

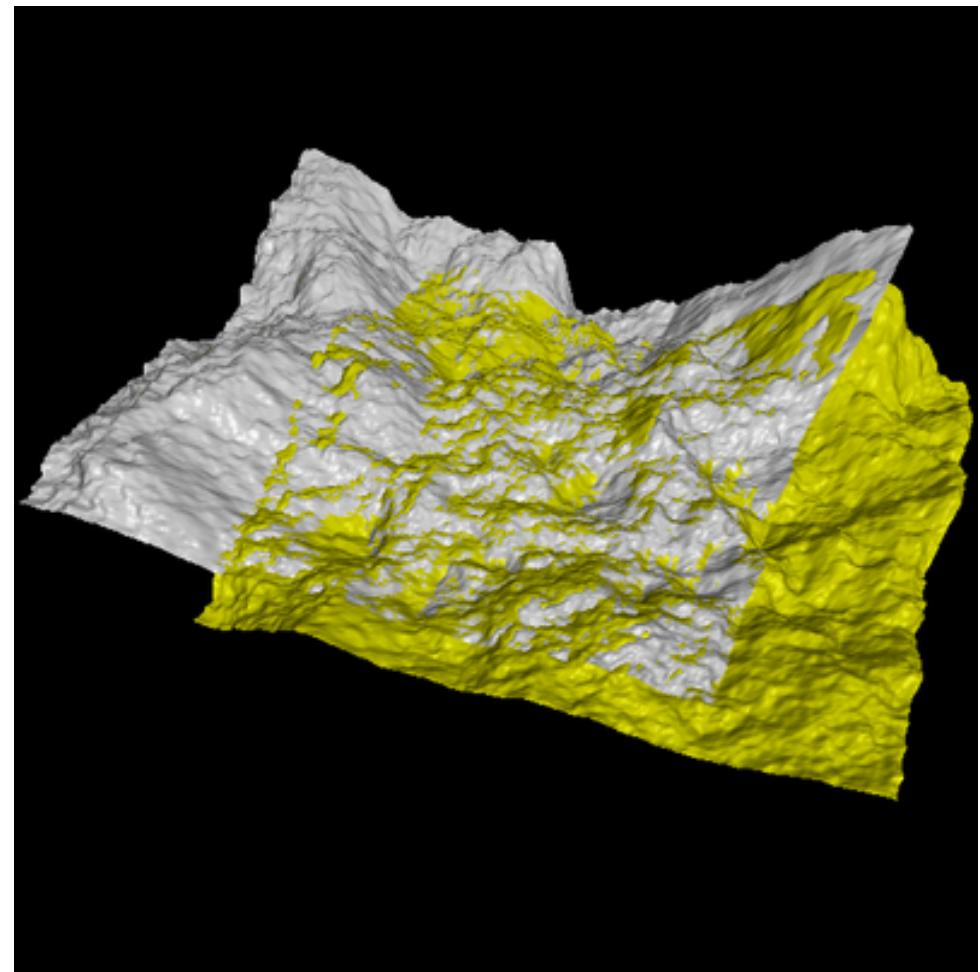
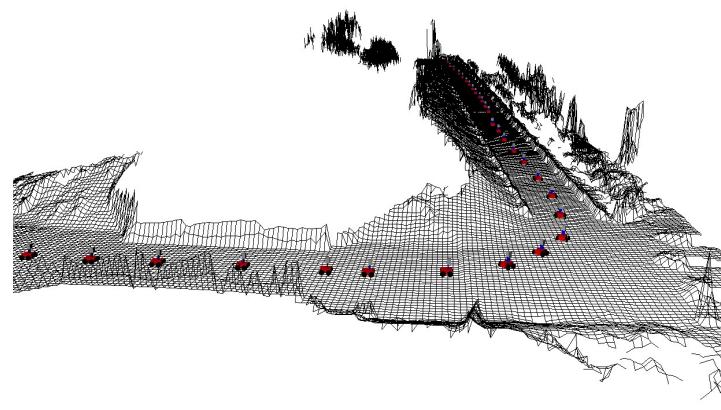
# Introduction to Mobile Robotics

## Iterative Closest Point Algorithm

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Slides adopted from: Wolfram Burgard, Cyrill Stachniss,  
Maren Bennewitz, Kai Arras and Probabilistic Robotics Book

# Motivation



# The Problem

- Given: two corresponding point sets:

