

On Zone Balancing of Peer to Peer Networks: Analysis of Random Node Join

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Overview

Motivation



Single-point

Random
Split

Center
Split



Multi-point

Random
 d -sampling

Deterministic
 d -sampling



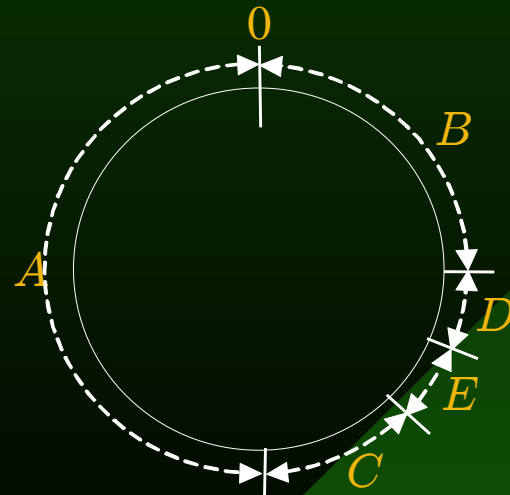
P2P Simulations



Conclusion

Motivation

- Structured P2P systems construct DHTs (Distributed Hash Tables) for efficient routing
 - Chord, CAN, de Bruijn
- Data objects are hashed into some virtual coordinate spaces
- Each user holds a zone in the DHT space
 - Stores data objects within its zone and answers queries for these objects



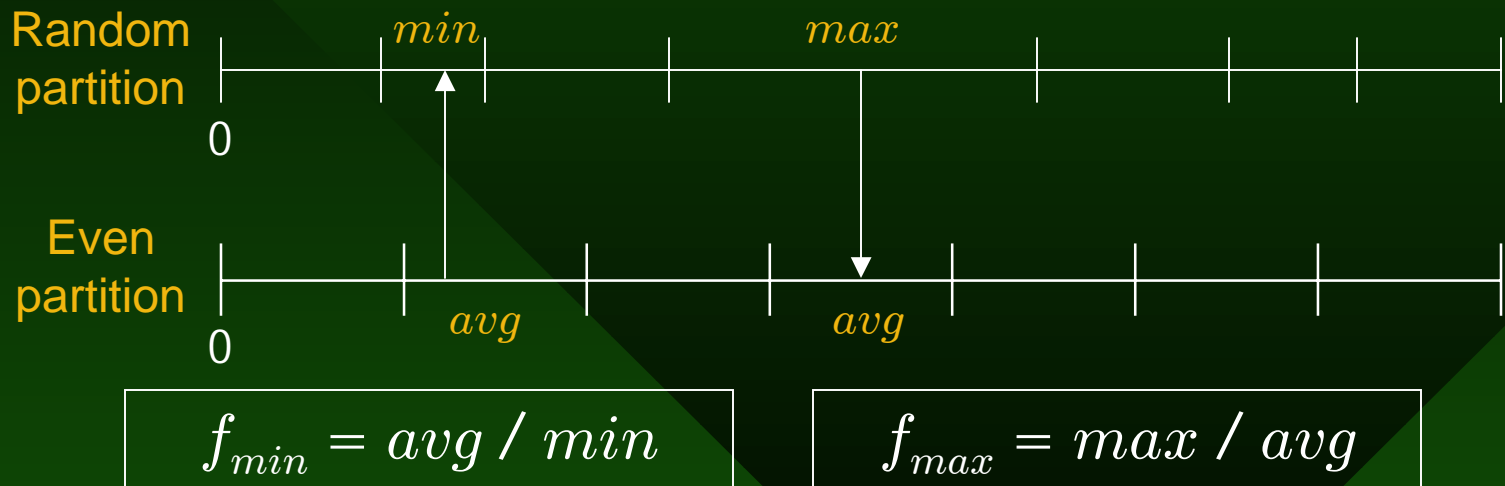
An instance of zone partition

Motivation 2

- Notice that the amount of user load is proportional to zone size
 - Imbalance can lead to “hotspots” and lower performance
- In addition, graph structure is unbalanced
 - Which leads to increased diameter, smaller node degree, lower bisection width
- Our paper studies how zone-balancing decisions during node join affect the resulting zone sizes
 - We derive the probability bounds on the maximum and minimum zone sizes

