

Reading

• Szeliski: Chapter 6.1.4

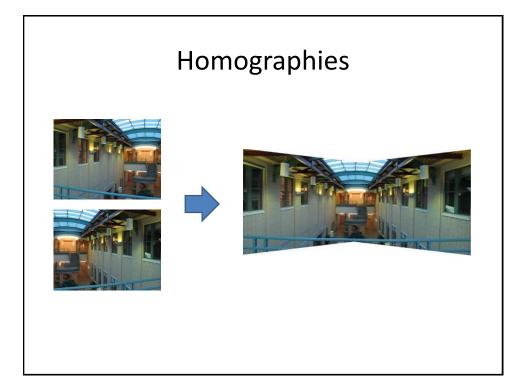
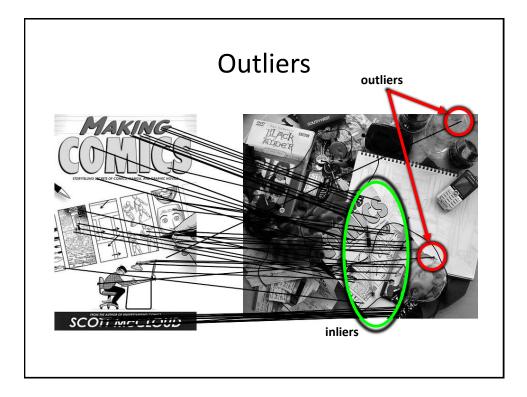


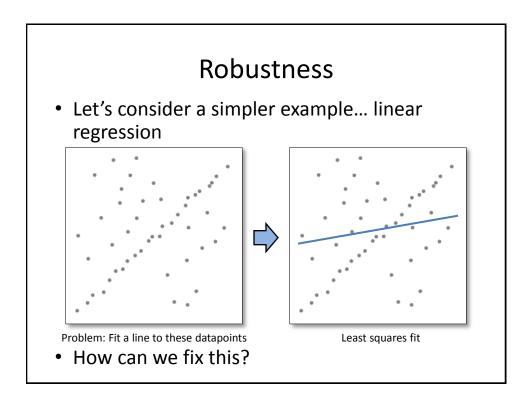
Image Alignment Algorithm

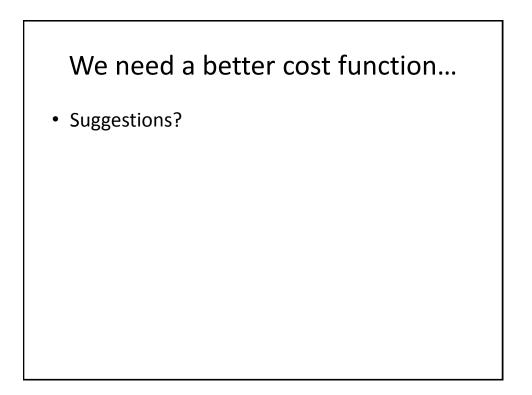
Given images A and B

- 1. Compute image features for A and B
- 2. Match features between A and B
- 3. Compute homography between A and B using least squares on set of matches

What could go wrong?