$$X = \{x_1, \dots, x_n\}$$

$$P = \{p_1, \dots, p_n\}$$

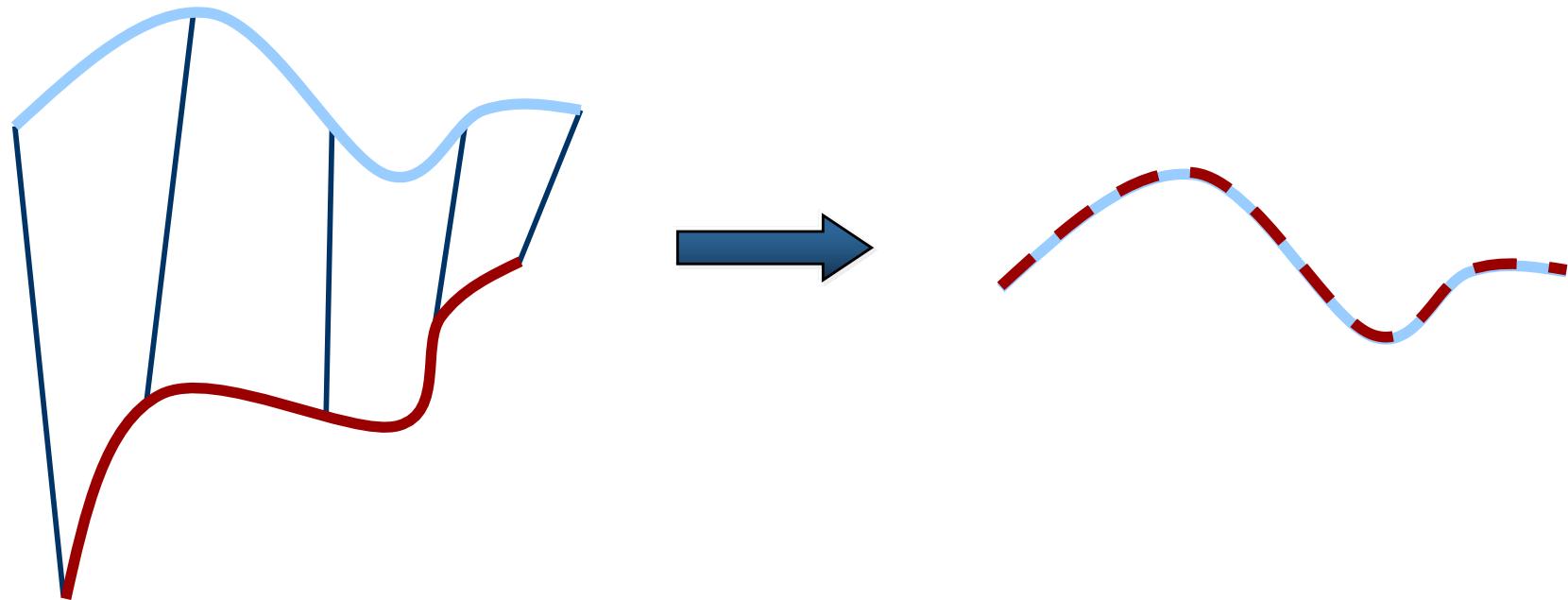
- Wanted: translation  $t$  and rotation  $R$  that minimizes the sum of the squared error:

$$E(R, t) = \frac{1}{N_p} \sum_{i=1}^{N_p} \|x_i - Rp_i - t\|^2$$

Where  $x_i$  and  $p_i$  are corresponding points.

# Key Idea

- If the correct correspondences are known, the correct relative rotation/translation can be calculated in closed form.



# Center of Mass

$$\mu_x = \frac{1}{N_x} \sum_{i=1}^{N_x} x_i \quad \text{and} \quad \mu_p = \frac{1}{N_p} \sum_{i=1}^{N_p} p_i$$

are the centers of mass of the two point sets.

## Idea:

- Subtract the corresponding center of mass from every point in the two point sets before calculating the transformation.
- The resulting point sets are:

$$X' = \{x_i - \mu_x\} = \{x'_i\}$$

and

$$P' = \{p_i - \mu_p\} = \{p'_i\}$$

# SVD

Let  $W = \sum_{i=1}^{N_p} x'_i p_i'^T$

denote the singular value decomposition (SVD) of W by:

$$W = U \begin{bmatrix} \sigma_1 & 0 & 0 \\ 0 & \sigma_2 & 0 \\ 0 & 0 & \sigma_3 \end{bmatrix} V^T$$

where  $U, V \in \mathbb{R}^{3 \times 3}$  are unitary, and  
 $\sigma_1 \geq \sigma_2 \geq \sigma_3$  are the singular values of W.

# SVD

**Theorem** (without proof):

If  $\text{rank}(W) = 3$ , the optimal solution of  $E(R,t)$  is unique and is given by:

$$R = UV^T$$
$$t = \mu_x - R\mu_p$$

Tr  $E(R, t) = \frac{1}{N_p} \sum_{i=1}^{N_p} \|x_i - Rp_i - t\|^2$  ( $R, t$ ) is:

$$E(R, t) = \sum_{i=1} (||x'_i||^2 + ||y'_i||^2) - 2(\sigma_1 + \sigma_2 + \sigma_3)$$

# Proof

$$P = \{p_1, \dots, p_n\} \quad Q = \{q_1, \dots, q_n\}$$

(1)

two sets of 3D points

$$E(R, t) = \arg \min_{R, t} \sum_{i=1}^n w_i \|Rp_i + t - q_i\|^2 \quad \text{for } w_i > 0$$

here we assume  $w_i = 1$

1. Assume  $R$  is fixed and solve for translation

$$\frac{\partial E}{\partial t} = 0 \quad \sum_{i=1}^n \alpha (Rp_i + t - q_i) = 2t \sum_{i=1}^n 1 + 2R \sum_{i=1}^n p_i$$

$$\text{denote } M_p = \frac{\sum p_i}{n} \quad \mu_q = \frac{\sum q_i}{n} \quad \text{mean of each point cloud}$$

