3DCV 2023

| | | Grade | TP1 | TP2 | TP3 | TP4 | Mean TP(/5) | Exam |
|---------------|-----------|-------|-----|-----|-----|-----|-------------|------|
| | MVA | | | | | | | |
| BAILLET | Victor | 16.5 | 4.5 | 4.5 | 4.5 | 5 | 4.6 | 13.5 |
| BORNE-PONS | Paul | 14 | 4.5 | 5 | 5 | 4.5 | 4.8 | 7 |
| DE SENNEVILLE | Adhémar | 12.5 | 5 | 3.5 | 4.5 | 4.5 | 4.4 | 5 |
| EVE | Célestin | 15.5 | 5 | 3 | 4 | 5 | 4.3 | 13.5 |
| GAUILLARD | Emma | 13.5 | 4.5 | 4.5 | 5 | 4 | 4.5 | 6.5 |
| GOLLIER | Maxence | 13 | 4.5 | 5 | 2 | 3 | 3.6 | 11 |
| JANVIER | Maya | 14 | 4.5 | 4.5 | 5 | 4 | 4.5 | 8 |
| JAUMAIN | Pierre | 12 | 3 | 5 | 5 | 4 | 4.3 | 5 |
| LABIAD | Ismall | 14.5 | 4 | 4 | 4.5 | 5 | 4.4 | 10.5 |
| LAFENETRE | Jamy | 14.5 | 4 | 4 | 4.5 | 4.5 | 4.3 | 11 |
| MARENGO | Matteo | 14 | 5 | 4.5 | 4.5 | 4 | 4.5 | 8.5 |
| MELO | Sébastien | 15 | 5 | 2 | 4 | 4.5 | 3.9 | 14 |
| POLLET | Florent | 13.5 | 5 | 4.5 | 4 | 4 | 4.4 | 8 |
| RANJBAR | Mahdi | | 4.5 | 1.5 | | | | |
| ROBERT | Hugo | 13.5 | 5 | 4.5 | 5 | 4 | 4.6 | 5.5 |
| TERRISSE | Theïlo | 16 | 5 | 5 | 5 | 5 | 5 | 9.5 |

| IMA | | Grade | TP1 | TP2 | TP3 | TP4 |
|---------------|------------------|-------|-----|-----|-----|-----|
| ARAOUI | Mohammed | 18 | 5 | 4.5 | 4 | 4.5 |
| BIENVENU | Marie | 16.5 | 5 | 4.5 | 4.5 | 2.5 |
| BIOSCA MARRON | Pau | 16 | 4 | 4.5 | 4.5 | 3 |
| CHEN | Yunqi | 10.5 | 4.5 | 4.5 | 0 | 1.5 |
| FODIL | Zine-Eddine | 18 | 4.5 | 4.5 | 4 | 5 |
| GUO | Youheng | 18 | 4.5 | 4.5 | 4.5 | 4.5 |
| LEONELLI | Caterina | 14.5 | 4.5 | 1 | 4 | 5 |
| LIU | Wenda | 13.5 | 4.5 | 4.5 | 3 | 1.5 |
| MADRAU | Maxime | 14.5 | 3 | 4 | 3.5 | 4 |
| MELLUL | Yona | 19.5 | 5 | 4.5 | 5 | 5 |
| MOEINI | Mina | 12 | 4 | 2 | 4.5 | 1.5 |
| MUSTROPH | Henryk Alexander | 5.5 | 2 | 1.5 | 1 | 1 |
| OMIDI | Mojan | 18 | 4.5 | 4.5 | 4 | 5 |
| PORTE | Leo | 12 | 5 | 1.5 | 3.5 | 2 |
| RICHARD | Tom | 8 | 3 | 1 | 3 | 1 |

3DCV 2023

| RIFFAUD | Hugo | 16.5 | 5 | 4.5 | 3 | 4 |
|------------|---------------|------|-----|-----|-----|-----|
| SADREDDINI | Kimia | 12 | 4 | 4.5 | 0 | 3.5 |
| SCHADE | Oskar Hartvig | 10.5 | 3.5 | 1.5 | 4.5 | 1 |
| THOMAS | Guillaume | 6.5 | 2.5 | 1 | 2 | 1 |
| XI | Yuxuan | 17.5 | 4.5 | 4.5 | 4 | 4.5 |
| YANG | Kai | 13 | 4.5 | 0 | 4 | 4.5 |
| ZHANG | Jacques | 9 | 4.5 | 2 | 2.5 | 0 |

| Audito | ors | Grade | TP1 | TP2 | TP3 | TP4 | Mean TP(/5) | Exam |
|-----------------|-------|-------|-----|-----|-----|-----|-------------|------|
| ADJEVI-NEGLOKPE | Ambre | 12.5 | 3 | 4 | 3.5 | 2 | _ | 4.5 |
| ZEINATY | Paul | 18 | 5 | 5 | 3.5 | 4.5 | _ | |

