A New Automated Redistricting Simulator Using Markov Chain Monte Carlo

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Joint work with Benjamin Fifield, Michael Higgins, and Alexander Tarr

Motivation

- Redistricting as a central element of representative democracy
- Redistricting may affect:
  - representation (Gelman and King 1994, McCarty *et. al* 2009)
  - turnout (Gay 2001, Baretto 2004)
  - incumbency advantage (Abramowitz *et. al* 2006)
- Substantive researchers simulate redistricting plans to:
  - detect gerrymandering
  - assess impact of constraints (e.g., population, compactness, race)
- Many optimization methods but surprisingly few simulation methods
- Standard algorithm has no theoretical justification
- Need a simulation method that:
  1. samples uniformly from a target population of redistricting maps
  2. incorporates common constraints
  3. scales to redistricting problems of moderate and large size

Overview of the Talk

1. Explain the difficulties of simulating redistricting plans
2. Propose new Markov chain Monte Carlo algorithms
3. Validate the algorithms on a small-scale data example
4. Present empirical analyses for New Hampshire and New Jersey

Characterizing the Distribution of Valid Redistricting Plans

- Scholars want to characterize the *distribution* of redistricting plans under various constraints
- Valid redistricting plans must have:
  - geographically contiguous districts
  - districts with equal population
- Other constraints of interest: compactness, community boundary, etc.
- Naive Approach 1: Enumeration
  - can't enumerate all plans (too many)
  - enumerating only valid plans is not trivial
- Naive Approach 2: Random assignment
  - too few plans will have equal population
  - too few plans will be contiguous

The Standard Simulation Algorithm

- Random seed-and-grow algorithm (Cirincione *et. al* 2000, Altman & McDonald 2011, Chen & Rodden 2013):
  1. randomly choose a precinct as a "seed" for each district
  2. identify precincts adjacent to each seed
  3. randomly select adjacent precinct to merge with the seed
  4. repeat steps 2 & 3 until all precincts are assigned
  5. swap precincts around borders to achieve population parity
- Modify Step 3 to incorporate compactness

- No theoretical properties known
- The resulting sample may not be representative of the population
- "Local" exploration is difficult

The Proposed Automated Redistricting Simulator

- Independent sampling is difficult
- Markov chain Monte Carlo algorithm
- Can sample uniformly from the target distribution
- Start with a valid plan and then swap precincts in a certain way

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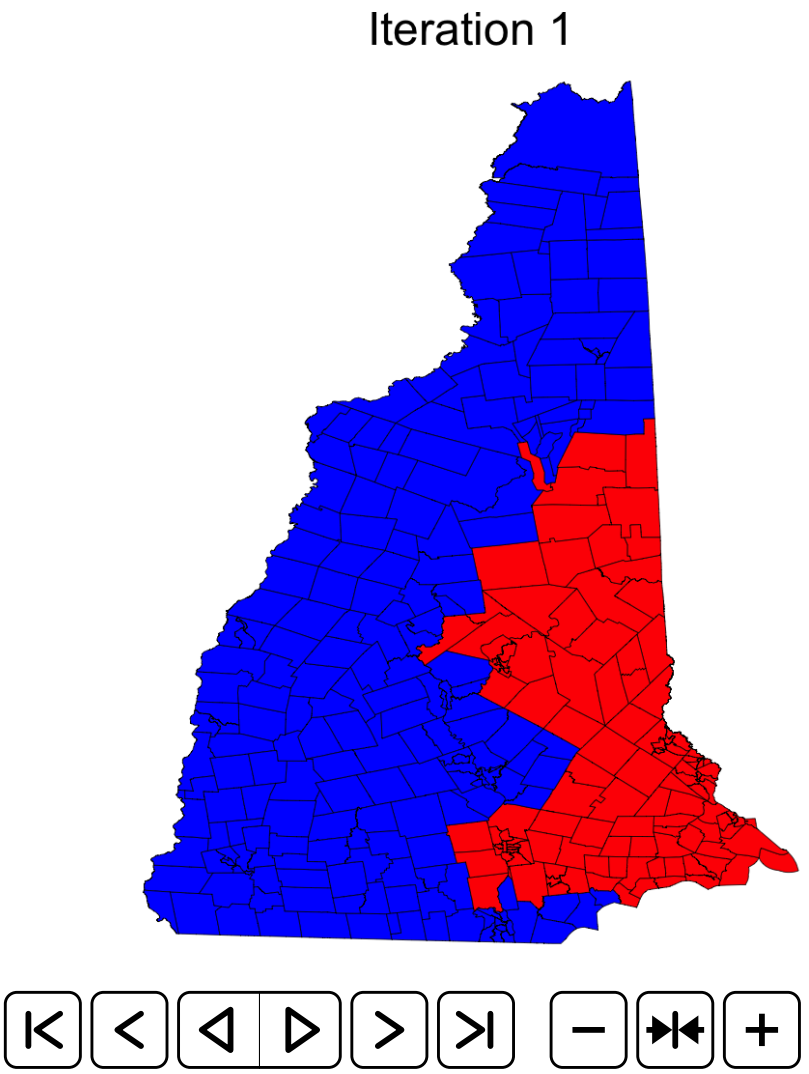
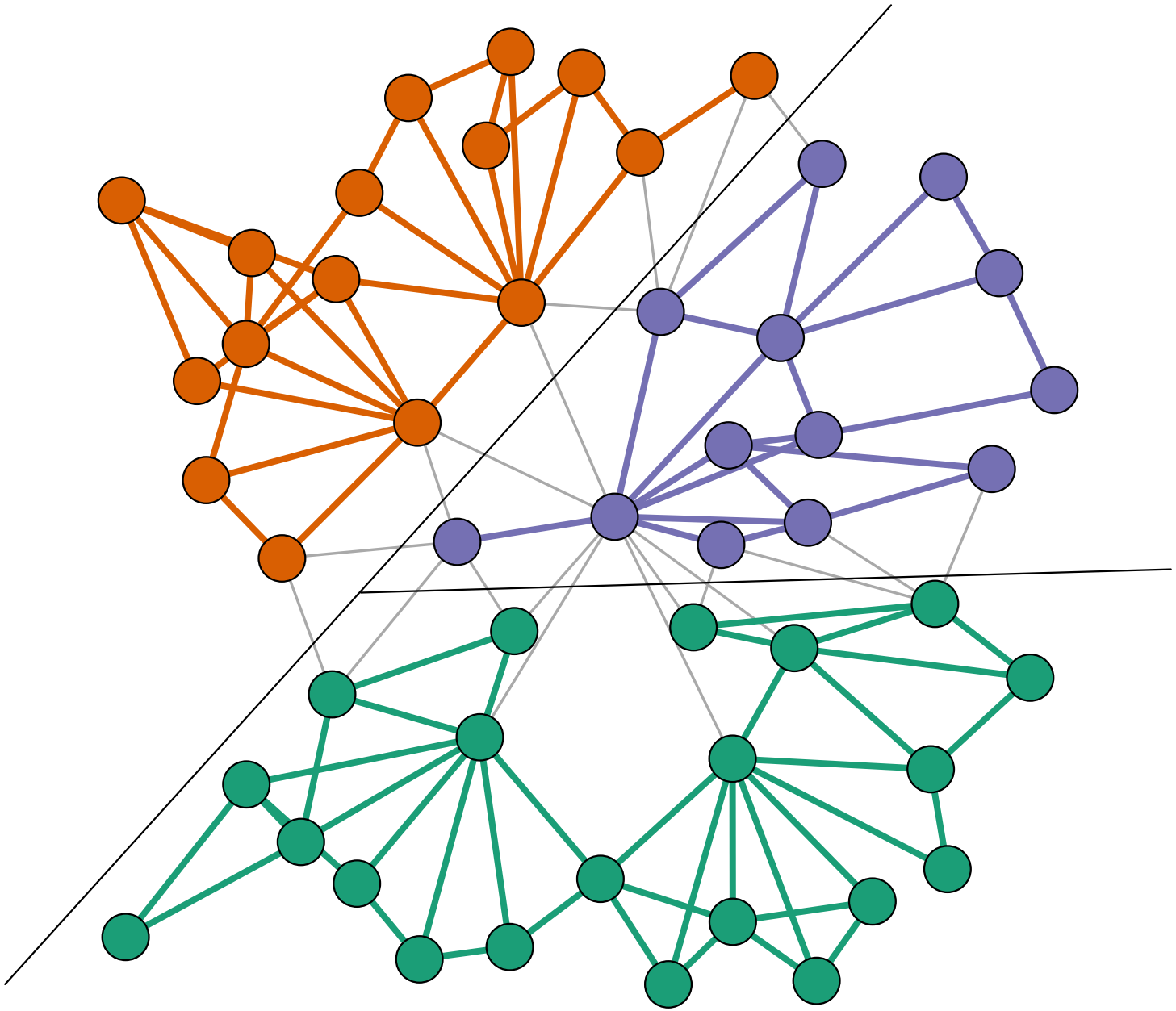
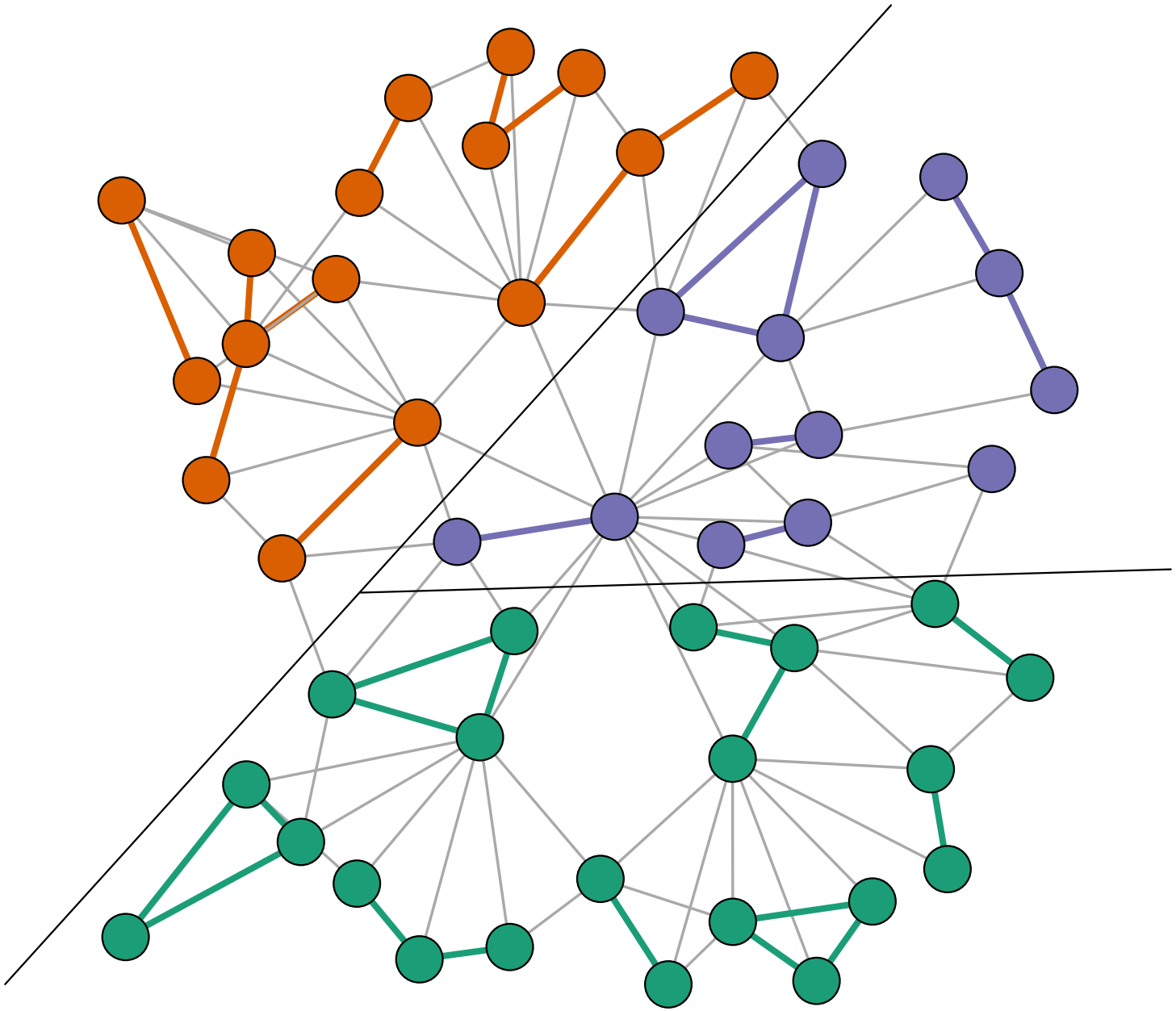