$$\text{then } t = \mu_q - R M_p$$

2. Substitute  $t$  to original  $E(R, t)$

$$\begin{aligned} \sum_{i=1}^n \|Rp_i + t - q_i\|^2 &= \sum_{i=1}^n \|Rp_i + \mu_q - R M_p - q_i\|^2 \\ &= \sum_{i=1}^n \|R(p_i - M_p) - (q_i - \mu_q)\|^2 \quad \text{denote } p'_i = p_i - M_p \\ &= \sum_{i=1}^n \|R p'_i - q'_i\|^2 \quad q'_i = q_i - \mu_q \end{aligned}$$

$$\|R p'_i - q'_i\|^2 = p'^T p'_i - 2 q'^T R p'_i + q'^T q'_i$$

$$\text{we have to solve } \arg \min_R (-2 \sum_{i=1}^n q'^T R p'_i)$$

This can be written as

$$\left[ \begin{array}{c} q_1^T \\ q_2^T \\ \vdots \\ q_n^T \end{array} \right] R \left[ \begin{array}{c} p_1 \\ p_2 \\ \vdots \\ p_n \end{array} \right] = \text{tr}(Y^T R X)$$

Trace properties  $Y^T X$

$$\text{tr}(AB) = \text{tr}(BA)$$

$$\text{tr}(Y^T R X) = \text{tr}(R X Y^T)$$

consider covariance matrix  $S = X Y^T$  take SVD of  $S$

$$S = U \Sigma V^T$$

$$\text{then trace } \text{tr}(R X Y^T) = \text{tr}(R U \Sigma V^T) = \text{tr}(\Sigma V^T R U)$$

$\Rightarrow$

trace is minimized if

$$V^T R U = I \Rightarrow R = V U^T$$

orthogonal matrices  
columns  
 $M_i^T M_i = 1$

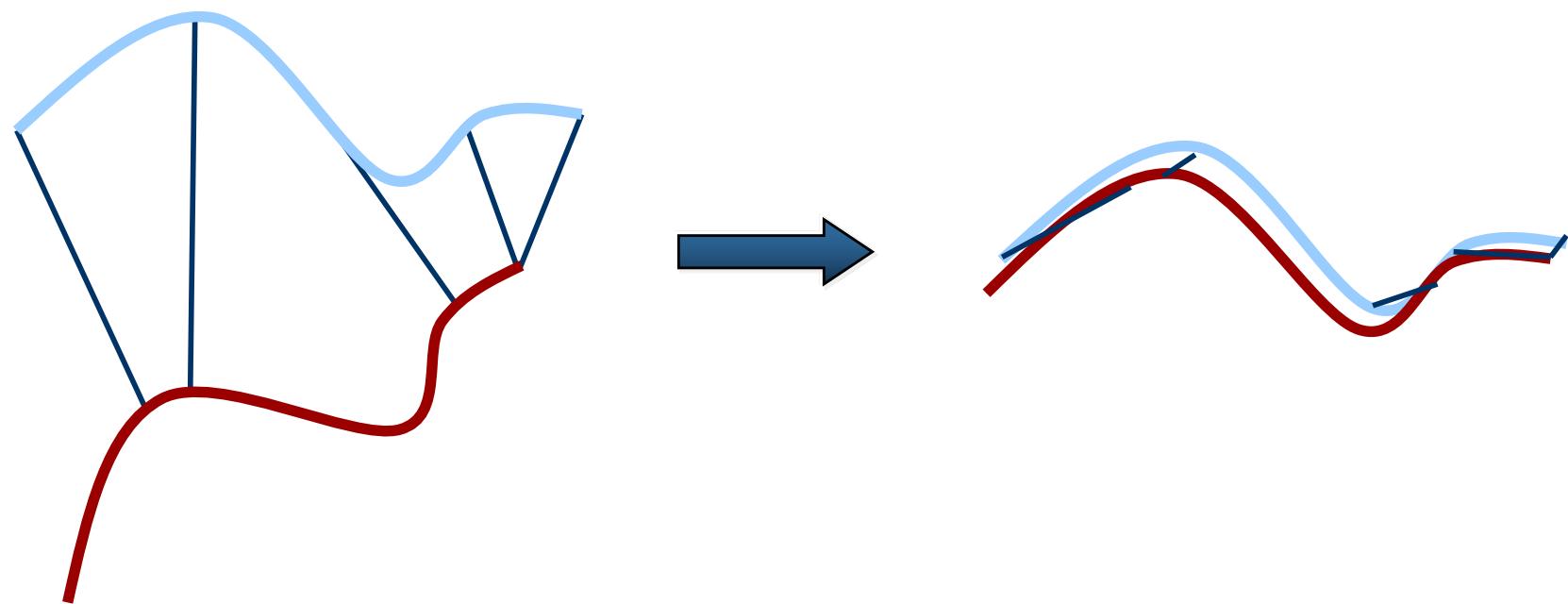
if  $\det(VU^T) = -1$  then it contains reflection it is not an orthogonal matrix

$\Rightarrow$  we need to invert one row of a rotation matrix

$$\Rightarrow R = V \begin{bmatrix} 1 & & \\ & \ddots & \\ & & \det(VU^T) \end{bmatrix} U^T$$