Panorama

Panorama(/5)

MVA

| | * * * | | |
|---------------|-----------|-----|---|
| BAILLET | Victor | 4.5 | Good work overall, but careful, 3 components for Color so at line 27 the loop condition must be I<3, not I<4. |
| BORNE-PONS | Paul | 4.5 | see comments for Emma Gauillard. Good for adding Doxygen comments. |
| DE SENNEVILLE | Adhémar | 5 | Great job! |
| EVE | Célestin | 5 | Good, code clean and tidy. |
| GAUILLARD | Emma | 4.5 | When pulling, the translation by (x0,y0) must be applied before the homography, not after. Do not recompute inverse of H at each iteration. Detecting overlap with white color is not the best idea. |
| GOLLIER | Maxence | 4.5 | Everything good, except the missing transparency at overlapping pixels. |
| JANVIER | Maya | 4.5 | see comments for Emma Gauillard. |
| JAUMAIN | Pierre | 3 | Disappointing results are due to one main problem: signs of B are wrong (should be +, not -). When fetching pixel in I1, point should be translated before applying the homography, not after. White pixel has values 255, not 0, anyway that is not the best way to check overlap. Do not recompute inverse of H at each pixel. Good for your experimental effort to assess the quality. |
| LABIAD | Ismall | 4 | You applied the push method, which yields missing pixels. Moreover detecting overlop through white color is not optimal. |
| LAFENETRE | Jamy | 4 | Your panorama looks fine because the upper left corner of I1 coincides with the one of the panorama, but it does not stand in general. |
| MARENGO | Matteo | 5 | Good, though the code for blending could have been shorter with factorization. |
| MELO | Sébastien | 5 | Good, though the clicking part could have been more friendly with anyGetMouse. |
| POLLET | Florent | 5 | Good job! |
| RANJBAR | Mahdi | 4.5 | It is somewhat wasteful to recompute the inverse of H at each pixel. Detection of overlap by white pixel is not quite satisfying. |
| ROBERT | Hugo | 5 | Good, but don't compute twice inverseH*pt2. |
| TERRISSE | Theïlo | 5 | Exactly as requested! |
| | | | |

IMA

| ARAOUI | Mohammed | 5 | Very good! However, take care of code presentation (indentation, mutlple consecutive blank lines with no logic). |
|---------------|----------|-----|--|
| BIENVENU | Marie | 5 | Very good! For the bilinear interpolation, you could have used directly Image::interpolate. |
| BIOSCA MARRON | Pau | 4 | Why is your point normalization after application of H commented out? It has to happen to get correct pixel coordinates. You did not implement transparency. |
| CHEN | Yunqi | 4.5 | Your transparency application in two steps could be simplified. |

