

# Motion/Path Planning II

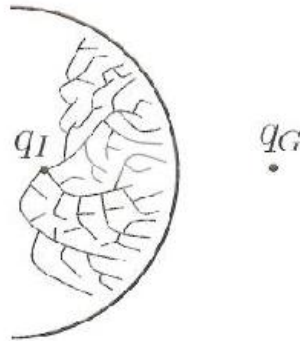
## Sampling techniques

### Approaches

- Cell decomposition
- Roadmaps
- Sampling Techniques  
(RRT, DRT, PRM,..)
- On-line algorithms  
D\*, ARA\*,..

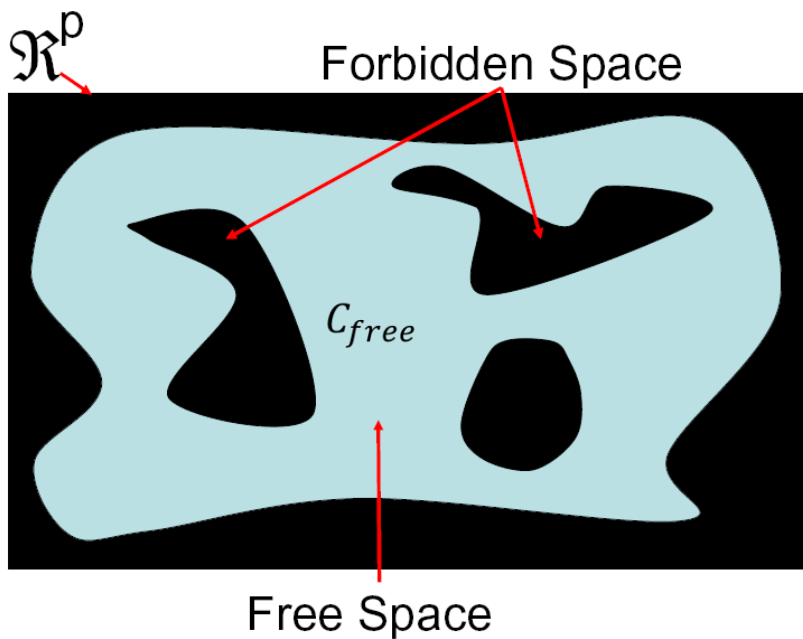
Completely describing and “optimally” exploring the C-space is too hard in high dimension + it is not necessary →  
Limit ourselves to finding a “good” sampling of the C-space

- This is why sampling of the entire space (rather than searching from start state) is necessary:



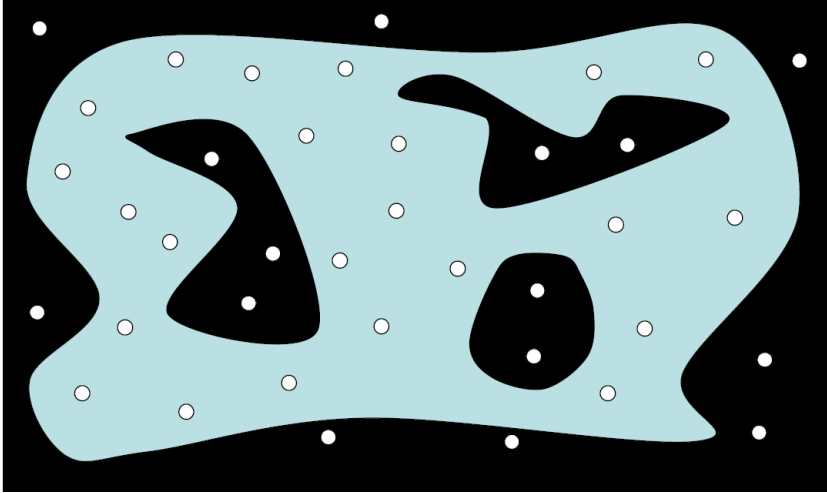
Example from Steve Lavalle

## Sampling Techniques



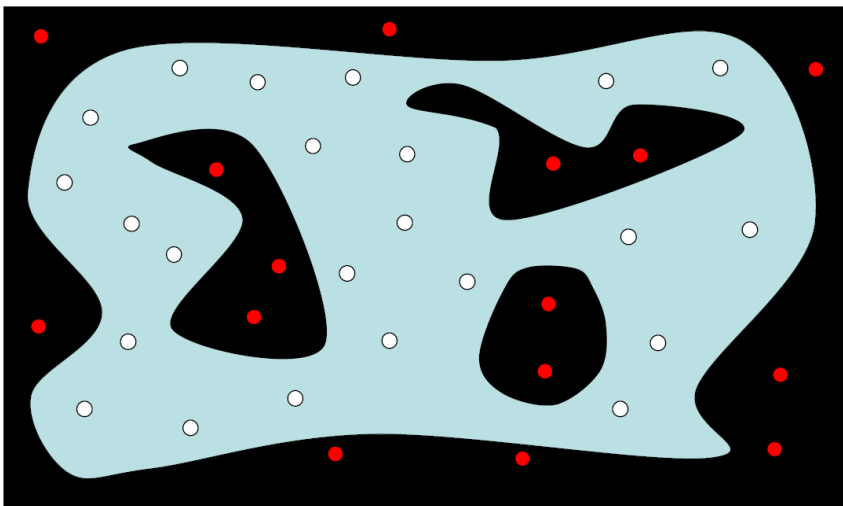
## Sampling Techniques

Sample random locations



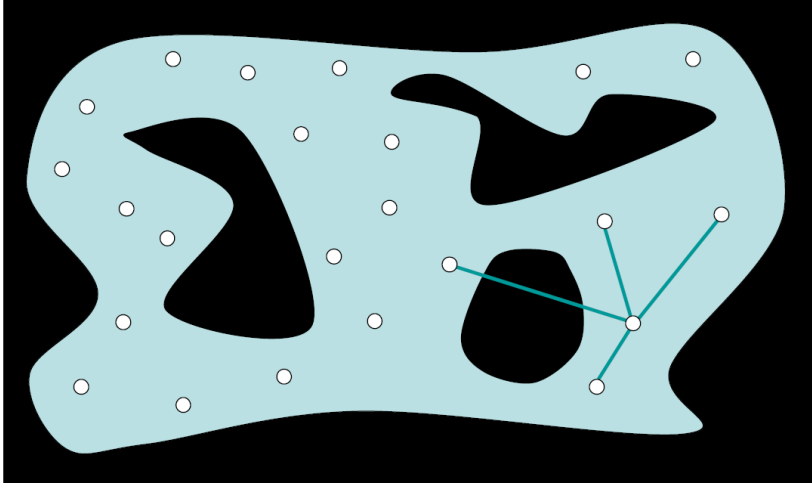
## Sampling Techniques

Remove the samples in the forbidden regions



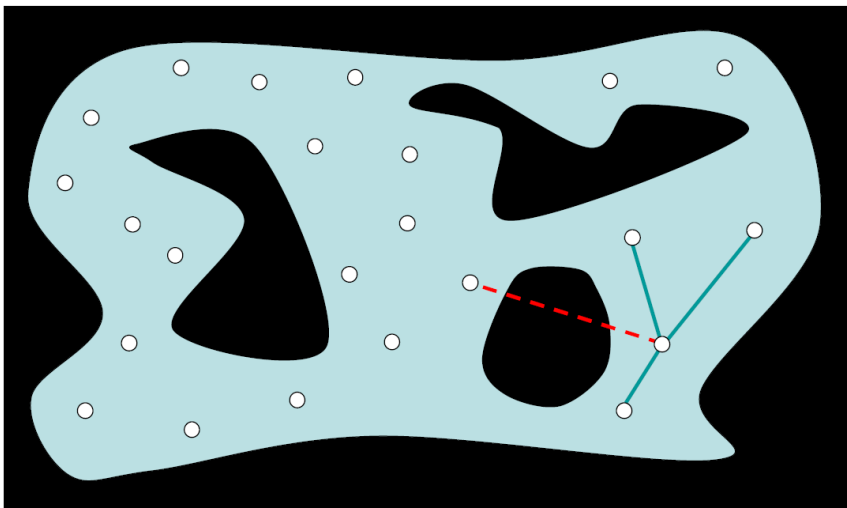
## Sampling Techniques

Link each sample to its  $K$  nearest neighbors



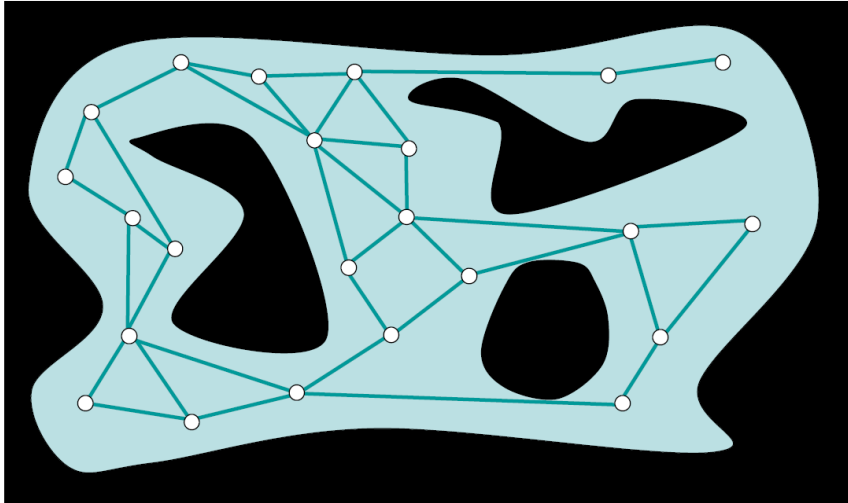
## Sampling Techniques

Remove the links that cross forbidden regions



## Sampling Techniques

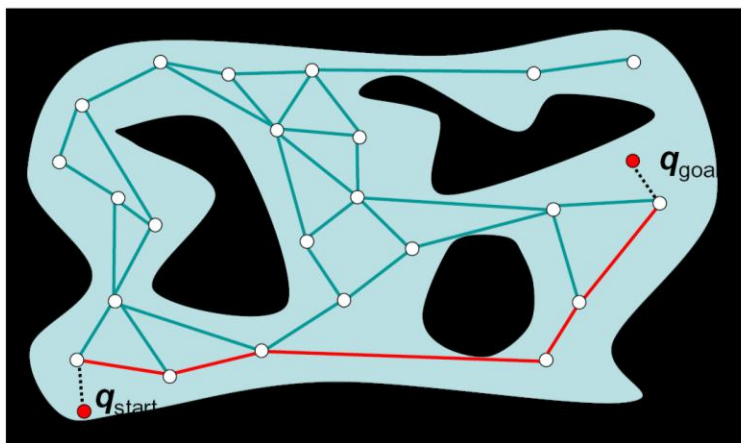
Remove the links that cross forbidden regions