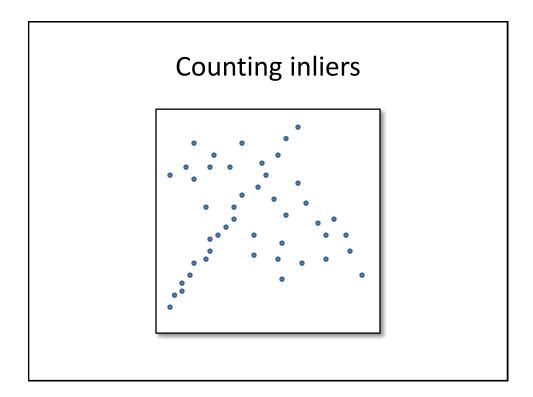


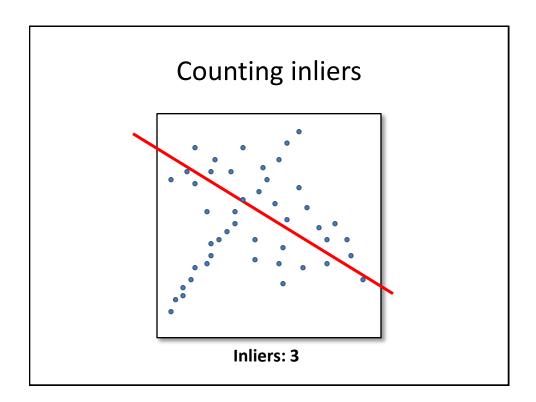


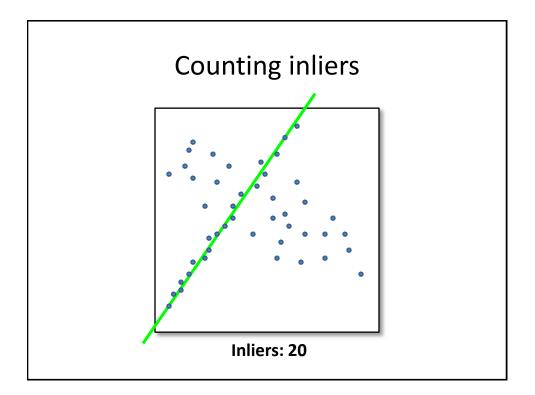


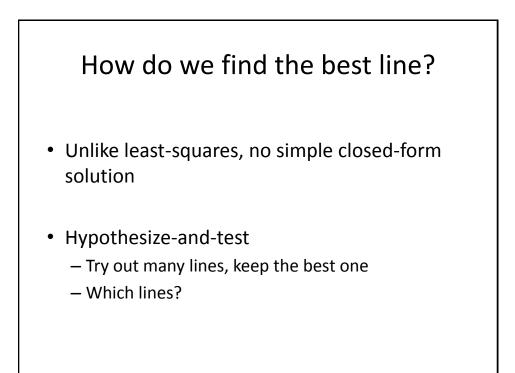


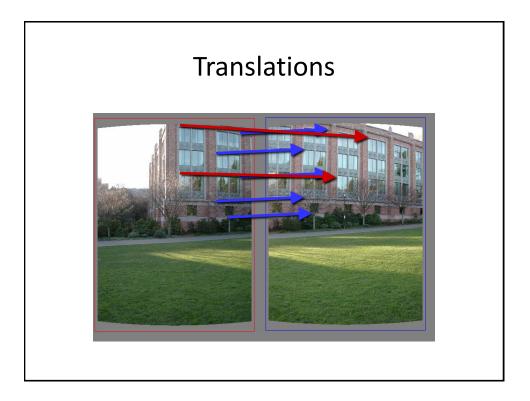
- Given a hypothesized line
- Count the number of points that "agree" with the line
 - "Agree" = within a small distance of the line
 - I.e., the inliers to that line
- For all possible lines, select the one with the largest number of inliers