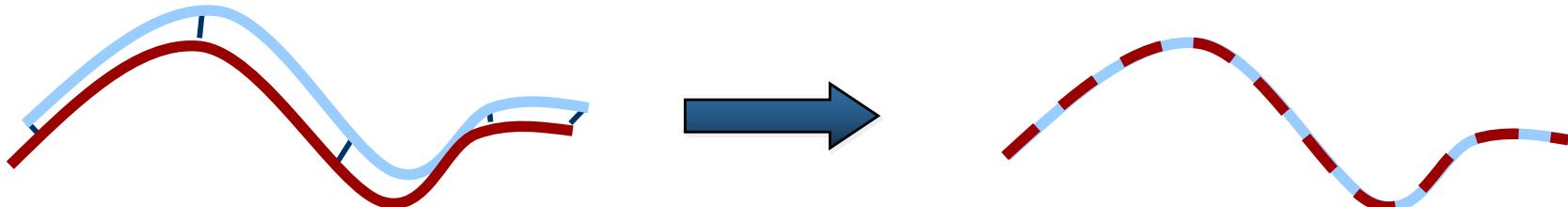
# ICP with Unknown Data Association

- If correct correspondences are not known, it is generally impossible to determine the optimal relative rotation/translation in one step

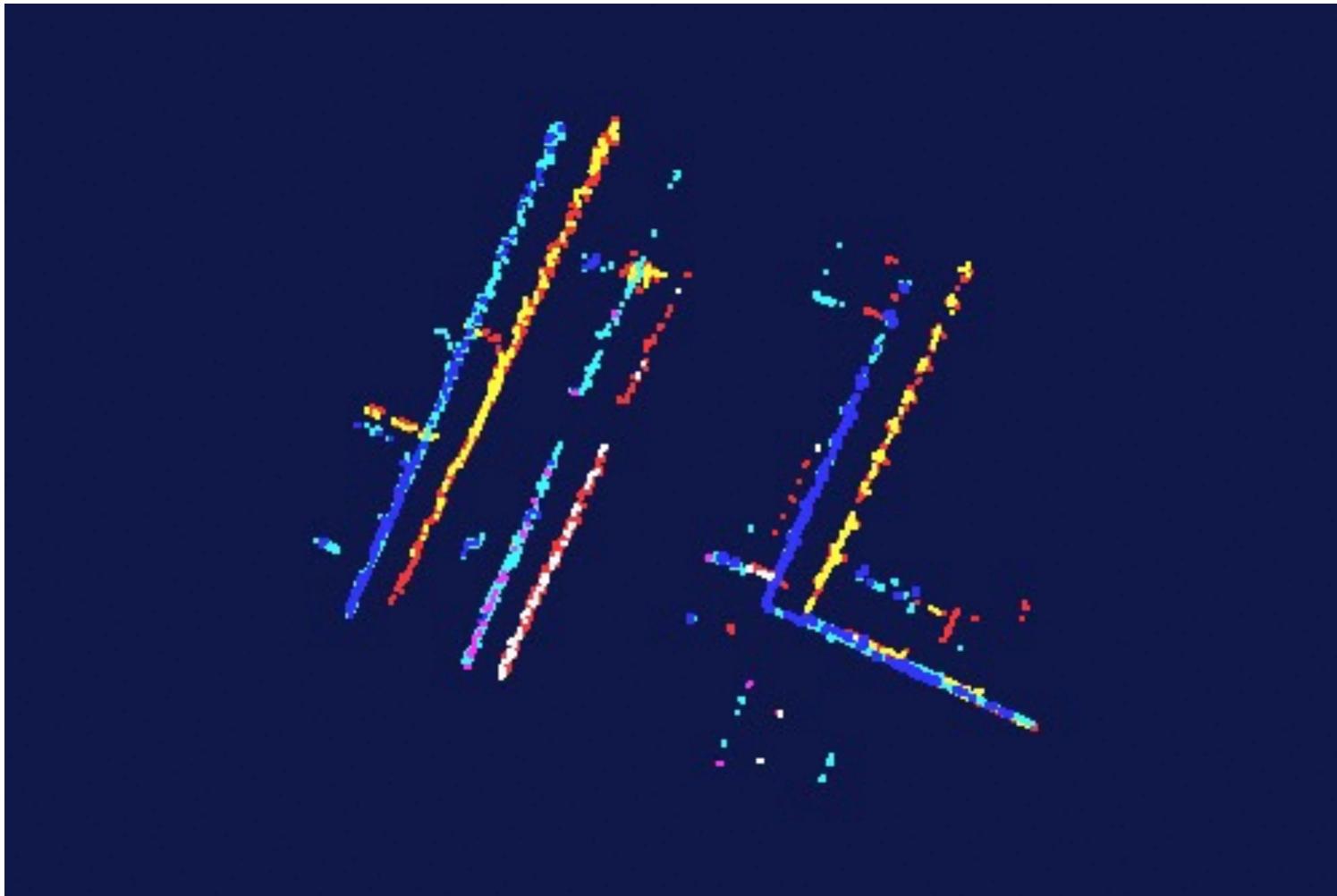


# ICP-Algorithm

- Idea: iterate to find alignment
- Iterated Closest Points (ICP)  
[Besl & McKay 92]
- Converges if starting positions are “close enough”



