

Supplementary Material for Adaptive Assignment for Geometry Aware Local Feature Matching

Anonymous CVPR submission

Paper ID 5298

1. Implementation Details

1.1. Architecture of Co-visible Area Segmentation Module

In order to obtain the co-visible area probability map, we borrowed the structure of DETR [3] and used a query to perform regression on the feature map. The specific network architecture is shown in the Fig 1. Firstly, the spatial attention map $F_{attn}^i \in \mathbb{R}^{1 \times h \times w}$ by the dot product operation of $Q^i \in \mathbb{R}^{1 \times 1 \times C}$ and $F_{1/8}^{i2} \in \mathbb{R}^{C \times h \times w}$, and then perform element-wise multiplication of F_{attn}^i and $F_{1/8}^{i2}$, followed by a shortcut connection to obtain $F_{co}^i \in \mathbb{R}^{C \times h \times w}$. Finally, a simple block with two convolution layers are used to obtain the co-visible area probability map, where the first convolution is followed by a ReLU activation and the second convolution is followed by a Sigmoid function.

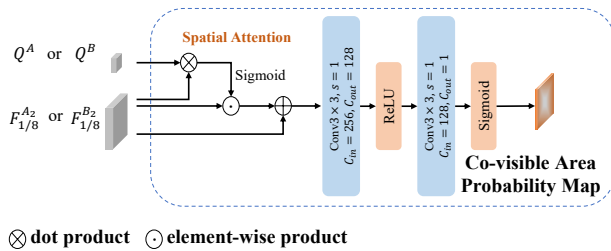


Figure 1. Architecture of co-visible area segmentation module.

1.2. Refine Network for SuperGlue

As stated in the main text, our adaptive assignment and sub-pixel refine module can be treated as a refinement network for other matching methods, such as SuperGlue [12]. Here, we will present in detail how to use our method as a refinement network for SuperGlue. To perform adaptive assignment, we simply remove the mutual nearest neighbours constraint, and obtain proposal matches by applying $\arg\max$ operation on each of the two dimensions of the matching matrix generated from SuperGlue, and then

filtering by adding a confidence threshold. The matching matrix can be the result of multiple iterations of the skinhorn algorithm, or the matrix obtained by decomposing the last iteration of the skinhorn algorithm. After obtaining the proposal matches, we sample the patch features $F^A, F^B \in \mathbb{R}^{n \times c \times w \times w}$ at the corresponding positions on the descriptor feature maps generated by SuperPoint [6] and feed them into our one-to-one refine module (Section 3.3 of the main text) to achieve scale alignment and sub-pixel location regression.

2. More Experiments

2.1. Additional Evaluation Metrics for HPatches

Metrics. Since the homography estimation accuracy contains the effect of the OpenCV-RANSAC, we use Mean Matching Accuracy (MMA) on the HPatches [1] dataset to evaluate different methods. We use the ratio of correctly matched features within thresholds of 1, 2, and 3 pixels, respectively, and the maximum amount of matches is limited to 1024. **Results** in Tab.1 show that Adamatcher outperforms other methods in terms of matching accuracy. Since adaptive assignment eliminates the ambiguity of matching in supervision and inference, AdaMatcher is able to generate more accurate matches when the viewing angle changes.

Methods	All	Viewpoint
	MMA@1px / 2px / 3px	MMA@1px / 2px / 3px
SIFT [9]+HardNet [10]	0.460 / 0.718 / 0.828	0.383 / 0.671 / 0.800
KeyNet [2]+HardNet [10]	0.422 / 0.701 / 0.837	0.342 / 0.661 / 0.811
R2D2 [11]	0.334 / 0.611 / 0.751	0.273 / 0.566 / 0.699
SP [6]+SG [12]	0.367 / 0.685 / 0.827	0.315 / 0.654 / 0.815
SP [6]+SG [12]+Ada	0.386 / 0.702 / 0.839	0.348 / 0.685 / 0.830
LoFTR-OT [13]	0.593 / 0.814 / 0.893	0.461 / 0.731 / 0.836
LoFTR-DS [13]	0.613 / 0.830 / 0.902	0.495 / 0.759 / 0.853
AdaMatcher-LoFTR	0.628 / 0.845 / 0.914	0.540 / 0.798 / 0.882
QuadTree [14]	0.642 / 0.844 / 0.911	0.517 / 0.777 / 0.870
AdaMatcher-QuadTree	0.640 / 0.850 / 0.917	0.550 / 0.799 / 0.881
ASpanFormer [4]	0.658 / 0.856 / 0.920	0.539 / 0.795 / 0.881
AdaMatcher-ASpan	0.657 / 0.857 / 0.920	0.552 / 0.802 / 0.887

Table 1. MMA metrics on HPatches.