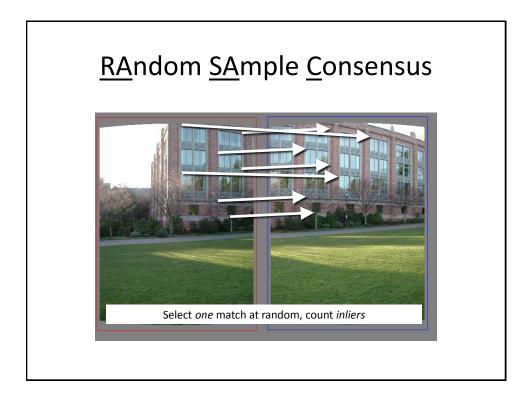


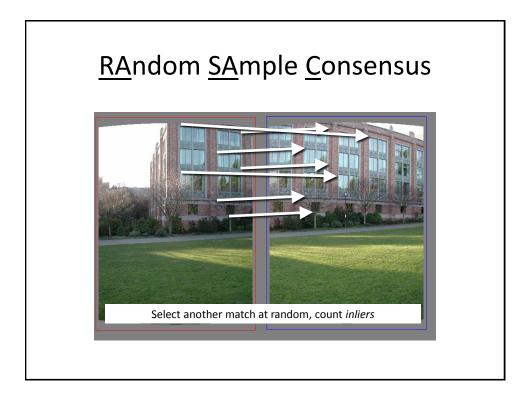


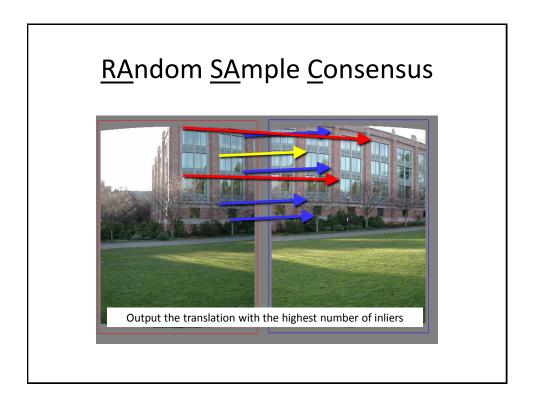


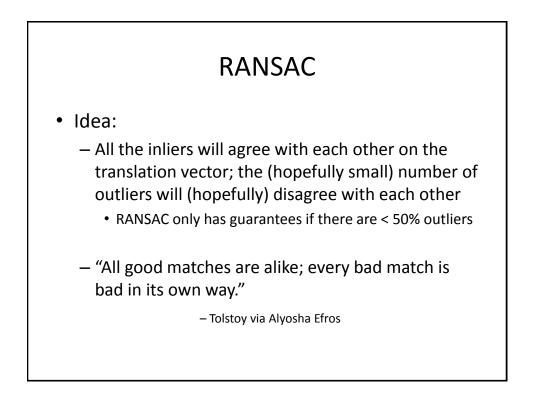


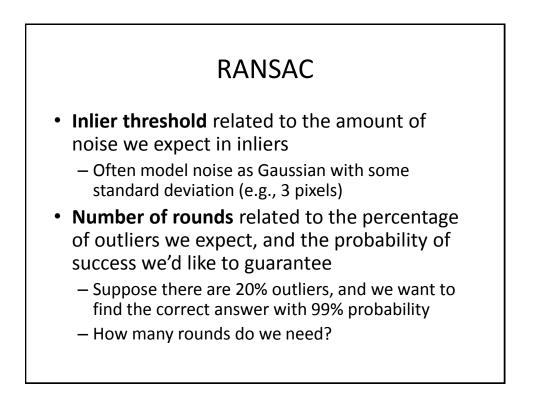


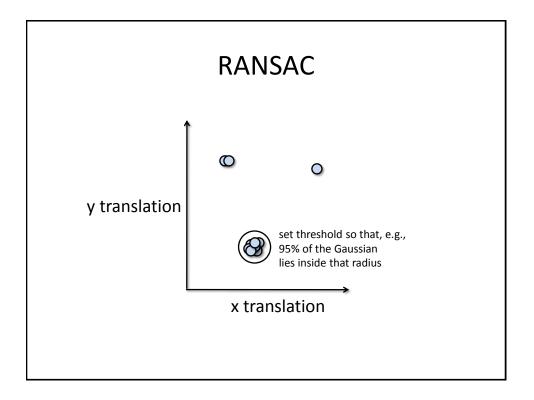


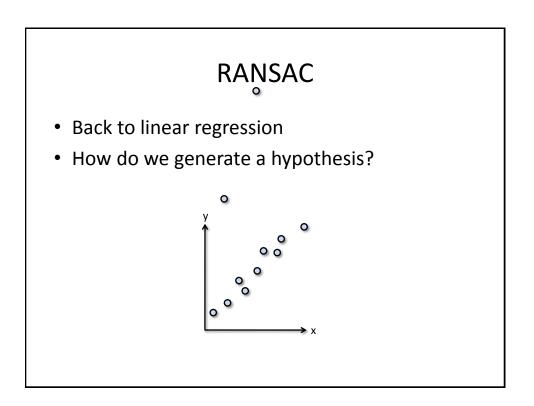


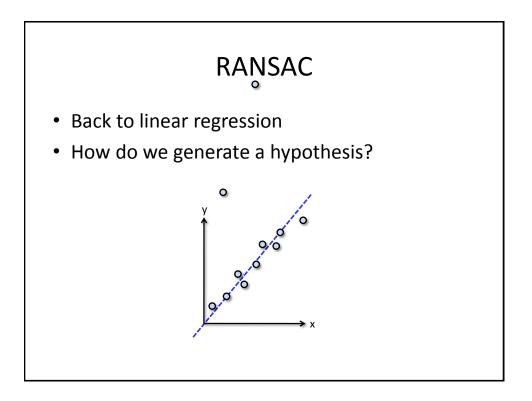












RANSAC

- General version:
 - 1. Randomly choose *s* samples
 - Typically *s* = minimum sample size that lets you fit a model
 - 2. Fit a model (e.g., line) to those samples
 - 3. Count the number of inliers that approximately fit the model
 - 4. Repeat N times
 - 5. Choose the model that has the largest set of inliers