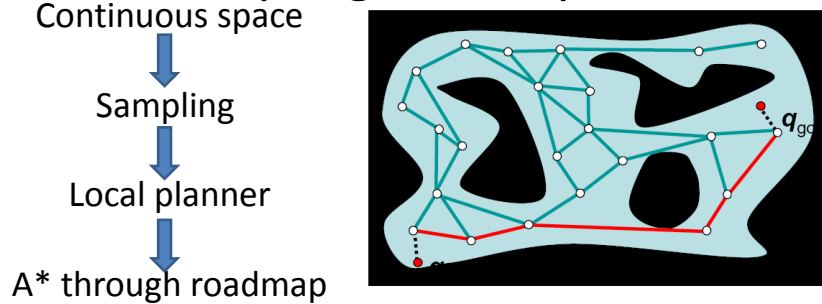
The resulting graph is a *probabilistic roadmap (PRM)*

## Sampling Techniques

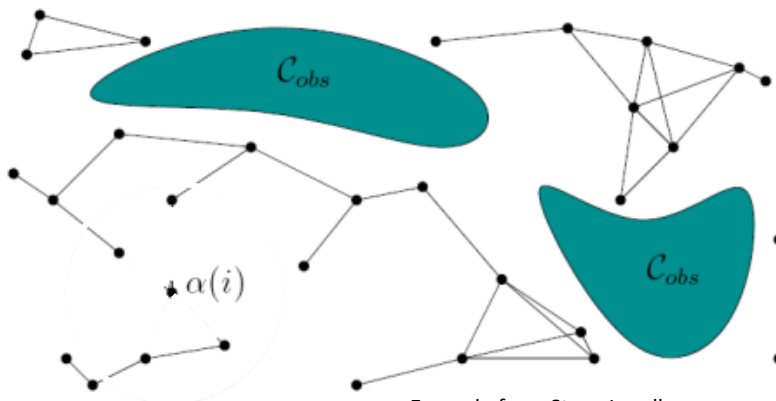
Link the start and goal to the PRM and search using A\*



# Sampling Techniques



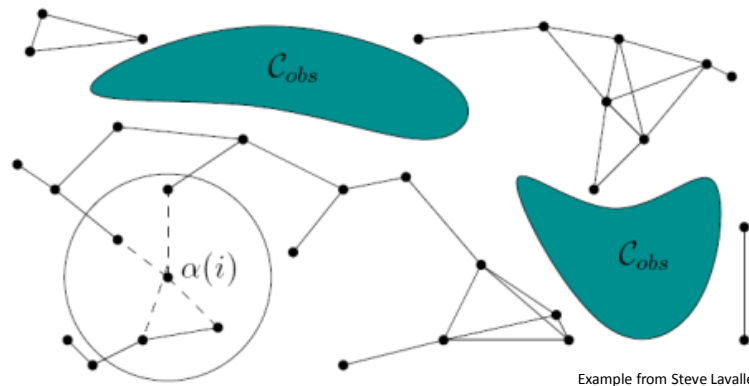
- How to connect the samples?
  - How to select the samples: “Good” sampling strategies are important
  - What are good strategy to maximize “completeness” and to minimize time?
- 
- Suppose that we have built a partial roadmap from  $i-1$  samples.
  - Let  $\alpha(i)$  be the  $i$ -th sample
  - **What strategy would you use to  $\alpha(i)$  connect to the existing roadmap?**
  - **What strategy would you use to select  $\alpha(i)$  ?**



Example from Steve Lavalle

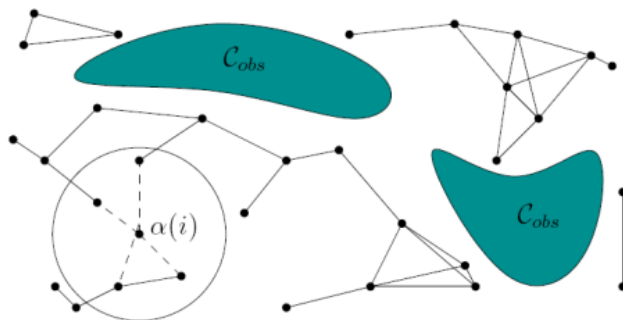
## Connecting samples

- Select  $K$  vertices closest to  $\alpha(i)$
- Select  $K$  (often just 1) closest points from each of the components in  $G$
- Select all vertices within radius  $r$  from  $\alpha(i)$