Basics

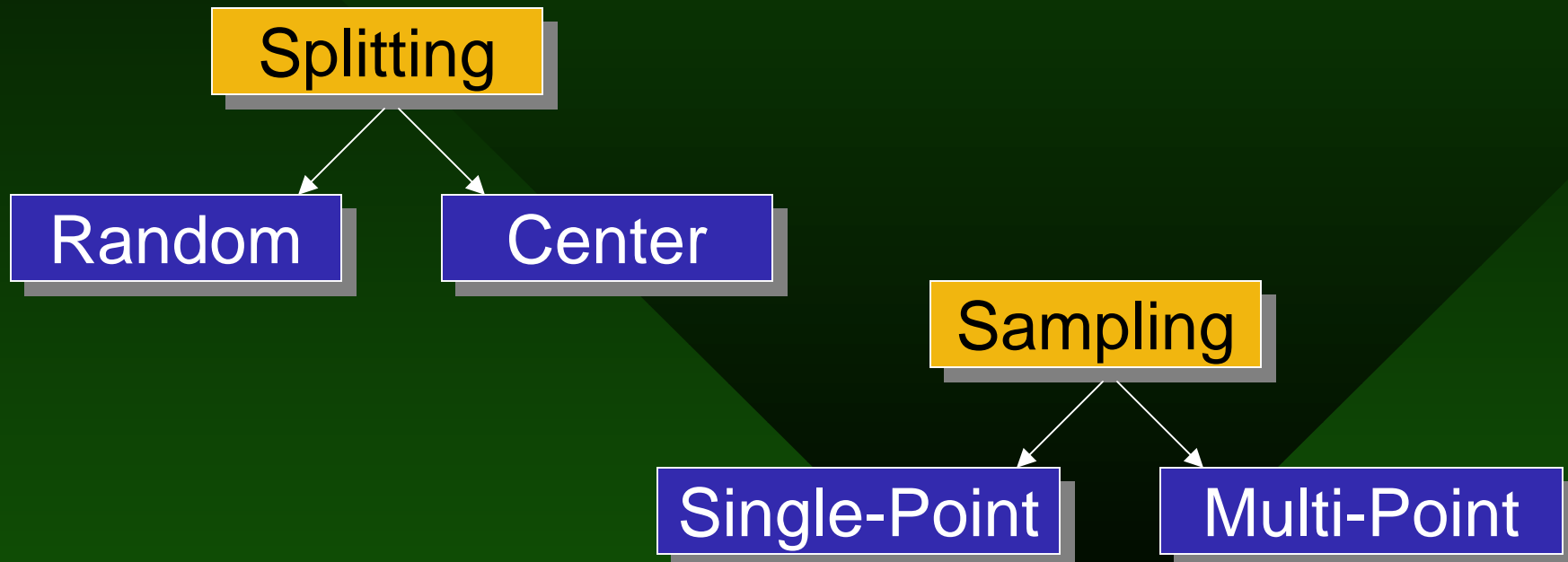
- Consider a system with n users
 - Assume a sequential join process
- Define two metrics for load balancing:



- We focus on the bounds of these two metrics that hold with probability $1 - n^{-\epsilon}$ ($\epsilon > 0$)

Random Join Process

- Each new user randomly samples one or more existing peers and splits one of their zones
- The join decision includes two factors:



Random Join Process 2

- We will compare these algorithms in terms of f_{max} and f_{min}
 - The optimal bound for the two metrics is 2
 - No method can achieve better load-balancing
- Due to the time limit, we skip the single-point algorithms
 - Summary for random and center splits:

Random	Center
$f_{max} \leq (1 + \epsilon) \log n$	$f_{max} \leq (1 + \epsilon) \log n$
$f_{min} \leq n^{1+\epsilon}$	$f_{min} \leq 3.246^{\sqrt{\log n}}$

A Big Map

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P2P Simulations



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Multi-Point Center-Split

- Next we examine multi-point schemes
 - We use center-split for the rest of the talk
- Greedy methods
 - Motivated by the “power of two choices”
- Idea: extend the center-split model to sample d random points before the actual join
- Intuitive observation:
 - The more points sampled, the better the graph is balanced, but what are the actual bounds?

Multi-Point Center-Split 2

- The extreme case is to sample every peer
 - The resulting f_{max} is always optimal and concentrates on the ideal value 2
- However, this method will suffer from huge traffic overhead
- Thus, the tradeoff is between:
 - The balancing performance of the algorithm, and
 - The amount of sampling traffic
- Next we study two multi-point schemes and present our analysis of this problem

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Purely Random d -sampling

- The method samples d independent uniformly random points X_1, X_2, \dots, X_d
 - Splits the largest zone among the d choices
- How does the performance improve as a function of d ?
- Based on the “balls-into-bins” model, we derive an asymptotic bound on f_{max}
 - The analysis is intractable when applying this model to f_{min}
 - We leave this direction for future work