# Iteration-Example



# ICP-Variants

- Variants on the following stages of ICP have been proposed:
  1. Point subsets (from one or both point sets)
  2. Weighting the correspondences
  3. Data association
  4. Rejecting certain (outlier) point pairs

# Performance of Variants

- Various aspects of performance:
  - Speed
  - Stability (local minima)
  - Tolerance wrt. noise and/or outliers
  - Basin of convergence  
(maximum initial misalignment)
- Here: properties of these variants

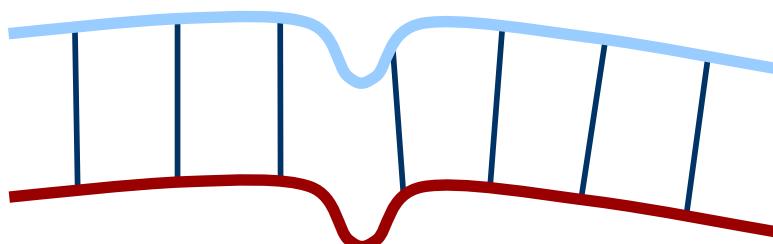
# ICP Variants

- 
1. Point subsets (from one or both point sets)
  2. Weighting the correspondences
  3. Data association
  4. Rejecting certain (outlier) point pairs

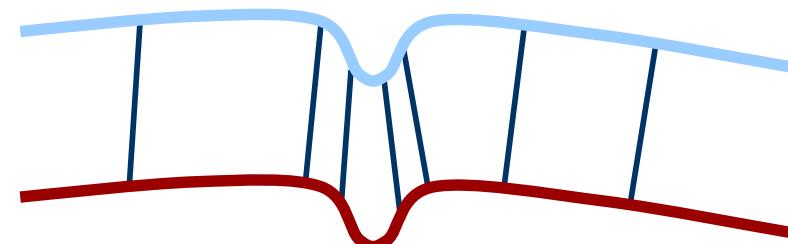
# Selecting Source Points

- Use all points
- Uniform sub-sampling
- Random sampling
- Feature based Sampling
- Normal-space sampling
  - Ensure that samples have normals distributed as uniformly as possible

# Normal-Space Sampling



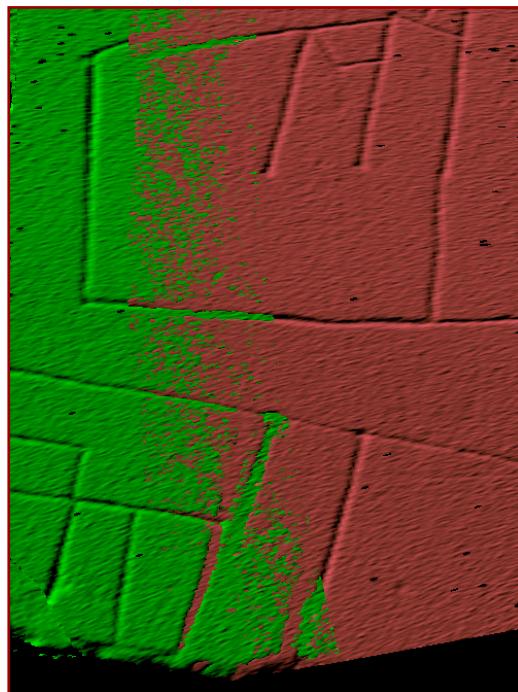
uniform sampling



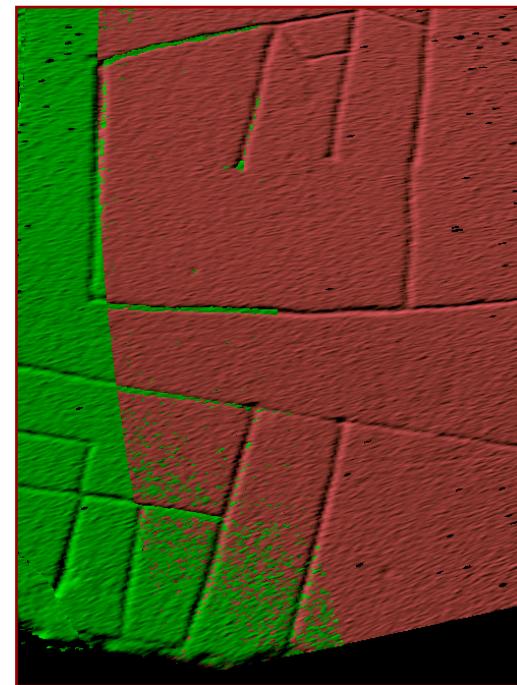
normal-space sampling

# Comparison

- Normal-space sampling better for mostly-smooth areas with sparse features  
[Rusinkiewicz et al.]



