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Optimierung von Lichtfeldmessungen

mithilfe des Strukturtensors

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ausgeführt am
Institut für wissenschaftliches Rechnen (IWR)
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born in Konstanz

2018

Optimizations of Light Field measurements using the structure tensor

This Master thesis has been carried out by Jacob Nieswand

at the

Institut für wissenschaftliches Rechnen (IWR)

under the supervision of

Herrn Priv.-Doz. Christoph Garbe

Optimierung von EPI-basierten Lichtfeldmessungen:

(Abstract in Deutsch, max. 200 Worte. Beispiel: ?)

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1 Introduction (TODO)

2 Theory

2.1 Light Field Parametrization

The earliest introduction to Light Fields in literature can be found in Adelson et al. [1991], where they parametrize the field of light as a so-called „plenoptic function“. If we assume that every point in space emits a light ray in a given direction which is characterized by a intensity value P , the whole information in the light field is given as a 6-dimensional function

$$P(V_x, V_y, V_z, \theta, \phi, \lambda), \quad (2.1)$$

where θ and ϕ are the solid angles describing the direction of any light field, λ describes the wavelength dependence. $V_{\{x, y, z\}}$ describe the room coordinates. If we use the pixel of a pinhole camera picture as the coordinate system of our choice, the plenoptic function would be parametrized as

$$P(x, y, V_x, V_y, V_z, \lambda). \quad (2.2)$$

A more generalized model of the plenoptic function could also include a time dependence, leading to a 7-dimensional ray space. In general, the plenoptic function serves as the global function that gets mapped into a low-dimensional space in some form by any camera device, e.g. a pinhole camera mapping the whole ray space down to a 2-dimensional image.

A more detailed introduction to Lightfields can also be found in [19].

2.1.1 The Lumigraph

In the form of 2.2 the plenoptic function is a 7-dimensional function, which is difficult to record and to handle. As described by Wu et al. [2017] this plenoptic function is usually simplified in 3 steps: First, we neglect the time dependence (assuming a static scene), further we neglect the wavelength λ , instead we define the mapped value of the plenoptic function as a vector containing the color channels. Additionally the plenoptic function obtains redundant information due to the fact, that light rays that are lying on one line in space propagate in the same direction, as introduced by Bolles et al. [1987]. The 4D- representation is also known as the *Lumigraph*. Those 4 dimensions can be split to 2 angular dimensions describing the direction of each ray and 2 dimensions for the location of the ray: Since every ray would pass an image plane of infinite size once (if the propagation vector is not

parallel), two coordinates for localizing the ray are sufficient. Commonly the *two-plane-parametrisation* is used to describe the light field. Every ray is characterized by the intersection of two arbitrary parallel planes Π and Ω . Mathematically spoken the light field L is a function

$$L : \Omega \times \Pi \rightarrow R \quad x, y, s, t \rightarrow L(x, y, s, t), \quad (2.3)$$

where x, y are the coordinates in the first plane Ω and s, t are the coordinates in the second plane Π . If we move a camera in a plane and take pictures of a scene orthogonal to the camera plane, the image itself is described by the image coordinates x, y while the position of the camera is denoted as s, t . Both planes are parallel to each other; that way a light field can be measured in a straight forward manner.

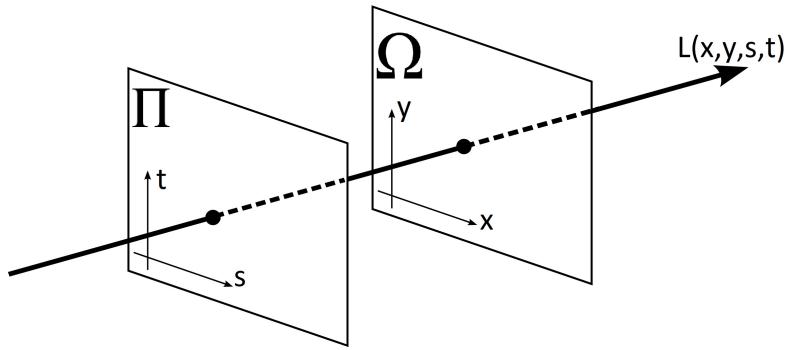


Figure 2.1: In a 4-dimensional two-plane parametrisation a light ray is characterized by the intersection with two parallel planes. we refer to the plane Π as the camera plane and the plane Ω as the image plane. From Xu et al. [2012]

2.1.2 Epipolar Plane Images

We measure the light field in order to obtain as much information about the scene we're looking at as possible. In order to obtain the threedimensional structure one needs to map the light field in a certain manner: We take a look at one 2-dimensional slice through the 4-dimensional space while keeping 2 coordinates constant: one horizontal coordinate x^* in the image plane and one vertical coordinate t^* in the camera plane (or vice versa). The slice Σ_{x^*,t^*} is a 2-dimensionnal image called Epipolar Plane Image (EPI). In figure 2.2 the extraction of an EPI from camera array data is visualized. As seen in figure 2.3 it consists of lines with different slopes, each point on one line with the same slope belongs to the same point in the scene und different angles. From the slope Δ of the line at each point we obtain the distance from the camera plane with the equation

$$\text{distance} = \frac{f \cdot b}{\Delta}, \quad (2.4)$$

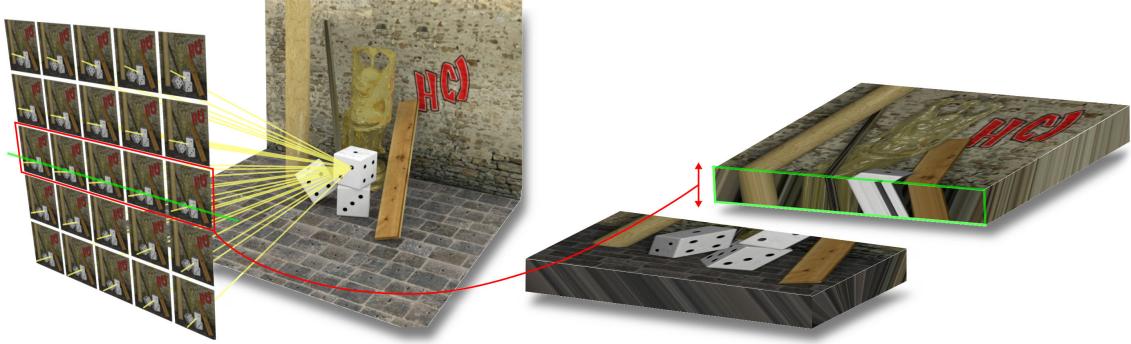


Figure 2.2: Visualization of an Epipolar Plane Image extractoin: A camera array takes images of the same scene from slightly different angles (left array). For a fixed image coordinate y^* (green) and a fixed camera coordinate t^* (red) the pixels are extracted and stacked up resulting in an EPI Σ_{y^*,t^*} (green box on the right). From [2]



Figure 2.3: An Epipolar Plane Image (EPI) that consists of 9 rows (9 equidistant views or sample points in the camera plane). Points with the same color correspond to the same scene point. Since the viewpoints are slightly shifted, the scene point is also shifted in each view by the disparity d . Marked in red one can identify the center position of the camera viewpoints.

where f is the focal length of the camera and b is the baseline between the camera positions. One may notices that this equation looks equal to distance estimation in stereo vision, where we replace the slope Δ by the disparity shift of a feature in 2 views. In fact, a stereo light field capture would result in an EPI with only 2 pixel rows, the slope of a line consisting of only two points is then defined as the disparity shift between the views.

2.2 Depth from Structure Tensor

Measuring depth from light field data can be done using various approaches. Wu et al. [2017] divide light field depth estimation approaches in three categories:

1. Sub-Aperture Image Matching-based Methods
2. EPI-based methods
3. Learning-based methods

In the following we focus on one specific EPI-based method on which is investigated as part of this work. However, the reader is encouraged to take a look at the cited paper „Light Field Image Processing: An Overview“ by Wu et al. [2017] for a state-of-the-art overview of different light field approaches.

Wanner [2014] makes use of the oriented structure of the EPI using image-processing techniques. This idea has first been introduced by Bigun [1987] in 1987.

The 2-dimensional structure tensor J on a function $g : \Omega \rightarrow R, \Omega \subset R^2$ is defined as

$$J = \begin{pmatrix} G * \frac{\partial g}{\partial x} \frac{\partial g}{\partial x} & G * \frac{\partial g}{\partial x} \frac{\partial g}{\partial s} \\ G * \frac{\partial g}{\partial s} \frac{\partial g}{\partial x} & G * \frac{\partial g}{\partial s} \frac{\partial g}{\partial s} \end{pmatrix}, \quad (2.5)$$

where G is a gaussian window function. The derivation can be found in the appendix. The first eigenvector of this tensor J indicates the preferred orientation in the local neighbourhood (defined by the window function). The second eigenvector is orthogonal to the first.

Jähne [2013] proposes a coherence value as a measurement for the anisotropy of the local environment:

$$C = \frac{\sqrt{(J_{11} - J_{22})^2 + 4J_{12}^2}}{J_{11} + J_{22}} \quad (2.6)$$

which obeys the value 1 in case of complete anisotropy, a value of 0 would indicate total isotropy.

In case of an EPI, the structure tensor provides an estimate for the slope of an EPI line, as defined by Bigun [1987]:

$$\Delta = \tan \left(\frac{1}{2} \arctan \left(\frac{J_{22} - J_{11}}{2J_{12}} \right) \right) \quad (2.7)$$

we refer to Δ as the disparity d in the following. Via equation 2.4 one obtains the depth in meters.

2.2.1 Implementation

The following implementation steps have been proposed by Wanner [2014]. The code uses *crosshair* light field data, that can be obtained e.g. by a camera as depicted in figure 2.4. The scope of the light field in the camera coordinates is significantly reduced, in the following the horizontal camera row and the vertical camera column are viewed separately as 1-dimensional light-field cameras.

From the input images one extracts the EPIS, which have the dimensions

vertical view \rightsquigarrow Nr of rows in image coordinates \times Nr of cameras

horizontal view \rightsquigarrow Nr of columns in image coordinates \times Nr of cameras

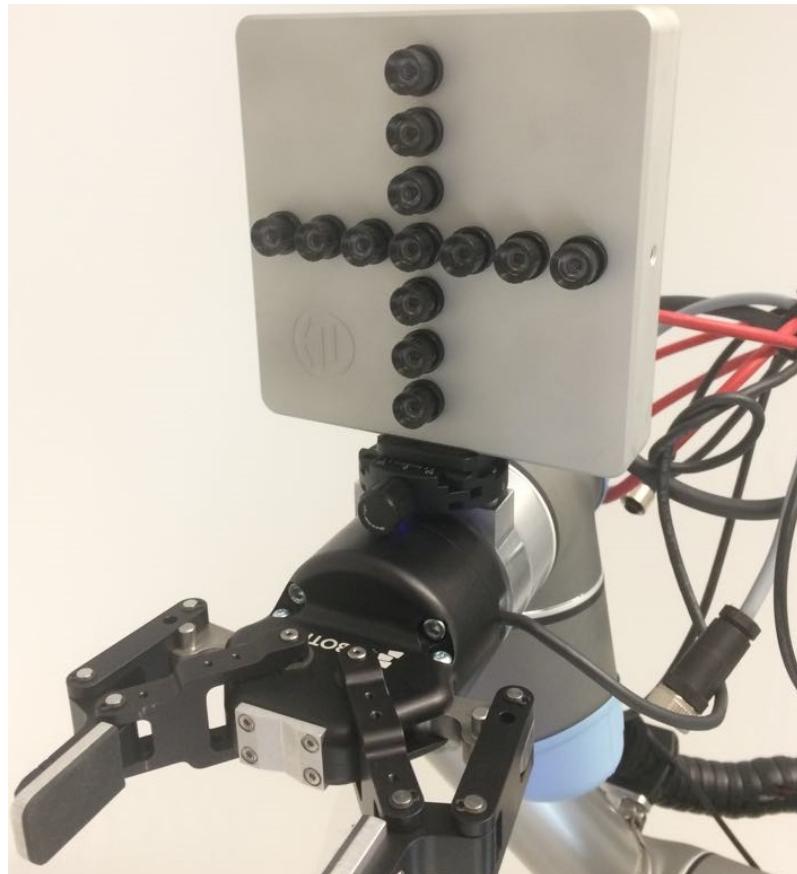


Figure 2.4: Instead of a complete camera array a crosshair camera reduces the number of cameras to one row and one column for faster processing. This *Lumi+* - Scanner from HDVisionSystems is mounted an a robot arm for industrial applications.

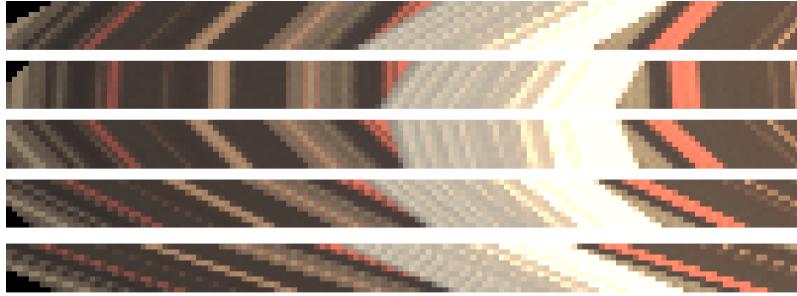


Figure 2.5: The EPI is refocussed by integer disparity steps. If the slope at a scene point is zero (vertical line) the EPI is perfectly focussed on that point. An integration of all views would still result in a sharp image at the corresponding depth.

For each EPI we calculate the structure tensor independently. This is done by first presmoothing the EPI with a 3×3 gaussian kernel to obtain reasonable results when calculating the gradient in the next step. Note that before calculating the gradient, the EPI is converted to grayscale format.

The gradient is calculated via the Scharr-filter, which has the form

$$\text{Scharr}_x = \begin{pmatrix} 3 & 0 & -3 \\ 10 & 0 & -10 \\ 3 & 0 & -3 \end{pmatrix} \quad (2.8)$$

$$\text{Scharr}_y = \begin{pmatrix} 3 & 10 & 3 \\ 0 & 0 & 0 \\ -3 & -10 & -3 \end{pmatrix}. \quad (2.9)$$

[Wanner \[2014\]](#) and [Diebold \[2016\]](#) both tested out different gradient filters and came to the conclusion that the Scharr-filter performs best. With the gradients the structure tensor and the disparity is obtained via equation 2.5 and 2.7.

Further Wanner found out that the structure tensor results are most accurate when the disparity is close to zero. Therefore the EPI is artificially refocussed by simple linear shifting of the rows such that the slope of any line in the EPI would be increased by the same amount. The disparity shift can simply be subtracted from the disparity later. The refocussing of the EPI is illustrated in 2.5. For each EPI the structure tensor is calculated at each disparity shift, so that the necessary disparity range is fully covered.

From all disparity shifts the value with the highest coherence measure (equation 2.6) is selected. Hence one obtains two depth estimations for each image coordinate in the center view: one from the vertical EPI, one from the horizontal EPI. The merging of both direction follows

$$d(x, y) = \begin{cases} d_{\text{horizontal}}(x, y) & C_{\text{horizontal}}(x, y) > C_{\text{vertical}}(x, y) \\ d_{\text{vertical}}(x, y) & C_{\text{horizontal}}(x, y) < C_{\text{vertical}}(x, y) \end{cases}. \quad (2.10)$$

2.2.2 The occlusion problem of the structure Tensor

Having a look at the results of the the benchmark test of Honauer et al. [2016] one realizes that most Light field depth estimation algorithms suffer from large errors near depth discontinuities. Since the center view pixels close to the edge of a depth discontinuity are at least partly occluded, this behaviour is to be expected. The ST almost always produces a systematic error near discontinuities, leading to a „magnification“ of the object closer to the camera in the depth map, see figure 2.7. We refer to this as „edge fattening“. The reason for this error has its origin in the smoothing of the EPI as part of the algorithm. At the occlusion, the edge between the fore- and background follows the same orientation as the foreground does. Furthermore the occlusion mostly comes with a color change, resulting in a high gradient which dominates the local environment orientation: Edge fattening is the consequence. In figure 2.6 one can recognize two effects leading to edge fattening: Firstly the EPI is smoothed with a 3×3 gaussian kernel which already will enlarge the foreground structure (b). This becomes clear when we illustrate the norm of the gradients (c), since the strong (white) gradients all measure the foreground (green) structure. A clear shift to the left is visible from (b) to (c). However, this bias error is small compared to the second error that is indicated in (d): The red dot lies clearly in the background structure. However, calculating the structure tensor components from the local environment (green square) one would obtain the foreground structure at the red dot, since the strong gradients in the square dominate. In (e) the total bias in labeling the two structures is illustrated.

2.3 Modifications on the structure tensor

In the following diffent methods will be explained which modify the Structure tensor algorithm proposed by Wanner [2014]. In section 3 implementations of those modifications will be evaluated and discussed.

2.3.1 Kernelsizes

The Structure tensor algorithm calculates the depth based on the preffered orientation of the EPI in a local neighbourhood. If one chooses a bigger neighbourhood, the resulting depth map is likely to be smoother, however the accuracy at edges in the scene will rapidly decrease.

Even though one often wants to smoothen the depth map in a postprocessing step to improve the overall result and cancel out outlyiers, the structure tensor algorithm by Wanner is forced to smoothen the depth map beforehand.

In the work of Wanner [2014] the effect of the kernel size has been examined extensively and he calculated the best parameters for different scenes by grid search.

Note that if the kernel is smaller then the height of the EPI, the outer cameras are neglected and do not enter in the calculation of the depth. To maintain the information of all cameras and vary the kernel size at the same time, Diebold [2016] proposes

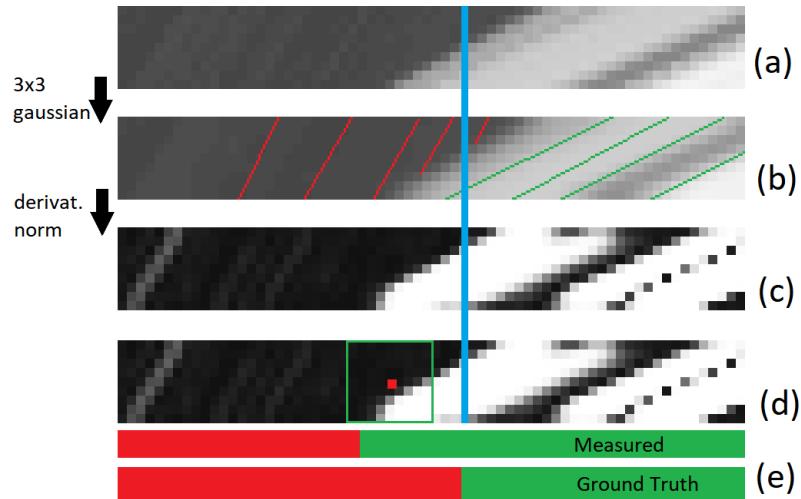


Figure 2.6: (a) The structure of an occlusion border in a gray-value EPI. (b) Before calculating the gradient, the EPI is smoothed with a 3×3 gaussian kernel. One sees the smoothed EPI with colored lines indicating the Ground Truth orientation. (c) shows the norm of the gradient calculated via the Scharr-filter. White signifies a high gradient, black signifies low gradient. In (d) the local environment around the red dot (x_r, y_r) as an example point is marked to show which gradient values go into the structure tensor components $J(x_r, y_r)$. (e) marks the orientation labeling measured and ground truth. The blue line marks the transition in the center view that corresponds with the ground truth labeling.

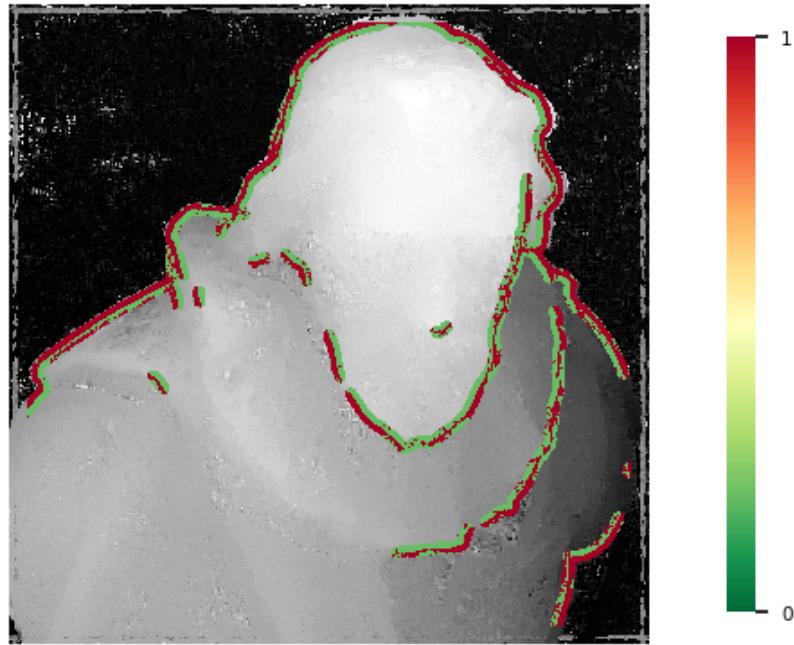


Figure 2.7: Evaluation of the deviation from Ground truth at the depth discontinuity for scene "cotton". The red border indicates that the depth map is erroneous at the outside of the edge.

asymmetrical kernelsizes to eventually reduce edge-fattening effects. However, asymmetrical kernels still need to find a trade off between denoising and avoiding edge fattening and do not significantly increase the quality of the depth map result. Another approach is to choose a custom kernel form which only weights the local gradients that could possibly be part of the „correct“ line in the EPI. Such a kernel is illustrated in figure 2.8, it has the form of a sandclock. in y-direction, it still follows a gaussian distribution.

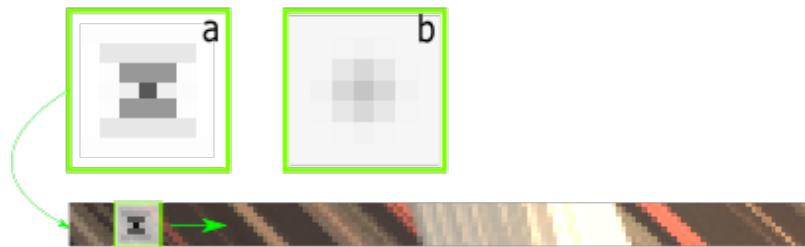


Figure 2.8: (a) shows a custom-shaped kernel in form of a sandclock. (b) shows a normal gaussian kernel.

2.3.2 Color-awareness with bilateral filtering

In its original form the structure tensor pipeline weights the square of all gradients in a local environment by the distance to the reference point. Since all local gradients in the EPI that belong to the same color in the EPI are very likely to belong to the same orientation, in theory one only has to find all points with the same color which then lie on the same line in the EPI. The preferred orientation of the gradients at those points will then correspond to their slope with high confidence. However, most objects are not perfectly lambertian such that not all points on the correct line have to be of the exact same color. Only filtering the exact color will lead to noisy results. Though one could implement a filter weighting not only the distance in a gaussian manner but also the distance from the EPI kernel origin in the color RGB space; such a filter is called „bilateral filter“ ([Tomasi and Manduchi \[1998\]](#)). Pixels with completely different color will be excluded from the Neighbourhood when calculating the structure tensor. In equation 2.5 the window function G is then defined as

$$G_P = \frac{1}{W_p} \sum_{q \in N} g_{\sigma_d}(\|p - q\|) g_{\sigma_c}(\|I_{p,EPI} - I_{q,EPI}\|), \quad (2.11)$$

where

p is the position of the pixel,

q is the position of the pixel in a neighbourhood N ,

g_{σ_d} is a gaussian function weighting the distance between p and q using σ_d as standard deviation,

g_{σ_c} is a gaussian function weighting the color distance in RGB space between the RGB values of the original EPI at p and q using σ_c as standard deviation,

W_p is a normalization factor so that the sum over all the neighbourhood always is 1.

It is important to notice that the color distance is extracted from the EPI, though we are filtering the components of the second derivatives, see equation 2.5. In fact, the EPI only serves as a „guide“ for the filtering.

2.3.3 Thresholding the gradients

In section 2.2.2 it is explained why the structure tensor struggles to serve with a good estimation at depth transitions. It is referred to two errors, one coming from the first smoothing, the other one as a result from the second gaussian smoothing. To avoid the second error, one could in principle normalize all local gradients such that in figure 2.6 (d) the foreground gradients in a local environment obey the same weight in the structure tensor calculation as the background gradients around the

red dot.

The gradient components of the EPI would be given by

$$\tilde{\vec{\nabla}}(x, y) = \frac{1}{\sqrt{\nabla_x(x, y)^2 + \nabla_y(x, y)^2}} \vec{\nabla}_x(x, y) \quad (2.12)$$

Nevertheless in some occasions one wants the stronger gradients to play a bigger role in the preferred orientation, since small gradients are more likely to be noisy. The only way to damp the gradients at occlusions while not strengthening the small noisy gradients in an EPI is to implement a truncation threshold. The gradient is then given by

$$\tilde{\vec{\nabla}}(x, y) = \begin{cases} \frac{\text{threshold}}{\text{norm}} \vec{\nabla}(x, y) & \text{if } \text{norm} > \text{threshold} \\ \tilde{\vec{\nabla}}(x, y) & \text{else} \end{cases} \quad (2.13)$$

$$\text{with } \text{norm} = \sqrt{\nabla_x(x, y)^2 + \nabla_y(x, y)^2}. \quad (2.14)$$

$$(2.15)$$

The threshold should be chosen the lowest possible in a manner that gradients higher than the threshold are barely affected by noise. In section ?? different thresholds are tested, if not mentioned otherwise a threshold of 0.1 is used assuming that the underlying EPI is a gray-value image with range 0-1.

2.3.4 Occlusion-awareness using segmentation of the EPI

In the section a method is proposed that attempts to handle both occlusion errors mentioned in section 2.2.2.