Panorama

| FODIL | Zine-Eddine | 4.5 | The color distortion in the transparency is due to an overflow of color components, stored in unsigned char. |
|------------|-----------------|-----|--|
| GUO | Youheng | 4.5 | Transparency is missing. Compute the inverse of H outside the loop. |
| LEONELLI | Caterina | 4.5 | The transparency on overlapping regions is missing, fine otherwise. |
| LIU | Wenda | 4.5 | Your way of applying the transparency is convoluted, it could be done directly. |
| MADRAU | Maxime | 3 | Results are not very good because you have some wrong values in matrix A: A(2i,7)=-y1*x2 and A(2i+1,6)=-x1*y2. Moreover, you forget to normalize by the third component when multiplying by the homography to retrieve pixel coordinates. |
| MELLUL | Yona | 5 | Good! NB: on a MacBook, clicking with two fingers simulates a right click. |
| MOEINI | Mina | 4 | Please send the source code next time, not only the report. What is missing is the transparency |
| MUSTROPH | Henryk Alexande | 2 | Something is wrong with the clicks, last points look suspect. In the panorama construction, you check if the point is inside image with the dimensions of I instead of I1 and I2. You do not implement the transparency. |
| OMIDI | Mojan | 4.5 | There is no blending or transparency, which would have helped assess the quality of the panorama. However, very good for trying on your own data. |
| PORTE | Leo | 5 | Very good! |
| RICHARD | Tom | 3 | To pull pixels from I1, you have to apply the inverse of H. Moreover, a translation by (x0,y0) must occur before, not after. Transparency is not handled correctly, white pixel is (255,255,255). |
| RIFFAUD | Hugo | 5 | Great, and congrats for having tested on your own data! |
| SADREDDINI | Kimia | 4 | Good, except the transparency is not implemented. |
| SCHADE | Oskar Hartvig | 3.5 | You did not get the interest of anyGetMouse, which indicates in which window the click occurred. To pull pixels from I2, you have to apply H, not its inverse. However, the bounding box was computed assuming I2 would stay fixed and I1 mapped. The homography should be applied to a single image, the other has just a |
| THOMAS | Guillaume | 2.5 | translation. Moreover, to get image coordinates, it is necessary to divide by the 3 rd |
| XI | Yuxuan | 4.5 | component. You did not apply the transparency effect, but otherwise it is fine. |
| YANG | Kai | 4.5 | Good, only the transparency on overlapping regions is missing. |
| ZHANG | Jacques | 4.5 | Good for testing on your own data. The number of clicked points should not be fixed in the program, right click should stop. Transparency is not impemented. (0,0) is a valid pixel. |

Auditors

Panorama

| ADJEVI-NEGLOKPE | Ambre | 3 | You should not expect unfilled coefficients of matrices to be zero. The code assumes that point (0,0) of I1 and panorama coincide. What you do is push pixels from I2 to the panorama, hence the missing pixels. Transparency has an overflow because channels are unsigned char and the addition may wrap around 0. |
|-----------------|-------|---|--|
| ZEINATY | Paul | 5 | Good, but some feedback would have been welcome while clicking. |

Fundamental

Fundamental(/5)

MVA

| | VA | | |
|---------------|-----------|-----|---|
| BAILLET | Victor | 4.5 | LS refinement is missing at the end. With thin SVD and A of size 8x9, we should have Vt of size 8x9, and the last column missing. Beware, 10e-3=1e-2=0.01. |
| BORNE-PONS | Paul | 5 | Very good, but the two functions for 8-point and n-point algorithms could be merged. |
| DE SENNEVILLE | Adhémar | 3.5 | Normalizing the matches inplace prevents their display in main function. You should not compute the SVD of A^t A but the one of A. In the refinement, you leave the last row of A undefined because of a loop bound off by 1. The user interaction is cumbersome because two clicks are expected. Good for trying on custom data. |
| EVE | Célestin | 3 | The vector inliers is not cleared at each iteration, in some runs I had more inliers than matches! No LS refinement at the end. |
| GAUILLARD | Emma | 4.5 | By default, the svd function should compute the thin SVD, hence Vt would be 8x9 and the last row is missing. Because of a bug, it computes the full SVD, hence you are spared. Niter may be become -infty if 1-(m/n)^8==1 (too few inliers). |
| GOLLIER | Maxence | 5 | Very good, though the fixed min number of inliers to catch the numerical problem could be better handled. |
| JANVIER | Maya | 4.5 | You should be careful, unset coefficients of Vector and Matrix are not initialized and may have random values. The formula for Niter update may suffer from numerical problem when the proportion of inliers is small (typically less than 10%). |
| JAUMAIN | Pierre | 5 | Good, though repeating very similar code for the minimal solver and the LS is not great. |
| LABIAD | Ismall | 4 | You should not compute A^t A, the SVD of A has the same V and it is numerically more favorable. After RANSAC, a least square minimization should be performed. |
| LAFENETRE | Jamy | 4 | This is good work, unfortunately results are disappointing due to a dumb mistake: y2 is computed from matches_sample[i].y1. Niter needs an update only when the number of inliers increases, not at each iteration. Very good for the gradient descent (which is not stochastic, by the way), but would be better with parameterization by the normalized F matrix. |
| MARENGO | Matteo | 4.5 | There can be a numerical problem with 1-(m/n)^8=1, hence a division by 0 in the formula for the update of Niter. The minimal solver and LS could have been merged. |
| MELO | Sébastien | 2 | Bad results because the vectors u and v have their last component uninitialized (should be 1) and some coefficients of N are left also uninitialized. Also 10e-3=0.01, you meant 1e-3, which is 0.001. Applying blindly the formula for Niter may lead to division by 0 because of numerical problems. |
| POLLET | Florent | 4.5 | Nothing wrong with your code, however Imagine++'s doc says that by default svd computes the thin SVD, hence if A is 8x9, Vt should be 8x9 and the last column of V would be missing. It happens there is a bug and the full SVD is always computed. |