Purely Random d -sampling 2

- Theorem 1: Under d -point sampling and center-splits, the following bound holds with probability at least $1 - n^{-\varepsilon}$

$$f_{max} \leq 2 + \frac{(1 + \varepsilon) \log n}{d} - \frac{\Theta(\log(d + \log n))}{d}$$

- For $d = 1$, it reduces to the single-point model

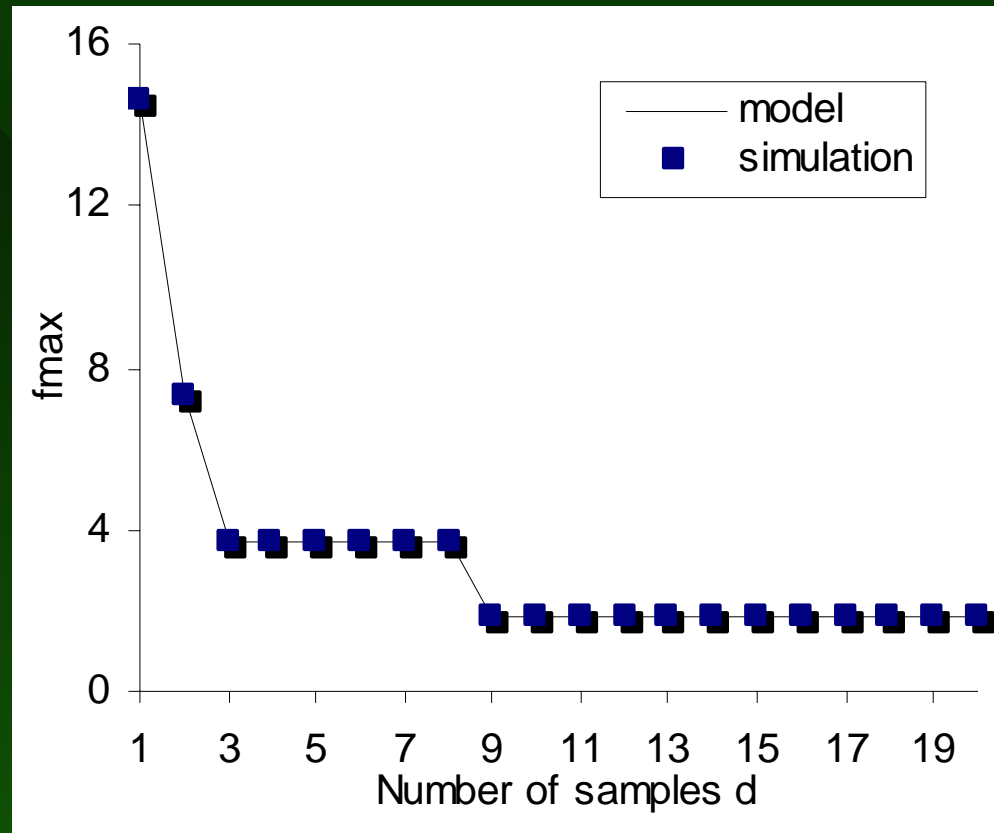
$$f_{max} \leq (1 + \varepsilon) \log n - \Theta(\log \log n)$$

- For $d \geq 2$, the term $(1 + \varepsilon) \log n$ is scaled down by a factor of d

- The “power of two choices” bound $\log \log n / \log d$ is not achieved here

Simulating Random d -sampling

- Each of the following simulations is run for 1,000 graphs with 30,000 nodes each



Further Discussion

- For $d = c \log n$,

$$f_{max} \leq 2 + \frac{1 + \varepsilon}{c} - o(1)$$

- For $c \rightarrow \infty$, the second term goes to zero
 - And f_{max} is bounded by 2 with high probability
- Recall from the single-point method,
 - $f_{max} \leq 28$ for $n = 10^6$ with probability $1 - 1/n$
- The improvement is significant
 - But results in additional traffic overhead

Reducing Traffic Overhead

- How to reduce the join overhead?
 - While keeping the graph balanced
- Idea:
 - Randomly sample a peer
 - Then deterministically sample its neighbors
 - Subsequently walk along the edges of the graph to find additional peers to sample
- Two walking strategies:
 - *Random walk* selects arbitrary (random) neighbors
 - *Biased walk* selects the largest neighbors

Reducing Traffic Overhead 2

- Intuition: “larger” nodes are more likely to know additional “large” nodes
- This reduces the join overhead by a factor of $\Theta(kD_{av})$
 - k is graph degree and D_{av} is the average distance
- The exact analysis is nontrivial since the walk process depends on the state of peers
 - We leave the exact model for future work
- Instead, we study a similar deterministic model
 - According to our analysis, it provides a lower bound on the performance of the other d -walk models