Bilateral filtering (section 2.3.2) and thresholding the gradients (section 2.3.3) may improve the depth map estimation, however they come with trade-offs in order to handle the occlusion problems. An alternative way to maintain sharp transitions is to find those transitions in the EPI and calculate them separately. By setting the gradient at the transition in the EPI to 0, background and foreground structure are calculated without being biased by the transition itself. Therefor the EPI is segmented binarily into occlusion transistions and continuous areas, whereby the norm vector is thresholded to create a binary mask on the EPI. Possibly one could think of a better criterion that is segmenting occlusion/non-occlusion areas, however this approach does not need to calculate the disparity values beforehand and can be implemented in the workflow directly. In principle the algorithm works as follows:

1. Segmentate the transitions and the rest of the EPI with a binary mask based on the gradient norm.
2. Calculate the structure tensor components on the masked gradient of the EPI. All masked gradients are zero and do not affect the local environment structure.

3. Calculate the structure tensor components again, now on the inverted masked gradient of the EPI.
4. Join the the disparity values from both masks.

In figure 2.9 the segmentation process is illustrated step-by-step. The gradient of the gray-valued EPI (a) is calculated in x- and y-direction (b), from which one calculates and thresholds the norm (c, d). The threshold itself is chosen to be 0.5 if not mentioned otherwise. It is essential to modify resulting mask from the binary thresholding (e) using morphological image-processing operators as explained in detail in [hom](#). It is made use of Morphological Closing (structured filling in of image region boundary pixels), which is explained in figure 2.10.

Having a look at the different scenarios in the given EPI that can happen when segmenting, the necessity for morphological closing becomes clear. The EPI has two visible disparity transitions, one at Pixel [133] and one at Pixel [185]. At [133] (a) it is seen that the transition affects multiple neighbouring pixels even though the EPI isn't smoothed yet. Our aim is to mask those transitions. However at Pixel [63] and [174] the threshold is also crossed, since big gradients do not necessarily have to lie on a transition. Still the false-positive masked transitions are less uncomfortable to handle then a false negative result. It is clearly visible at [63], that zero crossings in the gradient lead to holes in the mask which need to be closed by morphological closing. Since the local environment of each pixel is only given by near pixels in the same segment (masked/unmasked), small holes lead to erroneous results at those pixels. After morphological closing, a last postprocessing step has to be made to handle the smearing due to the 3×3 -gaussian presmoothing. In figure 2.10 (d) one sees that the transition peak is gaussian-formed. If one simply cuts out the segment at the threshold transition, the background segment gradient is still dominated by the gradients directly next to the segment transitions. This becomes visible in figure 2.11, where the inverted mask of the EPI gradient is shown. Thus morphological dilation with a 3×3 kernel is applied on the inverted filter, resulting in a complete separation of any peak crossing the threshold.

2.3.5 An alternative to the coherence as the confidence measure

The coherence (2.6) describes how strong the anisotropy of a structure in an image is. In the structure tensor pipeline (section 2.4) this value is used as a confidence measure for the corresponding depth value. In this section we try to define a more profound definition for the confidence based on the distribution of the gradients in a small environment. From the structure tensor components we obtain the disparity

$$d = \tan \left(\frac{1}{2} \arctan \left(\frac{J_{22} - J_{11}}{2J_{12}} \right) \right), \quad (2.16)$$

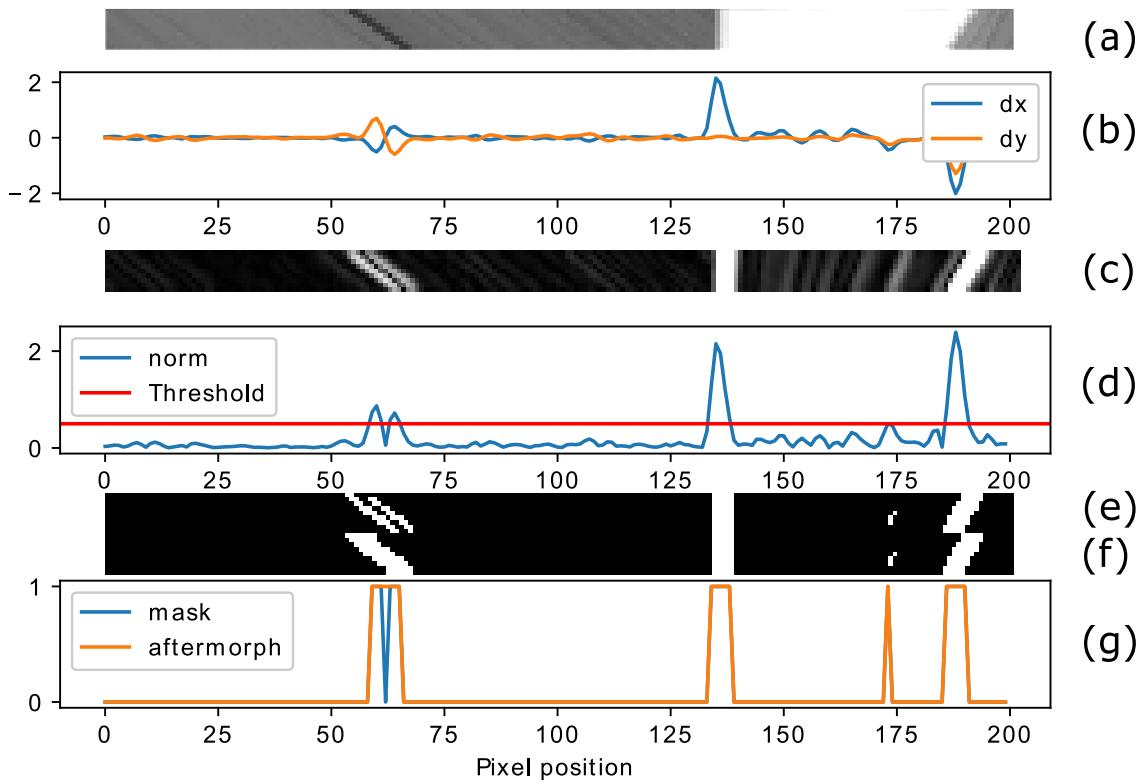


Figure 2.9: (a) Input EPI to be segmented. (b) show the local derivatives along the center view line. (c) shows the vector norm of the derivative. (d) shows the center view norm with the threshold that is applied to mask transitions. (e) shows the resulting mask. (f) shows the improved mask using morphological image processing. (g) shows the mask at the center view before and after using morph. image processing.

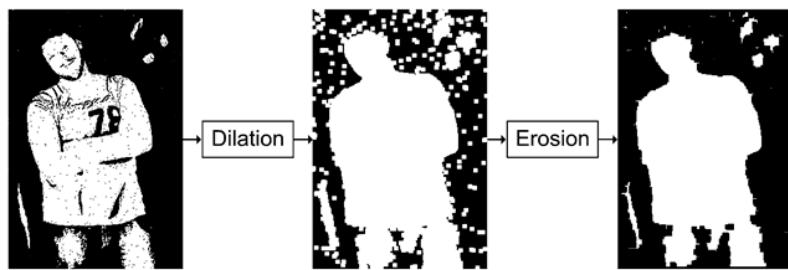


Figure 2.10: Morphological closing is a combination of Dilatation and Erosion. Dilatation uses a custom-sized kernel and turns any 0 to 1, if at least one 1 is found in the local environment. Erosion turns any 1 to 0, if at least one 0 is found in the local environment. Image from [wha](#)

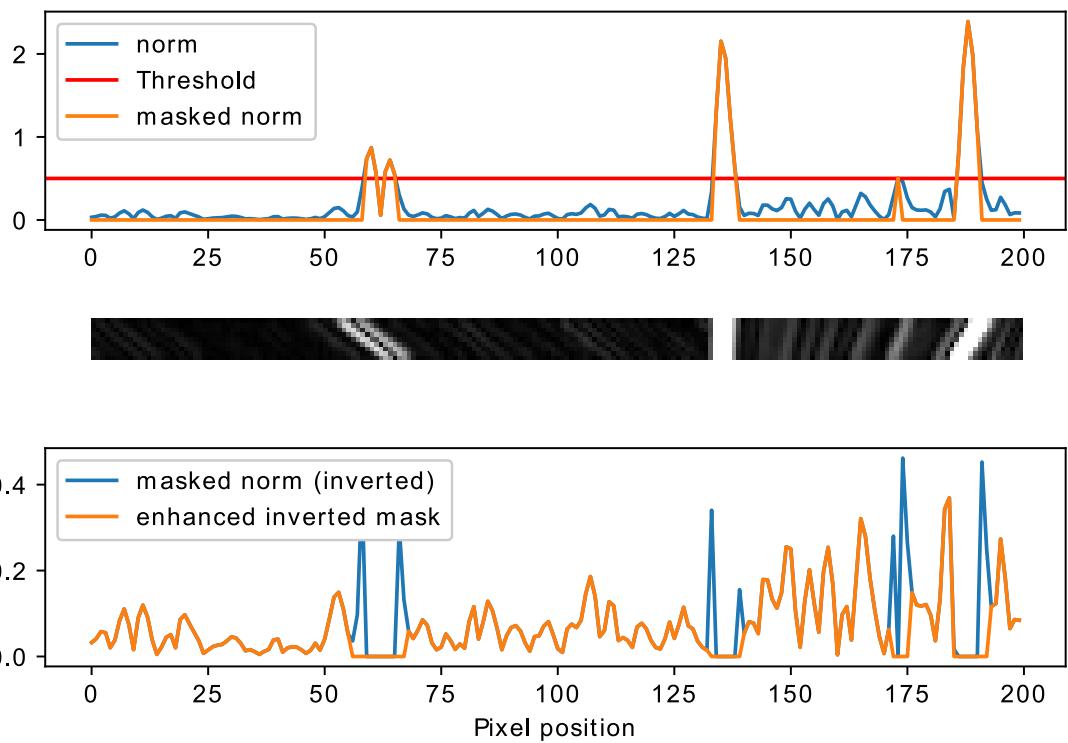


Figure 2.11: In the upper diagram the thresholded gradient norm of the EPI (middle) is seen. The norm (masked with the inverted mask) is depicted in the lower figure (blue), while the inverted mask with a dilation filter is depicted in orange.

whose error Δd is then given by first order error propagation:

$$\Delta d = \sqrt{\left(\frac{\partial d}{J_{11}}\Delta J_{11}\right)^2 + \left(\frac{\partial d}{J_{22}}\Delta J_{22}\right)^2 + \left(\frac{\partial d}{J_{12}}\Delta J_{12}\right)^2} \quad (2.17)$$

When substituting

$$d = \tan\left(\frac{1}{2}\arctan(x)\right), \quad \text{with } x = \frac{J_{22} - J_{11}}{2J_{12}} \quad (2.18)$$

we obtain

$$\frac{\partial d}{\partial x} = 0.5 \cdot \frac{d^2 + 1}{x^2 + 1}. \quad (2.19)$$

With

$$\frac{\partial x}{\partial J_{11}} = \frac{x}{J_{11} - J_{22}}, \quad \frac{\partial x}{\partial J_{22}} = \frac{-x}{J_{11} - J_{22}}, \quad \frac{\partial x}{\partial J_{12}} = \frac{x}{J_{12}} \quad (2.20)$$

inserting in 2.17:

$$\Delta d = \sqrt{\left(\frac{0.5 \cdot (d^2 + 1)}{x^2 + 1} \cdot x\right)^2 \cdot \left(\left|\frac{\Delta J_{12}}{J_{12}}\right|^2 + \left|\frac{\Delta J_{11}}{J_{11} - J_{22}}\right|^2 + \left|\frac{\Delta J_{22}}{J_{11} - J_{22}}\right|^2\right)} \quad (2.21)$$

The errors ΔJ_{11} , ΔJ_{12} and ΔJ_{22} can be obtained from the statistical deviations of the gradient:

$$\Delta J_{11} = G * |2\nabla_x| \Delta \nabla_x \quad (2.22)$$

$$\Delta J_{22} = G * |2\nabla_y| \Delta \nabla_y \quad (2.23)$$

$$\Delta J_{12} = G * \sqrt{\nabla_x^2 \Delta \nabla_y^2 + \nabla_y^2 \Delta \nabla_x^2} \quad (2.24)$$

Those equations are obtained from equation 2.5 using first order error propagation. We define $\Delta \nabla_x$, $\Delta \nabla_y$ as the statistical standard deviation of ∇_x , ∇_y in the neighbourhood and calculate it via

$$\Delta \nabla_x = \sqrt{E[\nabla_x^2] - E[\nabla_x]^2} \quad (2.25)$$

$$\Delta \nabla_y = \sqrt{E[\nabla_y^2] - E[\nabla_y]^2} \quad (2.26)$$

with the expectation value defined as the Gaussian weighted mean:

$$E[I] = G * I \quad (2.27)$$

However, equation 2.25 only makes sense under the assumption that the underlying distribution of the magnitude whose standard deviation will be computed is close to a normal distribution. This is in general not the case for the gradient components

since deviations in the magnitude of the gradient should not affect on the disparity error estimation. Given that the gradients are normalized, a gaussian distribution of the gradient around the mean normalized gradient is expected:

$$\Delta \nabla_{x,\text{normalized}} = \sqrt{E[\nabla_{x,\text{normalized}}^2] - E[\nabla_{x,\text{normalized}}]^2} \quad (2.28)$$

$$\Delta \nabla_{y,\text{normalized}} = \sqrt{E[\nabla_{y,\text{normalized}}^2] - E[\nabla_{y,\text{normalized}}]^2} \quad (2.29)$$

The interrelation between $\Delta \nabla_{y,\text{normalized}}$ and $\Delta \nabla_x$ is given by

$$\nabla_{x,\text{normalized}} = \frac{1}{\sqrt{\nabla_x^2 + \nabla_y^2}} \nabla_x \quad (2.30)$$

$$\Delta \nabla_{x,\text{normalized}} \approx \frac{1}{\sqrt{\nabla_x^2 + \nabla_y^2}} \Delta \nabla_x, \quad (2.31)$$

where we neglect the fact that the normalization itself is also erroneous. This simplifies and speeds up the calculation. Equation 2.30 leads to the expression

$$\Delta \nabla_x = \sqrt{\nabla_x^2 + \nabla_y^2} \cdot \sqrt{E[\nabla_{x,\text{normalized}}^2] - E[\nabla_{x,\text{normalized}}]^2} \quad (2.32)$$

$$\Delta \nabla_y = \sqrt{\nabla_x^2 + \nabla_y^2} \cdot \sqrt{E[\nabla_{y,\text{normalized}}^2] - E[\nabla_{y,\text{normalized}}]^2}, \quad (2.33)$$

which can be calculated from the EPI. If we insert the above equation 2.32 into 2.22 into 2.21, we obtain a deviation measure Δd for the disparity based on the gradient deviation in the EPI.

2.3.6 Alternative Merging of x- and y- orientation

In the old ST Pipeline, the coherence value as defined in equation 2.6 not only determines which depth value should be chosen from refocussing the EPI, it is also used to decide whether the local depth estimation is taken from the x-directed EPI or the y-direction. In the article of [Sheng et al. \[2018\]](#) a full Light field is evaluated such that multiple EPIS are available for each point. To handle occlusions, the EPI with less intensity variance is chosen to calculate the depth. In our case we are working with a crosshair light field only providing two EPIS for each scene point. Choosing the one with higher coherence seems logical, however at occlusions we are mostly provided with a clear structure in the EPI leading to a high coherence value. As a result, the EPI including edges is chosen in the close neighbourhood of a discontinuity edge, leading to **edge fattening**.

In his work, [Wanner \[2014\]](#) also proposes a TV-L minimization scheme to merge the disparity values resulting from x-directional and y-directional EPIS. Despite achieving good results, the improvement did not justify the calculation time of such optimization scheme. He thus sticks to choose the value with higher coherence. Instead one simply chooses the disparity value which relates to the a larger depth,

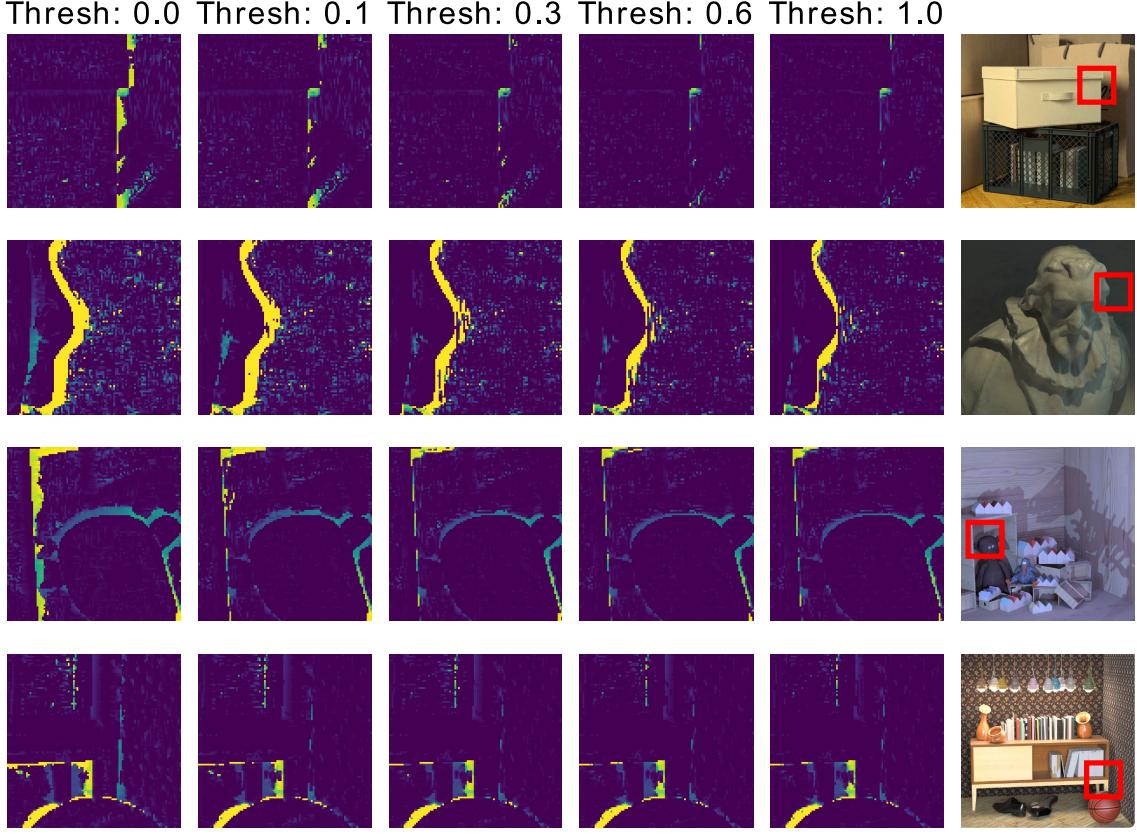


Figure 2.12: From left to right: Parameter dependent alternative merging. Up to down: scenes *boxes*, *cotton*, *dino*, *sideboard*.

and only chooses the closer disparity value, if the coherence difference is big enough. Therefor we introduce a threshold:

$$d(x, y) = \begin{cases} \min(d_x(x, y), d_y(x, y)) & \text{if } |c_x(x, y) - c_y(x, y)| < \text{threshold} \\ d_x(x, y) & \text{else, if } c_x(x, y) > c_y(x, y) \\ d_y(x, y) & \text{else} \end{cases} \quad (2.34)$$

Figure 2.12 shows the effect of varying the threshold introduced in equation 2.34. The occlusion edges shrink significantly with a higher threshold.

2.4 Depth from focus

One advantage of using lightfields for depth measure is its ability to get a two-dimensional mapping of the scene with focus at any depth. Integrating the views of the light field camera array has the same effect as the integration of a focussed lense camera, as the lense is simply integrating slightly different viewpoints of the same scene point when focussed on the correct depth.

Obtaining the refocussed integrated image is a synthetic process that only requires

shifting the view coordinates artificially. Given a full four-dimensional light field $L(u, v, x, y)$ we can refocus the light field as described in [15] :

$$L'(u, v, x, y) = L(u(1 - d'), v(1 - d'), x, y), \quad (2.35)$$

where d' describes the relative pixel shift. The disparity is directly related to the absolute depth of the focus (relate to PICTURE) if the relevant camera parameters are known. Given the baseline b in meters and the focal length f in pixels, the depth Z is given as

$$Z = \frac{f \cdot b}{d}. \quad (2.36)$$

We obtain

$$\bar{L}(x, y) = \frac{1}{N_{u,v}} \int \int L'(u, v, x, y) du dv = \frac{1}{N_{u,v}} \sum_u \sum_v L'(u, v, x, y) \quad (2.37)$$

Once we can focus at any range, one can adopt *depth-from-focus*-techniques as described in Watanabe and Nayar [1998] for depth measure. If the scene point at a given image coordinate (x, y) in the center view is in focus, the contrast in the integrated image $\bar{L}(x, y)$ is high, thus a contrast measure at each pixel combined with stepwise refocussing yields a depth map.

For measuring the contrast, one has different options: The most straight forward approach is calculating the first derivative of the grey-value image. At high contrast structure the local intensity changes are expected to be high. Alternatively one could measure the second derivative laplacian that eventually results in higher robustness. The implementation and tests of those techniques for the benchmark dataset can be found in section 3.1.

Using a pinhole camera array allows us to go further and find a response value that shows higher consistency. Taking the absolute difference between the center view of the camera array and the refocussed image yields to promising results as shown in Tao et al. [2017]. Under the assumption of lambertian surfaces the RGB- value of any scene point should be the same under all angles. Thus when refocussed on the correct depth, summing over all angles should result in a value that ideally is the same as in the center view alone. This is referred as *photo consistency*; for more information read Tao et al. [2017]. The response value at a given depth is obtained from

$$D'(x, y) = \frac{1}{|W_D|} \sum_{x', y' \in W_D} |\bar{L}(x', y') - P(x', y')|, \quad (2.38)$$

where $P(x, y)$ is the center view. For more robustness, it is averaged over a small window. We refer to this measuring technique as *photo consistency* in the following. Note that calculating the absolute results in a 1-channel-image while the input images are RGB-images.

Tao et al. propose another measure that they refer to as *angular correspondence*. It follows the same principle, but instead of integrating the refocussed lightfield followed by comparing it to the center view, they directly take the difference of each viewpoint to the center view and sum up those differences:

$$D'(x, y) = \frac{1}{N_{u,v}} \sum_u \sum_v |L'(u, v, x, y) - P(x, y)|. \quad (2.39)$$

We tested those methods against the common contrast measures mentioned above, the results are found in section 3.1.

2.5 Semi-Global Matching

2.5.1 Semi - Global Matching for Stereo Vision

In contrast to Light field depth estimation techniques Stereo systems often suffer from mismatching pixels between the left and right images. Many attempts have been made to smoothen bad pixels, resulting in blurred edges or long calculation times. One promising attempt to improve matching results was published in 2005 by Heiko Hirschmüller ([Hirschmüller \[2005\]](#)) that was described as „a very good trade off between runtime and accuracy“ ([Hirschmüller \[2011\]](#)): we speak of Semi-Global Matching.

In general, matching of two stereo images means shifting the disparity over the predefined disparity range and comparing both images (pixel- or blockwise) until we have a cost value at each image point for each discrete disparity. We assign to each pixel \vec{p} the disparity value $D_{\vec{p}}$ which is related to the lowest cost $C(\vec{p}, D_{\vec{p}})$. This matching does not have to be unique, resulting in erroneous pixel disparities. To overcome this one wants to minimize a global cost function of the form

$$E(D) = \sum_{\vec{p}} \left(C(\vec{p}, D_{\vec{p}}) + \sum_{q \in N_p} \begin{cases} P1 & \text{if } |D_{\vec{p}} - D_{\vec{q}}| = 1 \\ P2 & \text{if } |D_{\vec{p}} - D_{\vec{q}}| \geq 1 \\ 0 & \text{else} \end{cases} \right). \quad (2.40)$$

The first term sums all matching costs over the whole image, while the second term forces continuity by comparing the disparity of all neighbour pixels N_q to the disparity D_p ; if a small discontinuity is detected ($|D_{\vec{p}} - D_{\vec{q}}| = 1$), a small penalty is added to the global cost function. Since a small discontinuity can be found essentially at any tilted plane, only a small error is added. A bigger disparity difference indices a clear discontinuity in the disparity map. Note that the penalty $P2$ can be divided by the gradient of the original image to allow a disparity discontinuity when we find edges in the image; at these points we expect the disparity to be discontinuous.

However, minimizing the global cost function involves computational cumbersome algorithms as it is a NP-complete Problem ([Hirschmüller \[2011\]](#)). Semi-Global Matching however chooses another approach by minimizing the global cost function along one-dimensional lines – this can indeed be calculated in polynomial time.

The new smoothed cost function $S(\vec{p}, D_{\vec{p}})$ at pixel \vec{p} is then given as the sum of all 1D minimum cost paths that are ending in \vec{p} . The minimal cost L'_r along the path r is defined recursively as

$$L'_r(\vec{p}, D) = C(\vec{p}, D) + \min \begin{cases} L'_r(p_{\text{before}}, D) \\ L'_r(p_{\text{before}}, D + 1) + P1 \\ L'_r(p_{\text{before}}, D - 1) + P1 \\ \min_i L'_r(p_{\text{before}}, i) + P2 \end{cases} \quad (2.41)$$

By always adding the minimum path cost of the previous pixel on the scanline we are looking at, we solve equation 2.40 in one dimension. It is to mention that the rolling sum can reach quite high numbers that are unpleasant to handle on the computer; a normalization is implemented by subtracting $\min_D L'_r(p_{\text{before}}, D)$ from all pixel cost values $L'_r(\vec{p}, D)$. The position of the minimum cost function at pixel \vec{p} is unaffected by that normalization.

