

3D Computer Vision

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Open Informatics Master's Course

► Metropolis-Hastings (MH) Sampling

C, S – configurations (of all variable values)

e.g. $C = x$ and $\pi(C) = \pi(x)$ from $\rightarrow 117$

Goal: Generate a sequence of random samples $\{C_t\}$ from target distribution $\pi(C)$

- setup a Markov chain with a suitable transition probability to generate the sequence

Sampling procedure

1. given current config. C_t , draw a random config. sample S from $q(S | C_t)$

↗ collapse ↘

q may use some information from C_t (Hastings)

2. compute acceptance probability

the evidence term drops out

$$a = \min \left\{ 1, \frac{\pi(S)}{\pi(C_t)} \cdot \frac{q(C_t | S)}{q(S | C_t)} \right\}$$

$$q(S | C_t) = q(S) = \binom{q}{7}^{-1}$$

3. draw a random number u from unit-interval uniform distribution $U_{0,1}$

4. if $u \leq a$ then $C_{t+1} := S$ else $C_{t+1} := C_t$

'Programming' an MH sampler

1. design a proposal distribution (mixture) q and a sampler from q

2. write functions $q(C_t | S)$ and $q(S | C_t)$ that are proper distributions

not always simple

Finding the mode

- remember the best sample

fast implementation but must wait long to hit the mode

- use simulated annealing

very slow

- start local optimization from the best sample

good trade-off between speed and accuracy

an optimal algorithm does not use just the best sample: a Stochastic EM Algorithm (e.g. SAEM)

► The Nine Elements of a Data-Driven MH Sampler

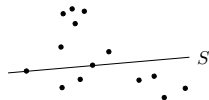
data-driven = proposals are derived from data

Then

1. **primitives** = elementary measurements

- points in line fitting
- matches in epipolar geometry or homography estimation

2. **configuration** = s -tuple of primitives minimal subsets necessary for parameter estimate



the minimization will be over a discrete set:

- of point pairs in line fitting (left)
- of match 7-tuples in epipolar geometry estimation

3. a map from configuration C to parameters $\theta = \theta(C)$ by solving the **minimal problem**

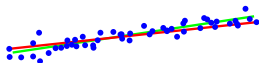
- line parameters \mathbf{n} from two points
- fundamental matrix \mathbf{F} from seven matches
- homography \mathbf{H} from four matches, etc

4. **target likelihood** $p(E, D | \theta(C))$ is represented by $\pi(C)$

- can use log-likelihood: then it is the sum of robust errors $\hat{V}(e_{ij})$ given \mathbf{F} (26)
 - robustified point distance from the line $\theta = \mathbf{n}$
 - robustified Sampson error for $\theta = \mathbf{F}$, etc
- posterior likelihood $p(E, D | \theta)p(\theta)$ can be used

MAPSAC ($\pi(S)$ includes the prior)

5. parameter distribution follows the **empirical distribution** of s -tuples. Since the proposal is done via the minimal problem solver, it is 'data-driven',



- pairs of points define line distribution $p(\mathbf{n} | X)$ (left)
- random correspondence 7-tuples define epipolar geometry distribution $q(\mathbf{F} | M)$

6. **proposal distribution** $q(\cdot)$ is just a constant(!) distribution of the s -tuples: $\frac{\tau(s)}{\pi(C_t)}$
- q uniform, independent $q(S | C_t) = q(S) = \binom{mn}{s}^{-1}$, then $a = \min \left\{ 1, \frac{p(S)}{p(C_t)} \right\}$
 - q dependent on descriptor similarity PROSAC (similar pairs are proposed more often)
 - q dependent on the current configuration C_t e.g. 'not far from C_t '

7. (optional) hard **inlier/outlier discrimination** by the threshold (27)

$$\hat{V}(e_{ij}) < e_T, \quad e_T = \sigma_1 \sqrt{-\log t^2}$$

8. **local optimization** from promising proposals

- can use the hard inliers or just the robust error (26) (more expensive but more stable)
- cannot be used to replace C_t (it would violate 'detailed balance' required for the MH scheme)

9. **stopping** based on the probability of proposing an all-inlier configuration →123

► Data-Driven Sampler Stopping

- The number of proposals N needed to hit the “true parameters” = an all-inlier config?
this will tell us nothing about the accuracy of the result

P ... probability that at least one proposal is all-inlier $1 - P$... all previous N proposals were bad