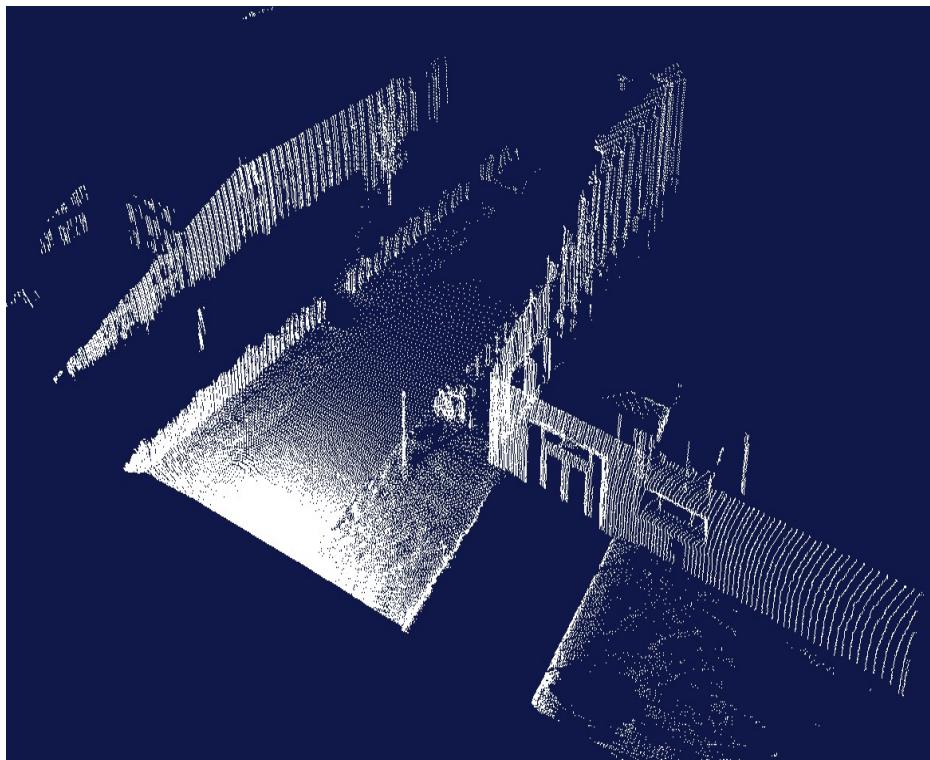
Random sampling



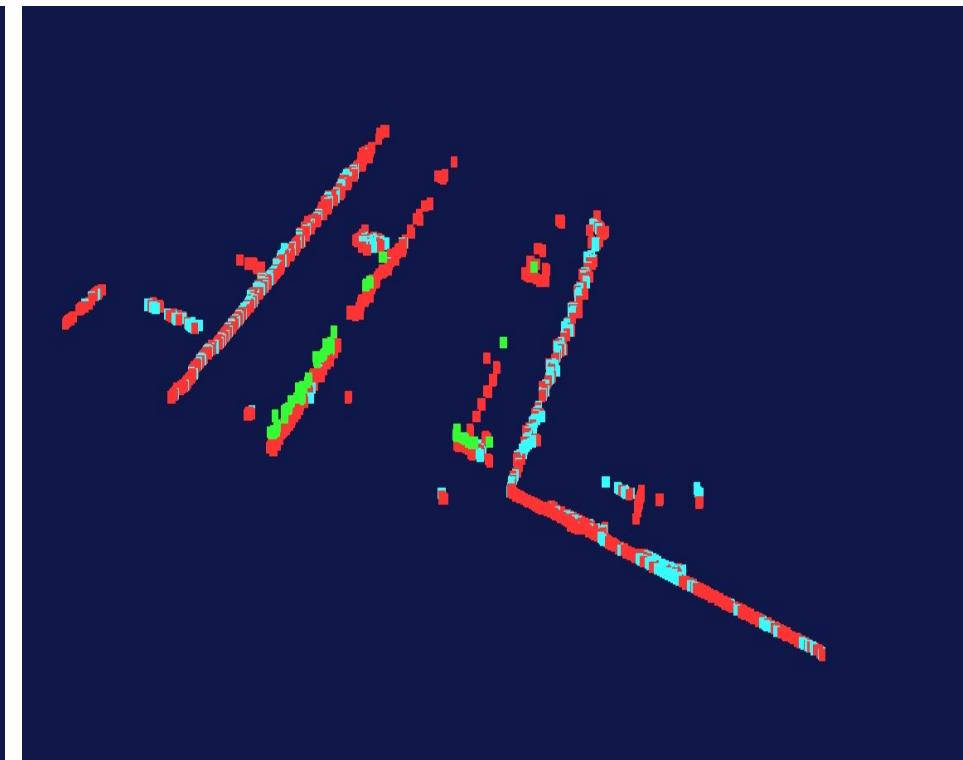
Normal-space sampling

# Feature-Based Sampling

- try to find “important” points
- decrease the number of correspondences
- higher efficiency and higher accuracy
- requires preprocessing