2.2. Results on YFCC100M

The YFCC100M [15] dataset is also used to conduct experiments to compare AdaMatcher with several baseline methods. To be fair, We use the same test pairs (a total of 4000 pairs) as in previous works [12, 16], using their evaluation metrics. The test set is derived from four selected landmark sequences, each sampling 1000 image pairs. All images are resized to 480×640 and all models are trained in MegaDepth [8]. The accuracy of pose estimation is measured by AUC under error thresholds (5° , 10° and 20°). The results are shown in Tab.2, where our methods all perform better than the corresponding baseline methods.

Methods	Pose Estimation AUC		
	@ 5°	@ 10°	@ 20°
LoFTR-DS [13]	43.06	62.21	77.26
AdaMatcher-LoFTR	44.06	63.04	77.61
ASpanFormer [4]	43.70	62.57	77.32
AdaMatcher-ASpan	43.93	62.92	77.54
QuadTree [14]	36.50	55.64	71.61
AdaMatcher-Quad	44.20	63.09	77.59

Table 2. The results of outdoor relative pose estimation on YFCC100M.

2.3. Indoor Pose Estimation

To validate the generalizability of different detector-free methods, we perform indoor pose estimation experiments on the ScanNet [5] dataset using models trained on the MegaDepth [8] dataset. We use the test split with 1500 image pairs following the experimental setting of [4, 12, 13]. To align with the existing methods [4, 13, 14], we resized all test images to 480×640 . We use the same evaluation protocols as in Sec. 2.2. As presented in Tab.3, AdaMatcher has a significant performance improvement on different baselines [4, 13, 14].

Methods	Pose Estimation AUC		
	@ 5°	@ 10°	@ 20°
LoFTR-OT [13]	15.46	31.28	47.87
LoFTR-DS [13]	17.26	33.93	50.16
AdaMatcher-LoFTR	18.60	35.00	50.75
ASpanFormer [4]	20.64	39.34	56.61
AdaMatcher-ASpan	21.33	39.93	56.69
QuadTree [14]	19.83	37.86	55.03
AdaMatcher-Quad	21.18	39.71	56.22

Table 3. The results of indoor relative pose estimation on ScanNet. All models are trained on MegaDepth dataset.

2.4. Computational Costs of Feature Interaction

We evaluate the computation and parameters between LoFTR’s feature interaction module [13] and our CFI module (using linear attention [7] as in LoFTR). The size of input tensor is $60 \times 80 \times 256$. As shown in Tab.4, compared to LoFTR’s feature interaction module (consisting of four sets of self- and cross-attention layers), our CFI module reduces about 38.79% of the computational costs and 14.29% of the parameters.

Method	Flops(G)	Param(MB)
LoFTR module	51.82	5.25
CFI	31.74	4.50

Table 4. Computational complexity of feature interaction module

3. D. Qualitative Results

We present more qualitative comparisons of AdaMatcher and baselines on Hpatches [1] dataset and MegaDepth [8] dataset. In Fig.2, we display inlier and outlier matches using different projection thresholds to compare the matching accuracy of different methods on the Hpatches dataset. Fig.3 presents more qualitative results on the MegaDepth [8] dataset and Fig.4 shows more qualitative results of the co-visible area estimation.

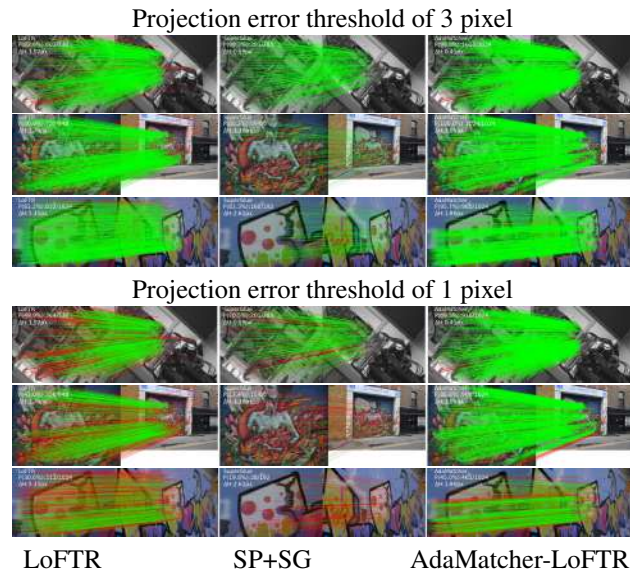


Figure 2. Qualitative image matches on Hpatches dataset. Matches with projection error less than the threshold are displayed in green, otherwise they are displayed in red.

