Supplementary Material: Machine Vision Guided 3D Medical Image Compression for Efficient Transmission and Accurate Segmentation in the Clouds

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In this supplementary material, we introduce four mapping functions and perform performance comparison in Section 1. In Section 2, we present more results with compression rate of 40. In Section 3, we then provide the image by image segmentation results to further demonstrate that the improvement by our method is statistically significant.

1 Mapping Function Exploration

We explored four types of mapping functions—reciprocal mapping (RM) (depicted as 'ours' in our paper), linear mapping (LM), discrete mapping (DM) and power mapping (PM), to search the best mapping between statistic index (SI) and quantization step (QS). Then we compared segmentation accuracy and compression rates using these mapping methods in Section 1.4.

1.1 Linear Mapping

We first assumed the relationship between SI and QS obeyed a linear mapping (LM) function:

$$Q_n = a \cdot \delta_n + b, \quad s.t. \quad Q_{min} \le Q_n \le Q_{max}, \tag{1}$$

where Q_n is the quantization step at subband n, Q_{min} and Q_{max} are the smallest and largest QS, and a and b are the fitting parameters.

1.2 Discrete Mapping

We further developed a seemingly more intuitive discrete mapping (DM) method defined as:

$$Q_n = \begin{cases} Q_{min} & \delta_n > T_1 \\ 2 \cdot Q_{min} & T_2 < \delta_n \le T_1 \\ 4 \cdot Q_{min} & T_3 < \delta_n \le T_2 \\ Q_{max} & \delta_n \le T_3 \end{cases}$$

$$(2)$$

where T_1 , T_2 and T_3 indicate the average intervals between the largest and smallest δ_n .

Table 1: Segmentation results of four candidate mapping methods (LM, DM, PM and RM (depicted
as "ours" in our paper)) using DenseVoxNet and HVSMR2016 dataset.

		LM	DM	PM	RM (Ours)
	Dice	0.771 ± 0.0438	0.821 ± 0.043	$0.834 {\pm} 0.0366$	$0.834{\pm}0.0386$
Myocardium	${f Hausdorff}$	$36.823{\pm}10.079$	32.372 ± 8.551	31.012 ± 7.911	$31{\pm}7.940$
	\mathbf{ASD}	$0.822 {\pm} 0.846$	0.669 ± 0.761	$0.651 {\pm} 0.724$	$0.652 {\pm} 0.671$
Blood Pool	Dice	0.903 ± 0.026	0.911 ± 0.027	0.913 ± 0.026	$0.914{\pm}0.024$
	Hausdorff	43.007 ± 11.358	41.792 ± 9.796	41.014 ± 9.769	$40.93 {\pm} 9.52$
	\mathbf{ASD}	$0.664 {\pm} 0.514$	$0.576 {\pm} 0.439$	$0.556{\pm}0.453$	$0.556 {\pm} 0.432$
Compression Rate		${\sim}40\mathrm{x}$	$\sim 35 x$	\sim 22x	$\sim 30 x$

Power Mapping 1.3

We adopted reciprocal mapping function in our paper (as Ours) to get QS for each subband from corresponding SIs. We also considered another non-linear mapping function, power mapping (PM),

$$Q_n = a \cdot \delta_n^b, \tag{3}$$

where a and b are fitting parameters. This mapping methods introduced more non-linearity between Q_n and δ_n .

Results Comparison

We compared segmentation accuracy and compression rate of these three mapping methods with the method in our paper. The experiment setup is the same as that in our paper for fair comparison. As shown in Table 3, we can observe that non-linear mapping methods (DM, PM and RM (ours)) can always achieve significant higher segmentation accuracy than linear mapping (LM) at all three segmentation measurement metrics, which demonstrates the advantage of the non-linear relationship between SI and QS. For the four mapping methods, it can be noticed that usually high accuracy corresponds to low compression rate except for RM (Ours). This clearly indicates that our adopted RM is the best option to balance the accuracy and compression rate, e.g. offering almost the best accuracy at a relatively high compression rate.

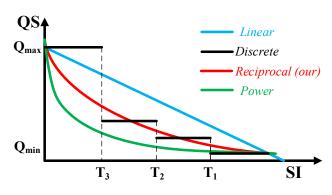


Figure 1: Mapping method comparison for result illustration.

The above phenomenon can be further explained with the illustration in Figure 1. The quantization steps in LM are always larger than non-linear mapping methods (PM and RM) at different SIs, thus resulting with a higher compression rate but a lower segmentation accuracy due to the more significant distortions of important features to DNN. Compared with RM (ours), PM has a steeper slope and a smaller QSs, thus decreasing compression rate $(22 \times v.s. 30 \times)$. However, as we can see from Table 3, the segmentation accuracy is not improved w.r.t. RM. This is because the additional information saved by PM has almost no contributions to DNN segmentation, i.e. redundant data. Although DM is similar with RM in terms of the magnitude of quantization steps (QS),

Table 2: Segmentation results of our method and JPEG-2000 using DenseVoxNet and HVSMR2016 dataset at different compression rates ($30 \times$ and $40 \times$).