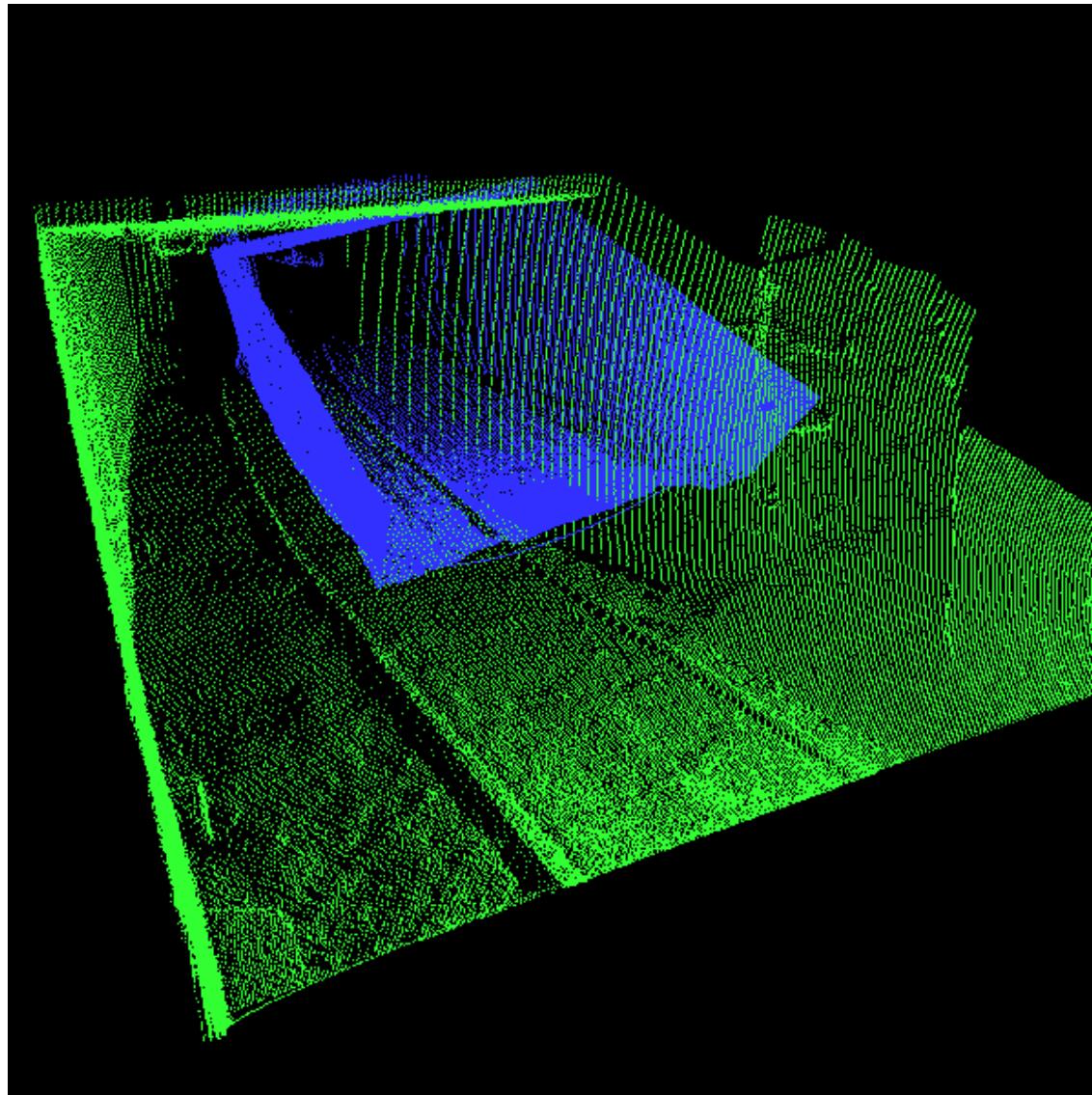


3D Scan (~200.000 Points)



Extracted Features (~5.000 Points)

# Application



[Nuechter et al., 04]

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# ICP Variants

1. Point subsets (from one or both point sets)
2. Weighting the correspondences
3. Data association
4. Rejecting certain (outlier) point pairs



# Selection vs. Weighting

- Could achieve same effect with weighting
- Hard to guarantee that enough samples of important features except at high sampling rates
- Weighting strategies turned out to be dependent on the data.
- Preprocessing / run-time cost tradeoff (how to find the correct weights?)

# ICP Variants

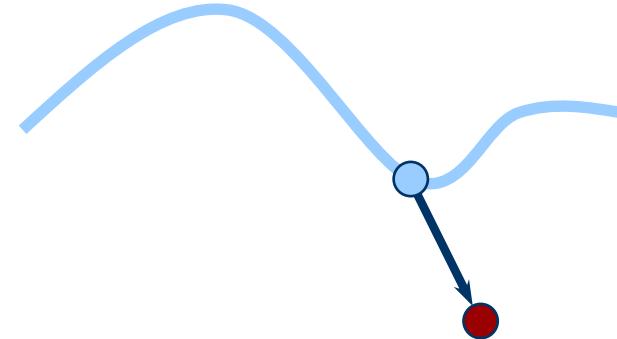
1. Point subsets (from one or both point sets)
2. Weighting the correspondences
-  3. **Data association**
4. Rejecting certain (outlier) point pairs

# Data Association

- has greatest effect on convergence and speed
- Closest point
- Normal shooting
- Closest compatible point
- Projection
- Using kd-trees or oc-trees