Summing along at least 8 path directions (crosshair + diagonals) results in disparity maps with reduced error pixel while maintaining clean edges.

2.5.2 Semi - Global Matching for Light fields

Even though Hirschmüller describes Semi - Global Matching (SGM) as a complete algorithm to obtain a disparity map from a stereo image input, we further refer to SGM as the true novelty of his work: the implementation of an approximation to the global solution of the cost function (equation 2.40). Indepent from the method one uses to calculate a disparity map, one needs a cost function defined in disparity space for each pixel to make use of SGM. Similar to the Stereo Matching depth estimation, the structure tensor depth estimation pipeline for Lightfield data sets produces a disparity map and a coherence value at each disparity shift. This implies, that the SGM algorithm can be adapted to improve the results of the structure tensor pipeline. However, there are some significant differences between those two methods:

1. The structure tensor algorithm is tuned to a much smaller disparity range. While in [Hirschmüller \[2005\]](#) Hirschmüller scans a disparity range of 32 pixels , The benchmark data sets for light fields mostly include close-up views of objects, with a disparity range between 2 and 10 pixels. In figure 2.13 one can see the different values that are allocated in memory for each pixel of the image.
2. The subpixel accuracy using the structure tensor is a lot higher than the stereo matching subpixel accuracy. A simple adaption of the algorithm to the structure tensor pipeline would require to give up the best feature that is provided by the ST, its subpixel accuracy.

To handle those problems, we do not throw away the subpixel accuracy: instead we use the float-value disparities to decide whether we penalize a disparity discontinuity

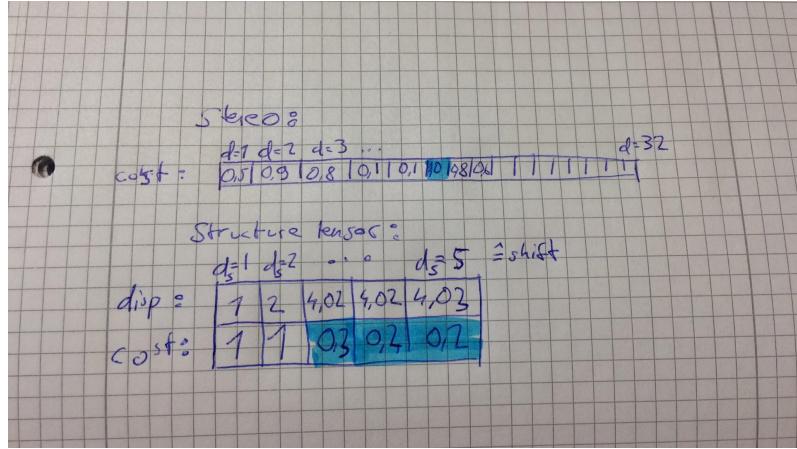


Figure 2.13: One point \vec{p} contains different values (values here: example values): For Stereo matching, the resolution is given by the discrete disparity steps. Each disparity value has a cost value assigned to it. Using the ST, we have a different subpixel accuracy for every disparity shift, while the subpixel accuracy can differ from the shift by up to 1.2

or not. As one can see in figure 2.13, we have to process an additional information, since the exact disparity value is not implicitly given by the index of the allocated cost value (in contrast to the original algorithm). Switching to a continuous space as depicted in figure 2.14 requires a new definition of the error propagation defined in equation 2.40.

In the following we refer to s as the disparity shift in the ST algorithm. Note that we replaced D by d to clarify that the disparity is no longer discrete:

$$E(d) = \sum_{\vec{p}} \left(C(\vec{p}, d_{\vec{p}}) + \sum_{q \in N_p} \begin{cases} P1 \cdot |d_{\vec{p}} - d_{\vec{q}}| & \text{if } |d_{\vec{p}} - d_{\vec{q}}| \leq 1 \\ P2 & \text{if } |d_{\vec{p}} - d_{\vec{q}}| > 1 \end{cases} \right). \quad (2.42)$$

The recursive 1-d form to solve the global constraint on a scanline then changes to:

$$L'_r(\vec{p}, s) = C(\vec{p}, s) + \min_i \begin{cases} L'_r(\vec{p}_{\text{before}}, s_i) + P1 \cdot |d_{\vec{p},s} - d_{\vec{p}_{\text{before}},s_i}| & \text{if } |d_{\vec{p},s} - d_{\vec{p}_{\text{before}},s_i}| \leq 1 \\ L'_r(\vec{p}_{\text{before}}, s_i) + P2 & \text{if } |d_{\vec{p},s} - d_{\vec{p}_{\text{before}},s_i}| > 1 \end{cases} \quad (2.43)$$

The biggest difference lies in the fact that the small factor that is smoothing the image linearly increases with the distance. This change is necessary under the assumption that the disparity space is continuous. In other words we cluster disparity differences between two neighbouring points as either part of one surface ($|d_{\vec{p},s} - d_{\vec{p}_{\text{before}},s_i}| \leq 1$) that gets smoothed by the linearly increasing penalty, or assume a real disparity discontinuity that is penalized regardless of the size of the

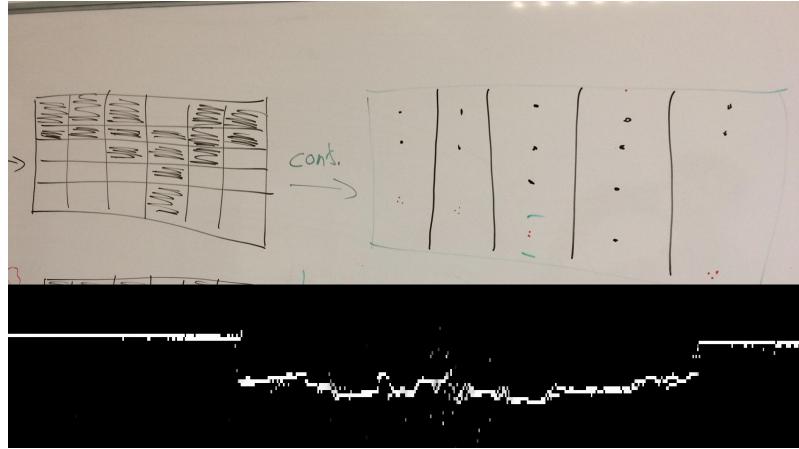


Figure 2.14: In this figure one can see the skizze of one arbitrary scanline of the structure tensor method. Under the Assumption that the ST algorithm recognizes the structure of the EPI perfectly, we either have two or three disparity shifts that have a high coherence (colored white) (a) and result in approximately the same final disparity. This can be seen if we plot the exact disparity values in a continuous space(b). In (c) real data scanline cost is plotted with a resolution of ca. 100 pixels.

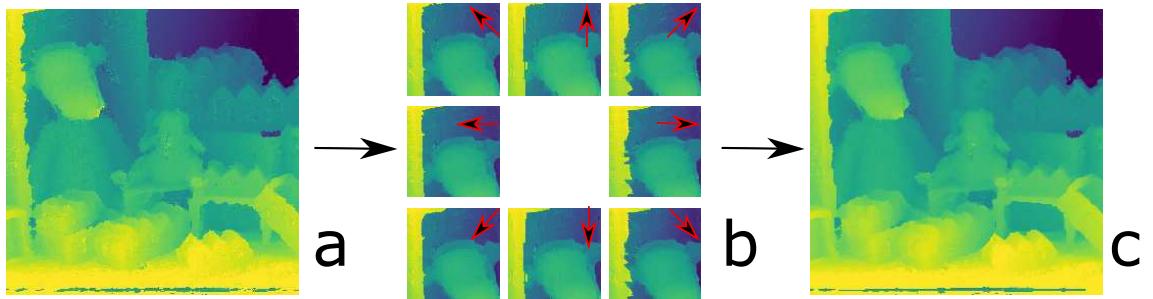


Figure 2.15: a: scene *dino* using the old pipeline,

jump - the second error remains constant as it is in Stereo matching. Note that in our implementation, $P2$ is modified by

$$P2' = \frac{P2}{\sqrt{(Im_b^2 + Im_r^2 + Im_g^2)}}, \quad (2.44)$$

with Im being the center view of the lightfield and $Im_{b,g,r}$ being the 3 color channels. If the color intensity changes, the penalty for a disparity discontinuity is lowered.

2.5.3 Occlusion awareness in SGM for light Fields

If we take a close look at figure 2.14, one can see that at least at some discontinuities the ST pipeline manages to calculate the depth of the background structure near

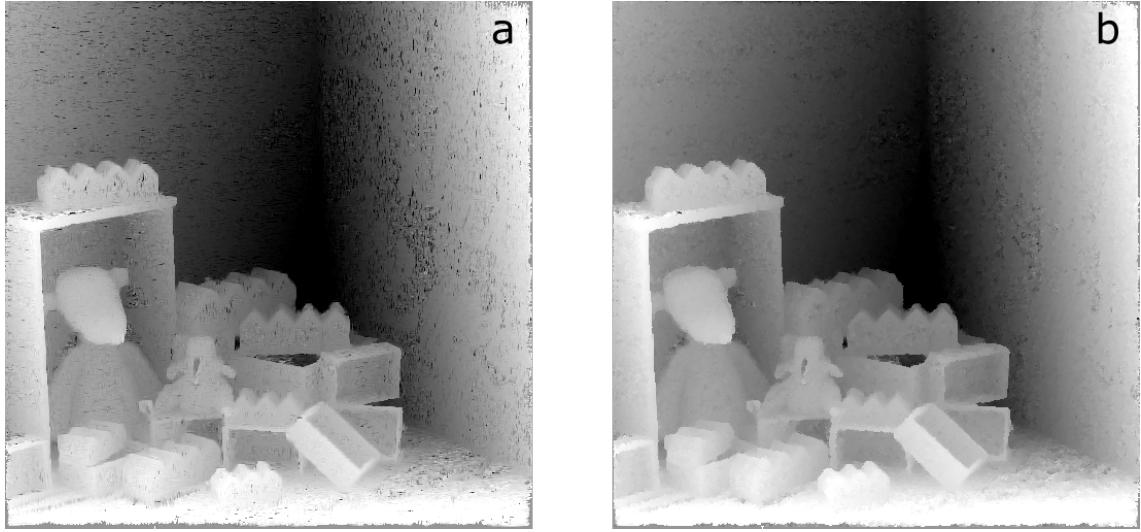


Figure 2.16: a: Errorneous disparity map before smoothing b: Same disparity map after smoothing

boundaries with good coherence, but the foreground structure is overlapping and quantitatively measured with higher coherence. Once we know that at least at some edges an improvement can be made by adapting the evaluation function in a sense that the highest coherence does not necessarily measure the right depth, we realize that SGM is doing the job already. The simple heuristic approach is to change the global minimization function 2.42 such that a positive disparity jump is less punished than a negative one. In fact, the function changes to

$$E(d) = \sum_{\vec{p}} \left(C(\vec{p}, d_{\vec{p}}) + \sum_{q \in N_p} \begin{cases} P1 \cdot |d_{\vec{p}} - d_{\vec{q}}| & \text{if } |d_{\vec{p}} - d_{\vec{q}}| \leq 1 \\ P2 & \text{if } d_{\vec{p}} - d_{\vec{q}} > 1 \\ P3 & \text{if } d_{\vec{p}} - d_{\vec{q}} < -1 \end{cases} \right). \quad (2.45)$$

2.5.4 SGM as postprocessing smoothing

In general Semi-Global Matching refers to minimizing an energy function in one dimension for different paths in the given space. In case of disparity mapping, the disparity candidate values and its initial cost values have to be given. To use SGM as an efficient tool for smoothing, these disparity candidate values have to be chosen wisely. In scope of this work we propose to choose for each pixel 5 neighbouring pixels that are randomly chosen from the close neighbourhood after constructing the disparity map with any algorithm. In this sense the SGM is used as a pure postprocessing step that can be used to improve any depth map. The effects of smoothing the disparity map while maintaining edges is depicted in figure 2.16 as an example.

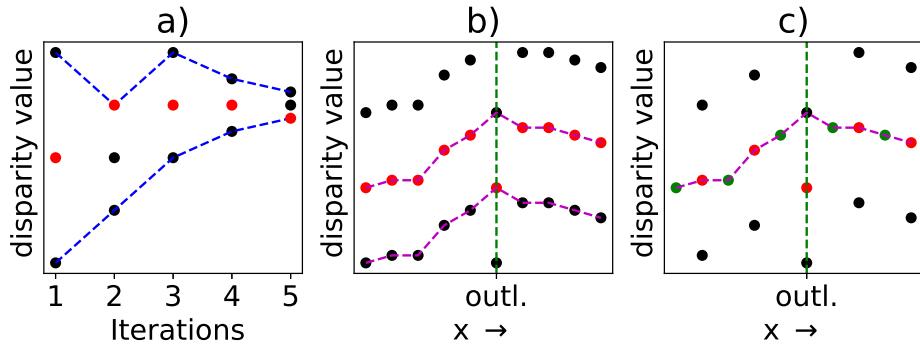


Figure 2.17: a) One exemplary pixel value in function of iteration steps. At each iteration, 3 candidate values for each pixel are chosen, the one with the lowest score is marked red. The candidate value window is marked with blue lines. b) One exemplary scanline through the depth map. Red marks the input values, the black points are the added candidate values. The green line marks the position of an outlyer. The magenta-colored lines show the pathes with the lowest cost. c) same as the diagramm b) with the difference of using a chess pattern initialisation.

2.5.5 Iterative postprocessing

To reach the best result possible the SGM- postprocessing is applied multiple times. This iterative process is inspired by ?. Instead of choosing random pixel from the close neighbourhood, the candidate values are found by adding/substracting a Δ from the original pixel depth value d_i , such that the candidate values are given by the triple $(d_i; d_i - \Delta; d_i + \Delta)$. The candidate with lowest cost is chosen and new candidates are formed for the next iteration. To reach convergence, this Δ shrinks by a factor of two each iteration as long as the chosen candidate was the central one. Otherwise Δ stays the same size. In figure 2.17 a) an exemplary disparity value is shown in dependence of the iteration step. The candidate window gets smaller each step, the stopping criterium is given by the overall iteration number. However, setting the same Δ at each pixel leads to problems along any SGM scanline, as sketched in figure 2.17 b). Since all candidates are initialized with the same Δ one finds 3 identical surfaces through the scanline that are shifted by Δ . Since they are almost identical, the cost of the disparity values lying either on one or the other surface is close to being identical. Hence the algorithm produces large errors along the scanline when choosing the wrong 'layer'. To overcome this problem one could apply a chess-pattern-like initialisation of the pixels. When proposing candidates to the SGM algorithm, every second pixel in the 2D-depth map is held constant to one value while the other pixels obtain 3 candidate values. After iterating, the second half of the pixels will be held constant. As depicted in figure 2.17 c) only one surface line will have a low cost value. However doing this iteration doubles the amount of time the code consumes.

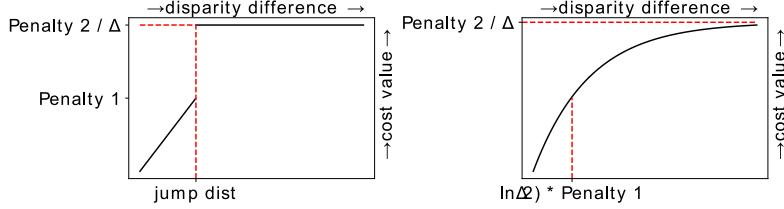


Figure 2.18: Sketch of different proposed Penalty functions. Left: Function as described in equation 2.43. Right function as described in 2.48.

2.5.6 A new penalty function

The iterative pipeline does not assume that any of its candidates at one pixel position have to be close to the ground truth disparity, e.g. having an outlier d_i at pixel $i(x, y)$ does not necessarily mean that either $d_i - \Delta$ or $d_i + \Delta$ are close to the ground truth. With punishing those candidates as described in equation 2.42, all 3 candidates would obtain the same cost function value. However one wants the candidate that is closest to its neighbours to be the one with the lowest cost. Thus the cost function should be monotonically increasing in function of the distance to the previous disparity value d_{i-1} . A simple linearly increasing cost function would lead to overly smoothed edges. The proposed global cost function that is to be minimized then looks like

$$E(d) = \sum_{\vec{p}} \left(C(\vec{p}, D_{\vec{p}}) + \sum_{q \in N_p} P2 \cdot \left(1 - \exp \left(\frac{-|d_{\vec{p}} - d_{\vec{q}}|}{P1} \right) \right) \right). \quad (2.46)$$

Note that the initial cost can be neglected since we assume the same initial cost for all candidates such that we obtain the form

$$E(d) = \sum_{\vec{p}} \sum_{q \in N_p} P2 \cdot \left(1 - \exp \left(\frac{-|d_{\vec{p}} - d_{\vec{q}}|}{P1} \right) \right) \quad (2.47)$$

with the 1D recursive form

$$L'_r(\vec{p}, s) = \min_i \left(L'_r(\vec{p}_{\text{before}}, s_i) + P2 \cdot \left(1 - \exp \left(\frac{-|d_{\vec{p}, s} - d_{\vec{p}_{\text{before}}, s_i}|}{P1} \right) \right) \right). \quad (2.48)$$

The equation 2.48 is plotted in figure 2.18 on the right side. In comparison to the old penalty function it has one less parameter, it is continuous and it always favours smaller disparity differences in order to get iterative processing to work. The error $P2$ is modified the same way it is like before, see equation 2.44. Large image gradients are more likely to lie on a disparity discontinuity, therefor the error is reduced. In figure 2.19 the effect of different penalty functions is depicted. Figure 2.19 a) shows the result of the scene *cotton* after smoothing with the exponential penalty function 2.48, while figure 2.19 b) shows the disparity map after iterative

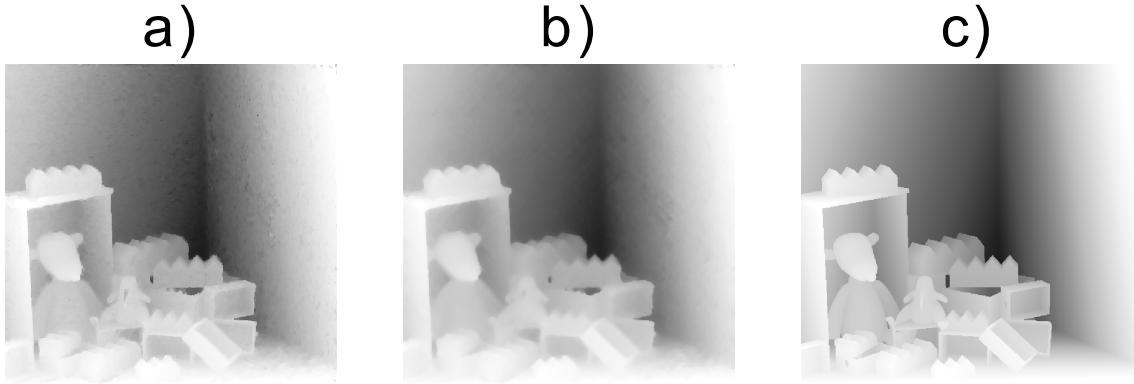


Figure 2.19: a) Depth map after Iterative SGM smoothing using an exponential penalty function. b) Instead a linear penalty function is used. c) Ground Truth

SGM processing with a linear penalty function. One sees that the linear function leads to smoother surfaces, however the smoothness constraint also oversmoothes the edges in the image. In a) the edges are maintained sharp. The figure 2.19 c) shows the ground truth for comparison.

2.5.7 SemiGlobal Matching implementation

The implementation for using SemiGlobal matching in our work is oriented on the opensource implementation published by [Moratto](#). This code however uses only 8 instead of 16 paths along the image, further it struggles with time efficiency as it is about 20 times slower than the implementation of [Hirschmüller \[2005\]](#) as published in his paper. Our implementation thus will also be slow. Further, [Ernst and Hirschmüller \[2008\]](#) implement the SGM algorithm on a GPU leading to an enormous increase of efficiency. Since all paths can be calculated on parallel, a GPU implementation can be implemented in a straight forward manner. However GPU programming exceeds the boundaries of this work and will be part of future research and development. Instead, we focus on the feasibility of implementing SemiGlobal Matching for lightfields.

2.6 Metrics for the evaluation of depth maps

In the following technical Dataset details and general evaluation measures are presented. The Datasets are provided by [Honauer et al. \[2016\]](#).

2.6.1 dense and sparse maps

The synthetic Lightfield datasets are designed with Blender ([Community \[2014\]](#)) and provide photos from the scene in a crosshair pattern, similar to the camera

configuration in figure 2.4. This allows to calculate a disparity map from only one point of view due to the restricted baseline of the cameras. In contrast to Stereo camera setups Light Field imaging enables a densely sampled reconstruction of the depth at each pixel. However a full pointcloud of a scanned object which is ideal for object detection applications will not be reconstructed due to occlusions. To evaluate the quality of the depth map, one could either try to optimize the overall error on the whole depth map (MSE) or count the number of pixels that are close to the Ground Truth with given tolerance Level (BadPixel Score). From thereon one either tries to provide a sparse depth map masking out all the pixel where the confidence is low or the disparity map is provided on the whole pixel space. While sparse disparity maps are the choice for exact measurements in industrial applications, the second method facilitates comparisons of different methods. For this reason the thesis mainly focusses on the Metric score on the dense disparity estimation maps using different algorithms. Only for the application on real data one has to face large regions with low confidence, thus masking becomes relevant again.

2.6.2 mean squared error

The Mean squared error, typically multiplied by a factor of 100, is defined as

$$\text{MSE} = \frac{\sum_x (d(x) - gt(x))^2}{\text{Nr of Pixels}} \cdot 100 \quad (2.49)$$

and describes the overall deviation from ground truth. $d(x)$ is the measured disparity value at pixel x , $gt(x)$ is the ground truth value at pixel x .

2.6.3 bad pixel score

The Number of pixels at which the disparity value $d(x)$ deviates more than the tolerance t from the ground truth $gt(x)$ is given by

$$\text{BadPixel}(t) = \frac{\{|d(x) - gt(x)| > t\}}{\text{Nr of Pixels}}. \quad (2.50)$$

2.6.4 bumpiness

The bumpyness score is quantified by the Frobenius norm of the hessian matrix H_f of a function f which is defined here as $f = d - gt$. Smooth surfaces lead to a better bumpyness score regardless of possible missorientations and offsets:

$$\text{Bumpiness} = \frac{\min(0.05, \|H_f(x)\|_F)}{\text{Nr of Pixels}} \cdot 100 \quad (2.51)$$

2.6.5 discontinuity and planar scores

In some scenes masks are applied to evaluate the score at different spots in the scene. This way it can be focussed quantitatively on challenges (planar surfaces or discontinuities) that are set by the scene. Given a mask M the BadPixel Score is then given as

$$\text{BadPixel}(t) = \frac{\{x \in M : |d(x) - gt(x)| > t\}}{\text{Nr of Pixels } x \in M}. \quad (2.52)$$

2.6.6 Mean Relative Error

In some cases we refer to the Mean Relative Error(MRE) instead of the Mean squared Error. The difference lies in the normalisation:

$$MRE = \frac{\sum_x (d(x) - gt(x))^2}{(\text{max. disp} - \text{min. disp}) * \text{Nr of Pixels}} \quad (2.53)$$

3 Evaluation

3.1 Depth from focus

The depth measure using epipolar plane analysis requires iterative calculation of the structure tensor for each EPI at each disparity. A way to overcome this is to generate a preestimate of the depth before actually calculating the correct depth. This could also help to prevent possible errors due to periodic scene characteristics which can lead to mismatch errors when calculating the structure tensor. Therefore the depth pre-estimate should fulfil the following criteria:

1. It should be *consistent*, meaning that the number of pixels with low confidence should be the lowest possible.
2. It should result in a *fast* measure, ideally faster than it would take to do the full iterative structure tensor algorithm.
3. It does not have to be subpixel accurate, since it only serves as a pre-estimate.

The methods that are tested are described in section 2.4. We test four different ways to obtain a depth map using depth from focus:

Photo consistency This measure takes advantage of the fact that the difference between the refocussed two-dimensional image and the center view is close to zero when refocussed to the correct depth. Response value:

$$D'(x, y) = \frac{1}{|W_D|} \sum_{x', y' \in W_D} |\bar{L}(x', y') - P(x', y')|, \quad (3.1)$$

Angular correspondence In contrast to the *Photo consistency* - measure, it first calculates the absolute difference between each camera array view and the center view followed by the summation of those deviations. The response value is given as in equation (2.39)

$$D'(x, y) = \frac{1}{N_{u,v}} \sum_u \sum_v |L'(u, v, x, y) - P(x, y)| \quad (3.2)$$

First derivative The first derivative is calculated for contrast measure by applying the sobel filter onto the refocussed image I :

$$G_x = \begin{bmatrix} -1 & 0 & 1 \\ -2 & 0 & 2 \\ -1 & 0 & 1 \end{bmatrix} \cdot I \quad G_y = \begin{bmatrix} -1 & -2 & -1 \\ 0 & 0 & 0 \\ 1 & 2 & 1 \end{bmatrix} \cdot I \quad (3.3)$$

The directional gradients are simply added up to the response value

$$D'(x, y) = |G_x(x, y)| + |G_y(x, y)| \quad (3.4)$$

Laplace Here we calculate the second derivative laplacian by applying the sobel operator twice:

$$D'(x, y) = \text{Laplace}(I)(x, y) = \frac{\partial^2 I}{\partial x^2}(x, y) + \frac{\partial^2 I}{\partial y^2}(x, y) \quad (3.5)$$

In the following we are going to compare the method qualitatively and quantitatively. In figure 3.1 one can see the pixel response value refocussed at different disparities for all the methods at example points in the testscene „complextestscene“. Points close to edges as well as points on clear surface with less structure on it are chosen. One can see that the pixel response of the Angular correspondence and the Photoconsistency method for those points show a more consistent behaviour, meaning that only one clear maximum can be seen. The derivative method as well as the Laplace method both seem to have trouble especially on pixel coordinates close to edges. This can be seen most clearly at the edge of the melon, where The first derivative shows 2 Maxima, while the Angular correspondence and Photo consistency measure find a clean maximum indicating at which depth the point can be found. However, on surfaces with less structure as for example on the potato, the Photo consistency measure struggles to find a clear maximum – still it has one at the right position. Having a look at the actual disparity maps produced by the different techniques (figure 3.2) we can already capture that the angular correspondence and photo consistency method produce the more consistent, smooth disparity maps. A quantitative evaluation confirms this impression: The diagram in figure 3.3 indicates that for all scenes either the Photo consistency or the Angular Correspondence method obtain the lowest mean relative error.