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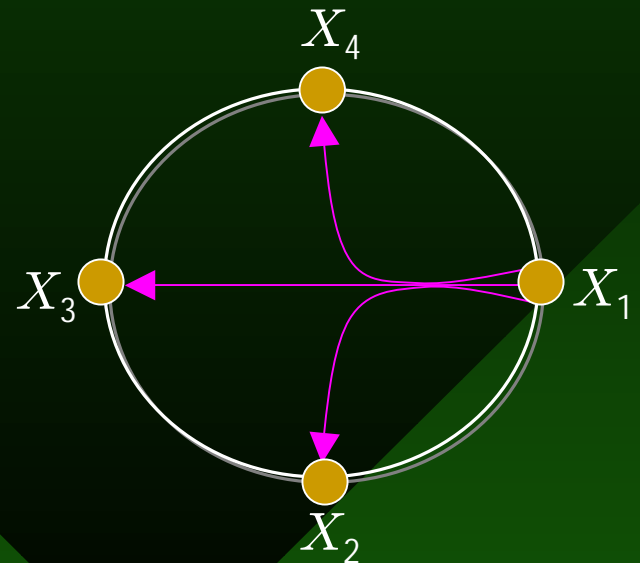


Conclusion

Deterministic d -sampling

- The model samples a random point X_1
 - Then checks $d - 1$ additional points according to a simple deterministic rule
 - Points $X_2, \dots, X_i, \dots, X_d$ are obtained by adding i/d of the total size of the DHT space to X_1

- An example of $d = 4$
 - X_1 is the first random sample
 - The points X_2, X_3, X_4 are found by adding $1/4, 1/2,$ and $3/4$ of the circle's circumference to X_1



Deterministic d -sampling 2

- Theorem 2: In deterministic sampling, the following bound holds with probability at least

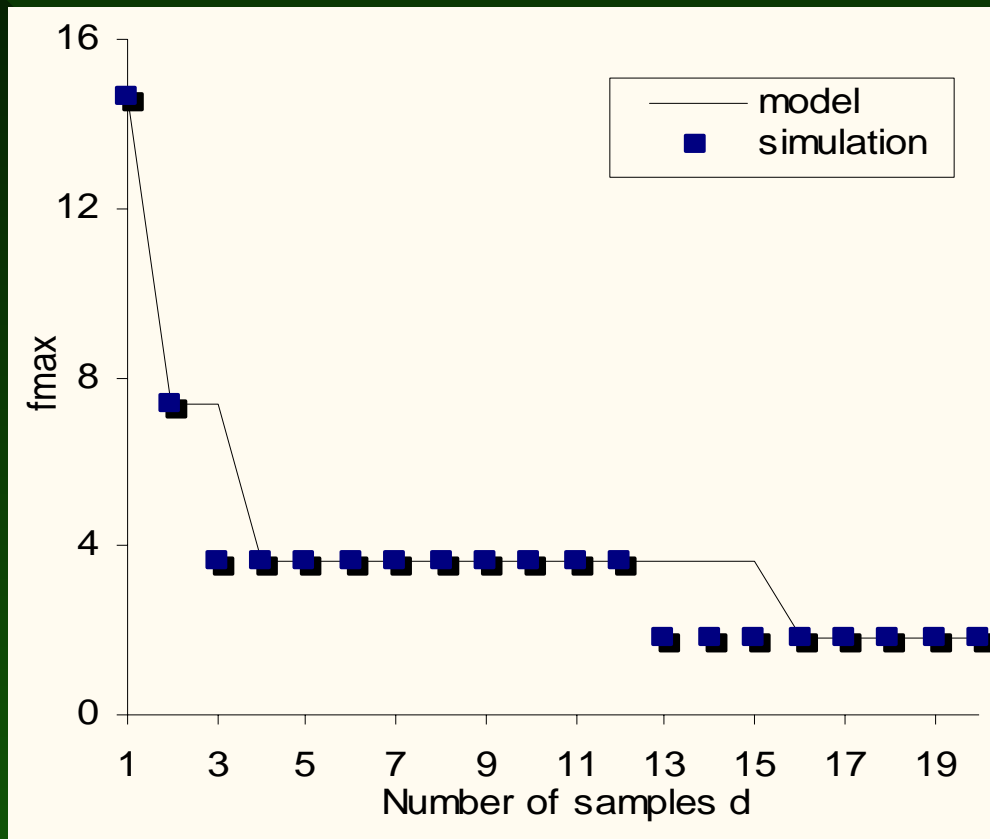
$$1 - n^{-\varepsilon} \quad f_{max} \leq 2 + \frac{(1 + \varepsilon) \log n}{d} + \eta - \frac{\Theta(\log \log n)}{d}$$

where $\eta = \log \left(1 + \frac{1+\varepsilon}{c} \right) + \log \left(1 + \frac{1+\varepsilon}{c} \right) + \dots$

- This result differs from that of random d -sampling by a constant η
- Notice that η is positive
 - Thus, the deterministic model is worse than the random model
 - But how much is the difference?