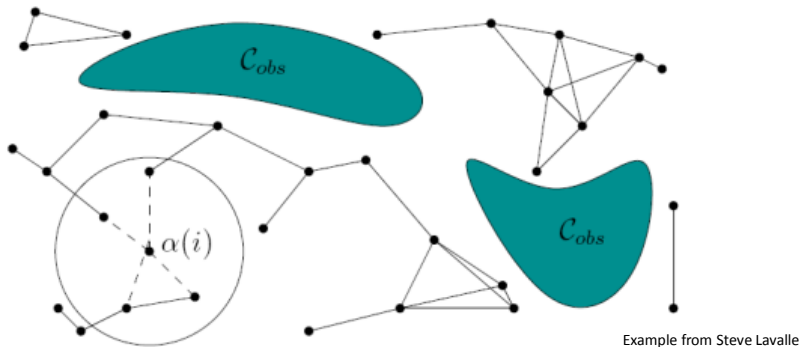
## Selecting samples I

- Sample uniformly from  $C_{free}$
- Select at random an existing vertex with a probability distribution inversely proportional to how well-connected a vertex is, and then generate a random motion from it to get a sample  $\alpha(i)$
- Bias sampling toward obstacle boundaries



## Selecting samples II

- Sample  $q_1$  and  $q_2$  from Gaussian around  $q_1$  and if either is in  $C_{obs}$ , then the other one is set as  $\alpha(i)$
- Sample  $q_1, q_2, q_3$  and set  $q_2$  as  $\alpha(i)$  if  $q_2$  is in  $C_{free}$  and  $q_1$  and  $q_3$  are in  $C_{obs}$
- Bias sampling away from obstacles



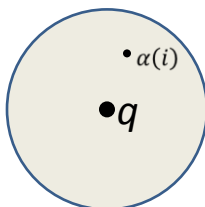
- Can we say something a little more formal about desired properties of  $\alpha(i)$  ?
- Can we say something a little more formal about “approximate completeness”?
- Can we avoid the (expensive) pre-processing step?



## Density

- A set  $Y$  is dense in  $X$  iff for any  $x$  in  $X$  and any  $\varepsilon > 0$ , there exist a  $y$  in  $Y$  inside the ball  $B(x, \varepsilon)$   
(e.g.,  $\mathbb{Q}$  dense in  $\mathbb{R}$ )
- A sequence  $\alpha(i)$  is dense in  $C$  if the corresponding set  $A$  is dense in  $C$
- A random sequence is *probably* dense:
  - For any  $q$  in  $C$  and any  $\varepsilon > 0$ , there exist a  $i$  such that  $\alpha(i)$  is inside the ball  $B(q, \varepsilon)$  with probability 1

Note: The distance could be or  $L_2$  or  $L_\infty$  ... the denseness property remains



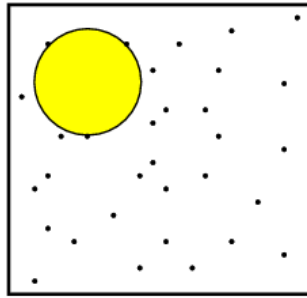
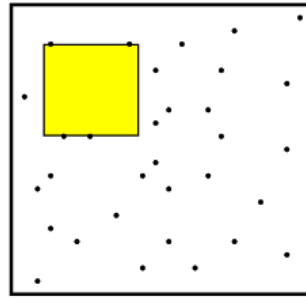
## Deterministic sequences

- All we need is asymptotic denseness
- We don't absolutely need randomness

$i$	Naive Sequence	Binary	Reverse Binary	Van der Corput	Points in $[0, 1]/ \sim$
1	0	.0000	.0000	0	●—————●
2	1/16	.0001	.1000	1/2	○—————●—————○
3	1/8	.0010	.0100	1/4	○——●——○—————○
4	3/16	.0011	.1100	3/4	○——○——○—————●
5	1/4	.0100	.0010	1/8	○——○——○——●——○
6	5/16	.0101	.1010	5/8	○——○——○——○——●
7	3/8	.0110	.0110	3/8	○——○——○——●——○
8	7/16	.0111	.1110	7/8	○——○——○——○——●
9	1/2	.1000	.0001	1/16	●——○——○——○——○
10	9/16	.1001	.1001	9/16	○○——○——○——●——○
11	5/8	.1010	.0101	5/16	○○——○——●——○○——○
12	11/16	.1011	.1101	13/16	○○——○○——○——○——●
13	3/4	.1100	.0011	3/16	○○○●○○○——○○○——○
14	13/16	.1101	.1011	11/16	○○○○○○○——○——○——○
15	7/8	.1110	.0111	7/16	○○○○○○○●○○○○○○○——○
16	15/16	.1111	.1111	15/16	○○○○○○○○○○○○○○○○○○●

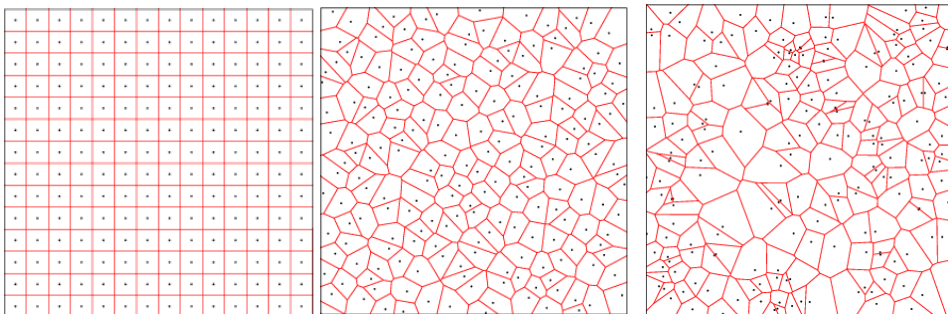
## Dispersion

- We may want the samples to become uniformly close to each other
- Dispersion:  $\delta(A) = \max_{q \in C} (\min_{\alpha \in A} d(q, \alpha))$