3.1.1 Using depth-from-refocus as a preestimate for the ST

The long-term aim of this work is to make the structure tensor pipeline more robust. We achieve this by using the depth-from-refocus method as a preestimate, therefore reducing the disparity window at each pixel. The Light field is now refocussed by integer disparity steps, resulting in disparity maps that obtain equidistant layers. All disparity maps with integer disparity stepsize are compared to the ground truth in figure 3.4. Again the Angular Correspondence and Photo consistency measures result in the lowest MRE (figure 3.5), though the Angular Correspondence cue is about 1.5 times more time-consuming than the photo consistency measure under the same conditions (figure 3.6). Further testing we stick to the Photo Consistency measure, as it produces the best results in terms of speed and precision.

The next step is to use the preestimate at each pixel to skip the iterative refocussing in the structure tensor pipeline. However, since not all points on an EPI are of

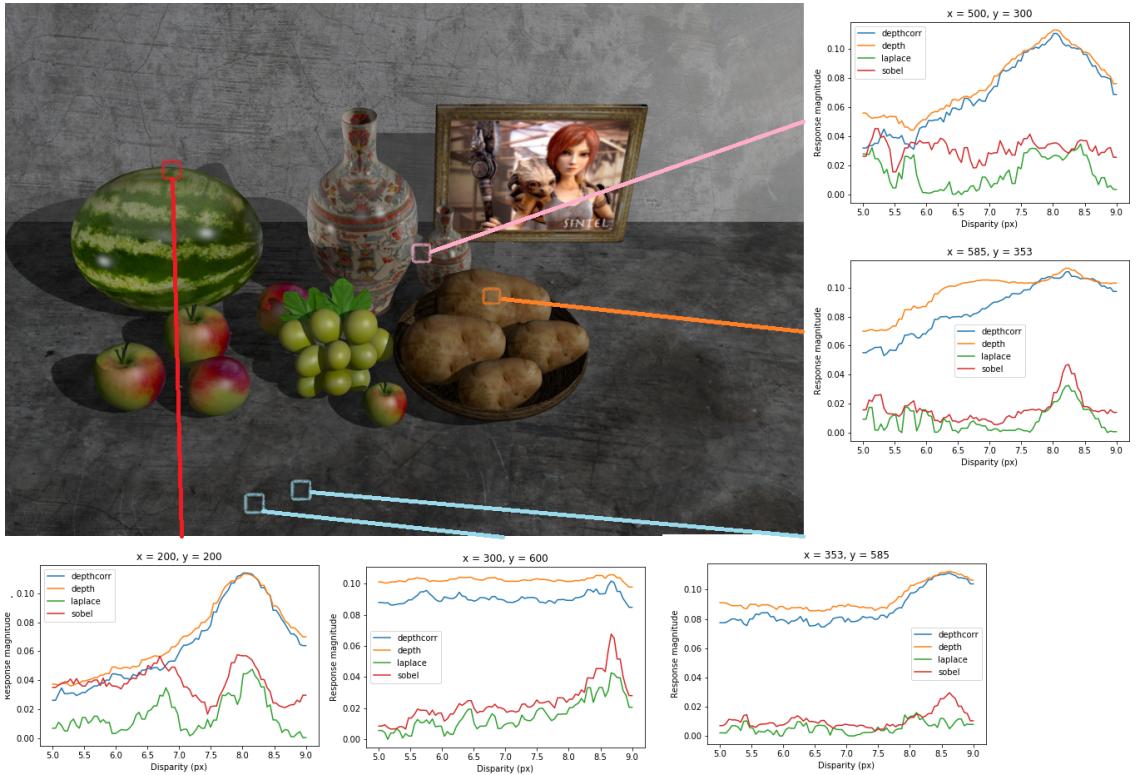


Figure 3.1: At different pixel positions we take a look on how the Pixel response value behaves for the four different depth-from-focus techniques: Photo consistency (orange), Angular correspondence (blue), First derivative (red) and The Laplace method (green) A high value means high confidence (low cost).

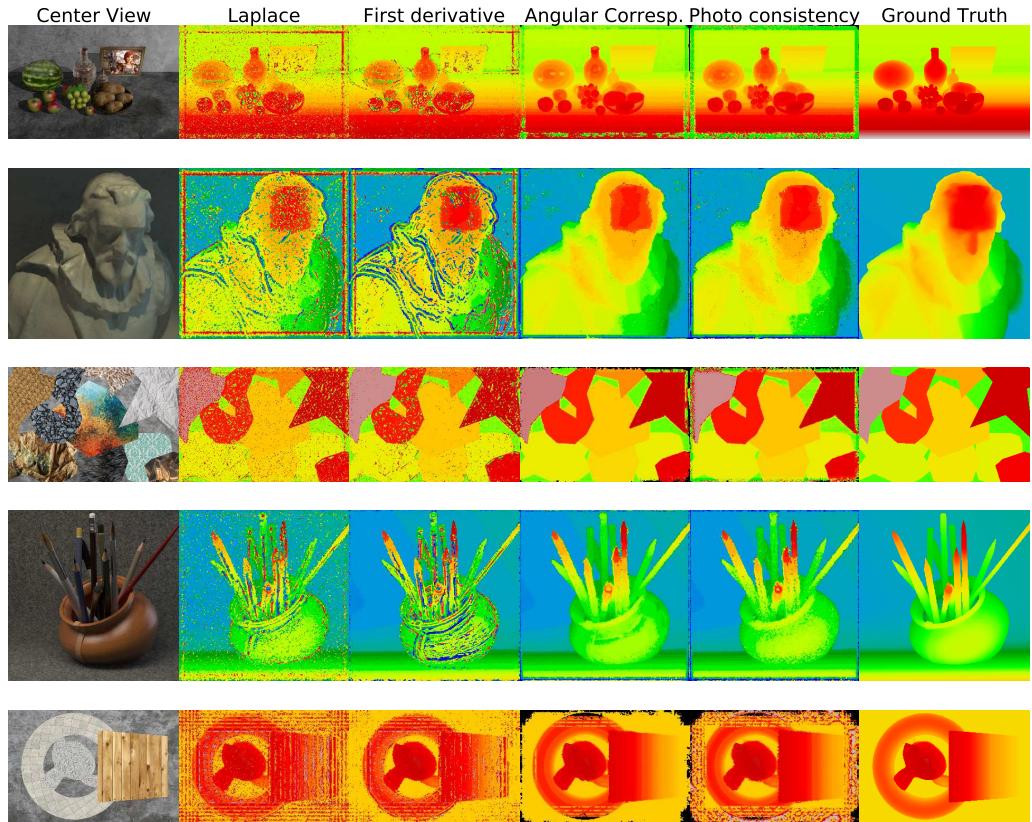


Figure 3.2: The depth maps of the scenes (from up to down) *complextestscene*, *cotton*, *occlusiontestscene*, *pens*, *testscene*, *tiltplane* is depicted for different depth-from-focus techniques. The Disparity stepsize is $1/20$.

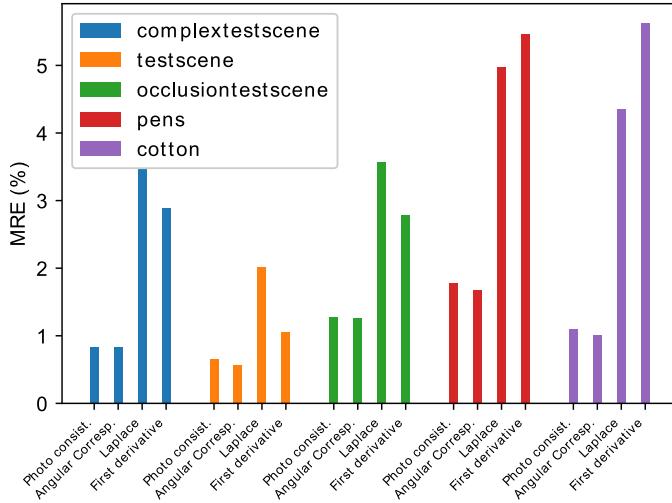


Figure 3.3: The mean squared error is depicted for depth-from-focus techniques. The stepsize is 1/20.

the same depth, the EPI is segmented in parts of the same pre-estimation depth. Those parts are refocussed separately to the estimated depth, results are illustrated in figure 3.7. Note that the edge of the depth images have been removed when calculating the MRE. Unfortunately the Photo Consistency pre-estimate pipeline yields worse results than the Structure Tensor itself. The reason is that artifacts from the preestimate maintain after the structure tensor pipeline has been applied to the EPI. Additionally one has to mention that only those pixels in the depth map that have a high coherence measure given by the ST pipeline are taken into account when calculating the errors. Hence a bias towards the Structure tensor pipeline in terms of optimal error results is expected.

Also the preestimation is even more time-consuming than the structure tensor itself. Because of the splitting of the EPI, the runtime is scene dependent. The time difference between the scene is therefore enormous, the results can be found in table 3.1.1. If one jumps the preestimation step and takes the Ground truth data as a „pre-estimate“, the pipeline is significantly faster than iteratively refocussing on each depth. The pre-estimate itself neither takes too long (about 2 seconds), but the EPI gets segmented into smaller parts when using the pre-estimate. A global smoothing scheme on the preestimate can possibly improve the runtime by minimizing the amount of disparity jumps, however this would result in even more artefact effects. We leave that aspect for future work.

In summary the obtained results using the Photo consistency measure as a pre-estimate for the structure tensor do not improve the obtained results regarding the Mean squared error and runtime-wise.

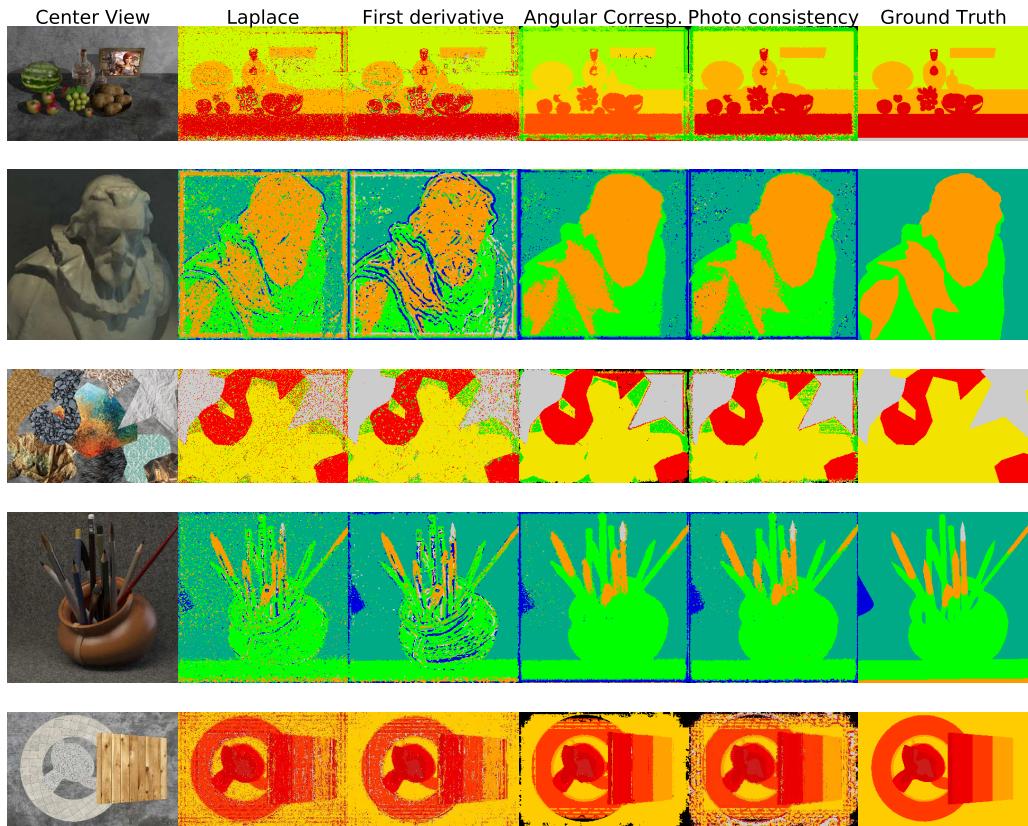


Figure 3.4: The depth maps of the scenes (from up to down) *complextestscene*, *cotton*, *occlusiontestscene*, *pens*, *testscene* is depicted for different depth-from-focus techniques. On the right the ground truth is depicted. The disparity stepsize is 1.

Scene	Structuretensor Pipeline	Preestimate+ ST	Use GT as preestimate
complextestscene	12.89 s	13.62 s	8.83 s
pens	5.1 s	5.2 s	4.7 s
cotton	5.09 s	6.78 s	3.97 s
tiltplane	22.25 s	17.46	9.3 s
occlusiontestscene	10.9 s	12.1 s	6.7 s

Table 3.1: Time for different pipelines

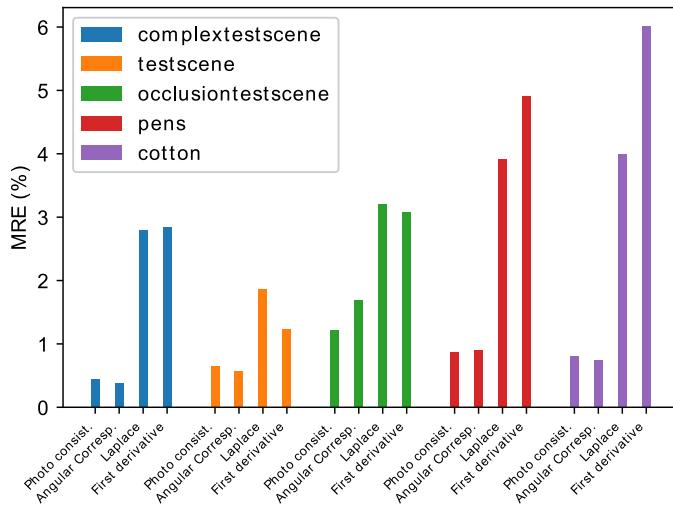


Figure 3.5: The mean squared error is depicted for depth-from-focus techniques. The stepsize is 1.

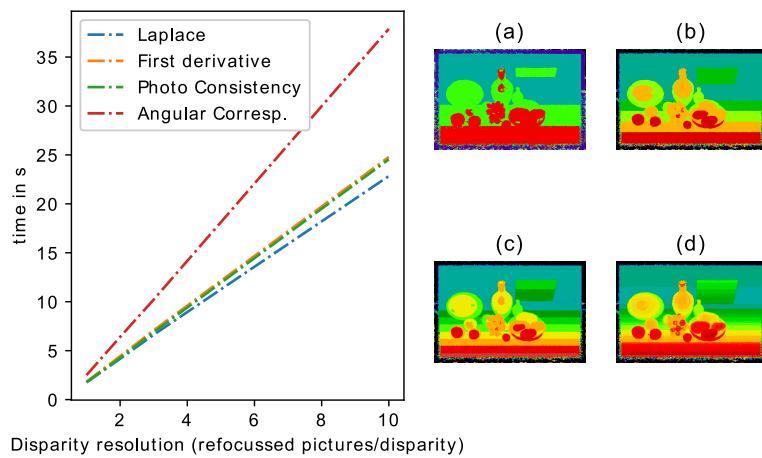


Figure 3.6: On the left the speed of different depth-from-focus techniques is depicted in function of the resolution chosen. (a)-(d) show the disparity map of the scene *complextestscene* for the resolutions 1, $\frac{1}{2}$, $\frac{1}{3}$, $\frac{1}{10}$ respectively.

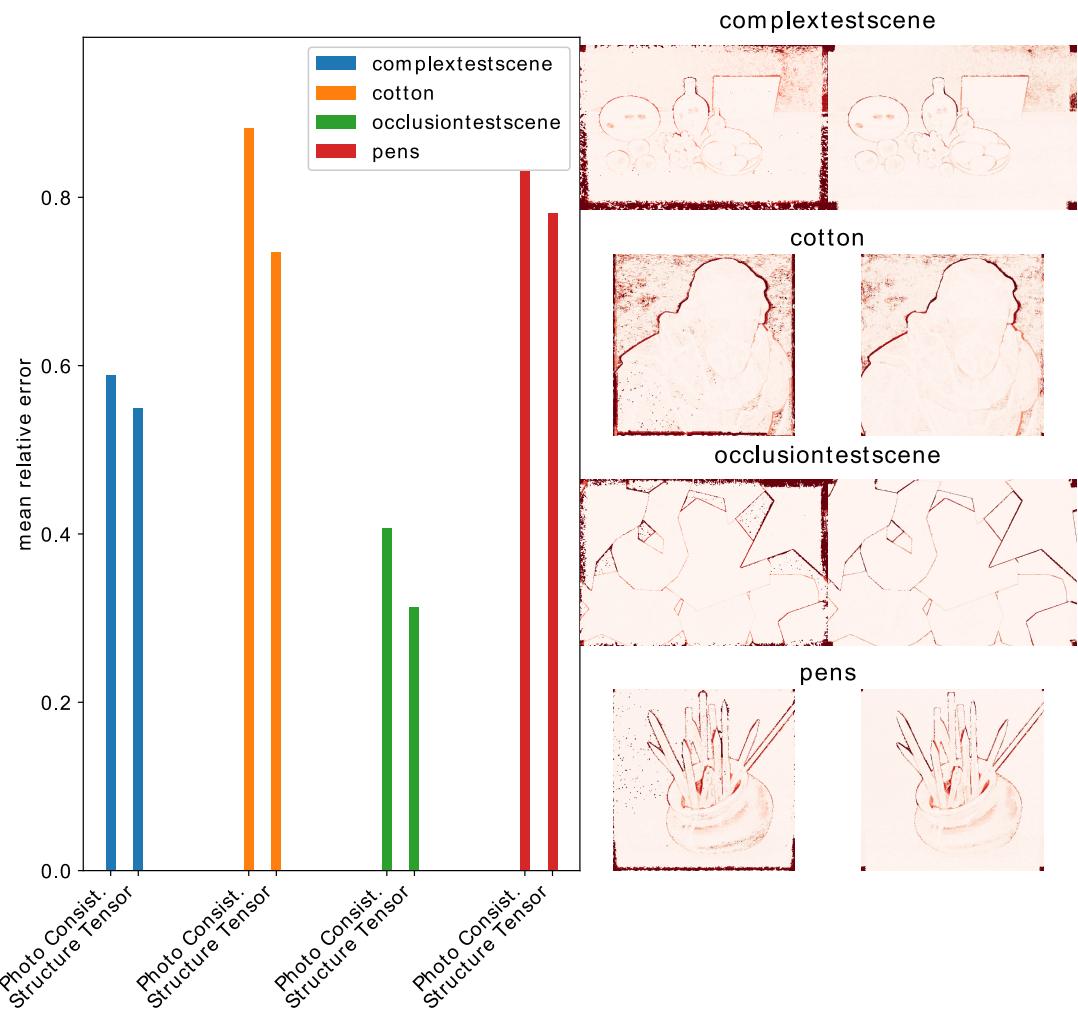


Figure 3.7: Comparison between the Structure Tensor Pipeline with and without a Photo Consistency pre-estimate. On the left the MRE is depicted for different scenes. The right images show the absolute difference of the depth map tho the ground truth, while the left column represents the results with the photo consistency preestimate.

3.2 Modification on the Structure Tensor

As depth from focus techniques didn't work out well for us, we examine different approaches by directly modifying the Structure Tensor Pipeline, with the aim on solving the occlusion problem while maintaining a dense disparity map.

In the following section we concentrate on benchmark data from the dataset of Honauer et al. [2016] as it not only provides a big data set, it also comes with a toolkit for evaluating scenes with different metrics. Whenever it is referred to a metric score, lower score means always better score.

3.2.1 Sandclock kernel

In figure 3.8 the results we obtain from using a sandclock-shaped kernel instead of a normal gaussian kernel are shown. Even though one could expect the custom-sized kernel to perform better at edges, the opposite is the case; the original gaussian kernel outperforms the sandclock kernel in all scenes. The motivation to define the kernel in form of a sandclock was to localize structure clearly to the pixel it belongs to. However it becomes clear that the sandclock kernel cannot handle the occlusion problem explained in section 2.2.2. On the surface regions less smoothing between neighbouring pixels lead to a higher noise resulting in a higher mean squared error.

3.2.2 Bilateralfilter

This section illustrates the effects of using a bilateral filter in the Structure Tensor pipeline instead of a Gaussian Filter. Figure 3.10 shows the Mean Squared Error as well as the discontinuity score when using a bilateral filter with the original EPI as a guide. The Scores are normed to the score of the Structure tensor pipeline. For large σ_{Color} both converge to 1, because in this limit the weighting based on the color space distance (see equation 2.11) is a constant function. Thus one again obtains the same result as with a normal gaussian filter. Note that the σ_{Space} remains the same as in the case of a normal filtering ($\sigma_{\text{Space}} = 7 \text{ pix}$). Unfortunately the MSE metric score and the discontinuity score is worse than with using the gaussian kernel for the complete parameter space. Especially surfaces with a lot of texture suffer from splitting the weights by their color values. This becomes clear as the scene *sideboard* with the fine patterned wallpaper reaches the worst results. (PICTURE). Any pixel that doesn't encounter enough neighboring pixels with an equal color results in a Stucture tensor that only takes into account the very local gradient in the EPI. Since the aim was to especially handle disparity transitions by not mixing up two regions in the EPI with different colors, we take a look at the results for the discontinuity metric defined by Honauer et al. [2016]. Even if one can see a slight improvement at some transitions looking at the disparity maps, the discontinuity score is still unaffected since the edge fattening is still visible. Since the results do not satisfy the expectations and the guided bilateral filter also takes about 5 times

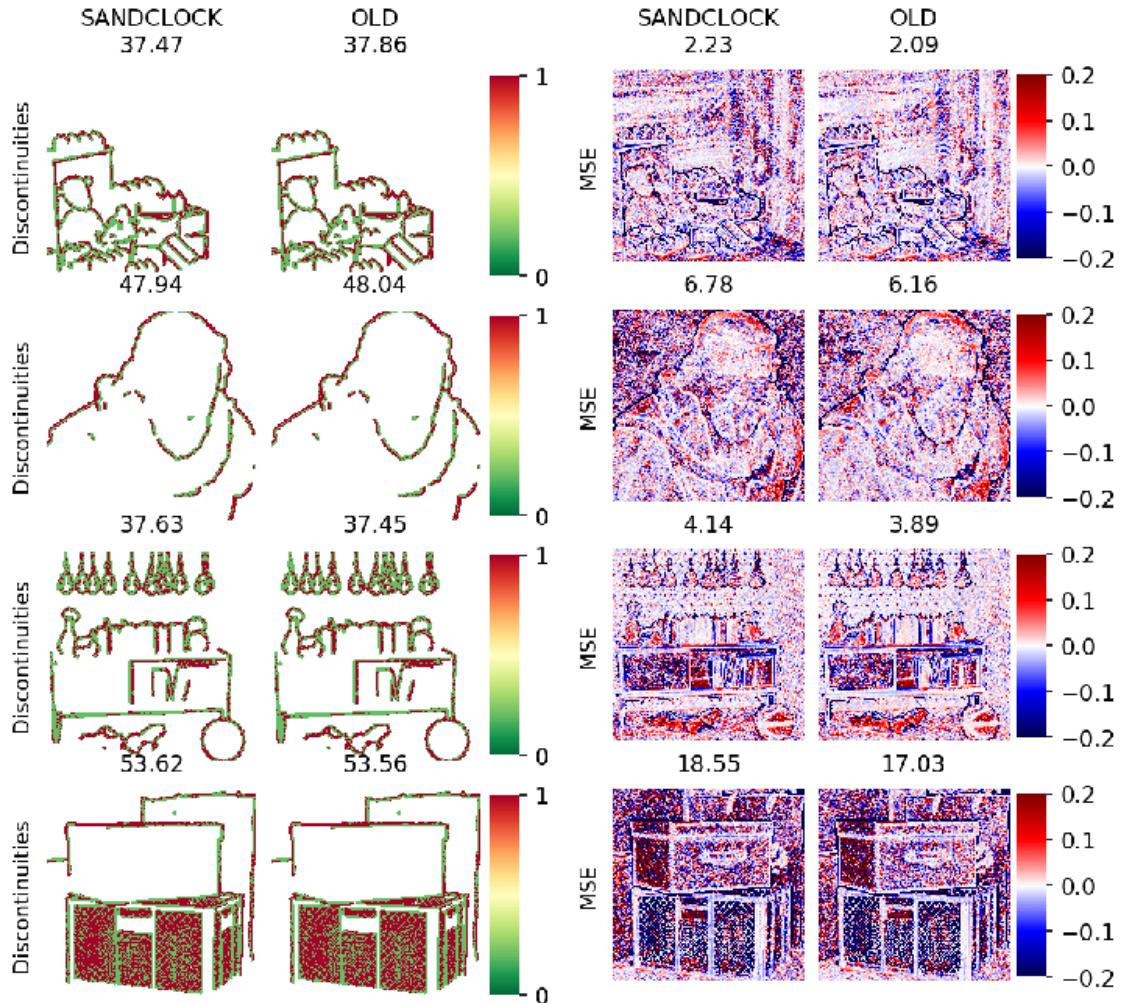


Figure 3.8: Comparison between the normal structure tensor and the sandclock-like kernel for the scenes *dino*, *cotton*, *sideboard*, *boxes*. The left side compares the mean squared errors in the disparity map, the right side shows the error where discontinuities in the disparity map are masked explicitly.