		Ours	JPEG2000	Ours	JPEG2000
	Dice	$0.834{\pm}0.0386$	0.816 ± 0.042	0.808 ± 0.043	0.802 ± 0.039
Myocardium	Hausdorff	$31 {\pm} 7.940$	33.513 ± 7.566	34.112 ± 8.551	$34.506 {\pm} 7.962$
	\mathbf{ASD}	$0.652 {\pm} 0.671$	$0.722 {\pm} 0.746$	0.767 ± 0.791	$0.802 {\pm} 0.824$
Blood Pool	Dice	$0.914{\pm}0.024$	0.912 ± 0.024	0.910 ± 0.023	0.909 ± 0.025
	Hausdorff	$40.93 {\pm} 9.52$	41.031 ± 9.648	41.192 ± 9.706	41.448 ± 9.671
	\mathbf{ASD}	$0.556{\pm}0.432$	$0.564 {\pm} 0.414$	$0.576 {\pm} 0.439$	$0.582 {\pm} 0.453$
Compression Rate		$\sim 30 x$	$\sim 30 x$	\sim 40x	$\sim 40 \mathrm{x}$

it only exploits QS with coarse-grained intervals. Therefore, DM can achieve a higher compression rate but lower segmentation accuracy due to the non-optimized (coarse-grained) quantization steps.

Therefore, we finally select RM for our compression framework which can achieve a high compression rate with the best segmentation accuracy.

2 Results with Different Compression Rates

The comparison of our method and JPEG-2000 with different compression rates is shown in Table 2. At $30\times$ compression rate, our method has remarkable segmentation accuracy improvement compared with JPEG-2000. However, at $40\times$ compression rate, the segmentation accuracy is improved. One reasonable explanation is that too aggressive compression will cause serious information loss resulting with low performance. When the compression rate is $30\times$, the necessary information for segmentation is still retained. However, when it comes to $40\times$, this information is seriously dropped which causes a large accuracy degradation. This phenomenon is common in image compression, and compression methods can only drop unnecessary information within their capacities for particular domains. Otherwise, performance loss will always happen.

3 Image by Image Segmentation Results Comparison

To better illustrate that our improvement is statistically significant, we further report the image by image segmentation results of the HVSMAR dataset by using our method and JPEG-2000 under two different DNN structures—DenseVoxNet and 3D-DSN. Table 3 and 4 clearly show that our method wins in almost every case w.r.t. JPEG2000 for Dice, Hausdorff and ASD characterized from various Myocardoum and Blood Pool images under the same compression rate for both networks. Moreover, compared with the original uncompressed images, our method always offers the similar or even better results due to filtering the unnecessary image features that can mislead the segmentation.

Table 3: Image by image segmentation results of original (uncompressed), our method, JPEG-2000 using DenseVoxNet and HVSMR2016 dataset. The compression rate of our method and JPEG-2000 is 30. The images compressed by ours can be segmented with almost the same accuracy, or sometimes even better than original ones, significantly beating those compressed by JPEG-2000.

	Myocardoum									
	Dice			Hausdorff			ASD			
	Original	Ours	JPEG-2000	Original	Ours	JPEG-2000	Original	Ours	JPEG-2000	
Img. 1	0.876	0.876	0.850	19.647	19.197	22.782	0.344	0.327	0.380	
Img. 2	0.786	0.774	0.755	37.229	37.815	39.434	1.426	1.453	1.53	
Img. 3	0.849	0.855	0.848	32.419	31.516	31.718	1.371	1.3	1.526	
Img. 4	0.851	0.846	0.839	27.258	28.083	29	0.122	0.089	0.083	
Img. 5	0.831	0.818	0.791	37.842	38.615	41.109	0.099	0.088	0.088	
				Blo	od Pool					
		Dice	:	Hausdorff			ASD			
	Original	Ours	JPEG-2000	Original	Ours	JPEG-2000	Original	Ours	JPEG-2000	
Img. 1	0.898	0.9	0.897	25.278	24.434	24.515	1.316	1.247	1.226	
Img. 2	0.9	0.9	0.898	45.200	44.878	44.922	0.323	0.311	0.341	
Img. 3	0.953	0.954	0.953	41.219	41.677	41.158	0.552	0.532	0.553	
Img. 4	0.895	0.897	0.895	49.669	48.363	48.518	0.683	0.62	0.649	
Img. 5	0.927	0.919	0.911	43.806	45.31	46.776	0.127	0.11	0.139	

Table 4: Image by image segmentation results of original (uncompressed), our method, JPEG-2000 using 3D-DSN and HVSMR 2016 dataset. The compression rate of our method and JPEG-2000 is 30. The images compressed by ours can be segmented with almost the same accuracy, or sometimes even better than original ones, significantly beating those compressed by JPEG-2000.

	Myocardoum								
	Dice			Hausdorff			ASD		
	Original Ours JPEG-2000			Original	Ours	JPEG-2000	Original	Ours	JPEG-2000
Img. 1	0.829	0.829	0.801	20.273	20.149	22.045	0.446	0.476	0.607
Img. 2	0.688	$\boldsymbol{0.692}$	0.703	39.712	38.652	38.026	0.401	0.386	0.39
Img. 3	0.799	0.802	0.781	35.567	34.929	35.185	0.451	0.5	0.526
Img. 4	0.834	0.839	0.836	25.199	24.880	25.160	0.093	0.091	0.088
Img. 5	0.773	0.769	0.748	40.976	40.902	44.744	0.157	0.171	0.164
Rland Paul									

Blood Pool									
	Dice			Hausdorff			ASD		
	Original	Ours	JPEG-2000	Original	Ours	JPEG-2000	Original	Ours	JPEG-2000
Img. 1	0.903	0.905	0.896	23.324	22.782	25.239	0.568	0.559	0.57
Img. 2	0.873	0.866	0.863	45.837	46.551	46.755	0.229	0.235	0.239
Img. 3	0.949	0.950	0.949	40.410	40.386	40.841	0.221	0.179	0.197
Img. 4	0.915	0.913	0.910	35.931	36.905	38.730	0.88	0.076	0.079
Img. 5	0.906	0.900	0.883	47.074	47.403	51.254	0.068	0.066	0.067