 $L_2$  $L_\infty$ 

## Dispersion

For  $L_\infty$  :  $\delta(A) \geq \frac{1}{2N^{1/p}}$  (dispersion for grids)



Grid

Halton

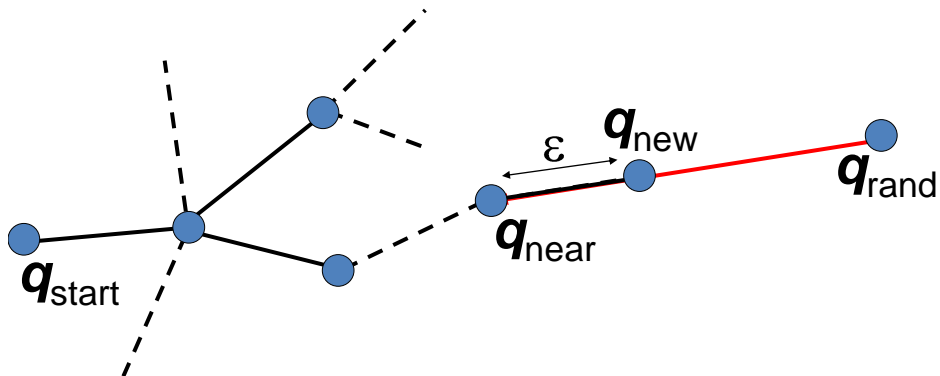
Random

- Can we say something a little more formal about desired properties of  $\alpha(i)$  ?
- *Can we say something a little more formal about “approximate completeness”?*
- Can we avoid the (expensive) pre-processing step?

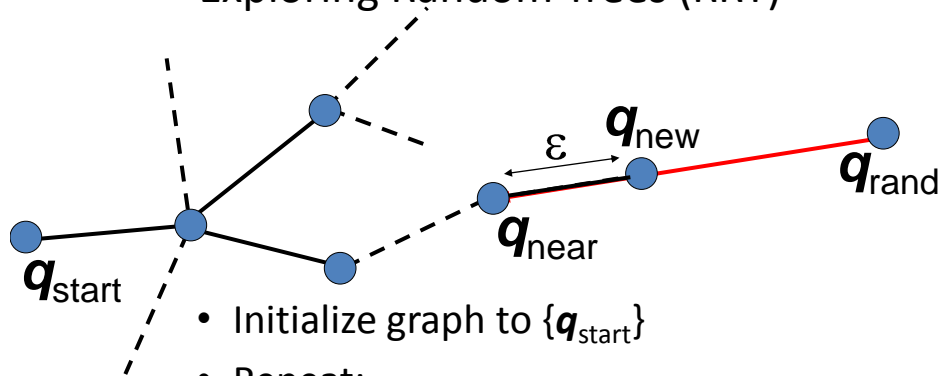
## “Completeness”

- Deterministic sampling:
  - If  $\alpha$  is dense sampling
  - Resolution complete  $\iff$  Guaranteed to find a path in finite time if one exists
- Random sampling:
  - If  $\alpha$  is probabilistically dense
  - Probabilistically complete  $\iff$  Guaranteed to find a path in finite time if one exists with probability 1

- Can we say something a little more formal about desired properties of  $\alpha(i)$  ?
- Can we say something a little more formal about “approximate completeness”?
- *Can we avoid the (expensive) pre-processing step?*

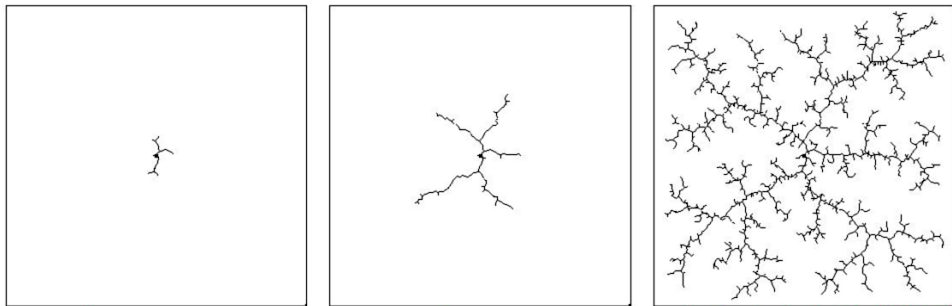


## No pre-processing: Rapidly Exploring Random Trees (RRT)



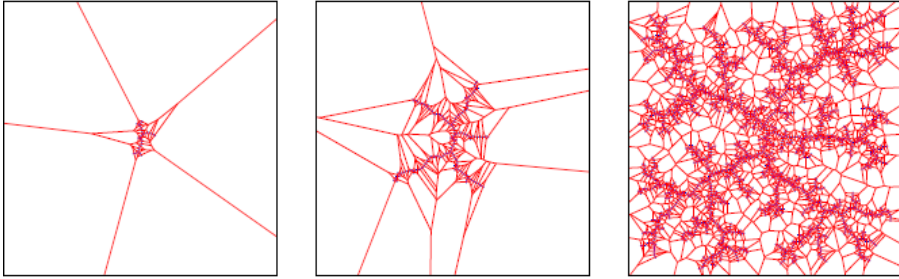
- Initialize graph to  $\{q_{start}\}$
- Repeat:
  - Select random new sample  $q_{rand}$
  - Find closest node  $q_{near}$  to  $q_{rand}$
  - Create edge  $(q_{near}, q_{new})$  if no collisions