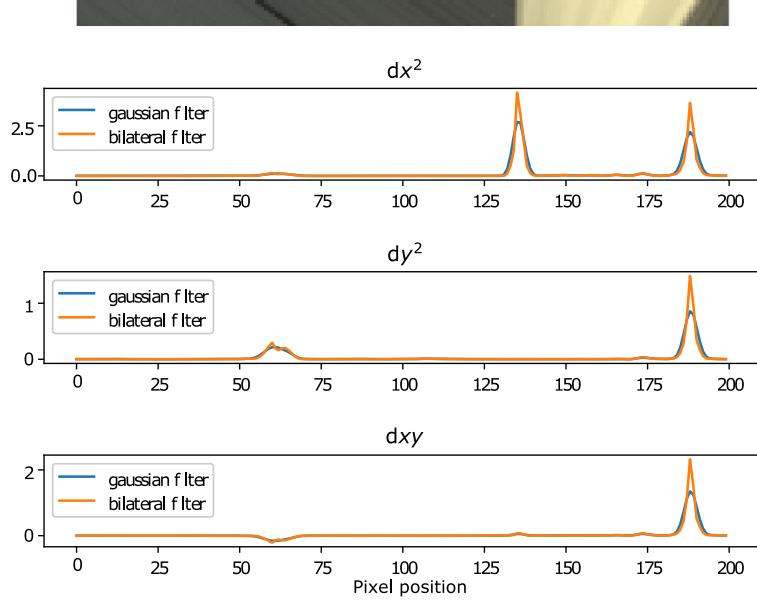


Figure 3.9: Bilateral filtering of an epi

longer then the normal structure tensor, we try an alternative approach. Instead of using the original EPI as a guide, one could directly take the structure tensor component vector

$$\vec{S} = \begin{pmatrix} \frac{\partial g^2}{\partial x \partial x} \\ \frac{\partial g^2}{\partial s \partial s} \\ \frac{\partial g^2}{\partial x \partial s} \end{pmatrix} \quad (3.6)$$

as the guide for the bilateral filtering of itself. The idea behind it is closely related to the EPI segmentation as described in section 3.2.4. Big gradients in the EPI are more consistently referring to the same transition. By separating small and big gradients, the influence of the structure in front is less significant than it is without using the bilateral filtering. As one can see in figure 3.11, using the ST components as a guide can actually result in a better MSE. However, for each scene the score is dependent on the parameter σ that is chosen. Also, the discontinuity score does not improve even though in figure 3.12 a clear improvement can be seen at occlusion borders. Actually the edge fattening is by far bigger then the masks used to determine the discontinuity score in figure 3.11. In figure 3.12 The depth maps zoomed at some disparity transitions in the given scenes *boxes*, *cotton*, *dino* and *sideboard* are shown, calculated with the gaussian and bilateral filter. One can observe that the bilateral filter (with the Structure tensor component as a guide) results in sharper edges that are significantly less „fattened“ than the gaussian equivalent. This becomes clear when looking at the head of the teddy bear in the scene *dino*. Even though the front object still occupies more space then it should (edge fattening), the edge has become smaller. This effect cannot be seen when using a bilateral filter with the

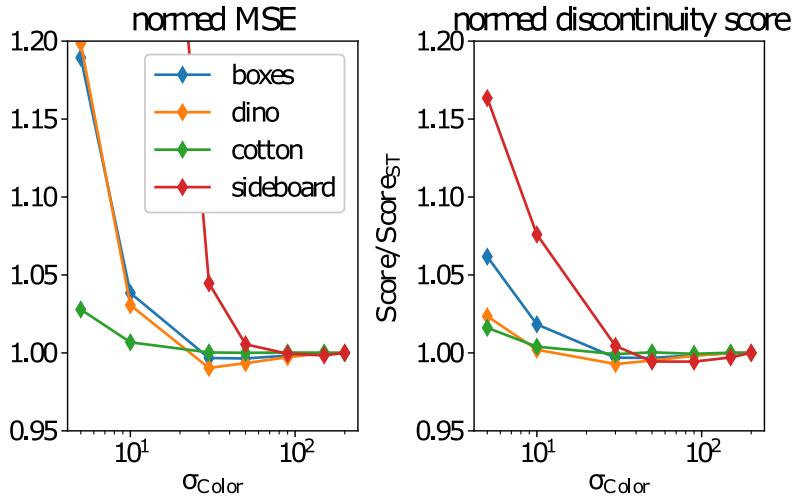


Figure 3.10: Bilateral filtering with original EPI as the guide: On the left the Mean Squared error for different scenes in function of the σ_{Color} of the bilateral filtering is shown, devided by the MSE with using a gaussian filter. On the right the Discontinuity score based on Honauer et al. [2016] is shown, also normed to the Score using the standard gaussian filter.

EPI as a guide however.

Additionally it becomes clear when looking at the scene *boxes* in figure 3.12 that the depth estimation using the bilateral filter with the structure tensor component as a guide is less affected by the by the color differences in the image. When using the gaussian ST, the error along the edge is dependent on the background structure as the transition to the lettering on the bag results in less error then above. The bilateralfilter however does produce a consistent error that may be better to handle for postprocessing.

3.2.3 Thresholding Gradients

When the gradient norm is thresholded as explained in section 2.3.3, the main aim is to change the weights of the gradients contributing to the structure tensor such that all gradients contribute the same. The smallest gradients however underly noise effects, therefor they are not upscaled. A threshold is set at which bigger gradient norms are normed to that threshold. As one can see in figure 3.13, for big threshold values the result is equal to the old ST pipeline results as all gradients in the EPI are below the threshold, thus they are unaffected. Setting an extremely low threshold has the same effect as normalizing all gradients to the same strength. Especially on planar surfaces which typically have less texture and therefor low gradient strengths in the EPI, the error score is worse when normalizing the gradient, as one can see in figure 3.13 in the right plot. However, the MSE is mostly affected by the edge fattening error as described in section 2.2.2. Therefor simple normalizing already

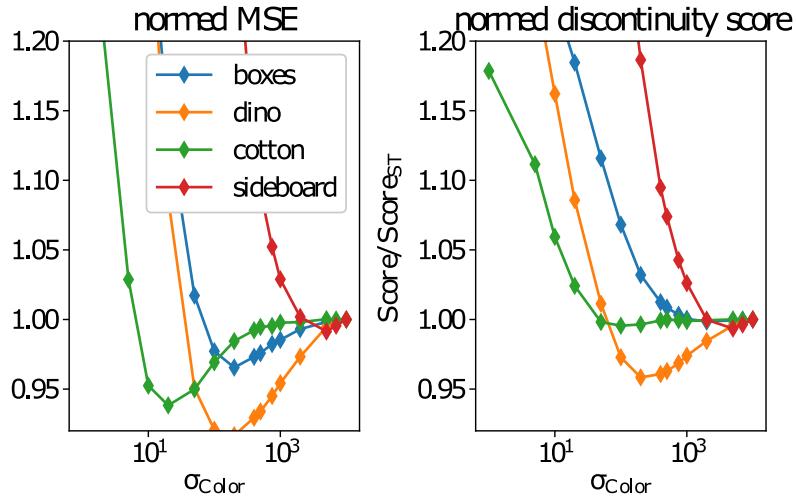


Figure 3.11: Bilateral filtering with Structure Tensor Components themselves as the guide: On the left the Mean Squared error for different scenes in function of the σ_{Color} of the bilateral filtering is shown, devided by the MSE with using a gaussian filter. On the right the Discontinuity score based on Honauer et al. [2016] is shown, also normed to the Score using the standard gaussian filter.

improves the MSE, since the gradient strength of the transition weights less into the structure tensor. The normed discontinuity score in the middle of figure 3.13 is counter-intuitive since it is even worse for low threshold values. The reason is that the mask used for calculating the discontinuity score by Honauer et al. [2016] is smaller than the edge fattening even after normalizing. Thus the discontinuity score cannot represent a better discontinuity behaviour if the edges do not lie in the mask.

Since the scene *boxes* is the noisiest one, the MSE is affected least by improving the disparity transitions. The MSE in the scene *cotton* on the other hand is mostly affected by the edge fattening error, thus its MSE is improved most by the Thresholding of the Gradients. All scenes show an improved MSE for the complete parameter space. Note that the threshold is plotted in a logarithmic scale. In figure 3.14 the thresholded gradient ST result is compared to the bilateral and gaussian ST result. It is obvious that with a gaussian kernel, transitions are fattened and erroneous. It seems that the bilateral filter at least for the upper 3 scenes performs best, while the thresholded ST still produces the same errors as the gaussian ST, as one can see in the *boxes*-scene (upper row). Nevertheless the additional noise (see scene *sideboard*, last row) produced by the bilateral filtering finally results in a worse MSE score.

Another advantage of using the Thresholded Gradient ST becomes clear in figure 3.15, where the gaussian outer kernelsize is varied for the old ST and for the Thresholded Gradient ST. Obviously the discontinuity score becomes worse for bigger gradients and the error score on planes improves due to bigger kernels being less

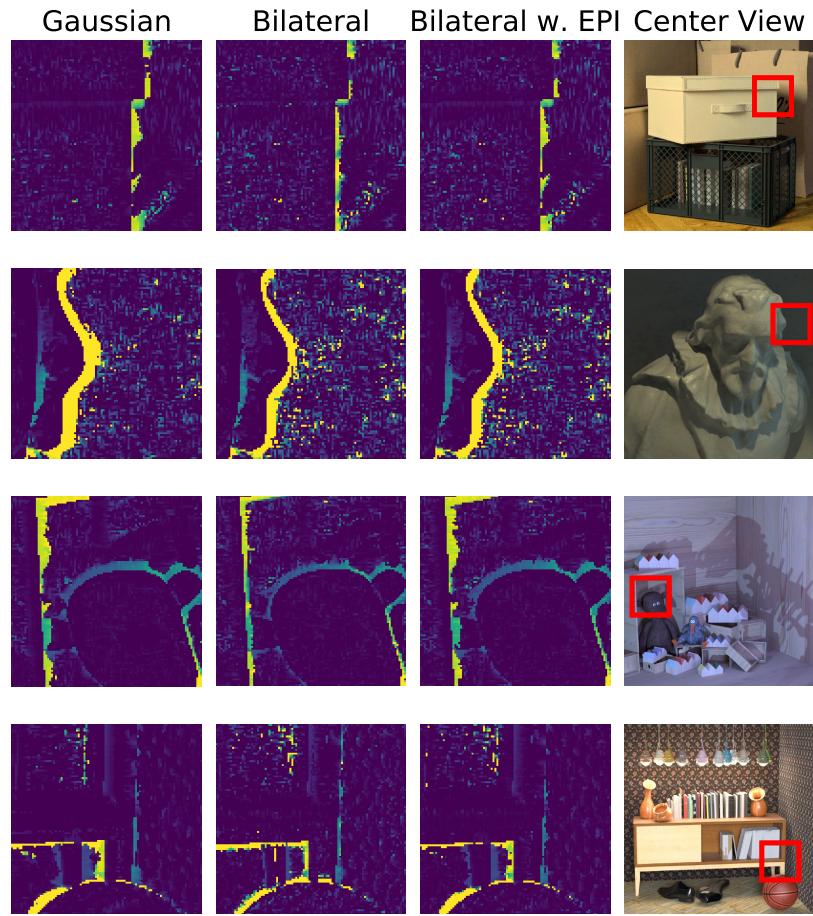


Figure 3.12: Results of Bilateral filtering for the scenes *boxes*, *cotton*, *dino*, *sideboard*.

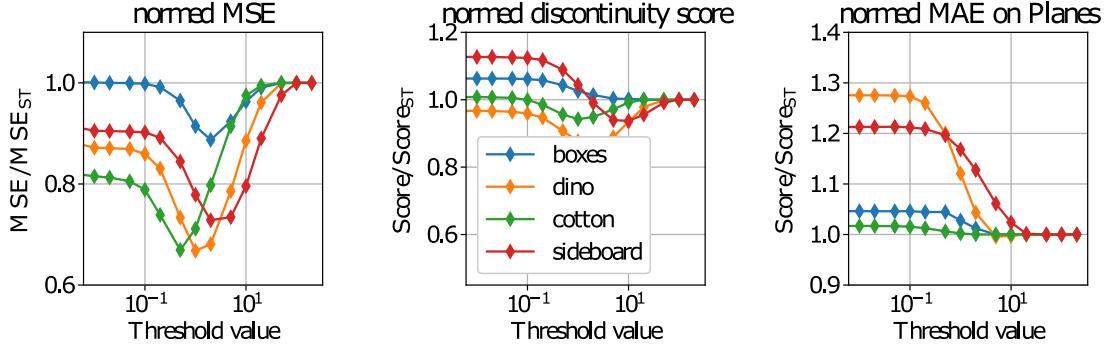


Figure 3.13: Results when thresholding the Gradients in the EPI are shown. All results are divided by the score obtained from the normal structure tensor algorithm. On the left the Mean Squared Error is shown, in the middle the discontinuity score is shown, on the right we see the Mean Average Error on planar surfaces in the scene. All metric scores follow the metrics from Honauer et al. [2016].

affected by noise in both cases. Still the Thresholded Gradient ST is less affected by the discontinuity problems when increasing the kernel size, leading to a quasi-independence from the kernelsize when looking at the MSE Score. This fact is to be enjoyed with care as the kernelsize does have a significant impact on the resulting image.

3.2.4 Occlusion awareness segmentation

In the next step we are trying to detect occlusions in the scan directly via the Gradient norm, as a high gradient is a good indicator for an occlusion edge. The result however is dependent on multiple parameters, two of them are the kernelsize of the morphology closing kernel. In figure 3.16 (upper row) the metric scores are plotted in function of the y -directional size of the kernel. The x -morphology kernel is held constant to the size of 7. From figure 3.16 (lower row) we can extract , that for all scenes we obtain the best result wne sticking to a 7-pix x -directional kernelsize. As the EPI is only 9 pixels wide, we do not go for bigger kernels.

The larger the kernel size is, the bigger the connected parts belonging to the same segment in the EPI are. More segmentations have negative effect on the Mean average error on planar surfaces with lots of texture, which we can see in the scenes *dino* and *sideboard*. This is approved in the upper right plot showing the MAE on planar surfaces for bigger and smaller closing kernels. A larger morphology size kernel results in a better planar surface score, however in the limit for very large kernels no segmentation is happening, thus the relative score is converging to 1. Furthermore the scenes *cotton* and *dino* are the ones whose mean squared error is mostly dependent on occlusions in the scene.

(FIGURE SHOWING PIXEL DISTANCES / ZOOMED IN EPI NORM)

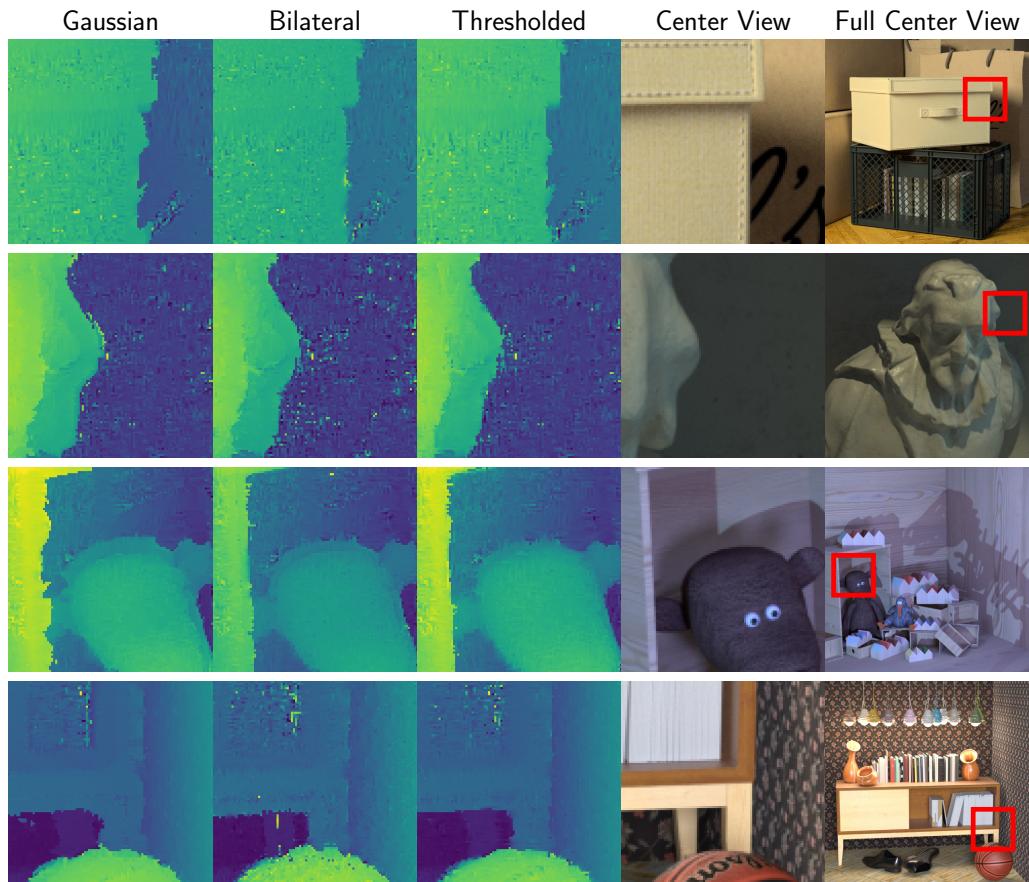


Figure 3.14: Disparity map Zoom-ins for different Structure Tensor filtering. From Left to right gaussian filtering, bilateral filtering and gaussian filtering when thresholding the gradients.

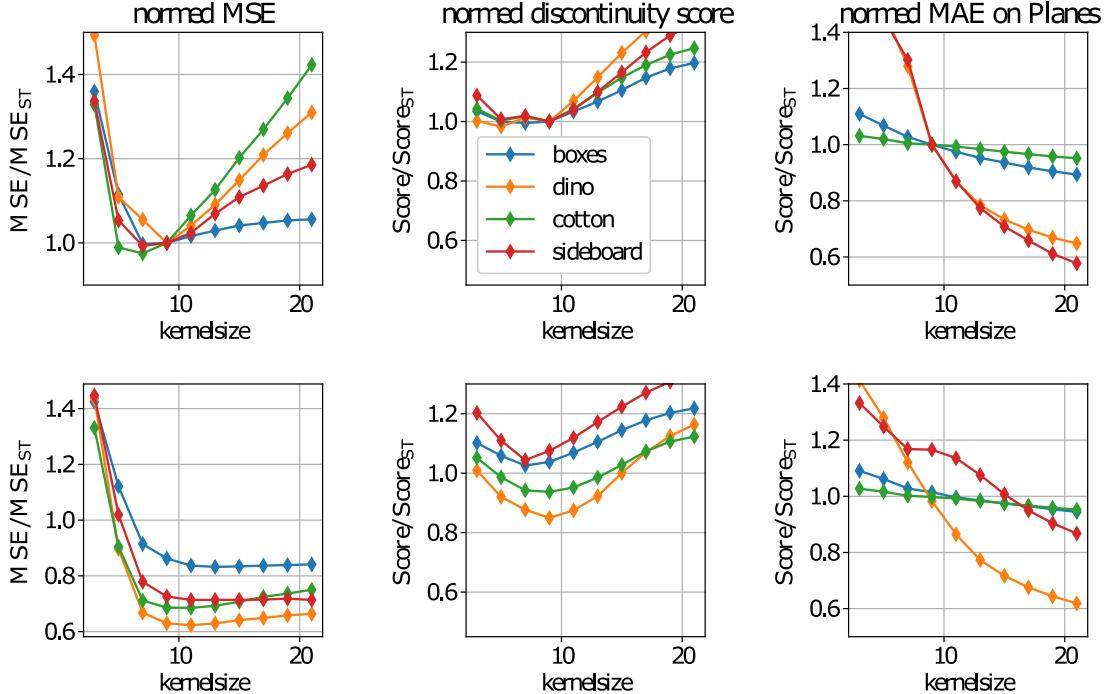


Figure 3.15: Top: Varying the kernelsize, using the old ST pipeline. Bottom: Varying the kernelsize with thresholding the gradients.

In figure 3.17 the segmentation algorithm is compared to the old pipeline as well as to the thresholded norm pipeline that is evaluated on the previous section. Again we have a look at the same scenes *boxes*, *cotton*, *dino* and *sideboard*. It becomes clear that both algorithms to some degree improve the occlusion handling, still the results differ from each other. In the Zoom-in on the scene *boxes* the Thresholded pipeline seems to achieve slightly better results, however on the occlusion in the scene *cotton* the Segmentation algorithm proves to produce the better results. The Segmentation especially works well with 'clean' edges, where the edge itself results in a clear peak in the norm of the plot (See figure 2.9 in section 2.3.4). However in the other scenes the segmentation does not seem to work well as the algorithm only achieves minor improvements.

3.2.5 Alternative coherence

The use of the alternative Cost function as explained in section 2.3.5 only works in combination with thresholding the gradient (section 2.3.3). For this reason we compare the error maps of the old pipeline, the threshold pipeline and the new error measure to each other in figure 3.18. From the given score above each subimage one can conclude that the new error measure does perform about equally as good as the thresholded pipeline. The alternative coherence measure was expected to result in a better disparity map, because in theory it should handle occlusion edges to some

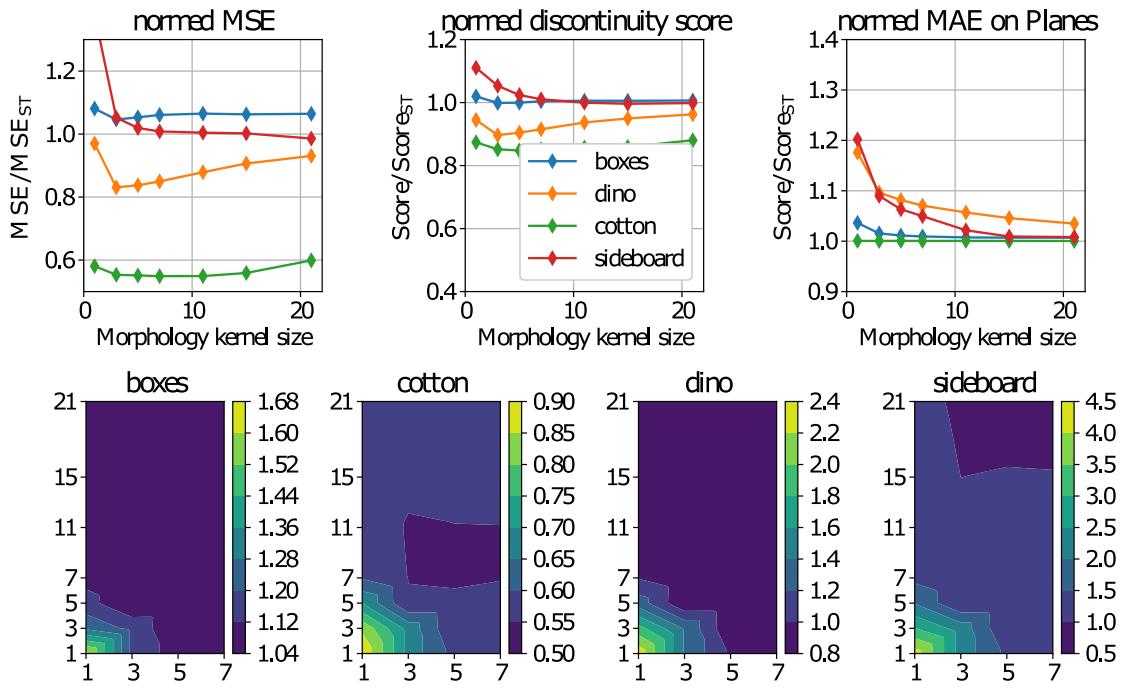


Figure 3.16: Upper row: Metric scores in dependence of the size in y -direction of the morphology kernel, x -direction size is 7.
 Lower row: dependence of the MSE for the size of the kernel. the y -size is depicted on the x -axis.

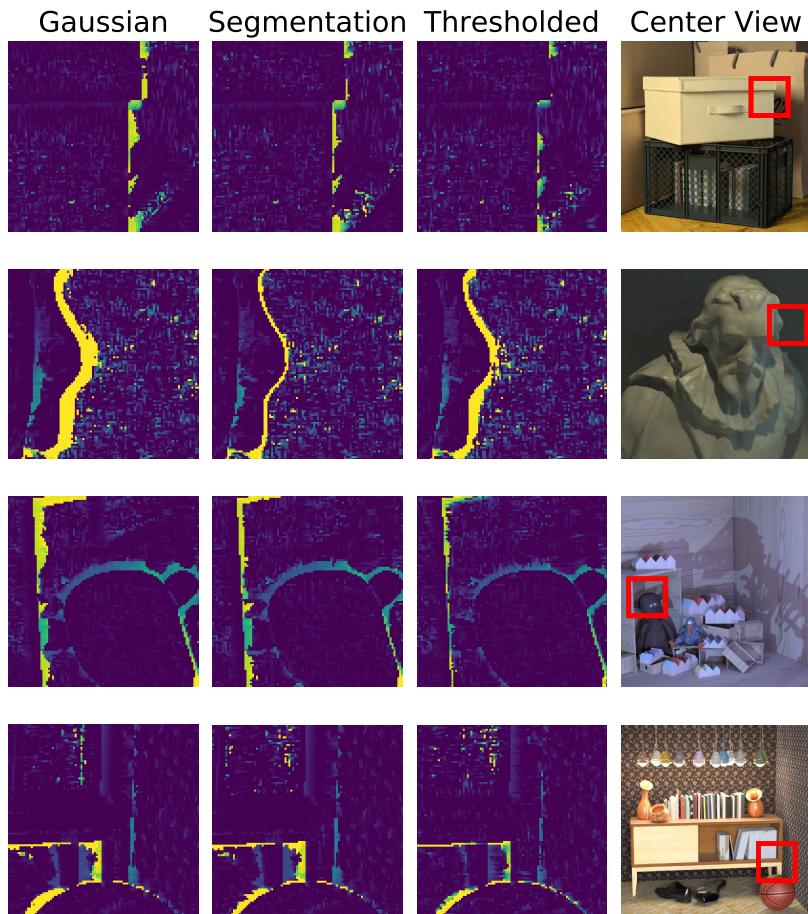


Figure 3.17: Disparity map Zoom-ins for different Structure Tensor filtering. From Left to right gaussian filtering, segmentation filtering and gaussian filtering when thresholding the gradients.

degree better than the old coherence measure does. Still, figure 3.18 shows that the errors remain roughly in the same spots.

