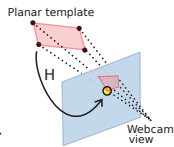


WHAT WE SHOW ? HOMOGRAPHY FITTING USING A THRESHOLD-FREE ROBUST ESTIMATOR

- A threshold-free model estimation method:
 - adapts to noise,
 - works in real time,
 - works with large outlier contamination.



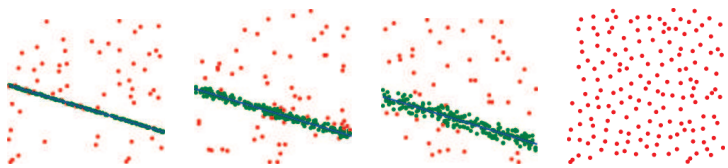
- Underlying mathematical tool: *a contrario* RANSAC (AC-RANSAC)
 - finds both a model and an associated confidence score.
- Used for:
 - F matrix [1], H matrix [2], Structure from Motion [3].

THE ROBUST ESTIMATION PROBLEM: THE THRESHOLD DILEMMA

RANSAC requires the choice of a threshold T , which must be balanced:

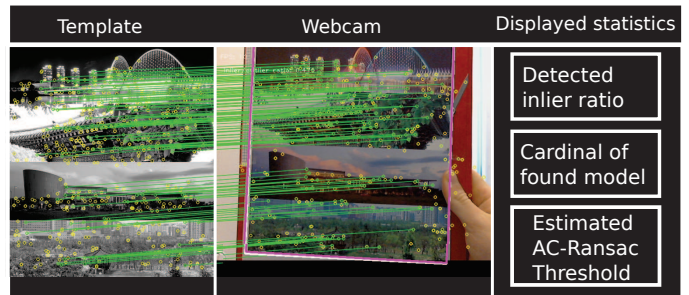
- Too small: too few inliers, leading to model imprecision,
- Too large: models are contaminated by outliers (false data).

Robust line estimation:



Goal: making T adaptive to data and noise.

Demonstration scenario: Homography estimation:



Validated inliers in green. Estimated homography is pictured by reprojecting template boundaries: RANSAC estimation in magenta, AC-RANSAC estimation in white.

From left to right: impact of RANSAC threshold (transfer error through homography):



$T = 0.5$ pixels: 6 points correspondences



$T = 2$ pixels: 19 points



$T = 6.8$ pixels: 50 points

$T = 6.8$ was automatically computed with the *a contrario* technique, that statistically determines a confidence threshold.

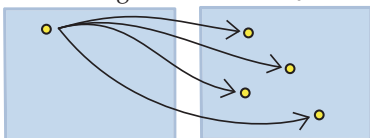
A CONTRARIO MODEL ESTIMATION : A THRESHOLD-FREE FRAMEWORK

AC-RANSAC. A threshold-free rigid model estimation framework.

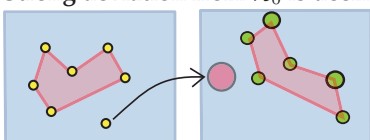
- The method answers the question: "Could the rigid set of data have occurred by chance?"
- The threshold T adapts for inlier/outlier discrimination.
- It provides a confidence score for each model.

A *contrario* criterion [1]:

- Use a background model \mathcal{H}_0 : uniform distribution.



- Strong deviation from \mathcal{H}_0 is deemed meaningful.



$$\mathbb{P}(\text{residual} \leq T | \mathcal{H}_0) = \frac{\pi T^2}{w \times h}$$

(ratio of radius T disk and image areas)

AC-RANSAC relies on the following definitions:

- Number of False Alarms (NFA) measures model fitness to data.
- Given model M , assuming k inliers among n correspondences, T_k denotes the k^{th} smallest residual.

$$NFA(M, k) = N_{\text{tests}}(n, k, N_{\text{sample}}) \mathbb{P}(\text{residual} \leq T_k | M, \mathcal{H}_0)^{k - N_{\text{sample}}}$$

Expectation: $NFA(M) = \min_{k=N_{\text{sample}}+1 \dots n} NFA(M, k) \leq 1$.

RANSAC maximizes inlier count wrt sample at fixed T .
AC-RANSAC minimizes NFA wrt sample with varying T_k .

- ACCV12 paper [3], Poster PF-36:

- AC-RANSAC for SfM. Released in the openMVG library.



REFERENCES

[1] L. Moisan and B. Stival. A Probabilistic Criterion to Detect Rigid Point Matches Between Two Images and Estimate the Fundamental Matrix. In *IJCV 2004*.
 [2] L. Moisan, P. Moulon and P. Monasse. Automatic homographic registration of a pair of images, with a contrario elimination of outliers. In *IPOL 2012*, <http://dx.doi.org/10.5201/ipol.2012.mmm-oh> (with online demo and source code)
 [3] P. Moulon, P. Monasse and R. Marlet. Adaptive Structure from Motion with a *contrario* model estimation. In *ACCV 2012*.