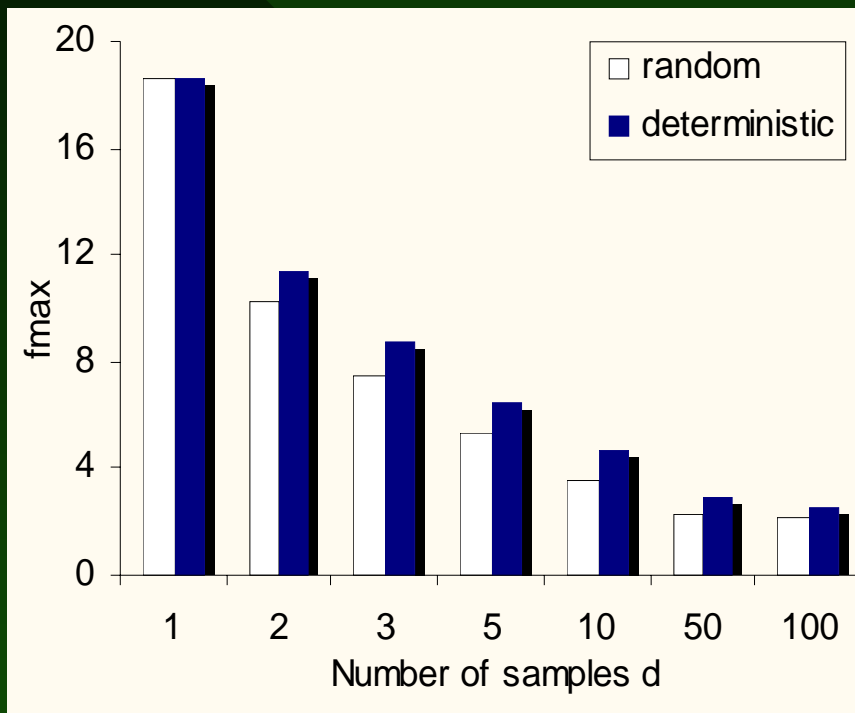
Simulating Deterministic Sampling

- The model is conservative on some points
 - Round-off errors at d not powers of 2

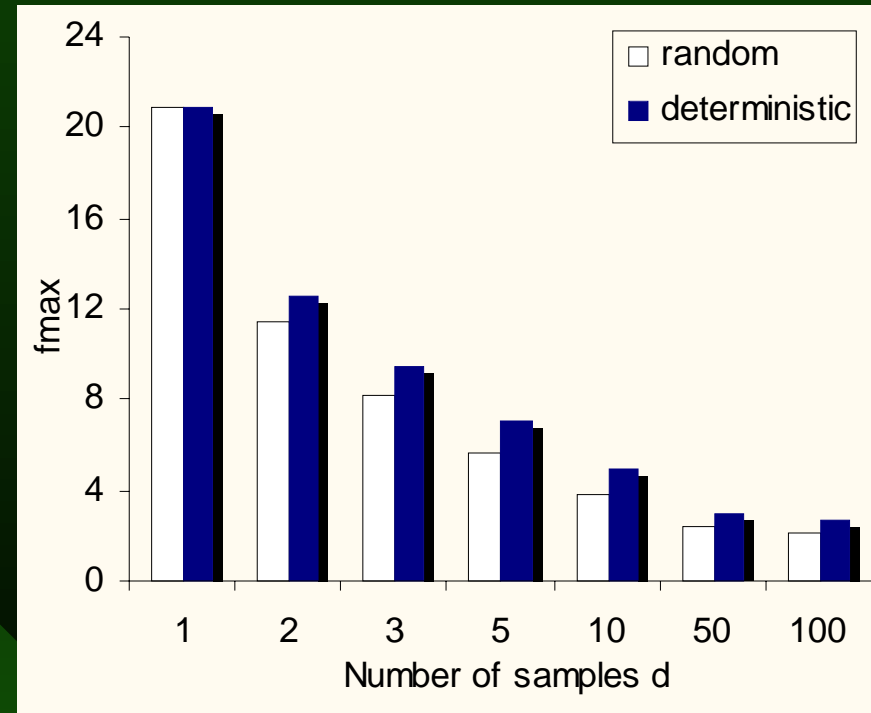


Purely Random vs Deterministic

- With the previous results on f_{max} , we compare the two multi-point models ($\varepsilon = 0.22$)