Both the old coherence and the new error measure calculate the respective error from the EPI in horizontal and vertical direction. Both views get merged based on the error measure of each view, resulting in a merged Coherence/Error map. This can be seen in figure 3.19, where for each scene the horizontal, vertical and merged Coherence/Error map is illustrated as well as the actual error that was produced by the pipeline. Since in figure 3.18 one recognizes that the disparity error maps for both pipelines don't differ much, only the error of the old pipeline is shown in figure 3.19. One can see that the merged error of the new error measure seems to have a bigger overlap with the actual error, making it a better tool for masking out bad pixels than the coherence value. Especially in the scene *cotton* the occlusion edge errors fit better to the error with the new error measure.

To quantify this result, we choose different error thresholds for both pipelines and set all disparity values whose related error is below the threshold to zero. The higher the amount of masked out pixels is, the higher we expect the MSE to be since the amount of disparity values being 0 increases linearly (those disparity values also account for the total MSE). This linear relation between the percentage of disparity values set to zero and the total MSE is plotted in figure 3.20 in a logarithmic manner. Using equal axes this would result in curves with linear form. The logarithmic scale however allows us to evaluate more easily how many pixels have to be set to zero until the result gets significantly worse. As long as the masked out pixels are already highly inaccurate, the resulting MSE doesn't change much by setting them to zero. Note that in all scenes zero is circa the mean depth of the scene. These plots confirm the observation that the error in the scene *cotton* is prognosticated better by the new error measure, since setting the upper 10 % of the highest error pixels to 0 results in a better score than doing the same with the old coherence measure. In the other scenes the quality of the error measure seems to be equal in both cases.

3.2.6 Alternative merging of x- and y-direction

When merging the x- and y- directional depth estimations giving preference to the depth value that is further away, one obtains better results on occlusion edges while the number of outliers rises. This trade-off is favourable, because statistically "random" outliers are easier to handle in a postprocessing step than systematic errors such as edge fattening. The trade-off is controlled via the threshold in equation 2.34 and is set to 0.7 in figure 3.23. In the close-up views of the deviation from the ground truth in all four scenes (column 3 and 4 in figure 3.23) one sees that occlusion edges are significantly improved, especially in scene *boxes* and *dino*. In scene *cotton* the edge remains, however it is smaller. The last scene does not show a significant improvement at occlusions. In all scenes however the noise has increased. On planar areas the higher coherence value is mostly the better one while on occlusions the coherence does not serve as a good measure to decide whether to take the x-directional or the y- directional depth estimation. As a result, one faces noise as

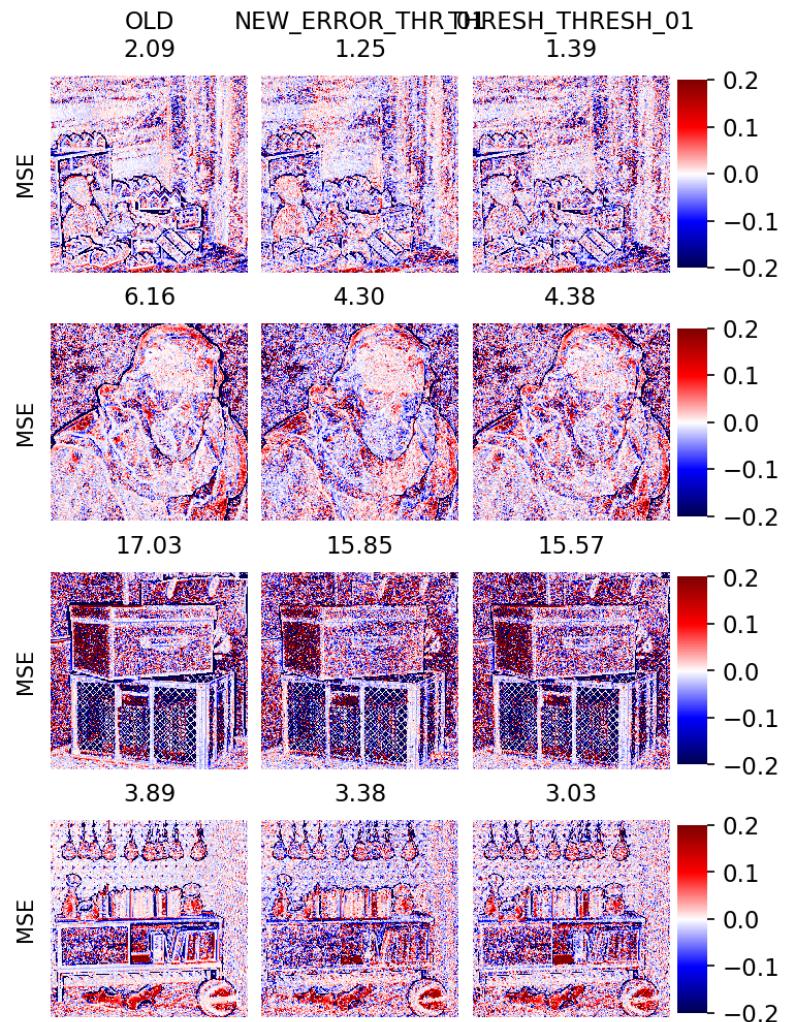


Figure 3.18: Left: The Mean squared error using the old ST pipeline. Middle: Using the alternative error measure + thresholding the gradients. Right: Using the old coherence measure + thresholding the gradients.

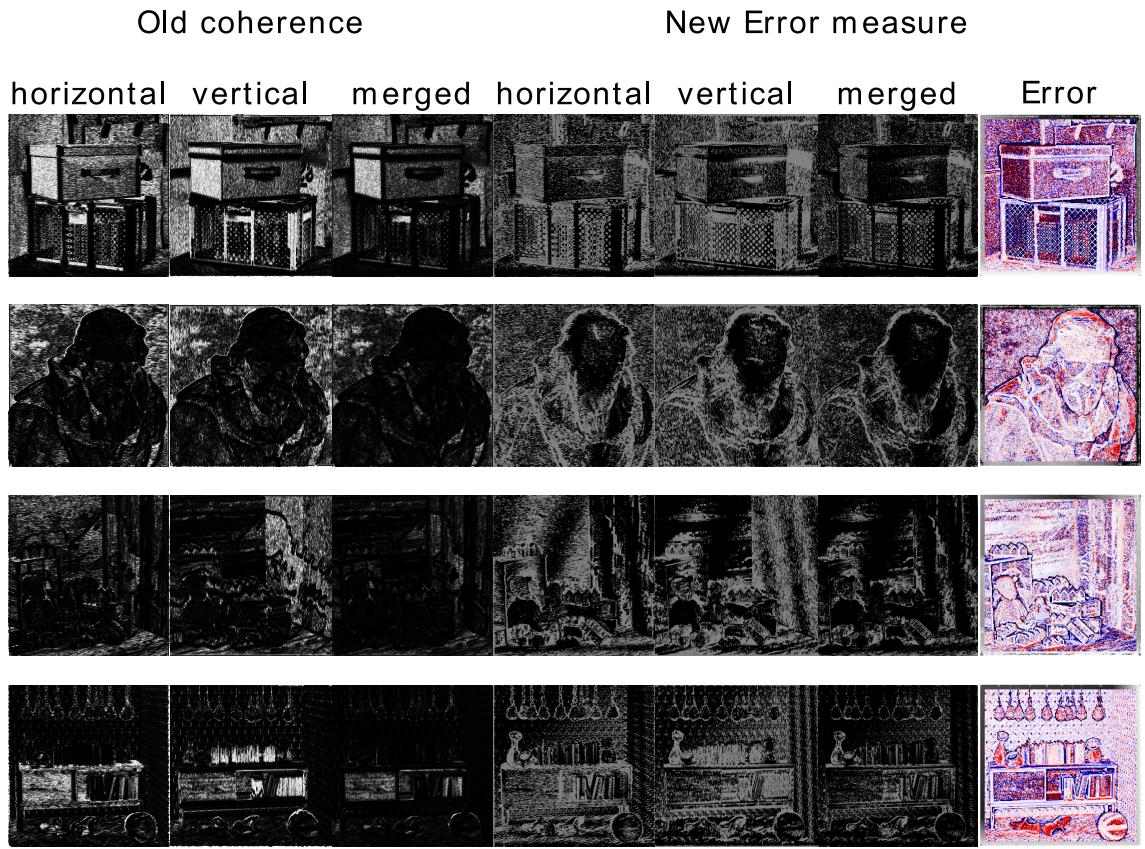


Figure 3.19: Columns on the left: The coherence value from the horizontal EPI, vertical EPI and the merged coherence map. Middle to the right columns: Alternative error measure based on error propagation. On the very right: Actual deviation from the ground truth.

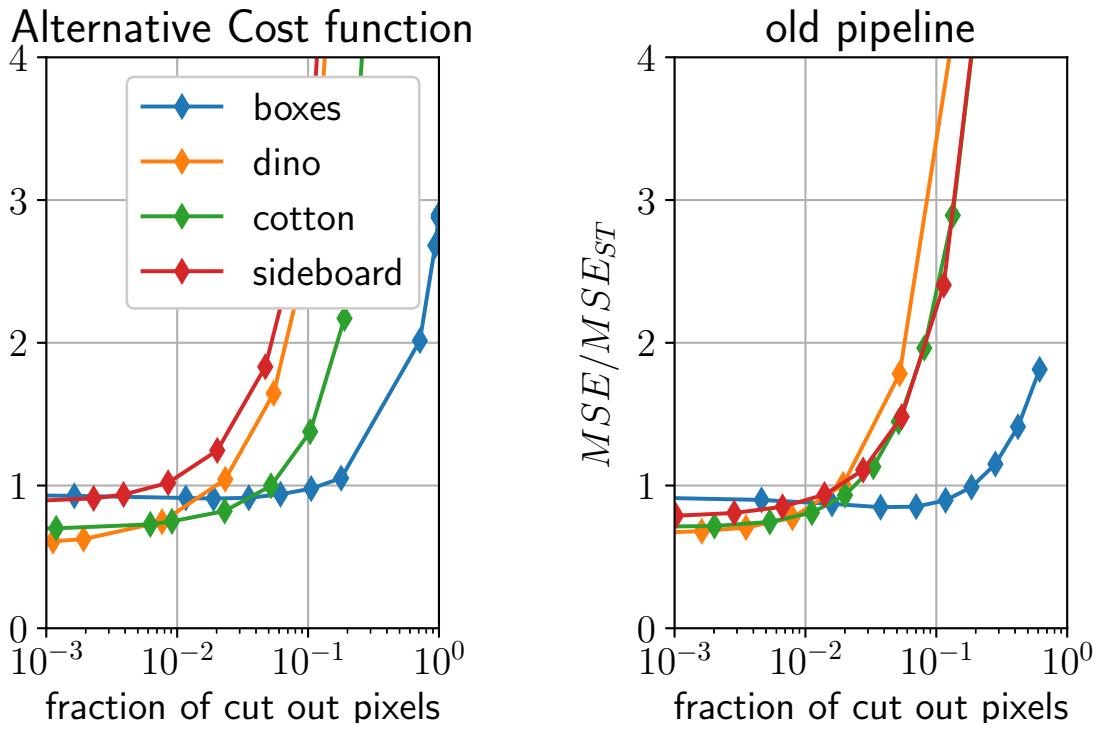


Figure 3.20: The relative MSE is plotted on the percentage of pixels that are set to 0 at different coherence values. Left the New error metric is used, right the old pipeline is used.

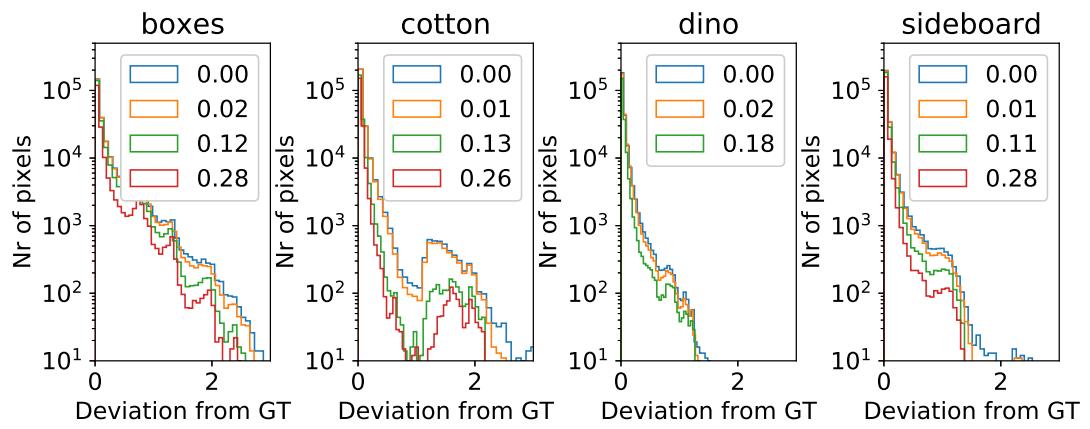


Figure 3.21: The error histogram for different masks based on thresholding the coherence map. The labeling indicates the fraction of masked out pixels.

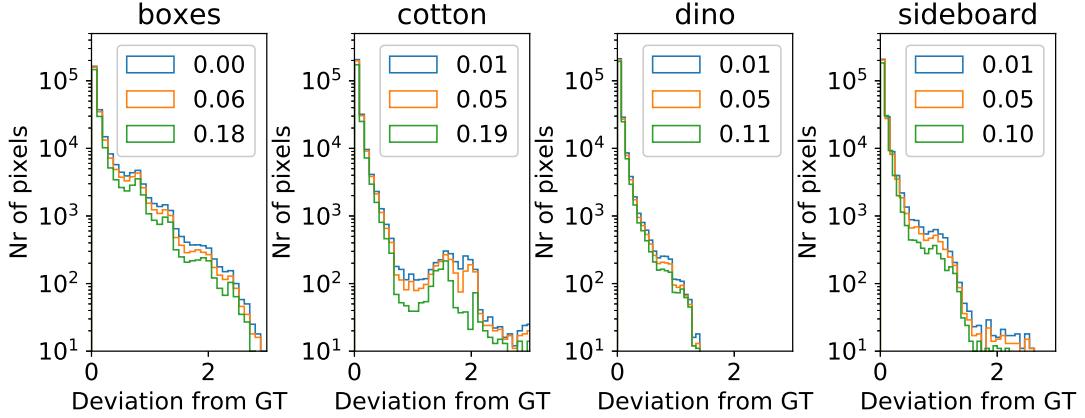


Figure 3.22: The error histogram for different masks based on thresholding the new error map. The labeling indicates the fraction of masked out pixels.

it can be observed well in scene *boxes* and *sideboard*.

Nevertheless the MSE is improved for all scenes if the threshold value is set between 0 and 0.5 as one can see from figure 3.24 on the left side. The threshold value 0 corresponds to the old pipeline, the threshold value 1 corresponds to always choosing the lower disparity value. Choosing big threshold values however leads to increased noise, thus worsening the MSE score. Note that we can see from figure 3.24 on the right that the BadPixel score is increasing, followingly the amount of pixels that are highly accurate decreases for a high threshold value. However the occlusion handling can compensate for the noise which is leading to the increased BadPixel score. Further we can smoothen out slight errors more easily than bigger deviations at occlusion edges.

The same plot can be found in figure 3.25 where the algorithm is combined with thresholding the gradients as described in section 2.3.3. One can see that we can see a slight improvement in the MSE for the scenes *dino* and *sideboard*, while the improvement of the occlusions in scene *boxes* cannot compensate the increasing noise. The scene *cotton* with big occlusion edges obtains the best result for the highest threshold value. Even though it doesn't improve the MSE Score it may be favourable to have this merging implemented if it can be handled by postprocessing smoothing. This is evaluated using the SemiGlobal Matching algorithm in section 3.3. Note taht the dependent parameter is the one referenced in equation 2.34, the gradient treshold is set to 0.1 . For this reason the MSE/MSE_{ST} Score does not start by 1. Even though in figure 3.24 and figure 3.23 it seems that occlusions can be handled by alternative merging, the gradient threshold does the same, such that both combined do not necessarily improve the result as seen in figure 3.25 on the left. For a higher Merging parameter the BadPixel Score is always increasing due to higher noise coming from low coherence disparity values.

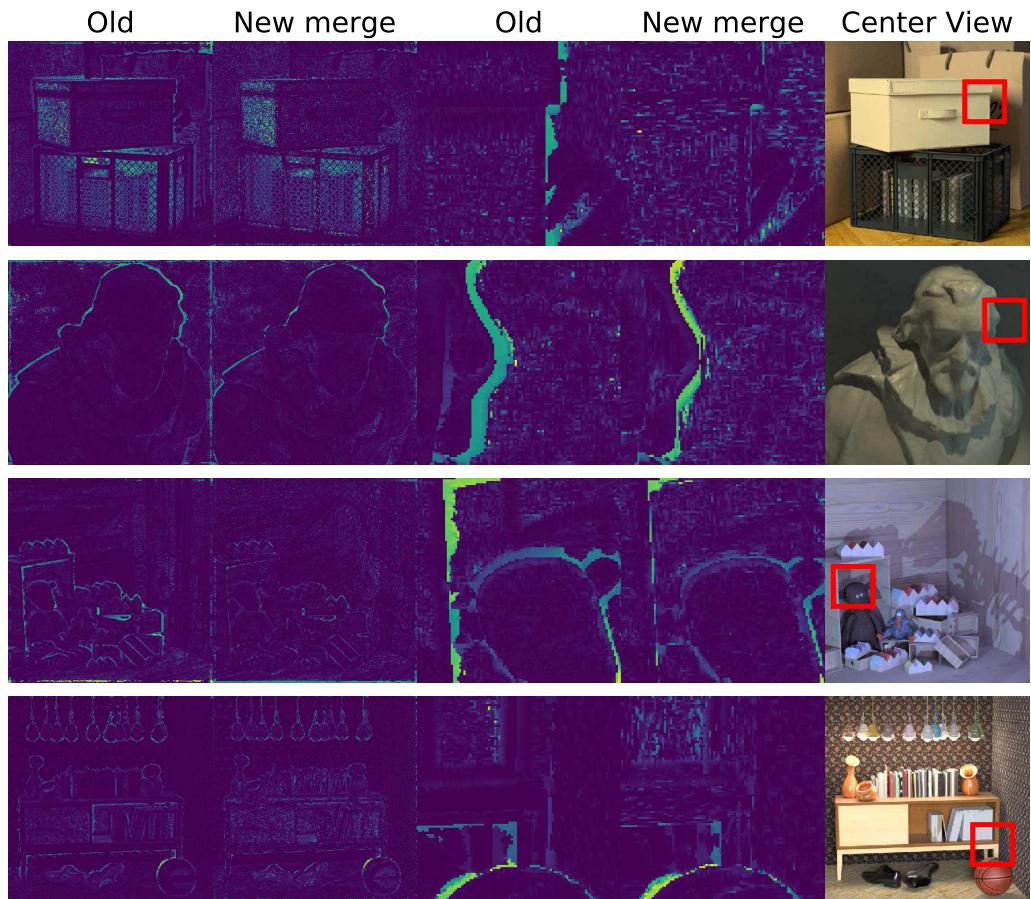


Figure 3.23: Left columns: Comparison between the old merging and the new merging for the whole depth map. The absolute deviation from the ground truth is shown. Middle columns: Close-up view on occlusion edges. Right column: Localisation of the close-up view in the scene. Parameter from equation 2.34 is set to 0.7.

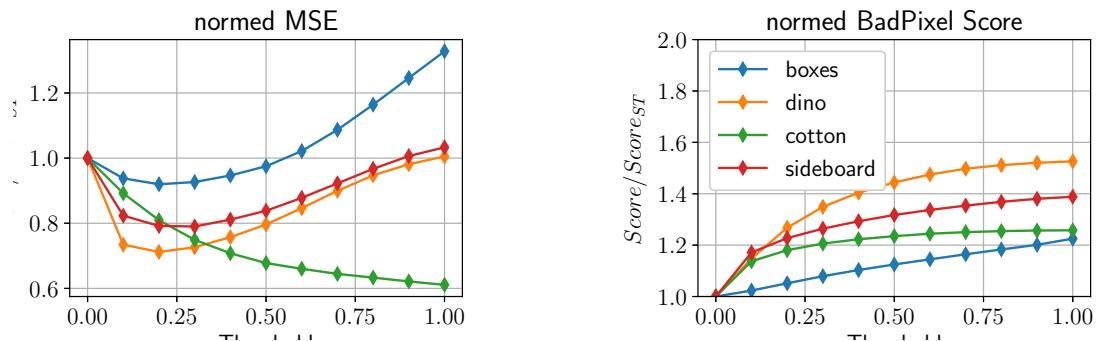


Figure 3.24: Alternative Merging: Parameter dependence. Left: Mean Squared Error. Right: BadPixel Score

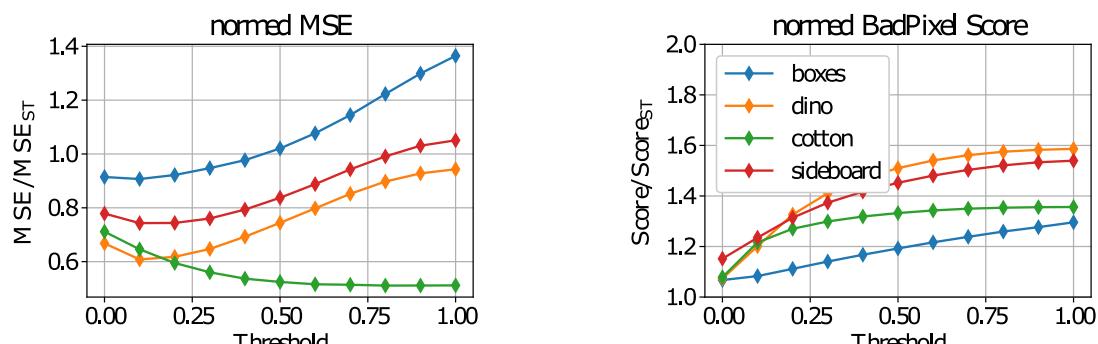


Figure 3.25: Alternative Merging + tresholded gradients: Parameter dependence. Left: Mean Squared Error. Right: BadPixel Score

3.3 Semi-Global Matching

First we evaluate the SGM algorithm as described in section 2.5.3. The Semi-Global Matching algorithm as implemented here is dependent on 3 parameters from equation 2.45.

We tried to perform a grid search to find the best parameters. The illustration of the grid search can be found in the appendix in figure B.2, B.3 and B.4 for different metrics. Even though one may conclude from figure B.4 that the score for planes is improved by setting a high parameter 1, the difference between the maximum and minimum score in the grid search is too small to see any dependence on any of the parameters. Note that we sampled the parameter space logarithmically to cover a wider space. Obtaining independence of the parameters is positive for working with the algorithm, the reason for the independence lies in the fact that at most pixels the candidate disparity values between whom the SGM algorithm decides are the same. If the Structure tensor only finds one unambiguous disparity value, this value is chosen regardless of the parameters. On planar surfaces this unambiguity is not given, thus the SGM algorithm is able to smoothen such surfaces. This can be seen in figure 3.26, where the old pipeline is compared to the result using with the sgm algorithm. The depth map is improved at surfaces with noisy values, thus the SGM has a smoothing effect. However at occlusion edges the SGM pipeline does not perform significantly better.

Improved Coherence Measure by using SGM

Even though the SGM pipeline does not handle occlusion edges well, it still does recognize disparity jumps as high costs in the cost function. In figure 3.27 the cost function is depicted in comparison to the actual error as well as in comparison to the old pipeline. Large disparity jumps are punished with a big penalty, thus they can be detected easily. In scene *cotton* and *sideboard* such disparity jumps are detected, while the other disparity jumps we see are less than 1 pixel shift and they are only punished by a small penalty. The cost function itself could be used to mask out possibly erroneous pixels. To test the quality of such a masking the error histogram for all 4 scenes is plotted in figure 3.28. It becomes obvious for scene *dino* and *sideboard* that large errors are masked out more often than it did with using the old coherence measure as depicted in figure 3.21. In scene *boxes* the disparity map is quite noisy which makes it hard to mask out those regions. However the usability of an error measure based on SGM is restricted due to the dependence on the penalty parameters mentioned in equation 2.45.