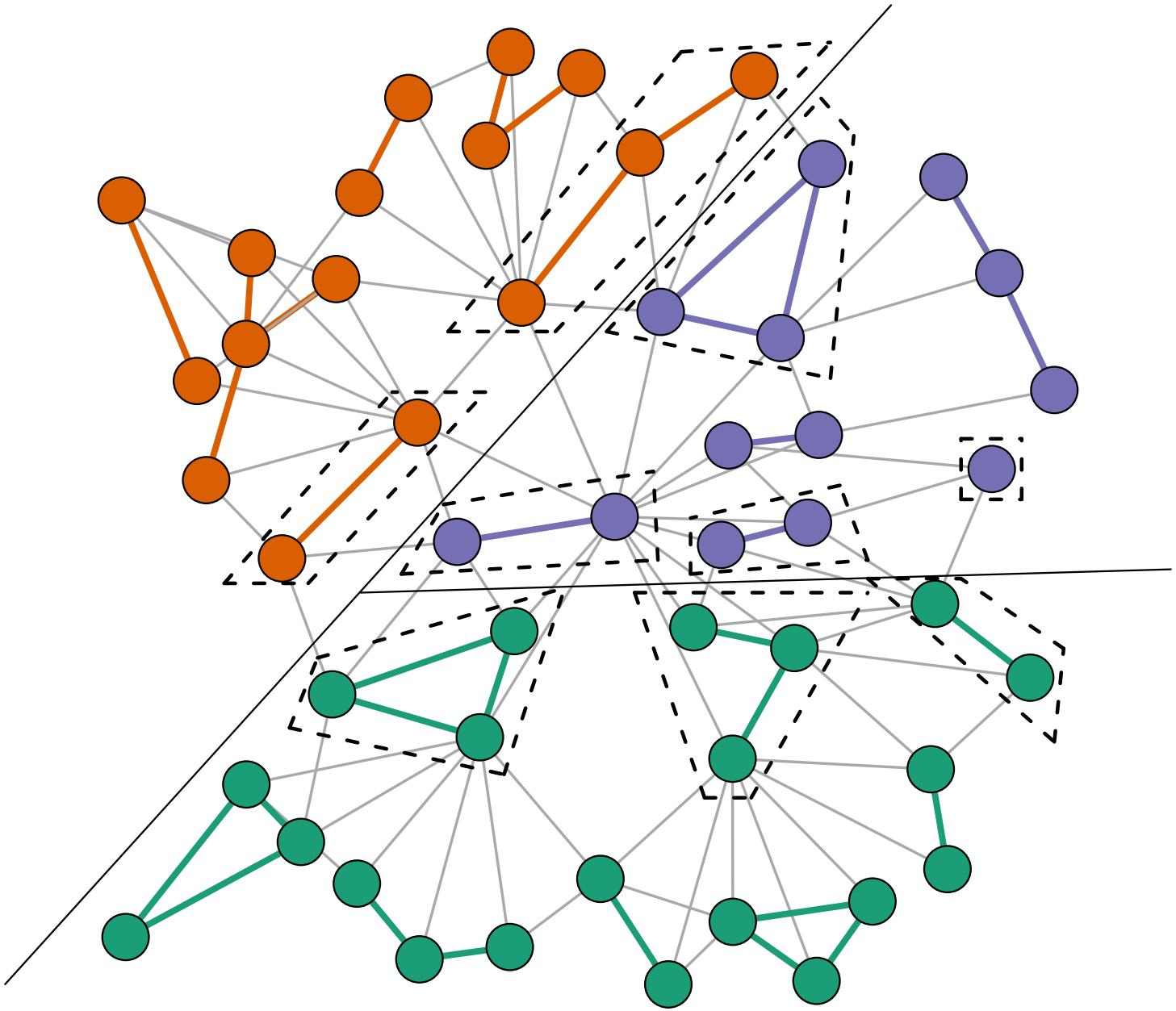
Redistricting as a **Graph-Cut** Problem