Fundamental

| RANJBAR | Mahdi | 1.5 | You don't compute the denominator for the point-line distance, and at the end compare with the squared maximum distance. The number of inliers is completely wrong (I got all matches). |
|---------------|-------------|-----|--|
| ROBERT | Hugo | 4.5 | Good! However, you should not compute A^t A, the SVD of A is enough and numerically better. |
| TERRISSE | Theïlo | 5 | Very good! |
| IMA | A | | |
| ARAOUI | Mohammed | 4.5 | Good but the update formula for Niter should be protected from numerical problems (potential division by 0). Indentation of the code is disastrous. Notice it makes no sense to try on panorama images, there is no translation. |
| BIENVENU | Marie | 4.5 | Very good, except the formula for update of Niter must be protected from numerical errors. And you inverted F and its transpose in displayEpipolar! |
| BIOSCA MARRON | Pau | 4.5 | Very good! LS estimation should still be performed after RANSAC and the formula for Niter update but be protected from numerical approximations. |
| CHEN | Yunqi | 4.5 | Very good, but refinement has a problem: the loop for filling A uses only the first 8 points. |
| FODIL | Zine-Eddine | 4.5 | No LS refinement. Moreover, if some model has too few inliers, you can get 1-(m/n)^8=1 and a division by 0, hence Niter=-infty and the loop stops. |
| GUO | Youheng | 4.5 | Good, would have been even better with least square refinement using all inliers. |
| LEONELLI | Caterina | 1 | Plagiarism Sadreddini for the code. Fortunately, you provided good experimental evaluation after my complaint. |

Missing LS at the end of RANSAC. Good otherwise, though the code is a bit LIU Wenda 4.5 verbose. No LS after RANSAC. The formula for Niter must be applied with precautions. When MADRAU Maxime 4 clicking on left image, the y of right point is wrong since its logical x is w, not 2*w. The LS refinement is not applied, and should actually involve all inliers, not just 9. MELLUL Yona 4.5 Off-diagonal coefficients of N are not initialized. Two fatal bugs: (1) line[2] is missing in formula for point-line distance. (2) Two 2 MOEINI Mina counters, samplingN and samplingNb, one is incremented but the other is tested. Your normalization of points makes no sense, they should be all normalized with the same factor. A(8,8) is left uninitialized. What is the role of x values and y values MUSTROPH Henryk Alexande 1.5 (which are unitialized, by the way)?

4.5

1.5

Mojan

Leo

OMIDI

PORTE

inliers for update of Niter is too conservative.

normalized space, the threshold distMax must be adapted.

Good job! The only missing part is the LS refinement. Also, your threshold at 50%

V(V.nrow(),...) is always out of bounds, you mean V(V.nrow()-1,...). Anyway, you

apply thin SVD and V is 9x8, V9 is missing. If you compute line-point distance in

Fundamental

| Tom | 1 | There is a confusion with H computation. Here, matrix A is 8x9 and each match gives a single equation. |
|---------------|---|--|
| Hugo | 4.5 | Good, only LS estimation is missing at the end of RANSAC. |
| Kimia | 4.5 | Good! What is missing is the LS refinement with all inliers. The safety against numerical problem in Niter update is not flexible with its fixed threshold. |
| Oskar Hartvig | 1.5 | When refining the estimation with all inliers, matrix A has the wrong size and does not require the addition of a line of zeros. There is no enforcement of rank-2 constraint. When drawing lines, line should be displayed on right part when click on left part, and thickness is wrong. |
| Guillaume | 1 | Confusiont with computation of H. For F, each match gives a single equation, hence a single row of matrix A. |
| Yuxuan | 4.5 | Missing LS refinement at the end of RANSAC and lack of precaution when applying the update formula for Niter. |
| Kai | 0 | Plagiarism |
| Jacques | 2 | Several bugs: A(i,2)=x1, not x2. Off-diagonal coefficients of N not initialized. Reshaping Vt as F actually computes its transpose. |
| | Hugo Kimia Oskar Hartvig Guillaume Yuxuan Kai | Hugo 4.5 Kimia 4.5 Oskar Hartvig 1.5 Guillaume 1 Yuxuan 4.5 Kai 0 |