$n=10^6$



$n=10^7$

Purely Random vs Deterministic

- Further question:
 - How many samples does the deterministic model need to approximate the random model?
- Theorem 3: Assuming that the random method samples $c_1 \log n$ points and the deterministic method samples $c_2 \log n$ points, the corresponding upper bounds on f_{max} are equal if

$$c_2 = \frac{(1 + \varepsilon)c_1}{1 + \varepsilon - c_1 \log\left(1 + \frac{1 + \varepsilon}{c_1}\right)}$$

Pure Random vs Deterministic 3

- For $c_1 = 1$ ($f_{max} \leq 4$) and $\varepsilon = 1$ (probability $1-1/n$), the two methods are equivalent if
 - The deterministic model samples 2.2 times more points than the random model
- For $c_1 = 2$ ($f_{max} \leq 3.5$) and $\varepsilon = 2$ (probability $1-1/n^2$), the difference is by a factor of 5.1
- In summary:
 - Each model has its benefits (low overhead vs. performance)
- What about graph properties?

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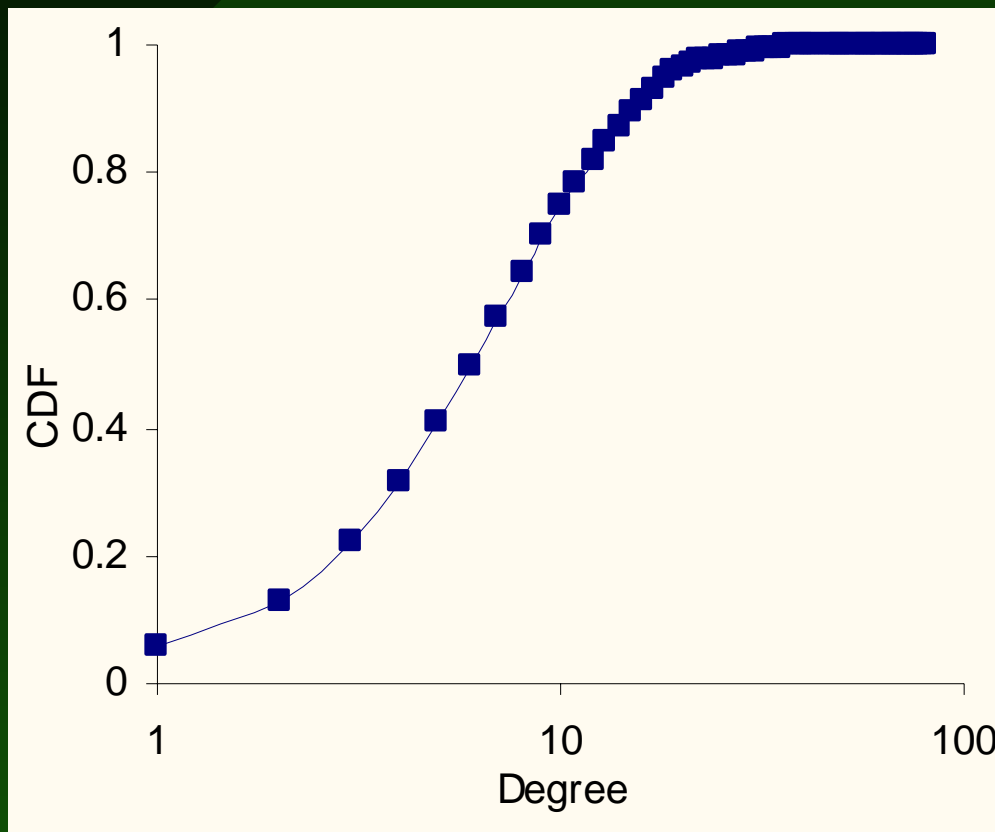
Conclusion

P2P Simulations

- We next compare the performance of multi-point methods in P2P simulations
 - Our main metric of interest is the degree distribution
- Three models
 - Purely random d -sampling
 - Random walk
 - Biased walk
- De Bruijn DHT (based on ODRI, SIGCOMM 2003) with $n = 30,000$ nodes and degree $k = 8$