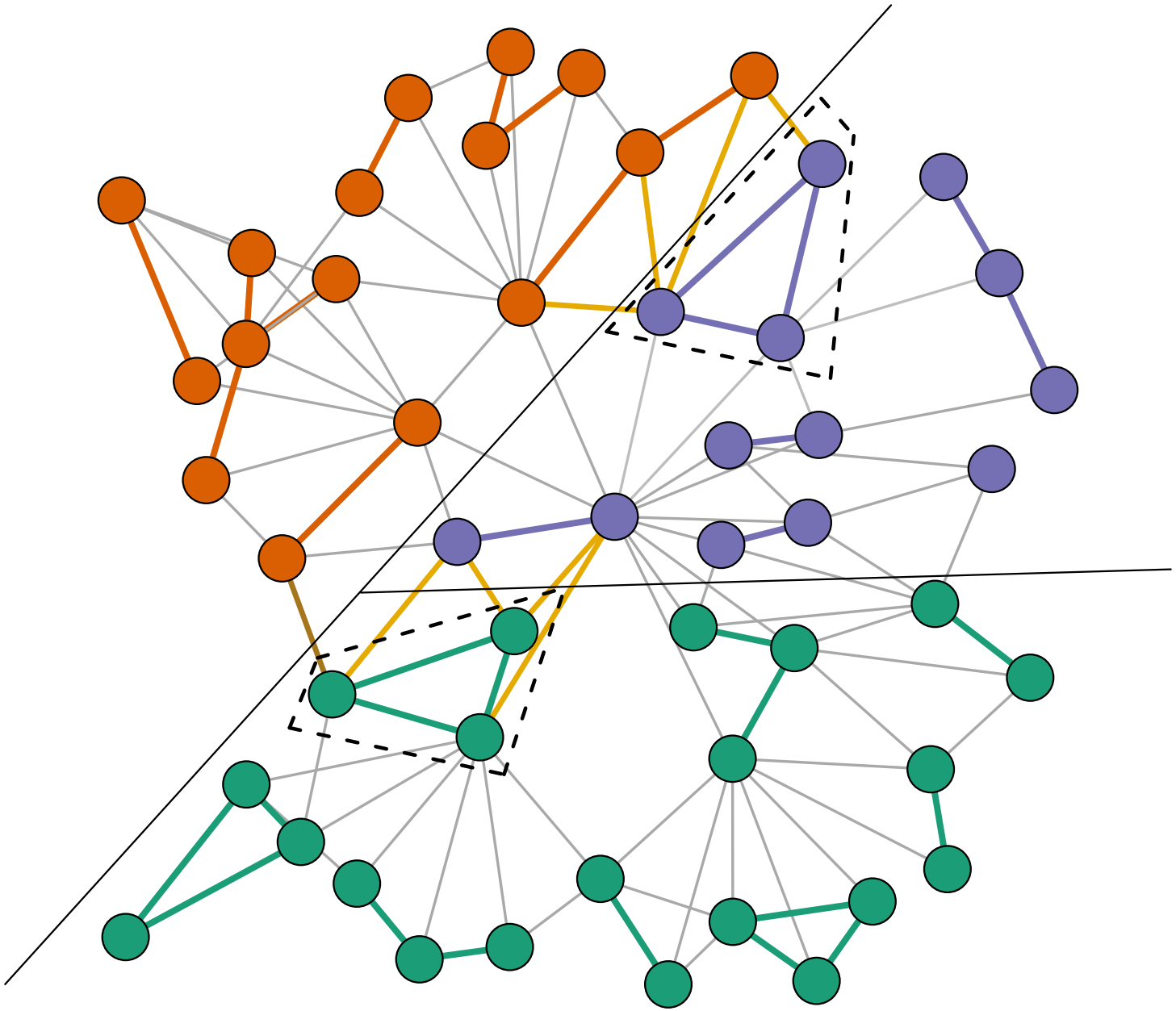
Step 1: Independently "Turn On" Each Edge with Prob. q



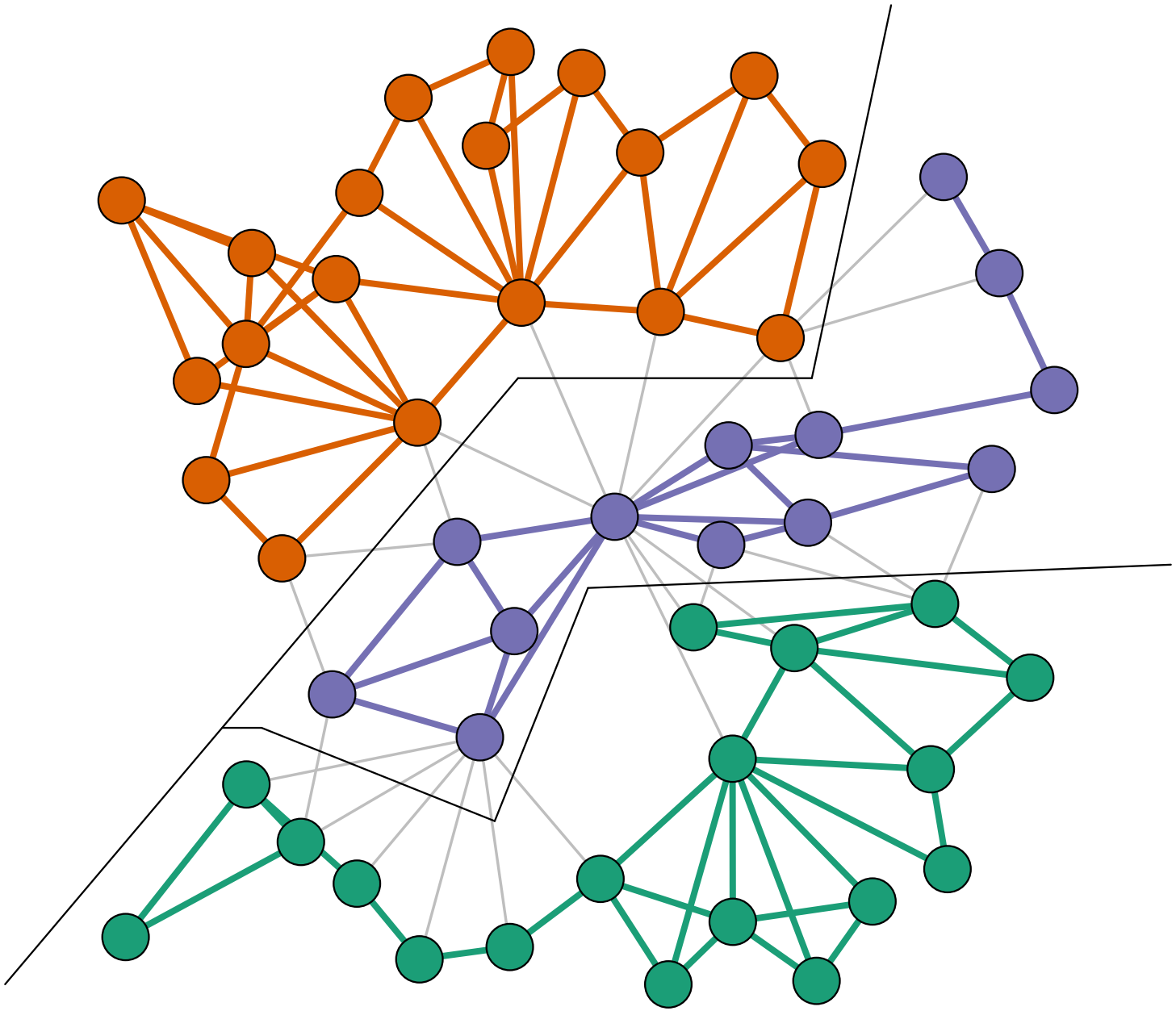
Step 2: Gather Connected Components on Boundaries