# Closest-Point Matching

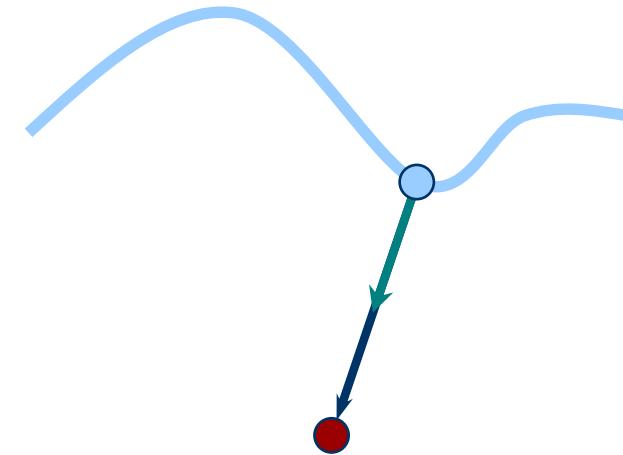
- Find closest point in other the point set



Closest-point matching generally stable,  
but slow and requires preprocessing

# Normal Shooting

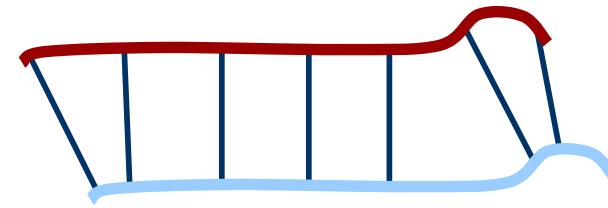
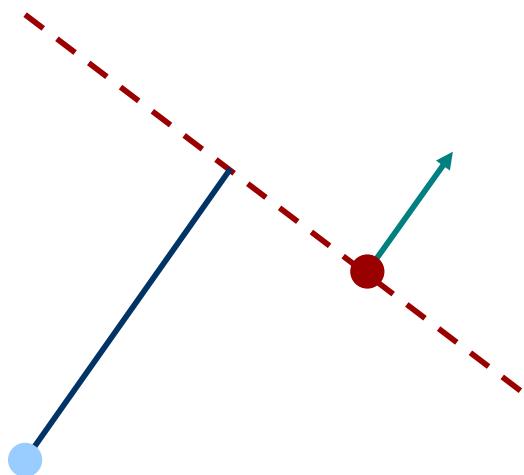
- Project along normal, intersect other point set



Slightly better than closest point for smooth structures, worse for noisy or complex structures

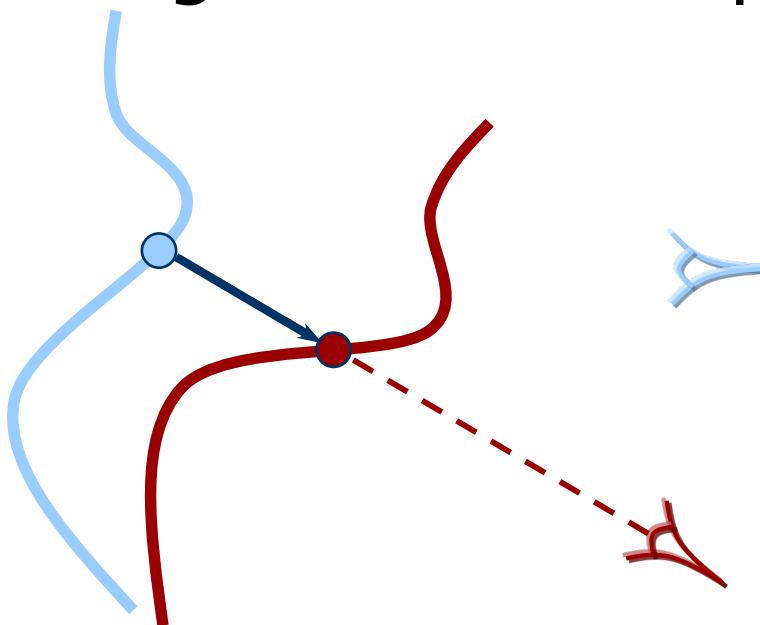
# Point-to-Plane Error Metric

- Using point-to-plane distance instead of point-to-point lets flat regions slide along each other [Chen & Medioni 91]



# Projection

- Finding the closest point is the most expensive stage of the ICP algorithm
- Idea: simplified nearest neighbor search
- For range images, one can project the points according to the view-point [Blais 95]



# Projection-Based Matching

- Slightly worse alignments per iteration
- Each iteration is one to two orders of magnitude faster than closest-point
- Requires point-to-plane error metric

# Closest Compatible Point

- Improves the previous two variants by considering the **compatibility** of the points
- Compatibility can be based on normals, colors, etc.
- In the limit, degenerates to feature matching

# ICP Variants

1. Point subsets (from one or both point sets)
2. Weighting the correspondences
3. Nearest neighbor search
4. Rejecting certain (outlier) point pairs

# Rejecting (outlier) point pairs

- sorting all correspondences with respect to their error and deleting the worst  $t\%$ , Trimmed ICP (TrICP) [Chetverikov et al. 2002]
- $t$  is to Estimate with respect to the Overlap



**Problem:** Knowledge about the overlap is necessary or has to be estimated

# ICP-Summary

- ICP is a powerful algorithm for calculating the displacement between scans.
- The major problem is to determine the correct data associations.
- Given the correct data associations, the transformation can be computed efficiently using SVD.