SGM as pure Postprocessing

For now we take as input the processed disparity map from the structuretensor and create the SGM candidates at each pixel from the local neighbourhood as described in section 2.5.4. Figure 3.29 shows how MSE, BadPixel Score and processing time

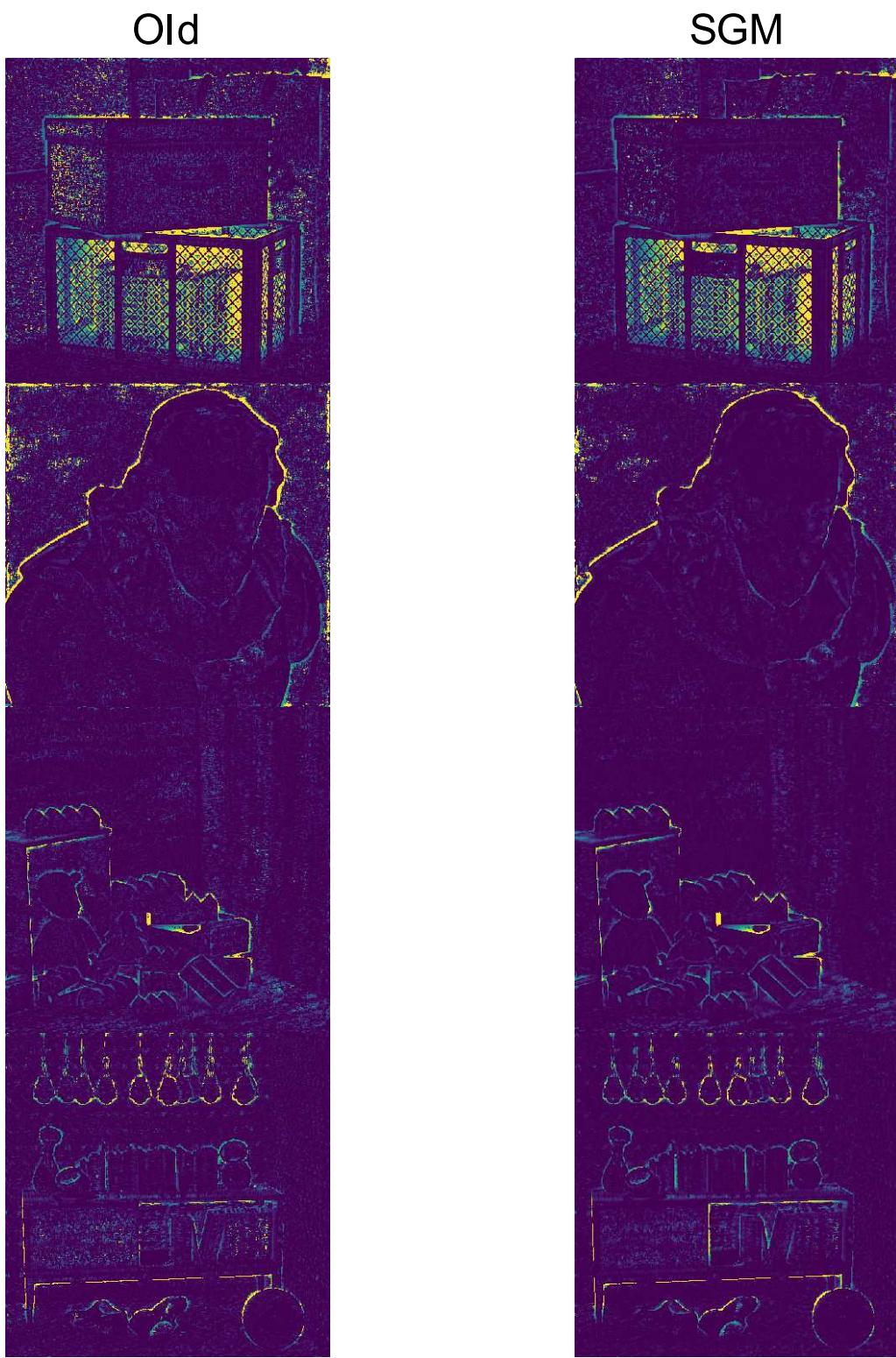


Figure 3.26: Disparity error maps using the old pipeline (left) and the Semi-Global Matching Algorithm (Right). Yellow is a big deviation, dark-blue is no deviation.

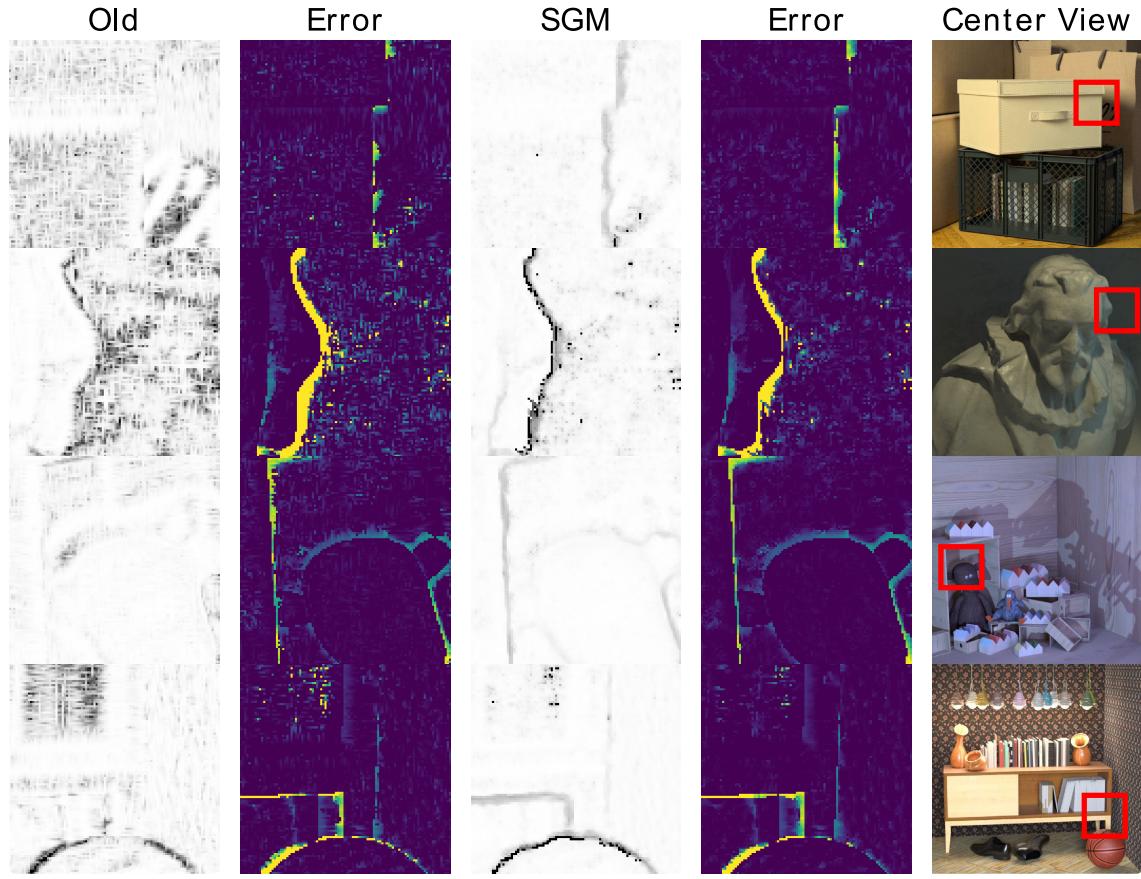


Figure 3.27: The assumed error (grey-valued) when calculating depth without using SGM (left) and with SGM (right). In Color the actual error for both pipelines is depicted.

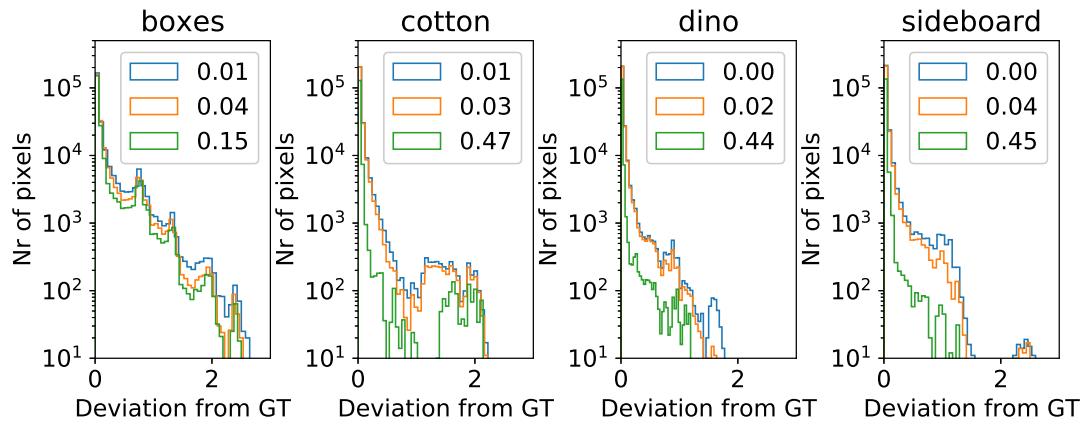


Figure 3.28: The error histogram for different masks based on thresholding with the cost function . The labeling indicates the fraction of masked out pixels.

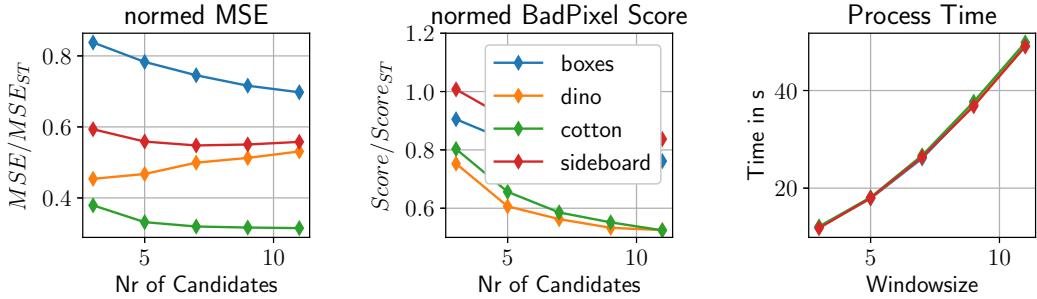


Figure 3.29: SGM as pure postprocessing combined with Alternative merging: The Number of candidates that is chosen from the direct neighbourhood is varied. Left/Middle: The MSE/BadPixel Score in dependence of the Nr of candidates. Right: the processing time of the algorithm dependent on the Nr of candidates. The window size is held constant at 5.

depend on the Number of candidates from the local neighbourhood which is defined as a 5×5 window around the relevant pixel. The candidates are chosen randomly from the neighbourhood. Note that the input depth map has been created with a modified structure tensor pipeline including **thresholded gradients** and **alternative merging**. The gradient threshold is constantly set to 0.1, while the merging parameter is set to 0.3 .

One sees that the MSE not necessarily improves given more candidates. In the scene *cotton* and *sideboard* we reach a fast saturation of the MSE while in the scene *boxes* one would even need more candidates to reach convergence. This has to do with the large noisy regions in the scene that make it impossible to get an acceptable candidate from the neighbourhood. In scene *dino* the MSE even gets worse with a higher number of candidates. On the other hand the BadPixel Score is improved for all scenes. One can conclude that the postprocessing SGM manages to improve the depth map at small regions of deviations (outliers), however larger erroneous regions can not be corrected because of the limited amount of candidate values. Using an iterative approach we try to overcome this problem in the next section.

Since the SGM algorithm is implemented here as a pure postprocessing step, one can analyse the effect of applying modifications on the structure tensor beforehand. Figure 3.30 shows the dependence on the merging parameter that is characterized by equation 2.34. No thresholding of the gradient is applied here. Larger parameters lead to better results at discontinuities, however as already shown in section 3.2.6 a higher merging parameter leads to an increased BadPixel Score, though the normed score stays below the 1.0 mark. Again scene *boxes* reaches a high MSE due to the large monochrome areas leading to noisy depth maps. One solution to handle such large deviations at surfaces is to mask out contiguous regions of low coherence. This has been implemented for the application on real data, see section ??.

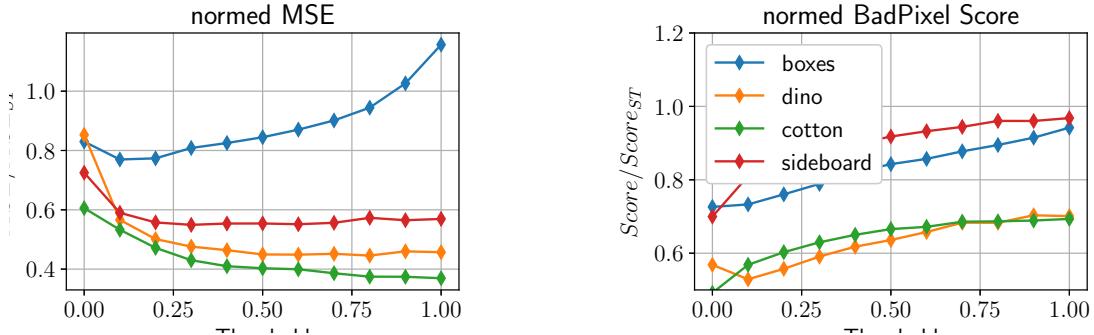


Figure 3.30: Alternative Merging with SGM Postprocessing. Left/Right: MSE/BadPixel Score as a function of the merging threshold.

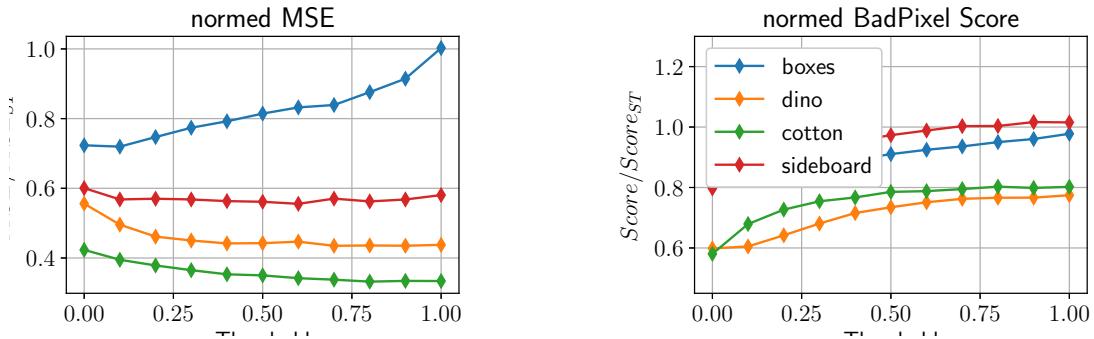


Figure 3.31: Alternative Merging with SGM Postprocessing while using thresholded gradients. Left/Right: MSE/BadPixel Score as a function of the merging threshold.

As we did before in section 3.2.6 we now take a look at the effect of implementing a gradient threshold additionally. In Figure 3.31 the MSE and the BadPixel Score are shown in dependence of the merging parameter. While the BadPixelScore behaves approximately as without a gradient threshold (figure 3.30), the MSE stays approximately constant for all scenes besides scene *boxes*. No improvement big improvement can be obtained, since the thresholding of the gradients already handles occlusions well. In figure 3.30 as well as in figure 3.31 the MSE does not rise for a larger merging parameter as it does in figure 3.24 and in 3.25, where no SGM has been applied. The SGM can handle the noisy effects which occur for a high merging parameter.

Using a grid search we tried to investigate the impact of varying the **Nr of candidates** as well as the **windowsize** on different Metrics. In figure 3.32 the MSE is shown in dependence of Nr of candidates and windowsize for all four scenes, in figure 3.33 the BadPixel Score is shown and in 3.34 the Discontinuity score is depicted. It becomes clear that the ideal kernelsize varies strongly from scene to scene as well as from metric to metric. Based on the shown results we fixed the windowsize for all calculations to 5x5 pixels. In general A higher Number of candidates promises

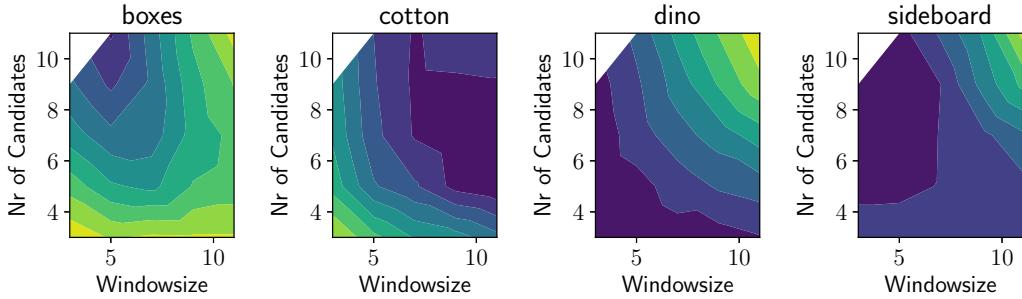


Figure 3.32: MSE Parameter dependence for Posprocessing SGM using alternative merging+thresholded gradients.

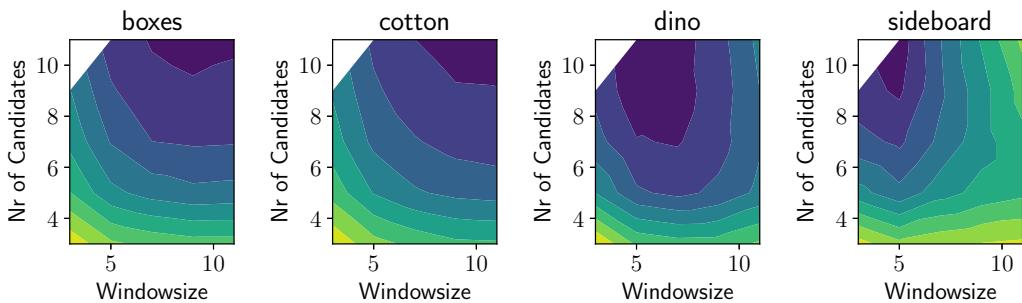


Figure 3.33: Badpixel score parameter dependence for Posprocessing SGM using alternative merging+thresholded gradients.

for better results, however the processing time scales linearly with the number of candidates. Thus we stick to Nr of candidates of 5, as shown in figure 3.29 on the right.

3.3.1 Iterative SGM postprocessing

In the following we evaluate the Iterative to sgm postprocessing as described in section 2.5.5. The iterative optimization should lead to a minimization of the global cost function. The global cost function we define as the mean cost value at all pixels. In figure 3.35 we confirm the iterative convergence of the Mean Cost Function on the example scene *cotton*. Parallel to the mean cost function the MSE also converges to a minimum. With the given set of parameters one can conclude that approximately 8 iterations are necessary to converge closely to the minimum value. In the upper row of figure 3.35 one can see how the iterative process smoothens outliers in the depth map.

The effect of the number of iterations on the MSE can also be extracted from figure 3.36. On the left the normed MSE is plotted. It becomes obvious that all

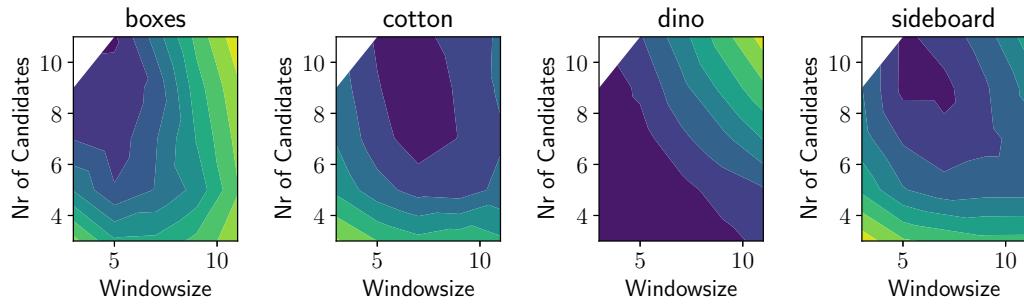


Figure 3.34: Discontinuity score parameter dependence for Postprocessing SGM using alternative merging+thresholded gradients.

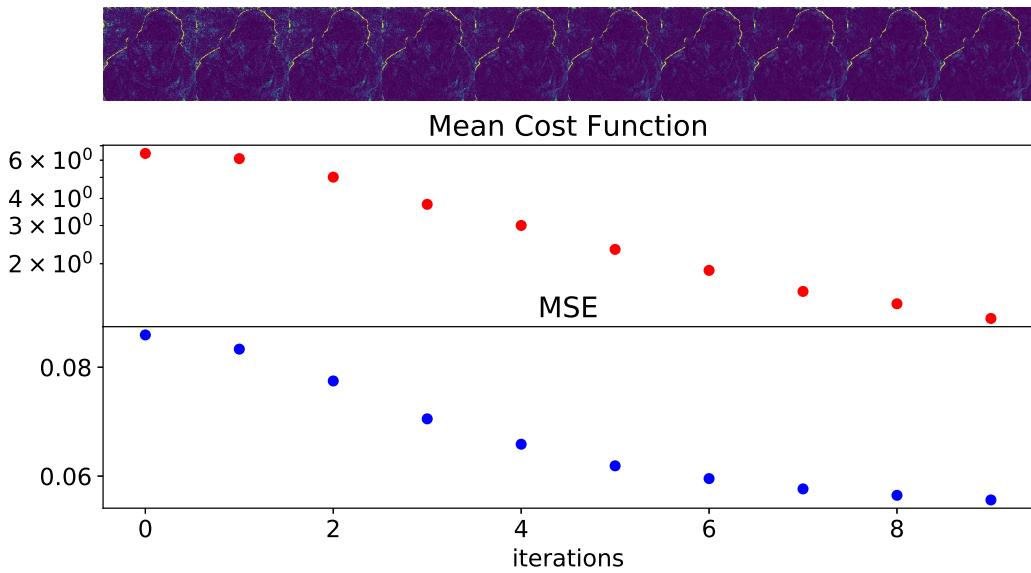


Figure 3.35: In the upper row the resulting depth map deviation from ground truth of the iterative SGM pipeline is shown for scene *cotton*. Below the Mean cost function averaged over all pixels as well as the MSE is depicted in dependence of the number of iterations. Note that the MSE is not normed as in figure 3.36.

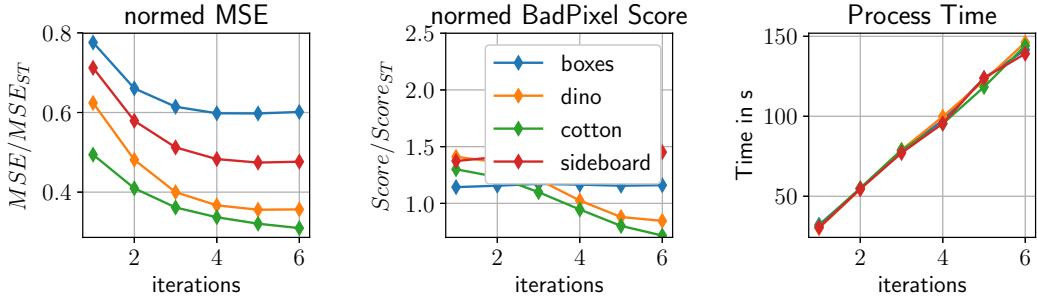


Figure 3.36: Iterative SGM pipeline dependence on Number of Iterations. Left: normed Mean Squared Error for all four scenes in function of Nr of iterations. Middle: Normed BadPixel Score in function of Nr of iterations. Right: Processing Time

scenes converge to a minimal MSE, however at different speed. Also for scene *cotton* and *dino* the BadPixel Score drops, while for scene *boxes* and *sideboard* it stays constant. However the process time increases linearly with the number of iterations, thus we stick to a maximum of 5 iterations in the upcoming calculation. This is still far from being comparable to the structure tensor pipeline, however as mentioned in section 2.5.7 we focus on the feasibility of such method and try to adapt the process time to a reasonable amount of time for our measuring purposes.

In the following we want to examine the effects of changing the initial Δ (see section 2.5.5) on the MSE as well as on the BadPixel Score. In Figure 3.37 The MSE and BadPixel Score are plotted in function of Δ_{initial} . For very low initial Δ no convergence can be reached, thus the MSE does stay high. For higher Δ the MSE is basically unaffected, given a the fixed number of 5 iterations. The BadPixel Score reaches a minimum for an initial $\Delta = 1$. To obtain the same result with higher Δ more iterations would be needed.

3.3.2 Application on real data

In this section we want to evaluate qualitatively the use of SGM postprocessing on real captured data instead of synthetic rendered images. Therefor we have a look at data from the Stanford light field archive ([Wiburn \[2004\]](#)) as well as captured light fields with the *Lumi+* sensor in course of an internship with HDVision Systems. In contrast to the synthetic scenes real captured light fields suffer from bad illumination, slight rectification errors and lense abberations. Thus the Structure Tensor coherence is mostly lower, leading to sparse disparity maps (after filtering out disparity values related to low coherence). One could simply run the SGM iterative postprocessing algorithm only on pixels with sufficient coherence, however we want to also smoothen outliers with low coherence instead of filtering them out. To include outliers for correction while excluding larger areas of bad coherence, we apply morphological opening to the binary mask which has been created based on

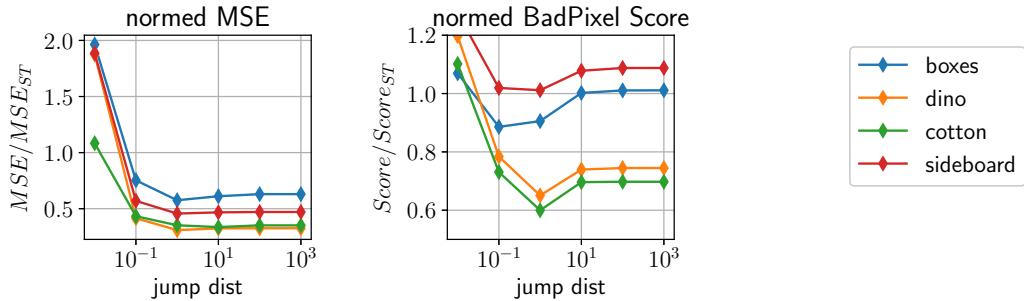


Figure 3.37: Iterative SGM pipeline dependence on Initial Δ with fixed Number of iterations (5). Left: normed Mean Squared Error for all four scenes in function of Δ . Middle: Normed BadPixel Score in function of Δ . Note that the x axis scale is logarithmic. (TODO: SAME PLOT WITH 10 ITERATIONS)

a coherence threshold. If not mentioned differently, the threshold is set to 0.85. We choose to use a rather small morphological kernel (3x3) to make sure every pixel has at least one neighbour initially which has been measured with sufficient coherence. In figure 3.38 one gets an impression on how SGM postprocessing changes the depth map. One can see the sparse depth map (a)) produced by the structure tensor (included gradient threshold already) of a glossy silicon fragmentary used in chemical industry to produce electrical devices. The accompanying mask can be seen in (b)), only pixels in black are evaluated. One can see a lot white pixels in between the black area, which we aim to include in the SGM pathwise optimization. Hence the masked is enhanced with a 3x3 morphological opening kernel, such that the new mask defines adjoint areas of low/high coherence (c)). We then run the SGM iterative postprocessing only on the pixels masked in black. The resulting depth map is more dense (d)). e) shows the center view of the camera in comparison .