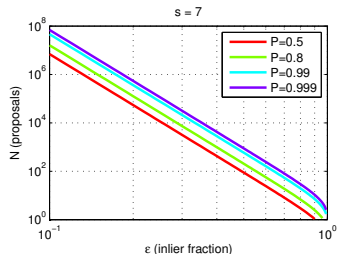
ε ... the fraction of inliers among primitives, $\varepsilon \leq 1$

s ... minimal configuration size 2 in line fitting, 7 in 7-point algorithm, 4 in homography fitting...

$$N \geq \frac{\log(1 - P)}{\log(1 - \varepsilon^s)}$$

- ε^s ... proposal does not contain an outlier
- $1 - \varepsilon^s$... proposal contains at least one outlier
- $(1 - \varepsilon^s)^N$... N previous proposals contained an outlier = $1 - P$

		N for $s = 7$	
		P	
ε	0.8	0.99	
0.5	205	590	
0.2	$1.3 \cdot 10^5$	$3.5 \cdot 10^5$	
0.1	$1.6 \cdot 10^7$	$4.6 \cdot 10^7$	



- N can be re-estimated using the current estimate for ε (if there is LO, then after LO)
the quasi-posterior estimate for ε is the average over all samples generated so far
- this shows we have a good reason to limit all possible matches to tentative matches only
- for $\varepsilon \rightarrow 0$ we gain nothing over the standard MH-sampler stopping rule

► Stripping MH Down To Get RANSAC [Fischler & Bolles 1981]

- when we are interested in the best config only... and we need fast data exploration...

Simplified sampling procedure

1. given C_t , draw a random sample S from $q(S|C_t)$ $q(S)$ independent sampling
no use of information from C_t

2. compute acceptance probability

$$a = \min \left\{ 1, \frac{\pi(S)}{\pi(C_t)} \cdot \frac{q(C_t | S)}{q(S | C_t)} \right\}$$

3. draw a random number u from unit-interval uniform distribution $\mathbb{U}_{0,1}$
4. if $u \leq a$ then $C_{t+1} := S$ else $C_{t+1} := C_t$
5. if $\pi(S) > \pi(C_{\text{best}})$ then remember $C_{\text{best}} := S$

Steps 2–4 make no difference when waiting for the best sample configuration

support

- ... but getting a good accuracy configuration might take very long this way
- good overall exploration but slow convergence in the vicinity of a mode where C_t could serve as an attractor
- cannot use the past generated configurations to estimate any parameters
- we will fix these problems by (possibly robust) 'local optimization'

► RANSAC with Local Optimization and Early Stopping

- initialize the best configuration as empty $C_{\text{best}} := \emptyset$ and time $k := 0$
- estimate the number of needed proposals as $N := \binom{n}{s}$ n - No. of primitives, s - minimal config size
- while $k \leq N$:

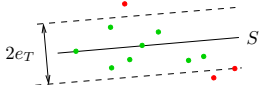
a) propose a minimal random config S of size s from $q(S)$

b) if $\pi(S) > \pi(C_{\text{best}})$ then

i) update the best config $C_{\text{best}} := S$

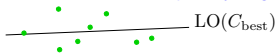
$\pi(S)$ marginalized as in (26); $\pi(S)$ includes a prior \Rightarrow MAP

ii) threshold-out inliers using e_T from (27)



$$\delta = \frac{1}{n\beta} \quad \frac{1}{2} \leq \beta \leq 1$$

iii) start local optimization from the inliers of C_{best} LM optimization with robustified (\rightarrow 114) Sampson error possibly weighted by posterior $\pi(m_{ij})$ [Chum et al. 2003]



iv) update C_{best} , update inliers using (27), re-estimate N from inlier counts

\rightarrow 123 for derivation

$$N = \frac{\log(1 - P)}{\log(1 - \epsilon^s)}, \quad \delta = \frac{|\text{inliers}(C_{\text{best}})|}{mn}$$

c) $n = k + 1$

4. output C_{best}

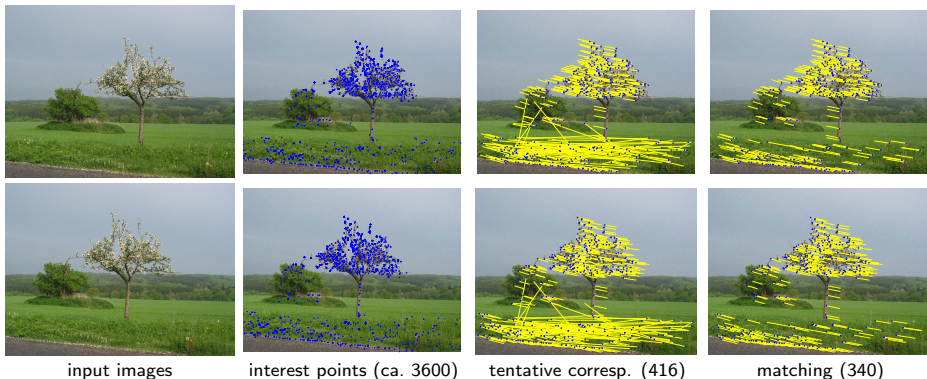
• see [MPV course](#) for RANSAC details