Auditors

| ADJEVI-NEGLOKPE | Ambre | 4 | If A is 8x9, Vt should be also 8x9 if thin SVD is applied (default for function svd), and the last column of V is missing. Beware, 10e-3=0.01. The code for displayEpipolar is hardly readable and i_min1 is left unitialized. Precautions should be taken when applying the update for Niter. |
|-----------------|-------|---|--|
| ZEINATY | Paul | 5 | Very good! |

Seeds(/5)

MVA

| IVIV | * * | | |
|---------------|-----------|-----|--|
| BAILLET | Victor | 4.5 | bestDisparity may be assigned unitialized to pixels. It happens at the left border, where every patch overflows the image. |
| BORNE-PONS | Paul | 5 | Very good! |
| DE SENNEVILLE | Adhémar | 4.5 | Good report! d_end should be excluded from the loop, as it would overflow if dmax were positive. In propagate, some pixels are assigned disparity 0 because they have no valid patch. Clipping the best disparity is not as good as just ignoring invalid disparities. |
| EVE | Célestin | 4 | In find_seeds, pixels with no valid patch are assigned dBest, which is unitialized. Same with ddBest in progagate. The patch may also overflow to the right if dmax were positive. No check of disparity bounds during propagation. |
| GAUILLARD | Emma | 5 | Very good! |
| GOLLIER | Maxence | 2 | Results are particularly bad due to a dumb bug: in correl, denominator is sqrt(var1)*sart(var2), not with pixel1 and pixel2! Pixels with no valid patch get assigned an uninitialized disparity. Right overflow of patch should also be taken care of. During propagation, some pixels have no valid patch and should not get a disparity. Clamping to [dmin,dmax] is worse than ignoring bad disparities. |
| JANVIER | Maya | 5 | Very good! |
| JAUMAIN | Pierre | 5 | Very good work! Just a little problem in propagate, where uninitialized bestDisparity is assigned to pixels with zero valid patch. |
| LABIAD | Ismall | 4.5 | During propagation, clamping to [dmin,dmax] is not as good as just ignoring wrong disparities. Moreover, it assigns dmin to pixels that have no valid patch. Notice also that the right overflow check should be stricter with >=im2.width(), not >. |
| LAFENETRE | Jamy | 4.5 | Good. No check for disparity bounds during propagation. The median filter does not help that much as wrong pixels are not initial seeds and are bound to have close disparity as their neighbors. |
| MARENGO | Matteo | 4.5 | Clamping the disparity to [dmin,dmax] after the minimization is not as good as just ignoring disparities out or range. Overflow of a patch to the right should also be guarded against, since in all generality dmax could be positive (no control over the crop or the rectification). |
| MELO | Sébastien | 4 | In find_seeds, pixels on the left with no valid patch are assigned argmaxd, which is unitialized. The same for pixels whose best NCC is negative. Overflow on the right should be checked with im2.width() (not im1), and the inegality should be >=. No check of disparity range [dmin,dmax] in propgate. Idem, argmaxd may be assigned while uninitialized. |

