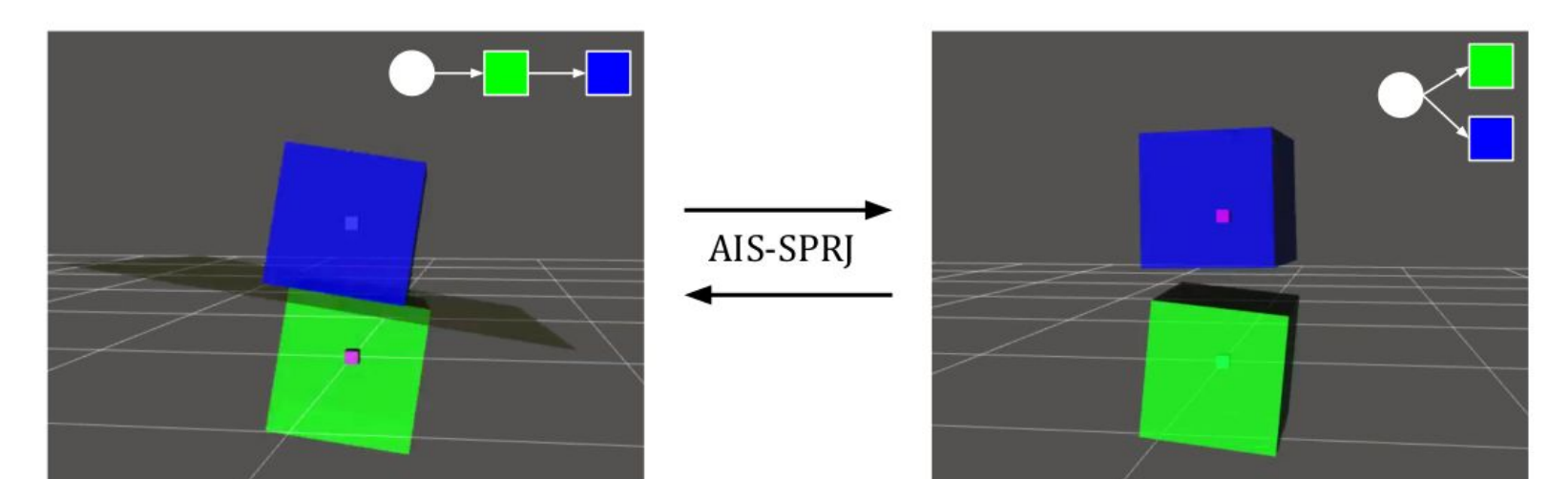


- Root node is the world coordinate frame
- Other nodes are objects' coordinate frames (poses)
- Edges from one object to another indicate sliding (3 degree-of-freedom) contact between objects
- Edges from root to object indicate the object has a full 6 degree-of-freedom pose

MCMC move on the tree (prune and regraft)



Use AIS to infer new continuous pose parameters during each move



Generalizes "subproblem" MCMC (Mansinghka et al., 2014):

- (1) Can use approximate instead of exact samplers
- (2) Applies generally to structure-changing moves

Related work: Georgios Karagiannis and Christophe Andrieu. 2013. Annealed importance sampling reversible jump MCMC algorithms. Journal of Computational and Graphical Statistics 22, 3 (2013), 623–648.