Step 3: Select Subsets of Components and Propose Swaps



Step 4: Accept or Reject the Proposal



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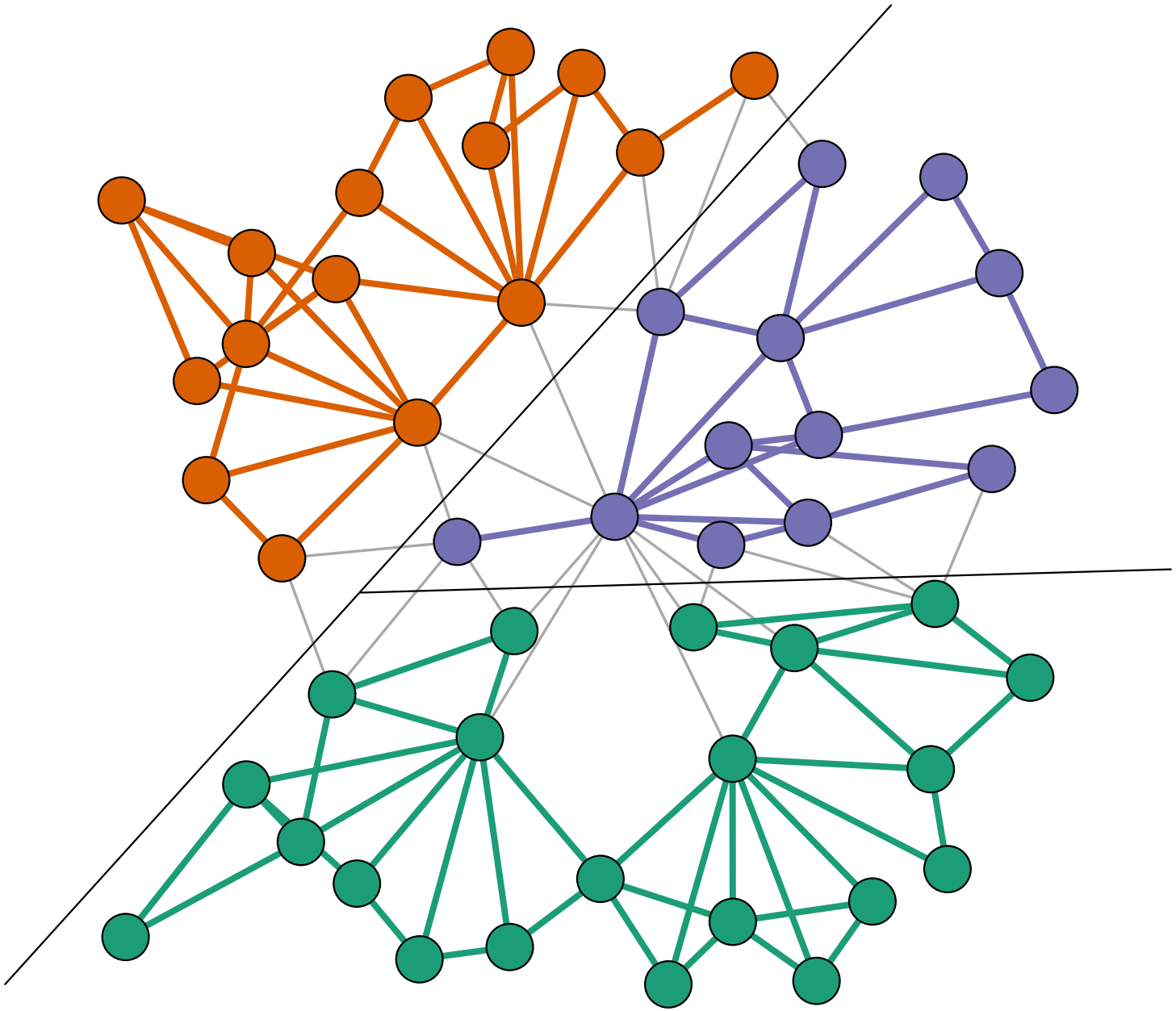


The Theoretical Property of the Algorithm

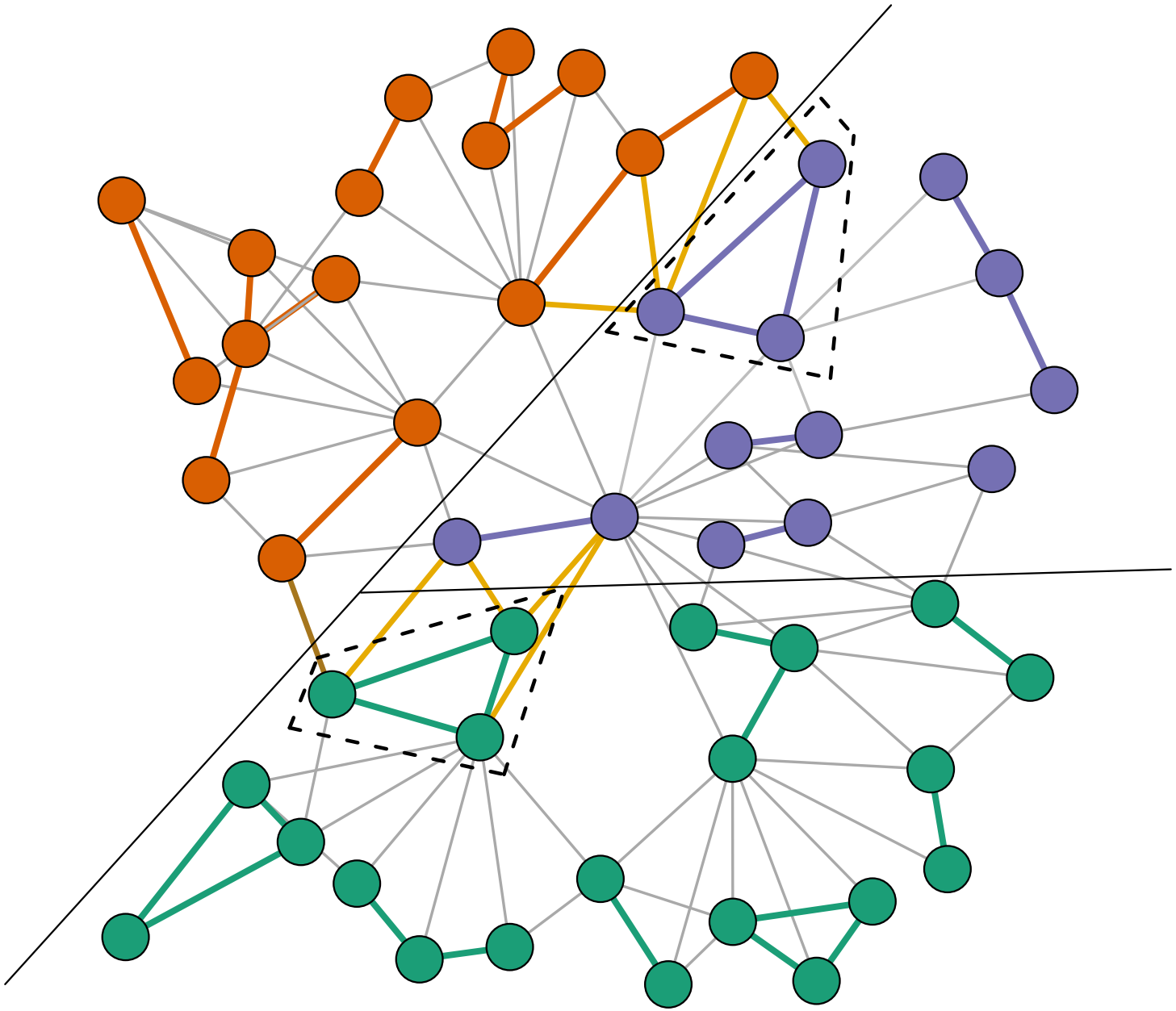
- We prove that the algorithm samples *uniformly* from the population of all valid redistricting plans

- An extension of the Swendsen-Wang algorithm (Barbu & Zhu, 2005)
- Metropolis-Hastings move from plan $\mathbf{v}_{t-1} \to \mathbf{v}_t^*$:

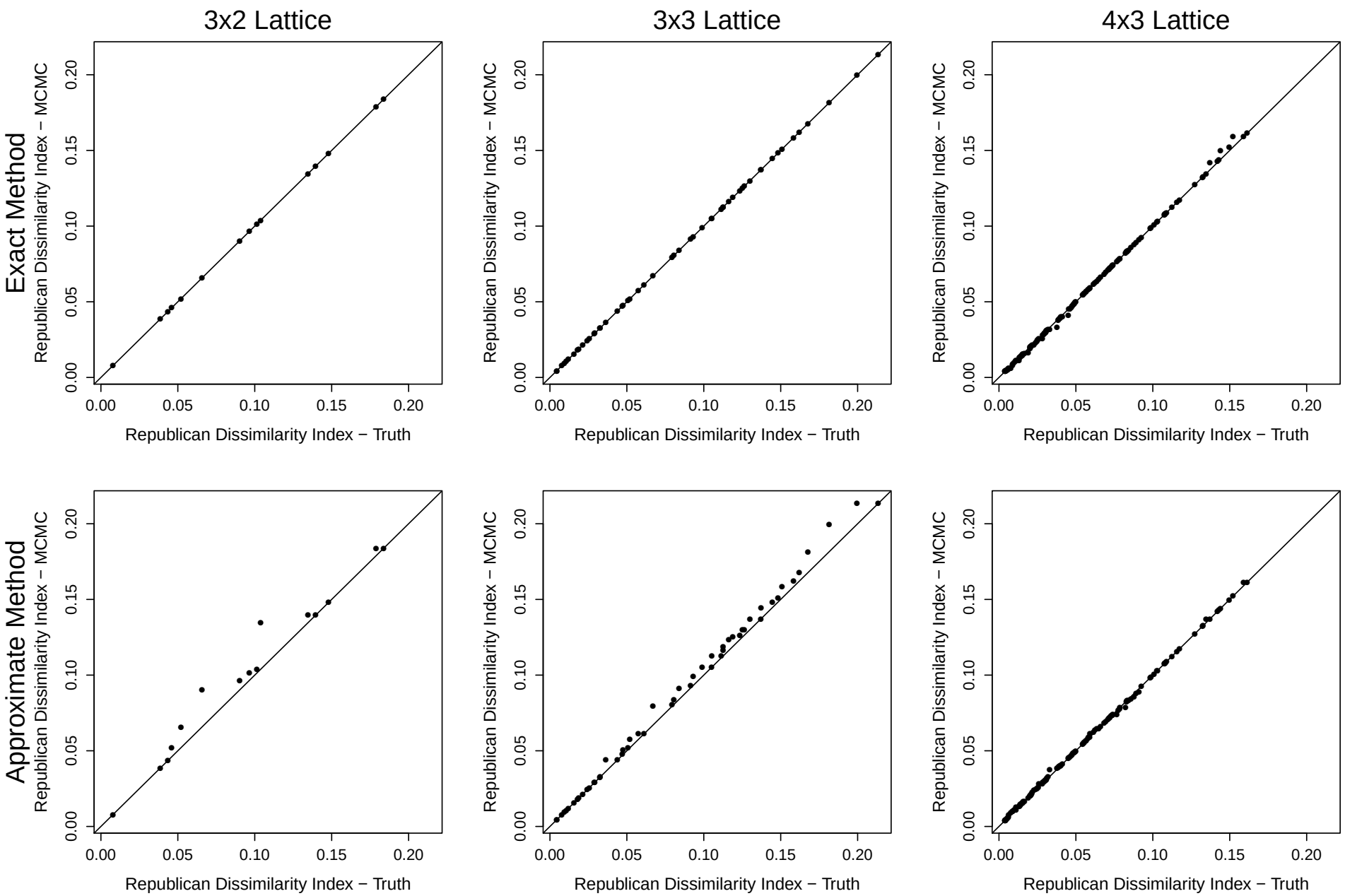
$$\begin{aligned} \alpha(\mathbf{v}_{t-1} \to \mathbf{v}_t^*) &= \min\left(1, \frac{\pi(\mathbf{v}_t^* \to \mathbf{v}_{t-1})}{\pi(\mathbf{v}_{t-1} \to \mathbf{v}_t^*)}\right) \\ &\approx \min\left(1,\ (1-q)^{|B(C^*,\mathbf{v})|-|B(C^*,\mathbf{v}^*)|}\right) \end{aligned}$$

  where q is the edge cut probability and $|B(C^*, \mathbf{v})|$ is # of edges between connected component and its assigned district in redistricting plan $\mathbf{v}$ $\rightsquigarrow$ Easy to compute
- Exact Metropolis ratio is too costly to evaluate (unless only a single unit is swapped at one time)

The Theoretical Property of the Algorithm



Accuracy of Approximation with a Long Chain



Incorporating a Population Constraint

- Equal population constraint:

$$\psi(V_k) \;=\; \left| \frac{p_k}{\bar{p}} - 1 \right| \leq \epsilon$$

  where p_k is population of district k, $\bar{p}$ is average district population, ϵ is strength of constraint (e.g., 2%)
- Strategy 1: Only propose "valid" swaps $\rightsquigarrow$ slow mixing
- Strategy 2: Oversample certain plans and then reweight
  1. Sample from target distribution f rather than the uniform distribution:

$$f(\mathbf{v}) \;\propto\; g(\mathbf{v}) \;=\; \exp\left(-\beta \sum_{V_k \in \mathbf{v}} \psi(V_k)\right)$$

     where $\beta \geq 0$ and $\psi(V_k)$ is deviation from parity for district V_k
  2. (Approximate) Acceptance probability is still easy to calculate,

$$\alpha(\mathbf{v} \to \mathbf{v}^*) \;\approx\; \min\left(1,\; \frac{g(\mathbf{v}^*)}{g(\mathbf{v})} \cdot (1-q)^{|B(C^*,\mathbf{v})| - |B(C^*,\mathbf{v}^*)|}\right)$$

  3. Discard invalid plans and reweight the rest by $1/g(\mathbf{v})$

Additional Constraints

1. Compactness (Fryer and Holden 2011):

$$\psi(V_k) \propto \sum_{i,j \in V_k, i<j} p_i p_j d_{ij}^2$$

where d_{ij} is the distance between precincts i, j

2. Similarity to the adapted plan:

$$\psi(V_k) = \left| \frac{r_k}{r_k^*} - 1 \right|$$

where r_k (r_k^*) is the # of precincts in V_k (V_k of the adapted plan)

- Any criteria where constraint can be evaluated at each district

Improving the Mixing of the Algorithm

- Single iteration of the proposed algorithm runs very quickly
- But, like any MCMC algorithm, convergence may take a long time

1. Swapping multiple connected components
   - more effective than increasing q
   - but still leads to low acceptance ratio
2. Simulated tempering (Geyer and Thompson, 1995)
   - Lower and raise the "temperature" parameter β as part of MCMC
   - Explores low temperature space before visiting high temperature space
3. Parallel tempering (Geyer 1991)
   - Run multiple chains of the algorithm with different temperatures
   - Use the Metropolis criterion to swap temperatures with adjacent chains

