3D Computer Vision

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Open Informatics Master's Course

How Difficult Is Stereo?



Centrum för teknikstudier at Malmö Högskola, Sweden

The Vyšehrad Fortress, Prague

- top: easy interpretation from even a single image
- bottom left: we have no help from image interpretation
- bottom right: ambiguous interpretation due to a combination of missing texture and occlusion

A Summary of Our Observations and an Outlook

- 1. simple matching algorithms do not work
 - the success of a model-free stereo matching is unlikely ${\rightarrow}153$
 - without scene recognition or use high-level constraints the problem seems difficult
- 2. stereopsis requires image interpretation in sufficiently complex scenes or another-modality measurement

we have a tradeoff: model strength \leftrightarrow universality

Outlook:

- 1. represent the occlusion constraint:
 - disparity in rectified images
 - uniqueness as an occlusion constraint
- 2. represent piecewise continuity
 - ordering as a weak continuity model
- 3. use a consistent framework
 - finding the most probable solution (MAP)

correspondences are not independent due to occlusions

the weakest of interpretations; piecewise: object boundaries

Binocular Disparity in a Standard Stereo Pair



• Assumptions: single image line, standard camera pair

$$b = z \cot \alpha_1 - z \cot \alpha_2 \qquad b = \frac{b}{2} + x - z \cot \alpha_2$$
$$u_1 = f \cot \alpha_1 \qquad u_2 = f \cot \alpha_2$$

• eliminate
$$\alpha_1$$
, α_2 and obtain:
 $X = (x, y, z)$ from disparity $d = u_1 - u_2$:

$$z = \frac{b f}{d}$$
, $x = \frac{b}{d} \frac{u_1 + u_2}{2}$, $y = \frac{b v}{d}$

f, d, u, v in pixels, b, x, y, z in meters

Observations

- constant disparity surface is a frontoparallel plane
- distant points have small disparity
- relative error in z is large for small disparity

$$\frac{1}{z} \frac{\mathrm{d}z}{\mathrm{d}d} = -\frac{1}{d}$$

• increasing the baseline or the focal length increases disparity and reduces the error

Structural Ambiguity in Stereovision

- suppose we can recognize local matches independently but have no scene model
- lack of an occlusion model
- lack of a continuity model



left image



structural ambiguity in the presence of repetitions (or lack

right image



of texture)

► Understanding Basic Occlusion Types



• surface point at the intersection of rays l and r_1 occludes a world point at the intersection (l, r_3) and implies the world point (l, r_2) is transparent, therefore

 (l,r_3) and (l,r_2) are <u>excluded</u> by (l,r_1)

- in half-occlusion, every 3D point such as X_1 or X_2 is excluded by a binocularly visible surface point such as Y_1 , Y_2 , Y_3 \Rightarrow decisions on correspondences are not independent
- in mutual occlusion this is no longer the case: any X in the yellow zone above is not excluded

 \Rightarrow decisions inside the zone are independent on the rest



► Matching Table

Based on scene opacity and the observation on mutual exclusion we expect each pixel to match at most once.





matching table

- rows and columns represent optical rays
- nodes: possible correspondence pairs
- full nodes: matches
- numerical values associated with nodes: descriptor similarities

see next

► Constructing An Image Similarity Cost

• let $p_i = (l, r)$ and $\mathbf{L}(l)$, $\mathbf{R}(r)$ be (left, right) image descriptors (vectors) constructed from local image neighborhood windows



• another successful (dis-)similarity is the Hamming Distance over the Census Transform related to local binary patterns

How A Scene Looks in The Filled-In Matching Table



- MNCC ρ used ($\alpha = 1.5, \beta = 1$) \rightarrow 176
- high-similarity structures correspond to scene objects

Things to notice:

constant disparity

- a diagonal in matching table
- zero disparity is the main diagonal assuming standard stereopair

depth discontinuity

horizontal or vertical jump in matching table

large image window

- similarity values have better discriminability
- worse occlusion localization

repeated texture

horizontal and vertical block repetition

Descriptors: Image points are tagged by their (viewpoint-invariant) physical properties:

- texture window
- a descriptor like DAISY
- learned descriptors
- reflectance profile under a moving illuminant
- photometric ratios
- dual photometric stereo
- polarization signature
- . . .
- similar points are more likely to match

• image similarity values for all 'match candidates' give the 3D matching table

[Tola et al. 2010]

[Moravec 77]

[Wolff & Angelopoulou 93-94] [Ikeuchi 87]

also called: 'disparity volume'

click for video

Marroquin's Winner Take All (WTA) Matching Algorithm

- Alg: Per left-image pixel: The most SAD-similar pixel along the right epipolar line
 - 1. select disparity range

 \rightarrow 169 this is a critical weak point

2. represent the matching table diagonals in a compact form



4 5 6

2 3

3. use an 'image sliding & cost aggregation algorithm'



- **4**. take the maximum over disparities *d*
- 5. threshold results by maximal allowed SAD dissimilarity

A Matlab Code for WTA

```
function dmap = marroquin(iml, imr, disparityRange)
%
        iml. imr - rectified grav-scale images
% disparityRange - non-negative disparity range
% (c) Radim Sara (sara@cmp.felk.cvut.cz) FEE CTU Prague, 10 Dec 12
 thr = 20:
                                                 % bad match rejection threshold
r = 2:
 winsize = 2*r+[1 \ 1];
                                                 % 5x5 window (neighborhood) for r=2
 N = boxing(ones(size(iml)), winsize);
                                                 % the size of each local patch is
                                                 % N = (2r+1)^2 except for boundary pixels
 % --- compute dissimilarity per pixel and disparity --->
 for d = 0:disparityRange
                                                 % cycle over all disparities
  slice = abs(imr(:.1:end-d) - iml(:.d+1:end)): % pixelwise dissimilarity (unscaled SAD)
 V(:,d+1:end,d+1) = boxing(slice, winsize)./N; % window aggregation
 end
 % --- collect winners, threshold, output disparity map --->
 [cmap,dmap] = min(V,[],3);
                                                 % collect winners and their dissimilarities
 dmap(cmap > thr) = NaN;
                                                 % mask-out high dissimilarity pixels
end % of marroquin
function c = boxing(im, wsz)
% if the mex is not found. run this slow version:
 c = conv2(ones(1,wsz(1)), ones(wsz(2),1), im, 'same');
end % of boxing
```

WTA: Some Results



- results are fairly bad
- false matches in textureless image regions and on repetitive structures (book shelf)
- a more restrictive threshold (thr = 10) does not work as expected
- we searched the true disparity range, results get worse if the range is set wider
- chief failure reasons:
 - unnormalized image dissimilarity does not work well
 - no occlusion model (it just ignores the occlusion structure we have discussed ightarrow167)

Thank You