Degree Distribution - CDF

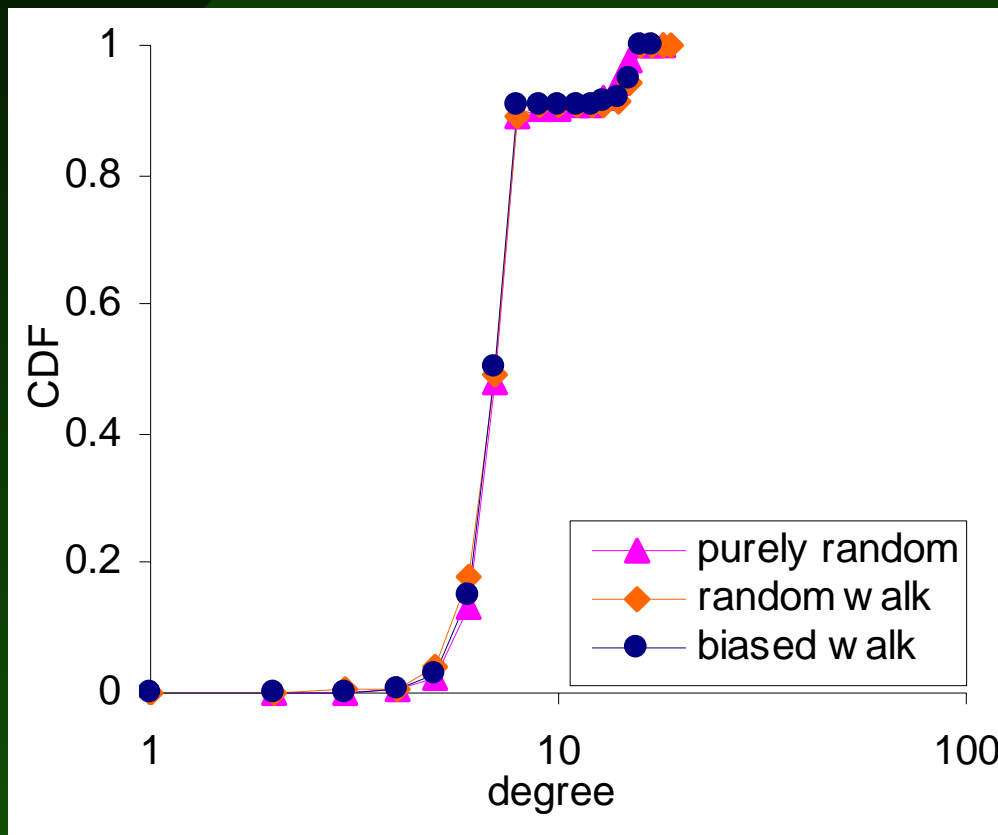
- Single-point, center-split scheme sets the basis for comparison



- 100 iterations
- Largest degree 81
- 5.7% of all nodes have degree 1
- 13% with degree 1 or 2