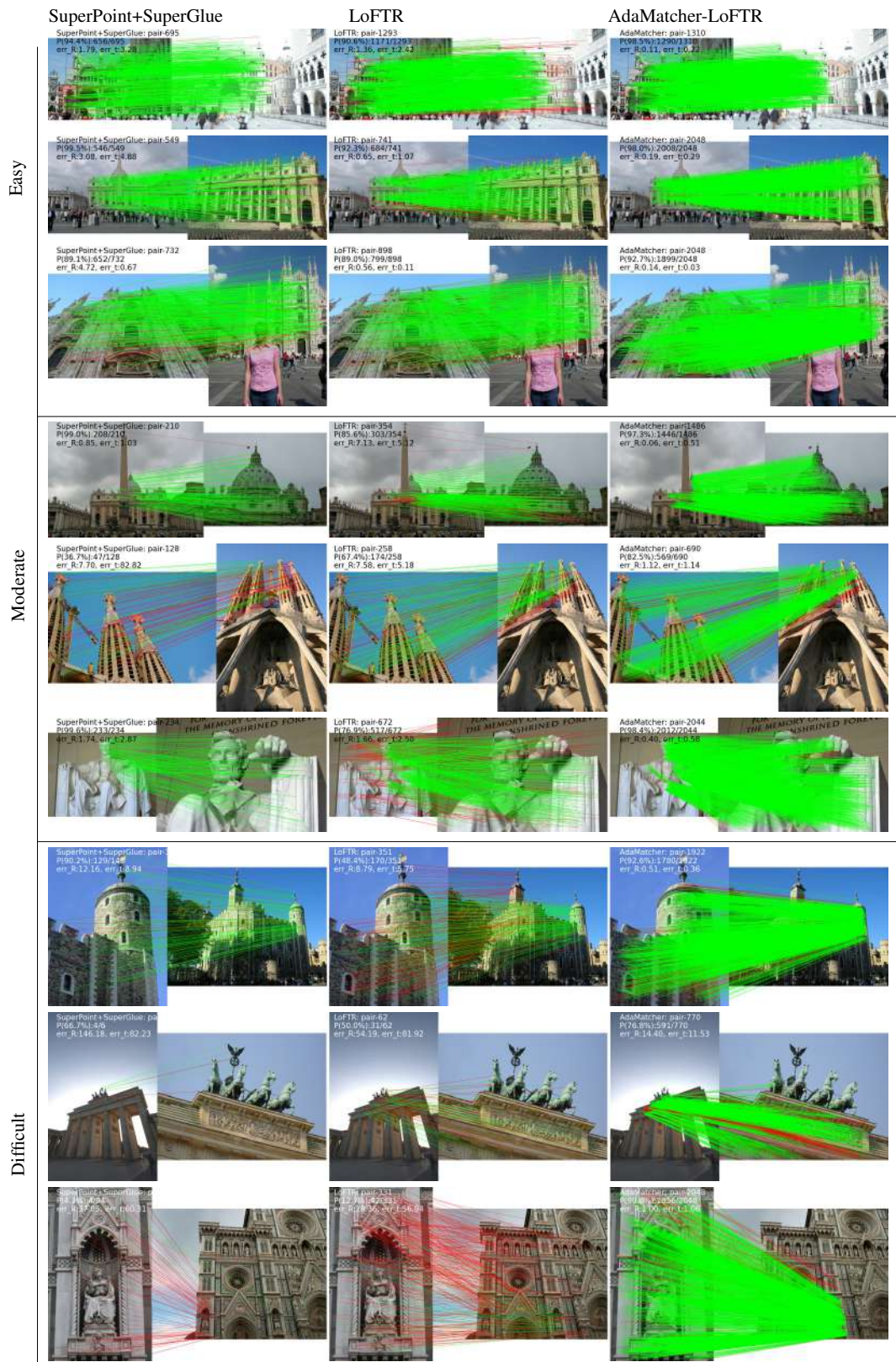


Figure 3. Qualitative image matches on MegaDepth dataset. Green indicates that epipolar error in normalized image coordinates is less than 1×10^{-4} , while red indicates that it is exceeded.