Another application example on real data is the Lego truck from the Stanford light field archive depicted in figure 3.39 (h)). We focus on a smal clipping of the scene which is depicted in figure 3.39 (g)). One sees the effects of different coherence thresholds, merging parameters and gradient thresholds in figure a) - f). Figure a) shows the Basic structure tensor result with a minimal coherence set to 0.85. In contrast figure b) shows the same result with a coherence of 0.5. Values with low coherence added to the depth map lead to a more dense map however the added pixels are mostly noisy. Obviously the wires in the image are fattened (edge fattening), also the holes on the blue lego part are not visible. In c) and d) we see the same depth maps as in a) and b) despite that we changed to alternative merging. Still the wires appear to be a lot wider than they actually are

3.4 Resumee

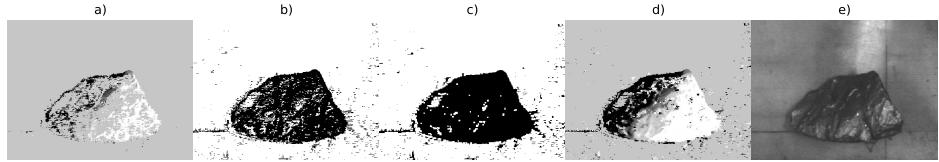


Figure 3.38: Glossy fragmentary scan captured with the *Lumi+* camera. a) Depth map obtained from structure Tensor b) Mask based on coherence value. All white pixels are masked out in [a)]. c) Enhanced mask using morphological opening d) depth map after applying Iterative SGM Post-processing on Pixels masked in [c)]. e) Centre view of the camera array.

Method	Time(s) (cotton)	MSE			
		boxes	cotton	dino	sideboard
Old Pipeline	4.78	17.03	6.16	2.09	3.89
Sandclock	6.81	18.55	6.78	2.23	4.14
epi-bilateralfilter ($\sigma = 90$)	43.96	19.26	6.41	2.22	4.16
bilateralfilter($\sigma = 750$)	20.18	18.95	6.38	2.10	4.32
Thresholding (thresh = 0.1)	6.06	15.57	4.38	1.39	3.03
occlusion segmentation (7x7 morph, thresh = 0.1)	11.28	18.08	3.39	1.78	3.91
alternative coherence + Treshold	14.15	15.85	4.30	1.25	3.38
SGM (4000,64000,1000)	8.90	14.98	3.84	1.75	3.18
SGM+ Threshold	12.75	13.31	2.96	1.11	2.43
pp Gauss 3		12.59	4.80	1.68	2.92
pp Gauss 5		11.95	4.54	1.59	2.72
pp Gauss 7		11.51	4.31	1.50	2.58
pp Gauss 9		11.34	4.16	1.46	2.52
pp sgm (4000, 256000, 400)		12.0	2.15	1.08	1.98
pp Median 3		13.21	4.96	1.83	3.35
pp Median 5		12.73	4.76	1.76	3.20
pp Median 7		12.73	4.76	1.76	3.20
pp Median 9		12.73	4.76	1.76	3.20
pp Median 11		12.73	4.76	1.76	3.20
Iterations 2					

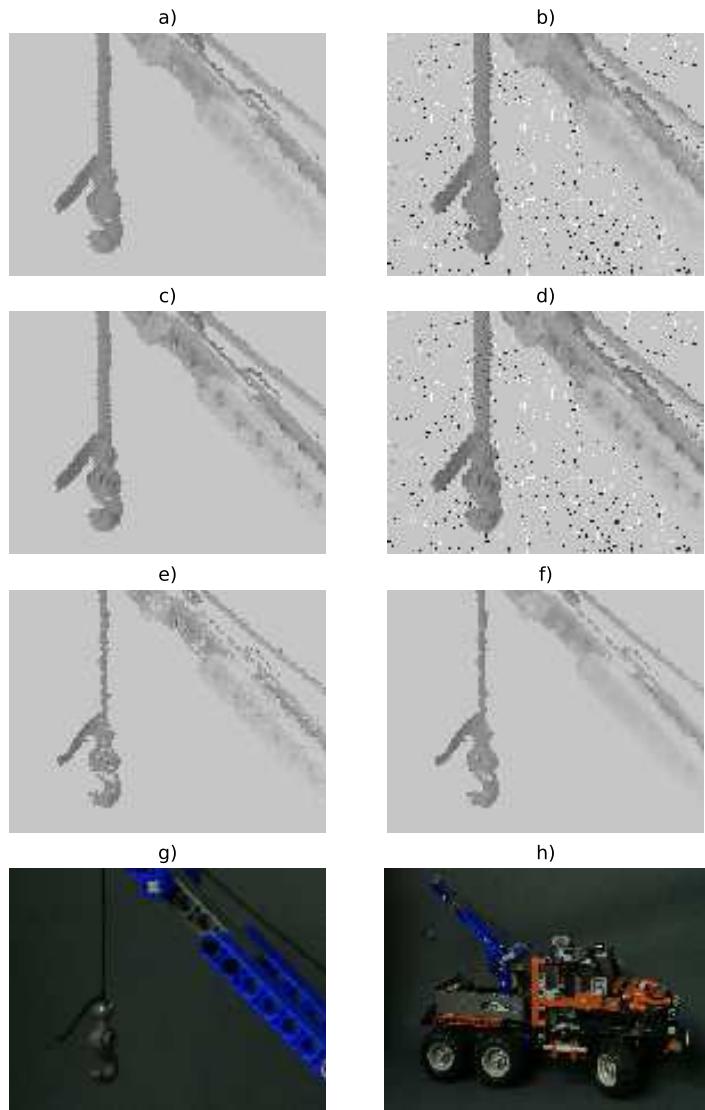


Figure 3.39: a) Depth map prod. by the structure tensor pipeline, with minimal coherence set to 0.85. b) Depth map prod. by the structure tensor pipeline, with minimal coherence set to 0.5 . c) Depth map when using alternative merging, minimal coherence set to 0.85. d) Depth map when using alternative merging, minimal coherence set to 0.5. e) Depth map when thresholding gradients, minimal coherence set to 0.85. f) e) Depth map when thresholding gradients and using iterative SGM Postprocessing, minimal coherence set to 0.85. g) Centre view clipping of the scene h) full centre view of the scene.

Part I

Appendix

A Derivation of the structure tensor

The derivation is taken from [Jähne \[2013\]](#). Taking a function $g : \Omega \rightarrow R, \Omega \subset R^D$, the preferred local direction $\vec{n} \subset R^D$ must satisfy the following equation:

$$(g^T \vec{n})^2 = |\nabla g|^2 \cos^2(\angle(g, \vec{n})) \quad (\text{A.1})$$

If ∇g is parallel or antiparallel to \vec{n} , the expression on the right side reaches a maximum. Therefore one needs to maximise the left hand expression in a local environment:

$$\vec{n}_{\text{preferred}} = \operatorname{argmax}_n \left(\int w(\vec{x} - \vec{x}') \left(\nabla g(\vec{x}')^T \vec{n} \right)^2 d^D x' \right), \quad (\text{A.2})$$

w is a window function defining the size of the local environment. Multiplying with \vec{n} we obtain:

$$\vec{n}_{\text{preferred}} = \operatorname{argmax}_n (\vec{n} J \vec{n}) \quad (\text{A.3})$$

$$J = \int w(\vec{x} - \vec{x}') \left(\nabla g(\vec{x}') \nabla g(\vec{x}')^T \right) d^D x' \quad (\text{A.4})$$

This results in a $D \times D$ tensor of the form

$$J_{pq} = \int_{-\infty}^{\infty} w(\vec{x} - \vec{x}') \left(\frac{g(\partial \vec{x}')}{\partial x'_p} \frac{g(\partial \vec{x}')}{\partial x'_q} \right) d^D x'. \quad (\text{A.5})$$

In two dimensions we can write

$$J = \begin{pmatrix} w * \frac{\partial g}{\partial x} \frac{\partial g}{\partial x} & w * \frac{\partial g}{\partial x} \frac{\partial g}{\partial s} \\ w * \frac{\partial g}{\partial s} \frac{\partial g}{\partial x} & w * \frac{\partial g}{\partial s} \frac{\partial g}{\partial s} \end{pmatrix}, \quad (\text{A.6})$$

where „*“ describes a convolution.

B SGM Parameter Testing

B.0.1 Real Data

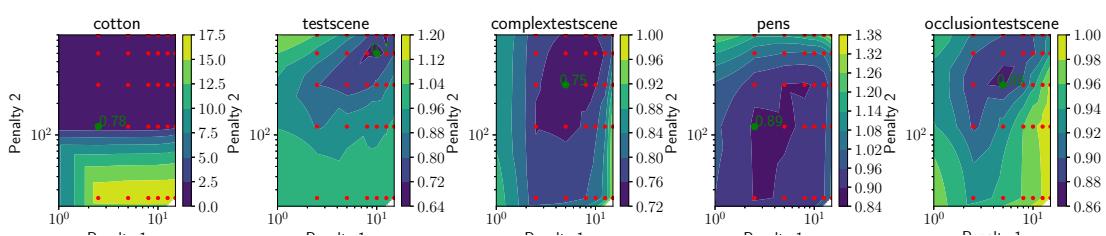


Figure B.1: parameter grid test for semi global matching

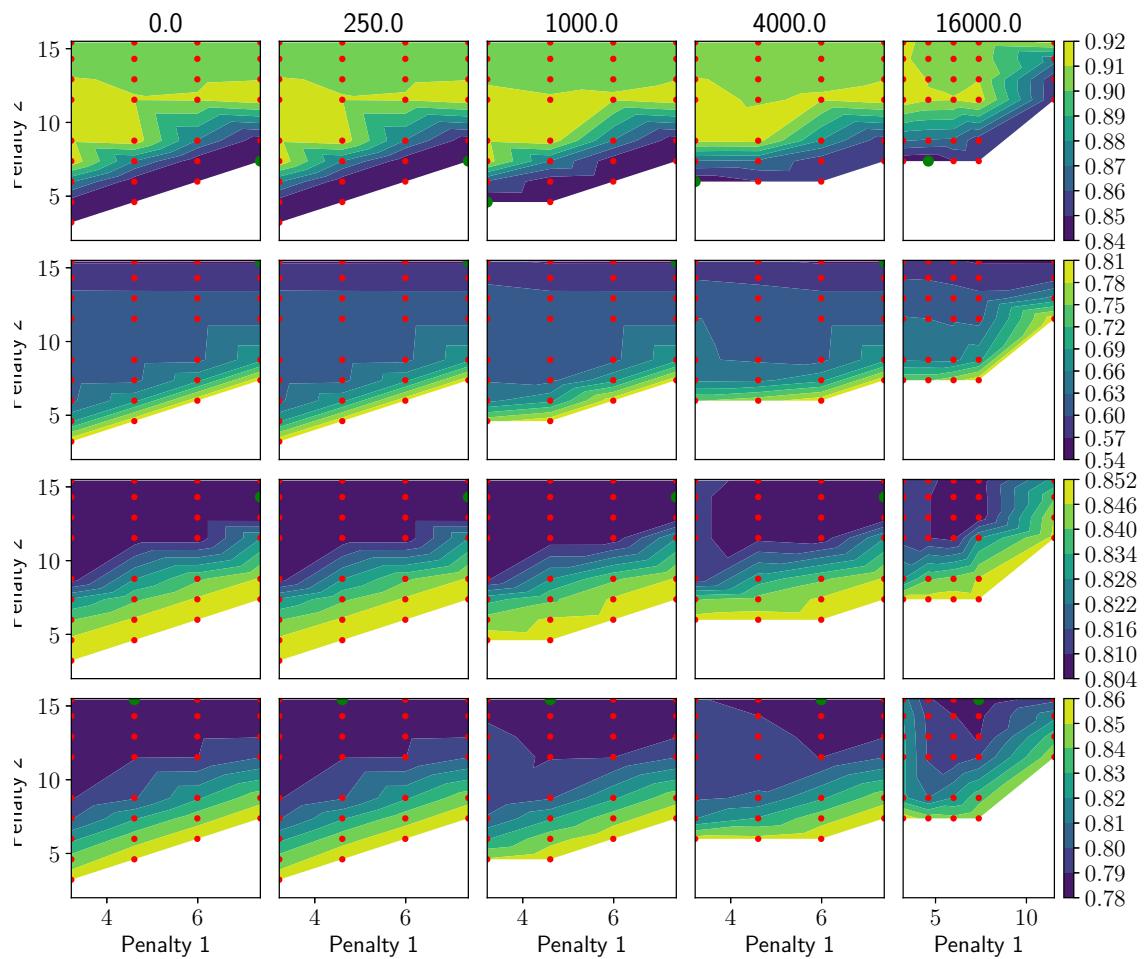


Figure B.2: Mean squared error for different (3) parameters

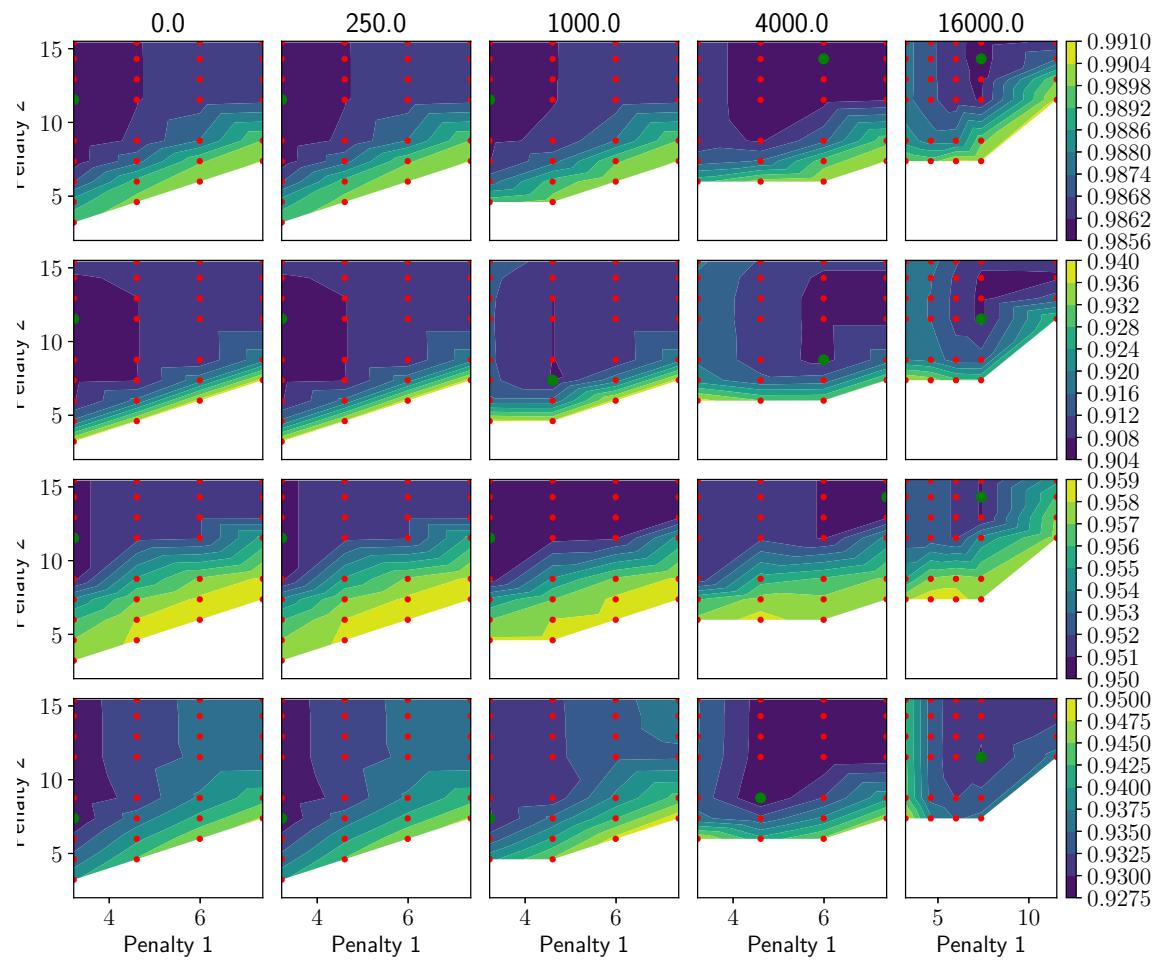


Figure B.3: discontinuity score for different (3) parameters

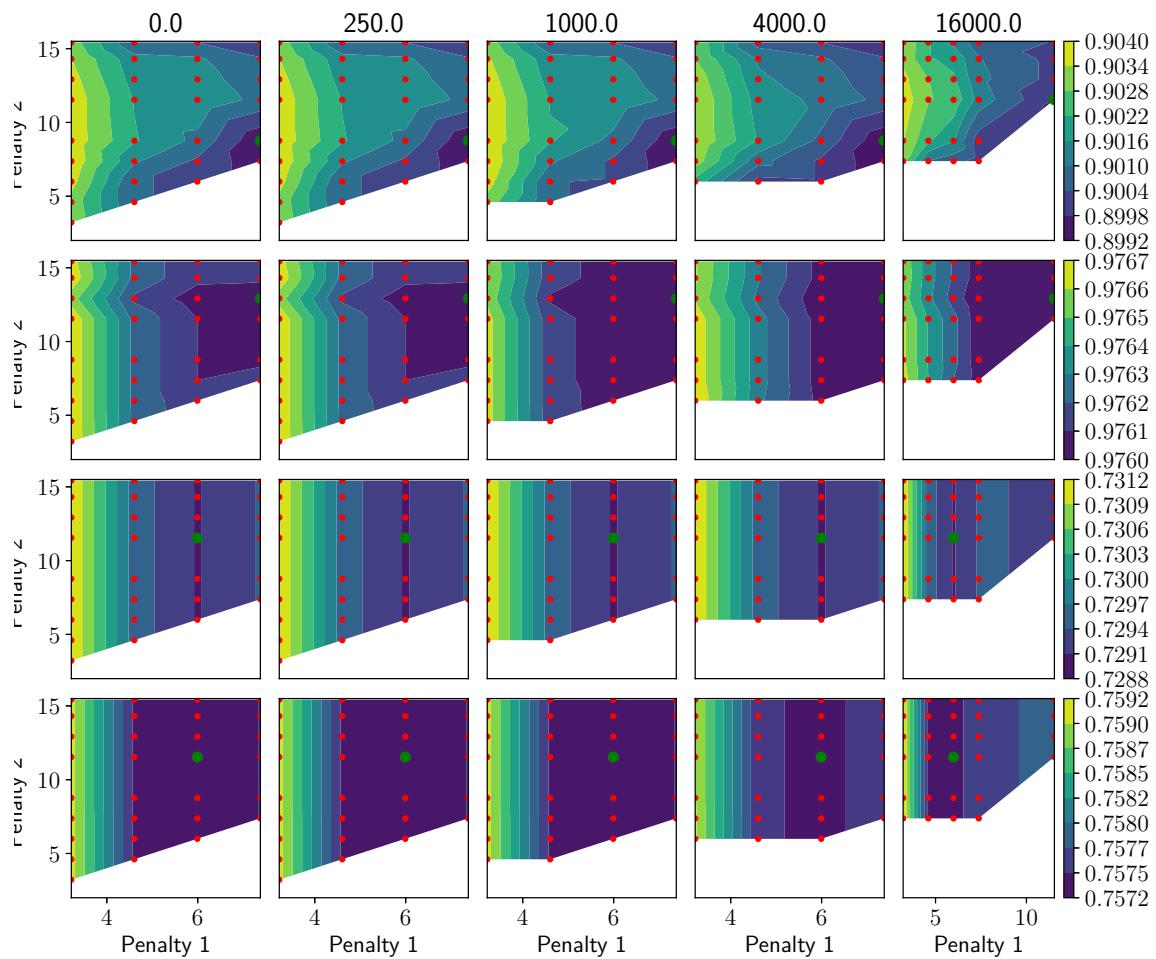


Figure B.4: planes score for different (3) parameters

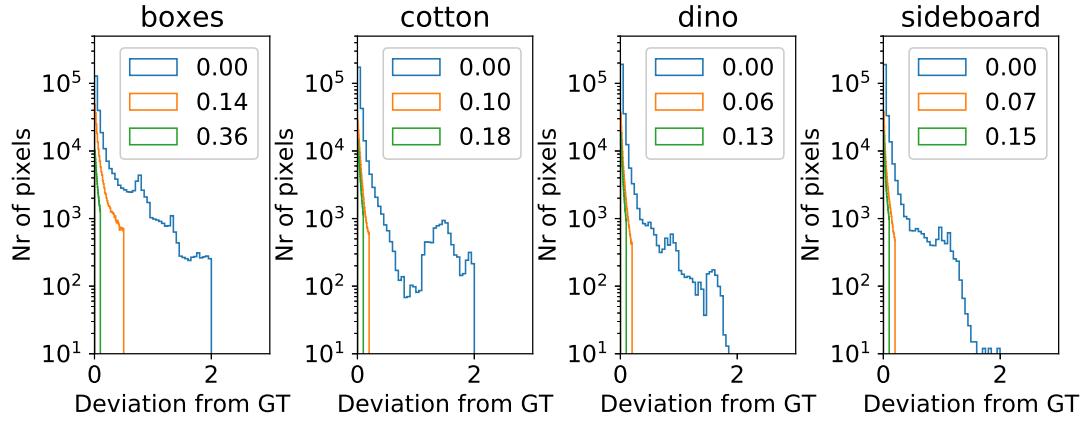


Figure B.5: The error histogram for different masks supposed one has found the ideal mask . The labeling indicates the fraction of masked out pixels.

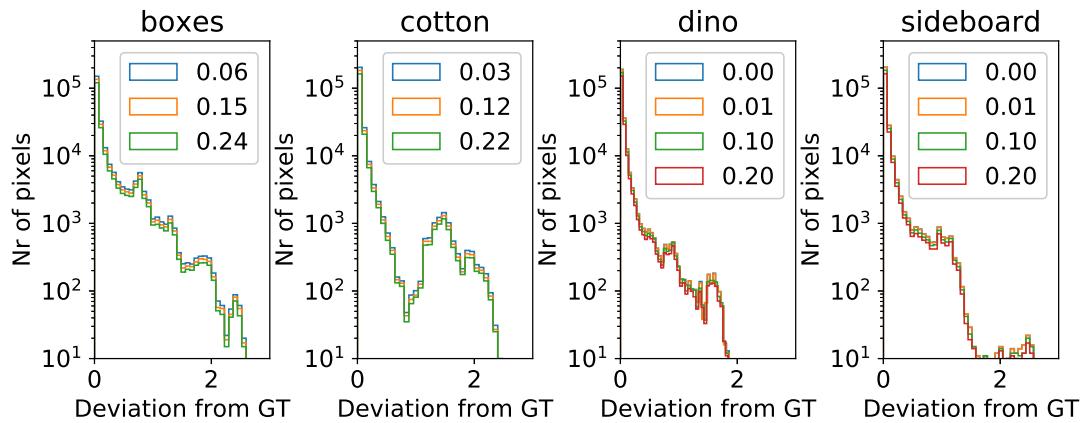
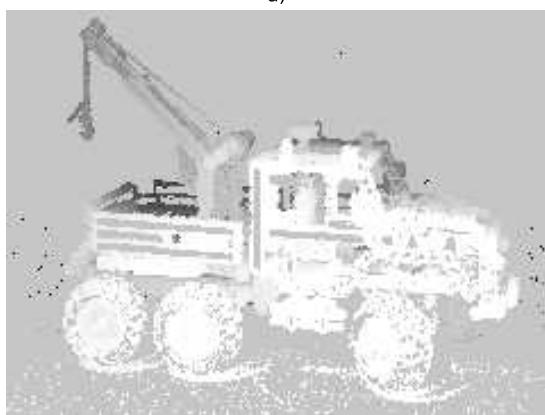
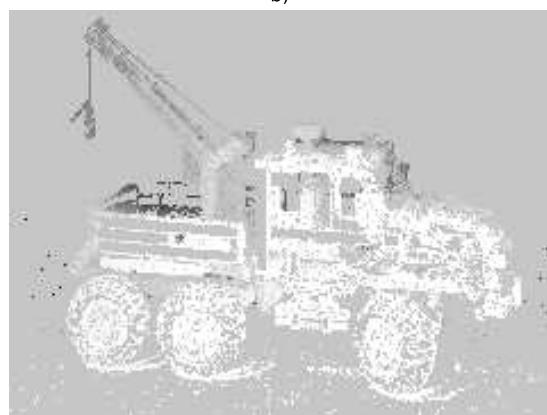


Figure B.6: The error histogram for different masks supposed the mask is completely random. The labeling indicates the fraction of masked out pixels.

a)



b)



C Lists

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Erklärung:

Ich versichere, dass ich diese Arbeit selbstständig verfasst habe und keine anderen als die angegebenen Quellen und Hilfsmittel benutzt habe.

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