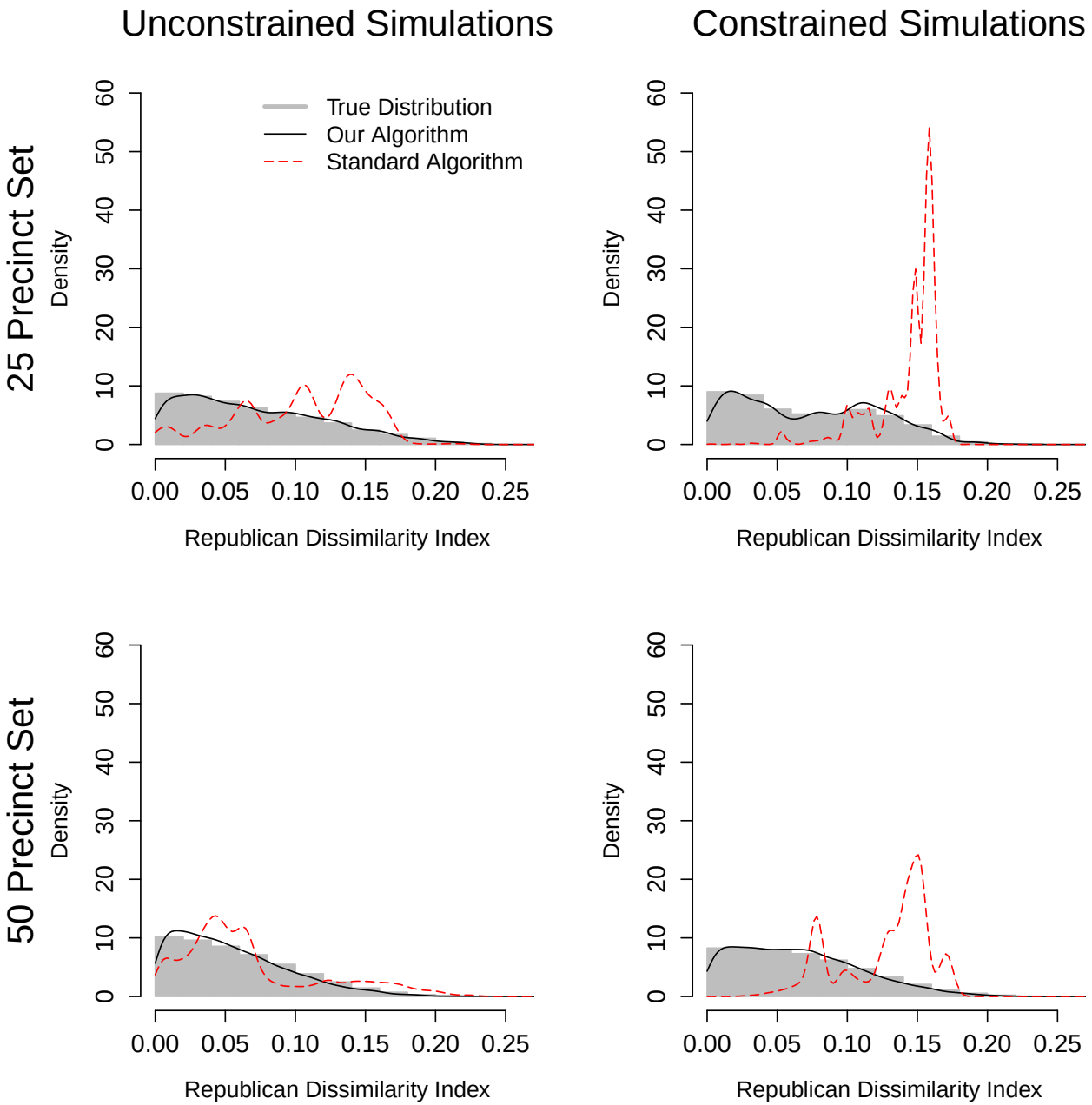
A Small-Scale Validation Study

- Evaluate algorithms when all valid plans can be enumerated

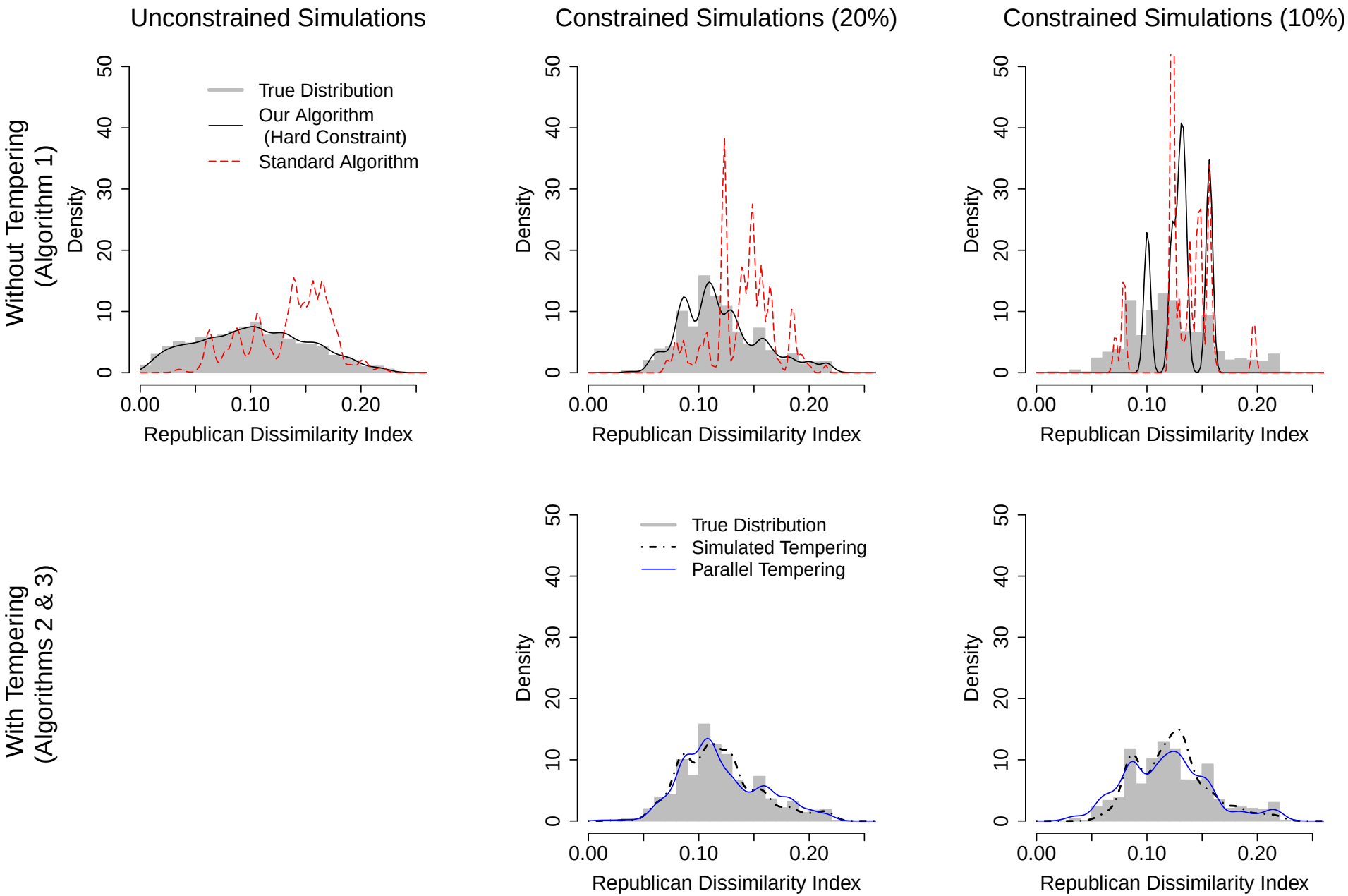
- # of precincts: 25 and 50
- # of districts: 2 and 3 for the 25 set, and 2 for the 50 set
- With and without a "hard" population constraint of 20% within parity
- Also, consider simulated and parallel tempering

- Comparison with the "random seed-and-grow" algorithm via the "industry-standard" `BARD` package (Altman & McDonald 2011)
- 10,000 draws for each algorithm