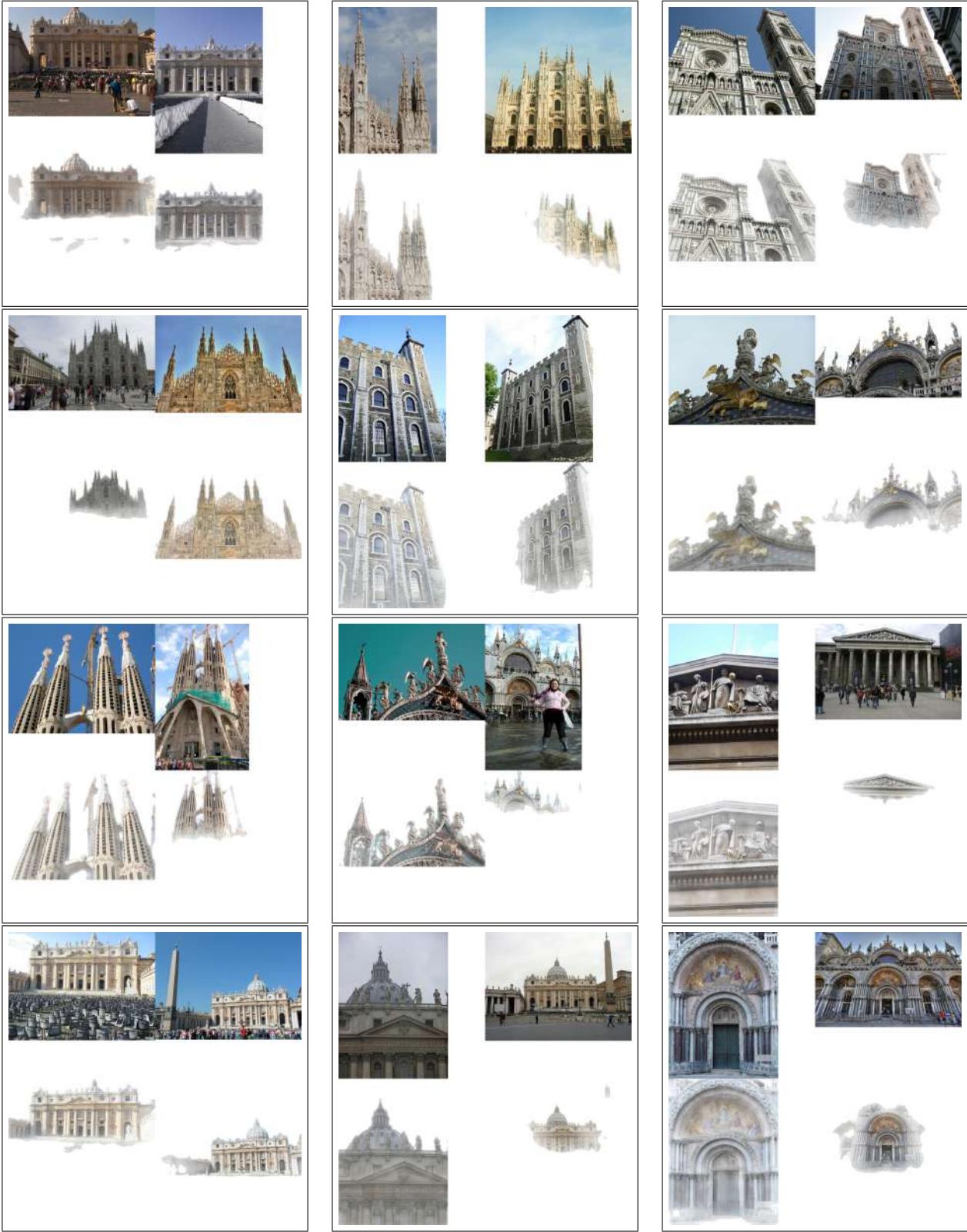


Figure 4. Qualitative co-visible area segmentation

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