$$\odot \quad \epsilon_m = (1 - \rho^{(m)}) \epsilon_{m-1} + \rho^{(m)} \delta$$

ROBUST-MONRO (1951)

see also [Fischler & Bolles 1981], [25 years of RANSAC]

Example Matching Results for the 7-point Algorithm with RANSAC



- notice some wrong matches (they have wrong depth, even negative)
- they cannot be rejected without additional constraints or scene knowledge
- without local optimization the minimization is over a discrete set of epipolar geometries proposable from 7-tuples

Beyond RANSAC

By marginalization in (23) we have lost constraints on M (e.g. uniqueness). One can choose a better model when not marginalizing:

$$\pi(M, \mathbf{F}, E, D) = \underbrace{p(E | M, \mathbf{F})}_{\text{reprojection error}} \cdot \underbrace{p(D | M)}_{\text{similarity}} \cdot \underbrace{p(\mathbf{F})}_{\text{prior}} \cdot \underbrace{P(M)}_{\text{constraints}}$$

this is a global model: decisions on m_{ij} are no longer independent!

In the MH scheme

- one can work with full $p(M, \mathbf{F} | E, D)$, then configuration $C = M$ \mathbf{F} computable from M
- explicit labeling m_{ij} can be done by, e.g. sampling from

$$q(m_{ij} | \mathbf{F}) \sim ((1 - P_0) p_1(e_{ij} | \mathbf{F}), P_0 p_0(e_{ij} | \mathbf{F}))$$

when $P(M)$ uniform then always accepted, $a = 1$

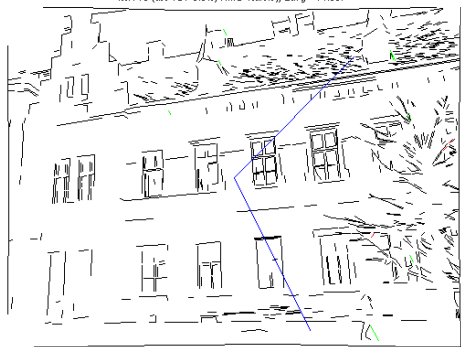
⊗ derive

- we can compute the posterior probability of each match $p(m_{ij})$ by histogramming m_{ij} from $\{C_i\}$
 - local optimization can then use explicit inliers and $p(m_{ij})$
 - error can be estimated for elements of \mathbf{F} from $\{C_i\}$ does not work in RANSAC!
 - large error indicates problem degeneracy this is not directly available in RANSAC
 - good conditioning is not a requirement we work with the entire distribution $p(\mathbf{F})$
 - one can find the most probable number of epipolar geometries (homographies or other models) by reversible jump MCMC and Bayesian model selection
- if there are multiple models explaining data, RANSAC will return one of them randomly

Example: MH Sampling for a More Complex Problem

Task: Find two vanishing points from line segments detected in input image. Principal point is known, square pixel.

iter: 10 (acc TOT=0.0%, HMC=NaN%); Eavg = 14.597



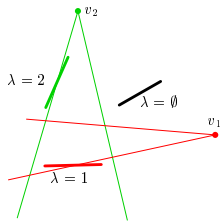
video

simplifications

- vanishing points restricted to the set of all pairwise segment intersections
- mother lines fixed by segment centroid, then θ_L uniquely given by λ_i , and the configuration is

$$C = \{v_1, v_2, \Lambda\}$$

- primitives = line segments
- latent variables
 1. each line has a vanishing point label $\lambda_i \in \{\emptyset, 1, 2\}$, \emptyset represents an outlier
 2. 'mother line' parameters θ_L (they pass through their vanishing points)
- explicit variables
 1. two unknown vanishing points v_1, v_2
- marginal proposals (v_i fixed, v_j proposed)
- minimal configuration $s = 2$



$$\arg \min_{v_1, v_2, \Lambda, \theta_L} V(v_1, v_2, \Lambda, \theta_L)$$

Thank You

$s = 7$