| POLLET | Florent | 4 | During propagation, out of bounds checks must still be performed. Clamping the disparity between [dmin,dmax] is worse than just ignoring wrong disparities. Finally, some pixels may have no valid patch and should be assigned a disparity. |
|---------------|-------------|-----|---|
| RANJBAR | Mahdi | | |
| ROBERT | Hugo | 5 | Very good! Still, during propagation, we should accept pixels with bestNCC negative. A better test is bestNCC>-1.0f. |
| TERRISSE | Theïlo | 5 | Very good, but the overflow should be checked wrt to im2.width() during propagation, not im1. |
| IM | A | | |
| ARAOUI | Mohammed | 4 | In find_seeds, initializing Max_ncc as 0 is a mistake, as pixels with no valid patch are still assigned Best_disparity, which is uninitialized. In the propagation, bestDisparity would rather never be tested if outside [dmin,dmax]. It is wasteful to add several times the same seed. |
| BIENVENU | Marie | 4.5 | Good work. During propagation, bounds [dmin,dmax] must still be observed. In find_seeds, the checks for bounds are useless, this is already ensured by the bounds of the for loop. |
| BIOSCA MARRON | Pau | 4.5 | In find_seeds, check for overflow to the right should be performed, it can happen with dmax>=0. In propagation, clamping the best disparity to [dmin,dmax] is not the best solution. Some pixels to the left with no valid patch still get bestDisparity, which is dmax due to clamping and dmax<0. |
| CHEN | Yunqi | 0 | Plagiarsim of Wenda Liu's work. The few changes introduced a bug: variable dp should vary around s.d, not s.x |
| FODIL | Zine-Eddine | 4 | In find_seeds, you are too conservative: if some disparities but not all in [dmin,dmax] yield a valid patch, they should be tested. Overflow to the right should be checked (could happen if dmax>0). During propagation, it is not bestCorr that should be compared to dmin and dmax, but dispar |
| GUO | Youheng | 4.5 | During propagation, clamping to [dmin,dmax] should not be performed after testing the 3 disparities, it is better to discard the ones outside the interval. Also, some pixels with no valid patch are still assigned dmin. |
| LEONELLI | Caterina | 4 | During propagation, some pixels have no valid patch associated, they should not be assigned a disparity. It is best to not consider disparities outside [dmin,dmax] then to clamp the result. There could be some overflow of the patch to the right (with different dmax), this should be checked. In functions correl and sum, you must go to win inclusive, otherwise the |
| LIU | Wenda | 3 | normalization in ccorrel is wrong. Moreover, in these functions you implicitly do 0-padding, which produces articial patches. It is better to not consider overflowing patches. In propagate, variable best_cur must be reset inside the for(i) loop, not |

before.

| MADRAU | Maxime | 3.5 | In find_seeds, a disparity is still set even though the NCC does not exceed nccSeed, which interferes with visualization of the seeds. During propagation, no check of patch overflow is performed. In non-Release mode, an assert stops the program. |
|------------|-----------------|-----|--|
| MELLUL | Yona | 5 | Very good! |
| MOEINI | Mina | 4.5 | In propagate, clamping to [dmin,dmax] a posteriori is not the best solution. Some pixels with no valid patch have best uninitialized compared to dmin. |
| MUSTROPH | Henryk Alexande | 1 | I think you have not understood at all the principle. For each possible disparity, we compute the NCC and finally select the argument of the maximum. |
| OMIDI | Mojan | 4 | In find_seeds, the test is too conservative, some disparities in [dmin,dmax] may be testable with all being so. During the propagation, you test also s.d+2. Clamping the disparity afterwards is not the best. When you test with other images, you must adust dmin and dmax. |
| PORTE | Leo | 3.5 | nccSeed has no role in propagation. There should also be a problem because at the bottom and the right you have disparities whereas the patch does not fit within the image. |
| RICHARD | Tom | 3 | Disparities should be tested only within the range [dmin,dmax]. A bit too conservative in the propagation, some pixels have one or two valid patches wihin [x+dp-1,x+dp+1]. Fatal mistake in propagate: conditions for if with = instead of ==. Infinite loop if new seeds are not marked during propagation. |
| RIFFAUD | Hugo | 3 | Some dumb bugs that ruin the results: (1) in correl, division by sqrt(cpt_im2)*sqrt(cpt_im2) instead of sqrt(cpt_im1)*sqrt(cpt_im2), (2) in sum, im(i+v,j+v) instead of im(i+u,j+v). Check for overflow of patch to the right should be performed. During propagation, for some pixels none of the three disparities yields an NCC, so best_disparity is compared to dmin without being set. |
| SADREDDINI | Kimia | 0 | Plagiarism. This is a copy-paste of a previous year's code. |
| SCHADE | Oskar Hartvig | 4.5 | During propagation, no check of disparity within [dmin,dmax] is performed. Also, some pixels with no valid patch still have a disparity assigned. Good otherwise. |
| THOMAS | Guillaume | 2 | You are mixing dmax, which is the maximal possible disparity, with an NCC. The range [dmin,dmax] is the set of possible disparities. Overflow on the right would be possible if dMax were positive. |
| XI | Yuxuan | 4 | In find_seeds, if some but not all of the disparities in [dmin,dmax] yield a valid patch, they should be tested. Overflow of a patch to the right should also be tested. Clamping to [dmin,dmax] is not the best. Moreover, for pixels with no valid patch, best_d is uninitialized and still compared to dmin and dmax. |
| YANG | Kai | 4 | In find_seeds, pixels with no valid patch are still assigned a disparity, which is actually bestD, left uninitialized. Similar problem in propagate. Patch inclusion should also check the right boundary. |