## Properties



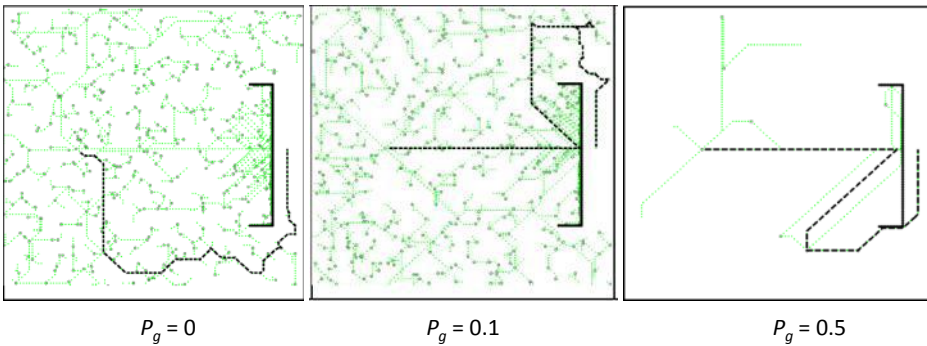
- Tends to explore the space rapidly in all directions
- Does not require extensive pre-processing
- Single query/multiple query problems
- Needs only collision detection test → No need to represent/pre-compute the entire C-space





uniform coverage of space:  
the growth is always biased by the largest Voronoi region

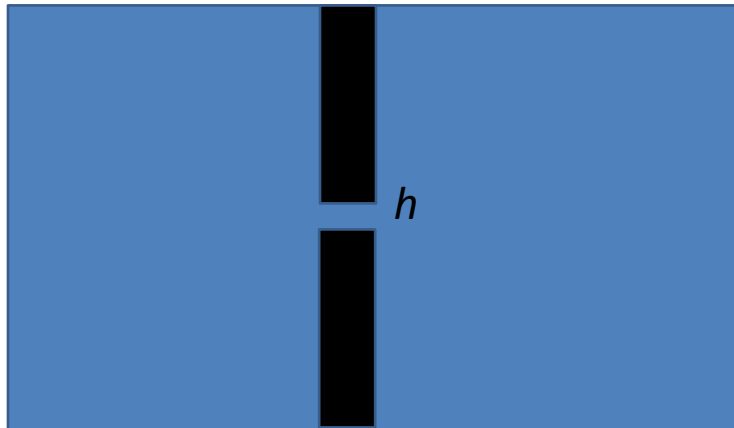
## How to connect the goal?



- With a probability  $(1-P_g)$ ,  $q_{rand}$  is chosen as a random sample in  $C_{free}$ , with probability  $P_g$ ,  $q_{rand}$  is set to  $q_G$
- Variations (ERRT): If known set of “preferred waypoint” locations  $W$ ,

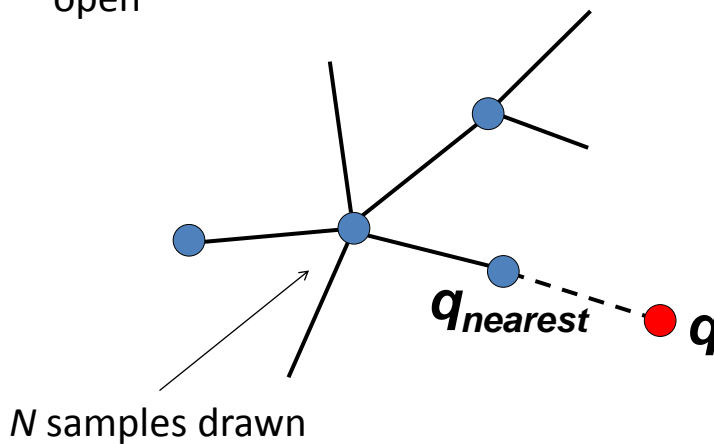
$$\begin{aligned} (1-P_g-P_w) &\rightarrow q_{rand} \\ P_g &\rightarrow q_G \\ P_w &\rightarrow \text{random } q \text{ from } W \end{aligned}$$

- In general: Probability of finding path after  $k$  samples depends on the smallest gap in  $C_{obs}$
- $P(\text{finding a path})$  after  $k$  samples assuming
  - Uniform sampling
  - Use nearest sample every iteration