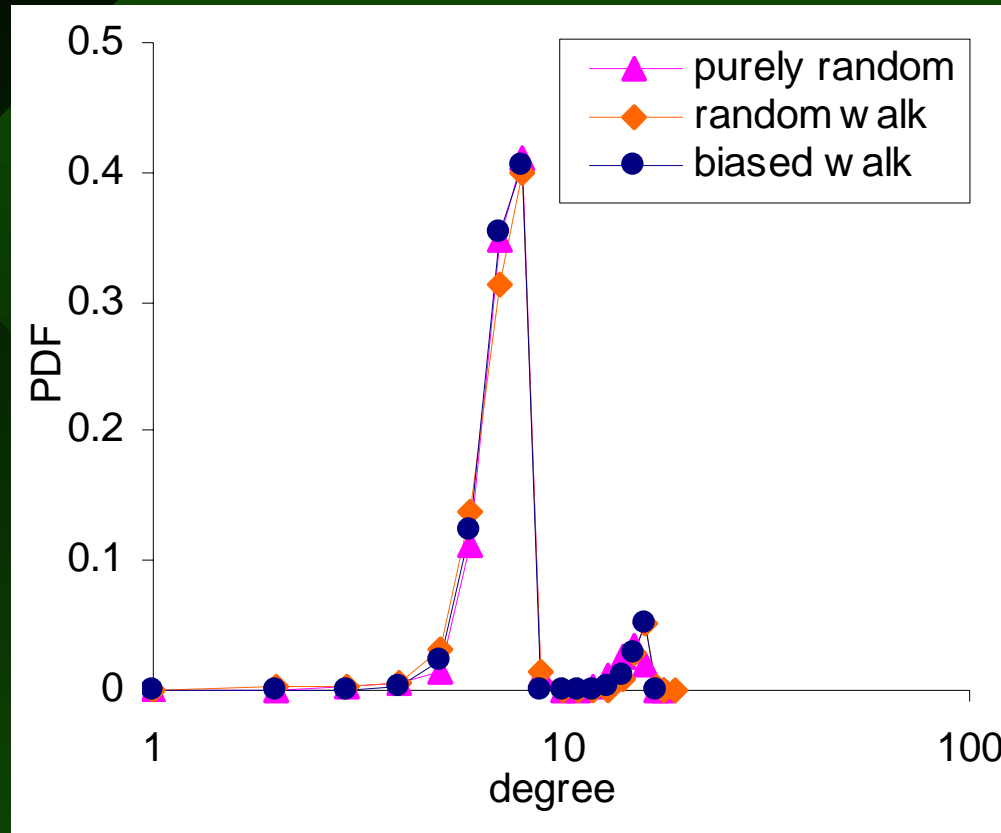
Degree Distribution - CDF

- Multi-point schemes perform much better



- Purely random
 - $d_1 = 11$
- Deterministic
 - $d_2 = 24 = 2.2d_1$
- 40% of the nodes have the ideal degree 8

Degree Distribution - PDF



- Overhead: random 55 messages per join and deterministic 7 per join, but performance is similar

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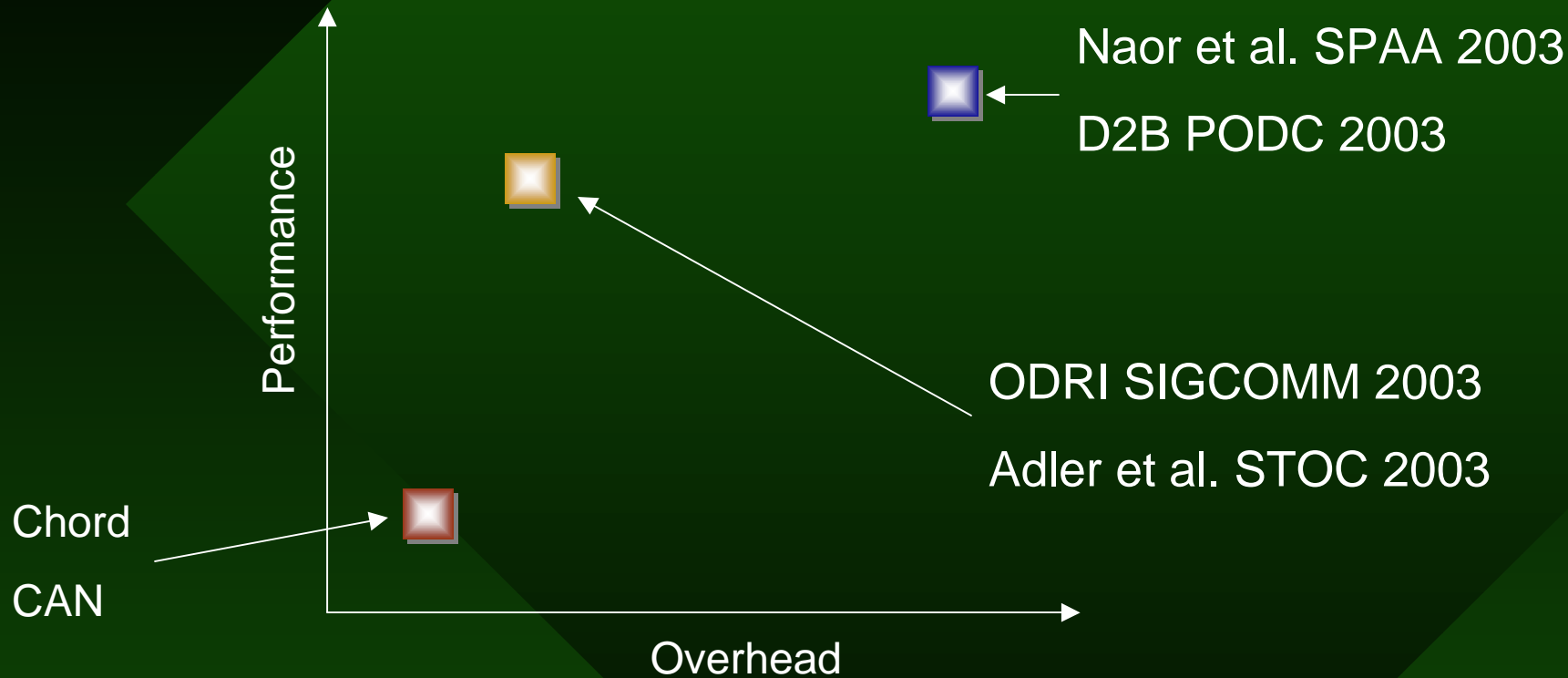


P2P Simulations



Conclusion

Conclusion



Legend:



Single-point center split



Deterministic



Purely random