GEOMETRY FOR COMPUTER VISION

> LECTURE 6B: SAMPLE CONSENSUS STRATEGIES

# LECTURE 6B: SAMPLE CONSENSUS STRATEGIES

LO-RANSAC

Preemptive RANSAC

**\* DEGENSAC** 

Today's paper: PROSAC

Not covered here: All the other variants MLESAC, NAPSAC etc.

# **RANSAC** ISSUES

In lecture 3 we introduced RANSAC (Fischler&Bolles 81).

It finds a model with maximal support in the presence of outliers

\*\* Approach: randomly generate hypotheses and score them.

Most novelties since 1981 covered in thesis by: Ondrej Chum, Two-View Geometry Estimation by Random Sample and Consensus, July 2005

# **RANSAC** ISSUES

Two problems with the original approach:



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# **RANSAC** ISSUES

\* Near degeneracies can be dealt with by sampling non-randomly, e.g.

DEGENSAC, for F estimation in plane dominant scenes. Chum et al., Two-view Geometry estimation unaffected by a Dominant Plane, CVPR05

Distance constraint for points used in E estimation. Hedborg et al., Fast and Accurate Structure and Motion Estimation, ISVC09 Reduces #iterations by 50% in forward motion.

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Inlier noise means that the heuristic for number of samples to draw:

 $N = \log(1-p)/\log(1-w^s)$ 

is overly optimistic.

\* A small modification makes the heuristic work again: Chum et al., Locally Optimized RANSAC, DAGM03

Small modification

#### RANSAC

loop:

- 1. Select random sample
- 2. Estimate model
- 3. Score model
- 4. If new high-score store model and score

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#### RANSAC

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LO-RANSAC

loop:

- 1. Select random sample
- 2. Estimate model
- 3. Score model
- 4. If new high-score
   run local optimisation
   then store model and score

Chum tries four variants of local optimisation:

 Linear estimation from all inliers
 Iterative linear estimation with decreasing inlier threshold.
 Inner RANSAC
 Inner RANSAC with #2.

#2 and #4 worked best, and came close to the heuristically expected #samples.

# The inner RANSAC step uses non-minimal sample sets. Errors for linear F estimation:



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Solution Structure and motion estimation, ICCV03

\*\* Total time for RANSAC is given by:  $t = k(t_M + E[m_S]t_V)$ 

% k- #iterations t<sub>M</sub>-model estimation time, t<sub>V</sub>-verification time. ms - #models/iteration

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# If many correspondences, ty will dominate.

# Idea: Do a probabilistic verification instead.

 $t = k(t_M + E[m_S]t_V)$ 

In a real-time system, t is fixed, so if we reduce tv we may increase k.

Preemptive RANSAC does this by evaluating all hypotheses in parallel.

#### **\*\* Preemptive RANSAC:**

Generate f(1) hypotheses in parallel.
 For n=1 to N

- 3. Evaluate f(n) hypotheses on a random correspondence
- Keep the f(n+1) best hypotheses according to accumulated score.

#### #f(1)=M and $f(n+1) \leq f(n)$

 f(n) - the preemption function  $f(n) = |M2^{-\lfloor \frac{n}{B} \rfloor}|$ B - block size (f only changes every B steps) M - number of models # Accumulated scoring  $L(m) = \sum \rho(n, m)$ n=1 Log-likelihood of sample *n* given model *m*  $\rho(n,m)$ 

#### Chum, et al., Two-view Geometry Estimation Unaffected by a Dominant Plane, CVPR'05

# Planar dominant scenes are also problematic



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In le5 we saw that if all seven points are in a plane, then

$$\mathbf{x}_k^T \mathbf{F} \mathbf{y}_k = 0, \ \mathbf{x}_k = \mathbf{H} \mathbf{y}_k, \quad k = 1 \dots 7$$

and  $\mathbf{F} = [\mathbf{e}]_{\times} \mathbf{H}$  for any epipole **e** (why epipole?)

\* If six points are in a plane  $\mathbf{x}_k^T \mathbf{F} \mathbf{y}_k = 0$ ,  $k = 1 \dots 7$   $\mathbf{x}_k = \mathbf{H} \mathbf{y}_k$ ,  $k = 1 \dots 6$  $\mathbf{F} = [\mathbf{e}]_{\times} \mathbf{H}$  for  $\mathbf{e} \in \mathbb{R}^3$ ,  $\mathbf{e}^T (\mathbf{H} \mathbf{x}_7 \times \mathbf{y}_7) = 0$ 



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# If *six* points are in a plane  $\mathbf{x}_k^T \mathbf{F} \mathbf{y}_k = 0, \quad k = 1 \dots 7 \quad \mathbf{x}_k = \mathbf{H} \mathbf{y}_k, \quad k = 1 \dots 6$  $\mathbf{F} = [\mathbf{e}]_{\times} \mathbf{H}$  for  $\mathbf{e} \in \mathbb{R}^3$ ,  $\mathbf{e}^T (\mathbf{H} \mathbf{x}_7 \times \mathbf{y}_7) = 0$ For *five* points in the plane  $\mathbf{x}_6 \times (\mathbf{H}\mathbf{y}_6)$  and  $\mathbf{x}_7 \times (\mathbf{H}\mathbf{y}_7)$ define two lines that intersect in e. F will have all points consistent with H as inliers.

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$$\mathbf{H} = \mathbf{A} - \mathbf{e}_1 (\mathbf{M}^{-1} \mathbf{b})^T$$

where  $\mathbf{A} = [\mathbf{e}_1]_{\times} \mathbf{F}$   $\mathbf{M} = [\mathbf{x}_1 \ \mathbf{x}_2 \ \mathbf{x}_3]^T$ 

and  $b_k = (\mathbf{x}_k \times \mathbf{A}\mathbf{y}_k)^T (\mathbf{x}_k \times \mathbf{e}_1) ||\mathbf{x}_k \times \mathbf{e}_1||^{-2}$ 

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This H is now checked for two additional inliers. If found, F is said to be H-degenerate

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There are  $\binom{7}{5} = 21$  ways to pick five points from 7.

\*\* But, if we pick the 3 points that define H as {1, 2, 3}, {4, 5, 6}, {1, 2, 7}, {4, 5, 7}, {3, 6, 7}
\*\* We will have covered all 21 permutations.

Thus at most five H need to be computed and tested to find out if F is H-degenerate.

#### DEGENSAC algorithm

- 1. Select 7 random correspondences and estimate F
- 2. IF best support this far
- 3. IF H-degeneracy
- 4. Do inner RANSAC and estimate F
   from H and 2 correspondences
   (Plane+Parallax algorithm)
   that are inconsistent with H
- IF new F has even bigger support, store F
   ELSE store H

## DISCUSSION

Discussion of the paper: Ondrej Chum and Jiri Matas, Matching with PROSAC -- Progressive Sample Consensus, CVPR'05

#### FOR NEXT WEEK...

#Hartley&Zisserman, Appendix A4.3

**% K. Shoemake**, Animating Rotation with Quaternion Curves, SIGGRAPH85