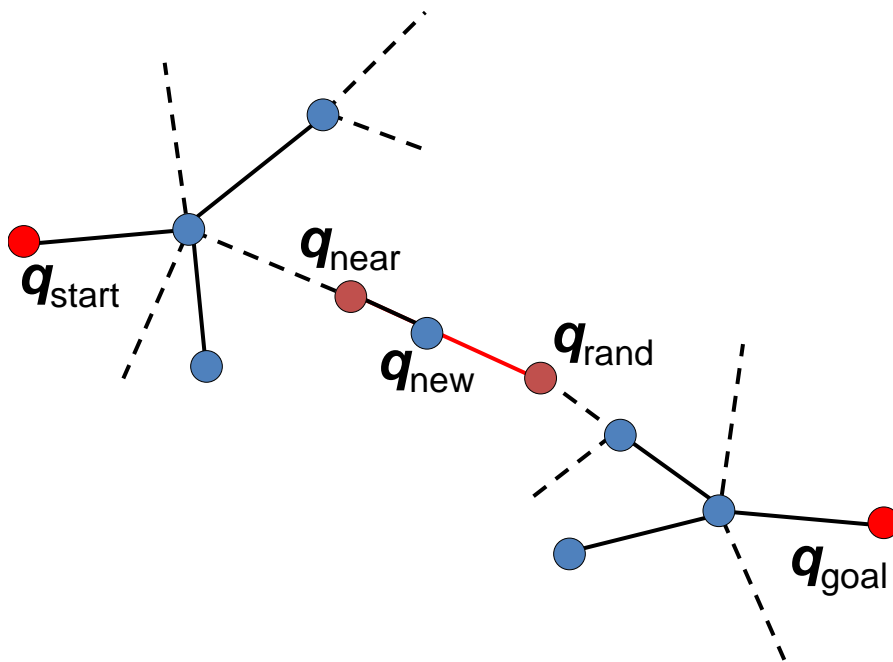
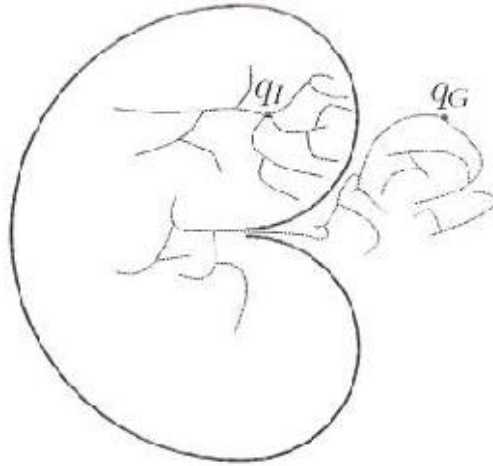


## Convergence?

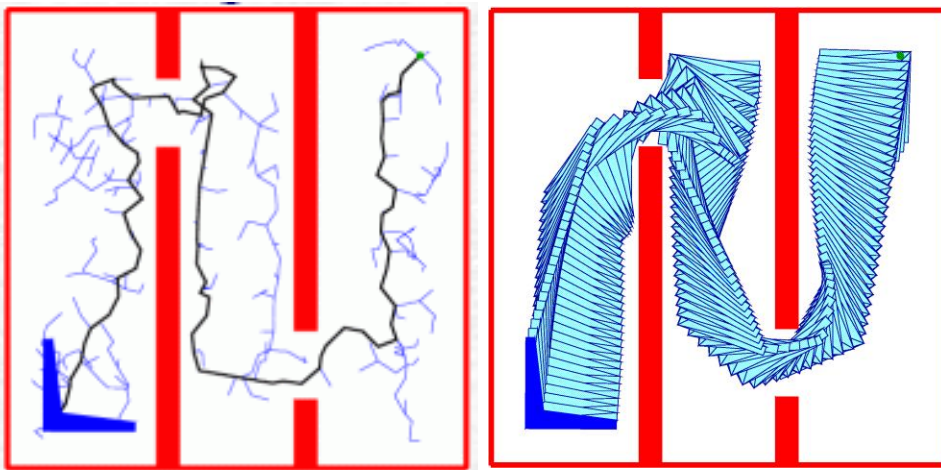
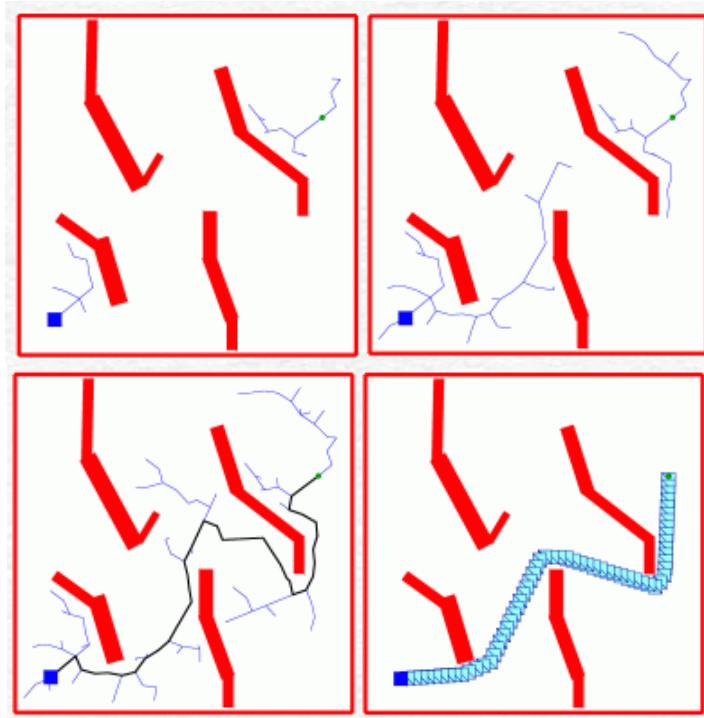
- For any  $q \in C_{free}$ ,  $\lim_{N \rightarrow \infty} P[d(q, q_{nearest}) < \epsilon] = 1$
- Assumptions:  $C_{free}$  is connected, bounded and open



- Should we use 2 trees?