| ZHANG | Jacques | 2.5 | nccSeed is unused in find_seeds. In the propagation, why is disps>s.d significant? We test s.d-1 and s.d, which are perfectly fine. When you test on other images, dMin and dMax must be adjusted. |
|-----------------|---------|-----|---|
| Auditors | | | |
| ADJEVI-NEGLOKPE | Ambre | 3.5 | Why divide the grey levels by 256 (through and RGB <float> on top of that)? NCC is invariant to such a factor. Disparities should be restriced to the interval [dmin,dmax]. In propagate, x+d+win should be compared to im2.width(), not im1.</float> |
| ZEINATY | Paul | 3.5 | In find_seeds, some pixels have their best NCC negative. In that case, since bestNcc is 0, they are assigned bestDisp, which is whatever happened before. Moreover, the right overflow check should be x+d>= im2.width()-win. In propagate, simliar problem and the interval [dmin,dmax] is not used. |

GCDisparity

GCDisparity(/5)

MVA

| BAILLET | Victor | 5 | Great job! |
|---------------|-----------|-----|---|
| BORNE-PONS | Paul | 4.5 | Good, though you add lambda twice by using 4 neighbors instead of 2. By putting INF as terminal weights, you are excluding dmin and dmax. Zncc at disparity d should feed the weight from d-1 to d, not d to d+1. Great gif animations! |
| DE SENNEVILLE | Adhémar | 4.5 | You forgot to put weight to y+1, hence the apparent horizontal stripes in the results. By putting infinite weight to sink, you are forbidding disparity dmax, which should actually depend on zncc. It you authorize it back, decode_graph should be adapted to it. Great report! |
| EVE | Célestin | 5 | Very good! |
| GAUILLARD | Emma | 4 | Disparities dmin and dmax are excluded as you put infinite weights to terminal nodes. Using the four neighbors amounts to adding twice lambda. |
| GOLLIER | Maxence | 3 | In call to zncc, disparity must not be multiplied by zoom as it is computed in original images. Put INF as weight if the patch is not in the image. In decode_graph, the result should be dmin+d, no normalization has to be performed. |
| JANVIER | Мауа | 4 | Putting (INF,0) and (0,INF) as terminal weights excludes disparities dmin and dmax. ZNCC at dmin+d should be used with nodes at d-1 and d, not d and d+1. If you consider 4 neighbors, you are adding twice lambda to the edges. In decode_graph, the test d <nd call="" good="" precede="" report.<="" should="" td="" the="" to="" what_segment.=""></nd> |
| JAUMAIN | Pierre | 4 | You are forbidding dmin and dmax as disparity because you put infinite weight from the source and to the sink. All is fine otherwise. The vertical streaks in the result are due to adding twice lambda for vertical smoothness but no horizontal smoothness. |
| LABIAD | Ismall | 5 | Excellent, both code and report! |
| LAFENETRE | Jamy | 4.5 | Very good, but considering the 4 neighbors amounts to adding lambda twice because (x,x+1) are neighbors and (x+1,x) also. All calls to zncc must be preceded by checks that the patch is within the image. The |
| MARENGO | Matteo | 4 | weight to sink should depend on zncc with disparity dmax. decode_graph does not detect correctly disparity dmax (all nodes should be linked to the source). Good |
| MELO | Sébastien | 4.5 | report. Good, just a minor mistake: you are still using the zncc at dmax-1 to feed the weight to sink. |
| POLLET | Florent | 4 | Function zncc cannot be called before checking the patch is within the image. In Release mode, assertions are disabled and it can go unnoticed. The weight for d=0 should correspond to disparity dmin, not dmin+1. Good report. |
| RANJBAR | Mahdi | | |
| ROBERT | Hugo | 4 | You cannot call zncc without guarding against patch overflow (assertion triggered in non-Release mode). In decode_graph, you would rather put dmax by default, as if the cut happens just before the sink, this indicates dmax. Very good report! |