Our Algorithm vs. Standard Algorithm

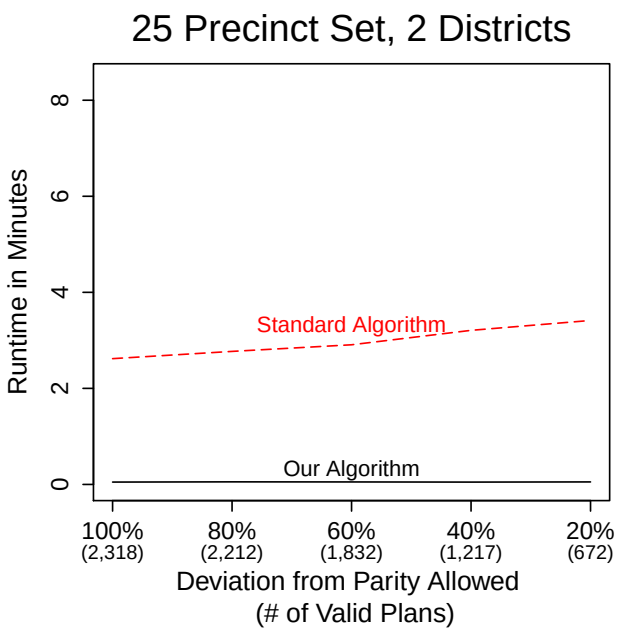


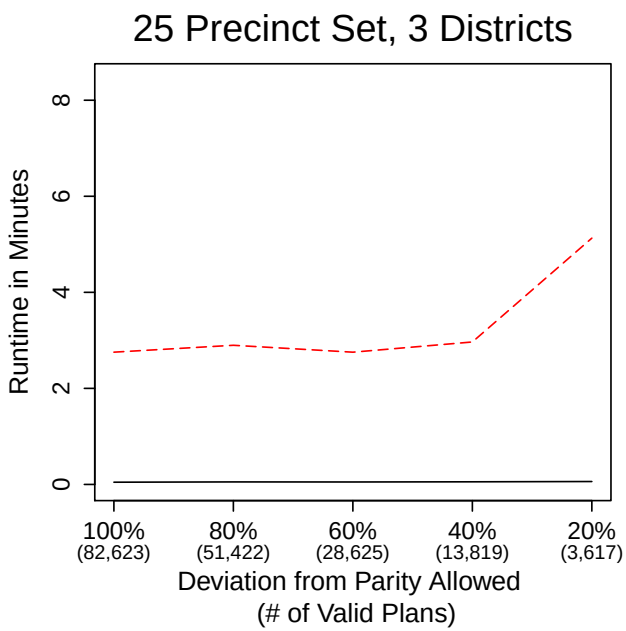
Simulated and Parallel Tempering

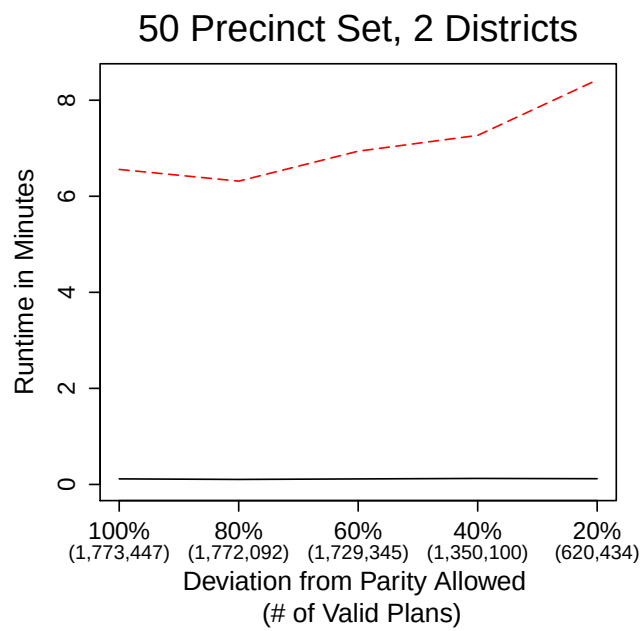


Runtime Comparison

- Run each algorithm for 10,000 simulations under different population constraints



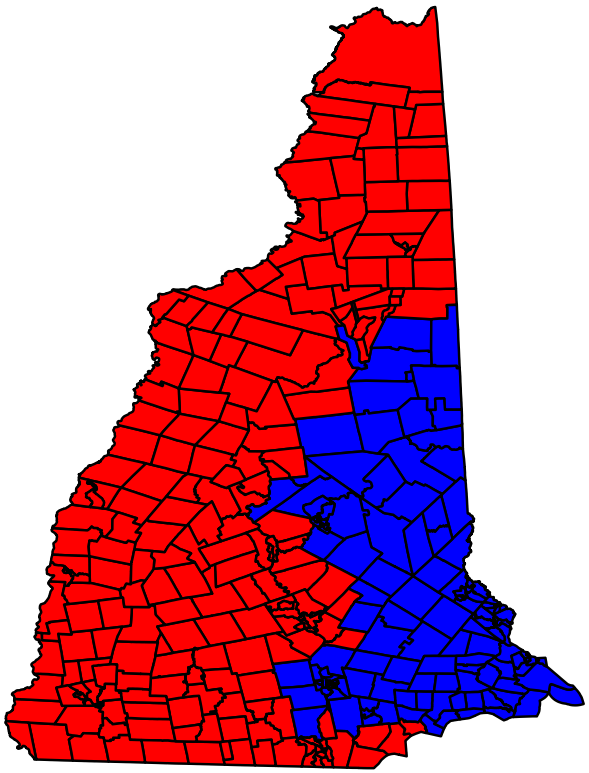




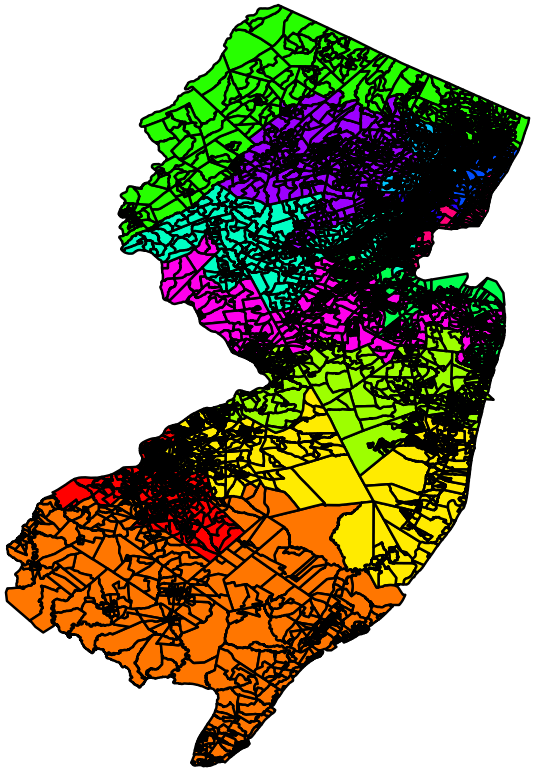
An Empirical Study

- Apply algorithm to state election data:
  1. New Hampshire: 2 congressional districts, 327 precincts
  2. New Jersey: 4 congressional districts, 6,344 precincts
  3. 5% deviation from population parity