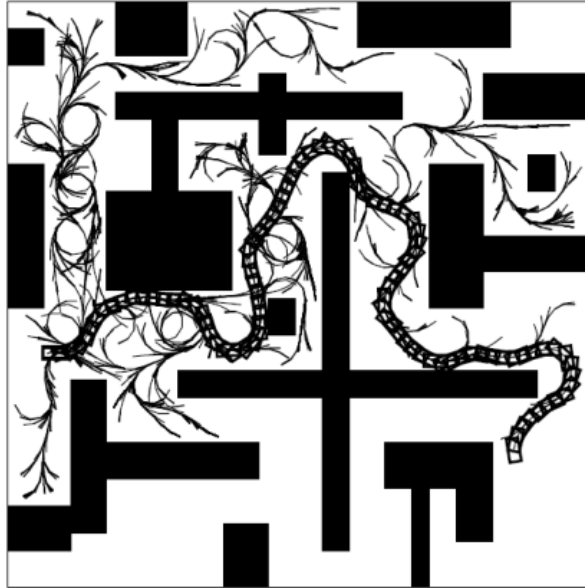




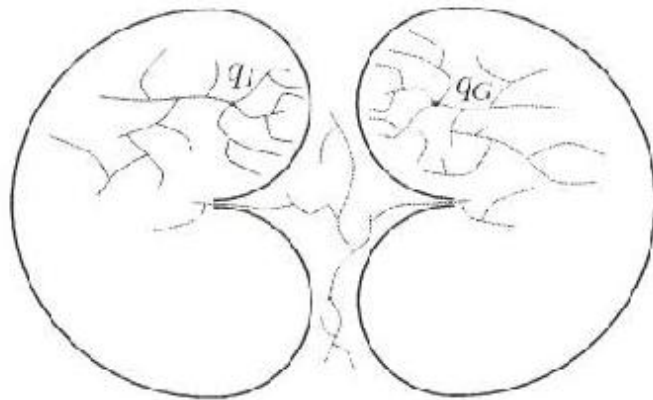


From Kuffner et al.



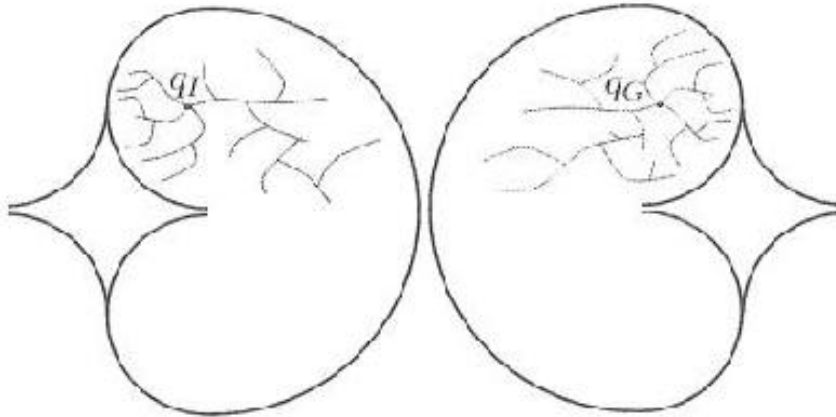


- Should we use more trees?



Example from Steve Lavalle

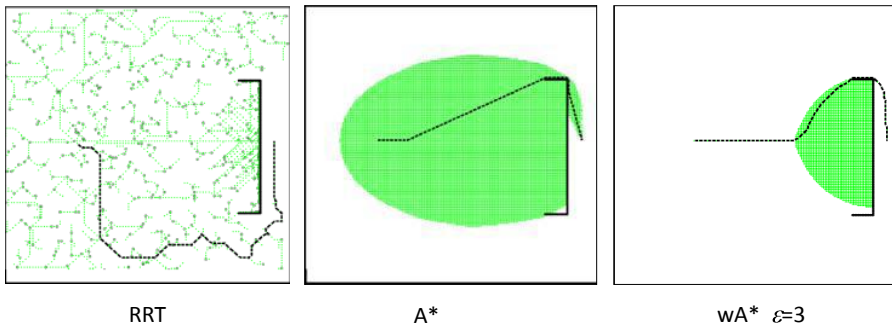
- Good luck!



Example from Steve Lavallo

## Comparison

- Even for easier problems that can be solved by A\* directly, needs much fewer node expansions
- Smaller memory
- Works well in (very) high dimensions
- (Very) sub-optimal solutions. Needs post-processing.
- More difficult to incorporate complex cost functions



Note:  $wA^* = A^*$  with heuristic  $wh(\cdot)$  where  $h(\cdot)$  is an admissible heuristic. Sub-optimal, guaranteed to find path at most  $\epsilon C^*$

Example from Max Likhachev