GCDisparity

| TERRISSE | Theïlo | 5 | Excellent, clean code and nice report! |
|---------------|-----------------|-----|---|
| IM | A | | |
| ARAOUI | Mohammed | 4.5 | Good, but zncc at d should feed weight from (d-1) to d, not d to (d+1). Problem in decode_graph if the disparity is actually dmax. |
| BIENVENU | Marie | 2.5 | There is some confusion about indices which are incremented by 1. This being fixed, decode_graph should add dmin and put dmax to unassigned pixels. |
| BIOSCA MARRON | Pau | 3 | You do not add K. Terminal weights are wrong. Decoding should put dmin+d. |
| CHEN | Yunqi | 1.5 | It does not work because you put INF weights to terminal links, which prevents cutting them. |
| FODIL | Zine-Eddine | 5 | Very good! Still, you should not rely on d>=0 and ignore the possibility of left overflow of patch. Beware that computation time is with zoom=2. |
| GUO | Youheng | 4.5 | Good overall, but you forget to add K to the cost, overflow check should not multiply d+dmin by zoom and weight to sink should depend on ZNCC at dmax, not dmax-1. Good! However, you should put weight INF instead of wcc for overflowing patch. |
| LEONELLI | Caterina | 5 | When testing on kitchen images, results cannot be good if the images are not |
| LIU | Wenda | 1.5 | rectified. Your weights to terminals are wrong. Why put (nd+20)*wcc? |
| MADRAU | Maxime | 4 | The code must not call zncc unguarded against patch overflow, which happens here. Otherwise, everything fine. |
| MELLUL | Yona | 5 | Very good! Do not ignore the possibility of left overflow in case d<0. |
| MOEINI | Mina | 1.5 | You cannot compute zncc without checking the patch is within the image. In decode_graph, you must put D=dmin+d, the integer disparity. |
| MUSTROPH | Henryk Alexande | 1 | Your node-number formula is wrong, x*y should not appear inside. You do not use ZNCC as unary weights. Why do you have a double for loop on depth? |
| OMIDI | Mojan | 5 | Very good! When testing on other images, dmin and dmax must be adjusted. |
| PORTE | Leo | 2 | You forgot to add K to the weight, which changes everything. When decoding, you must add dmin. Your function rho is not exactly the proposed one. |
| RICHARD | Tom | 1 | Everything should be changed: node numbers are wrong, weight does not care of zoom factor, decoding should just assign d. |
| RIFFAUD | Hugo | 4 | ZNCC with d must be used to link edge d-1 to d, not d to d+1. In decode_graph, result should be dmin+d, not d. |
| SADREDDINI | Kimia | 3.5 | Terminal weights are wrong: should be (w,0) for d=0 and (0,w) for d=nd-1. You do not add K. Weight should be INF for out-of-bounds patch. You did not understand the role of zoom: nx and ny already take into account the |
| SCHADE | Oskar Hartvig | 1 | zoom factor and the window radius. decode_graphs should be fine except that dmax must be excluded as disparity dmax is encoded by the cut occurring just before the sink. |

GCDisparity

| THOMAS | Guillaume | 1 | In build_graph, x and y should start from 0 and stop at nx and ny, win is already taken care of. Terminal node weights are wrong, do not put infinity. neighbor_node_id is wrong, should be with d-1. Moreover, the zncc must be mapped through the function sqrt(1-x) as weight. |
|--------|-----------|-----|---|
| XI | Yuxuan | 4.5 | The weight to the sink should depend on zncc at dmax=d+nd, not d+nd-1. Very good otherwise. When comparing on other images, dmin and dmax must be adjusted. |
| YANG | Kai | 4.5 | You should add K to the weights. You should check also overflow of patch by the left side (in case dmin<0). Default value for C in build_graph should be INF. |
| ZHANG | Jacques | | |
| | Auditors | | |
| | | | You do not take into account that there is a zoom factor to speed up the |

| ADJEVI-NEGLOKPE | Ambre | 2 | You do not take into account that there is a zoom factor to speed up the computation, so the real x and y are x*zoom+win and y*zoom+win. Calls to zncc must be done only after checking the patch is within the image. |
|-----------------|-------|-----|--|
| ZEINATY | Paul | 4.5 | Good! Still, the weight of (nd-1) to the sink should depend on zncc at disparity dmax, not dmax-1. The x and y increments are not very intuitive. Good report. |