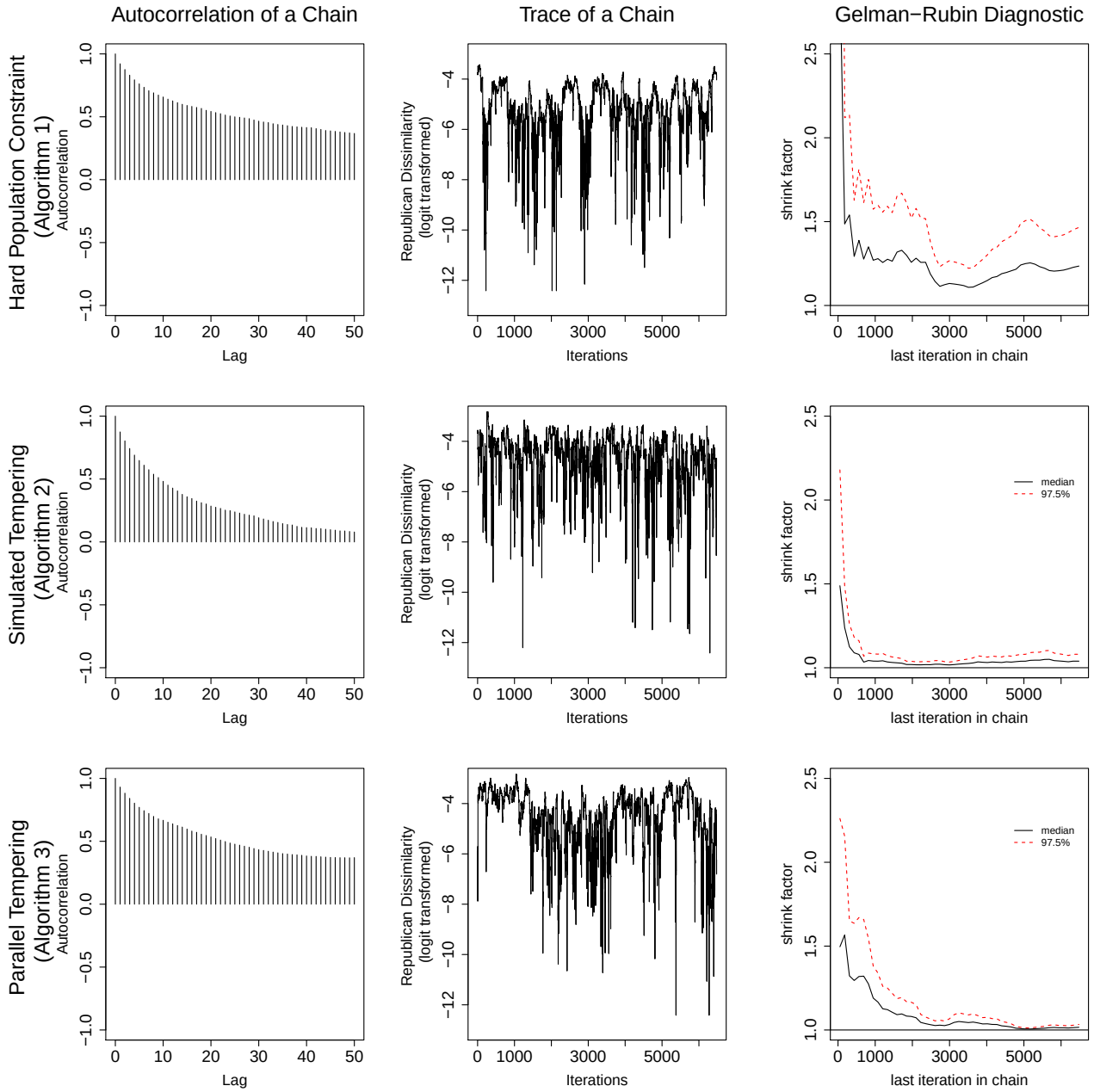
New Hampshire



New Jersey



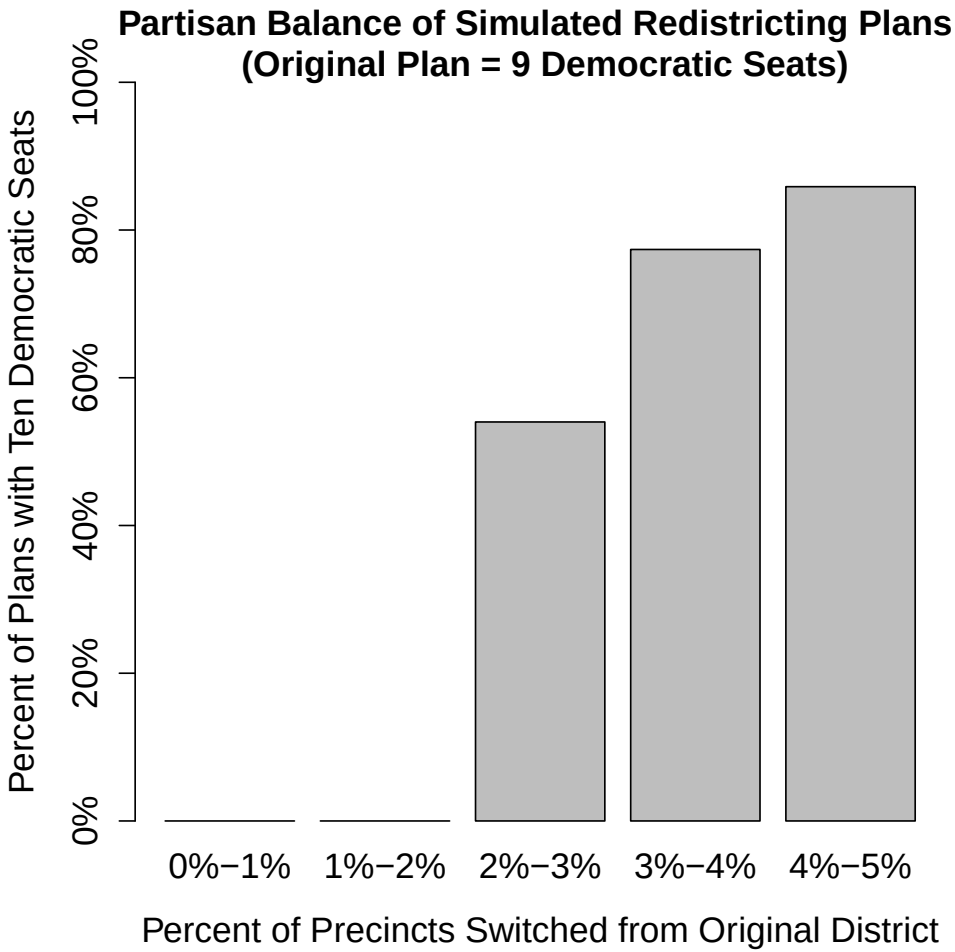
New Hampshire: Tempering Works Better

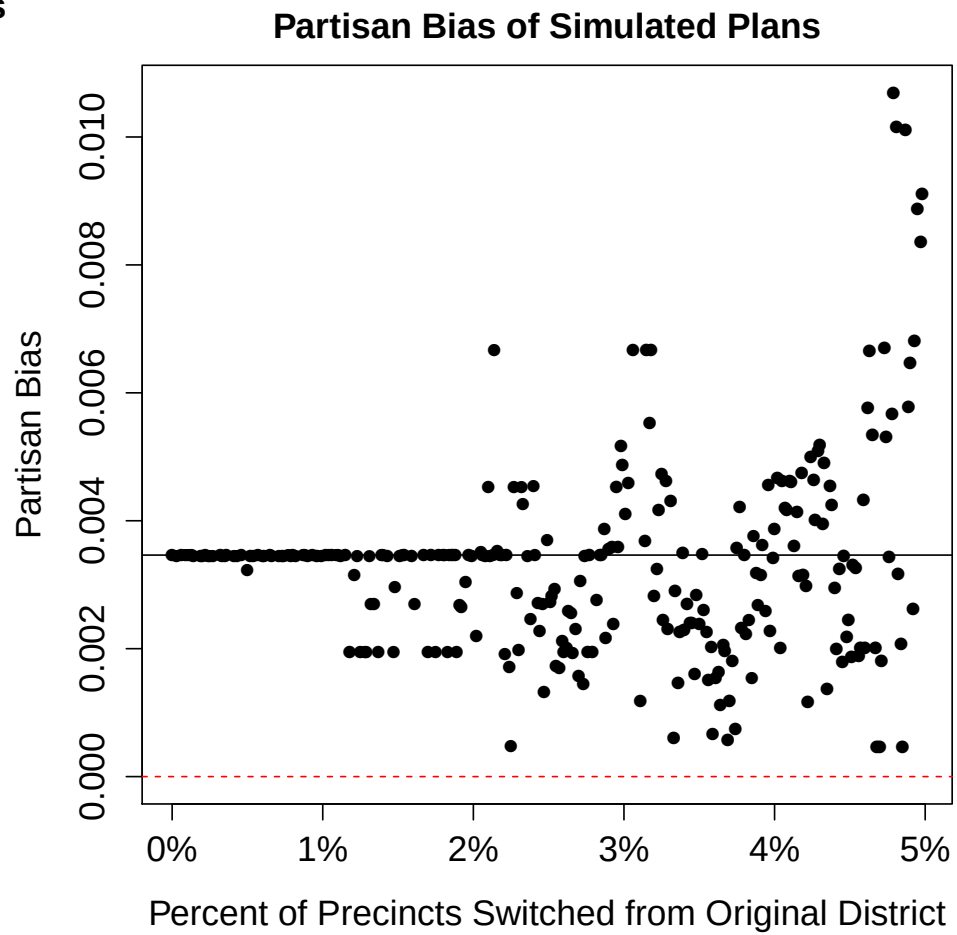


Redistricting Plans that are Similar to the Adapted Plan

- Question: How does the partisan bias of the adapted plan compare with that of similar plans?
- Two measures:
  1. Number of Republican winners under each plan
  2. Partisan bias (Gelman & King, 1994): Deviation of the seats-votes curve from partisan symmetry under each plan
- Geweke diagnostics: comparison of means within a single chain

Partisan Implications of "Local Exploration"





Concluding Remarks

- Scholars use simulations to characterize the distribution of redistricting plans
- Many optimization algorithms but very few simulation methods
- Commonly used algorithms are adhoc
- We propose a new MCMC algorithm that has:
  - better theoretical properties
  - superior speed
  - better performance in validation and empirical studies
- Future research:
  - Apply the method to historical redistricting data
  - Explore alternative ways to tackle large scale redistricting problems

References

1. Paper at http://imai.princeton.edu/research/redist.html
2. R package at https://github.com/redistricting/redist
3. Comments and suggestions: send them to kimai@princeton.edu