# Supplementary Material: A Morphology Focused Diffusion Probabilistic Model for Synthesis of Histopathology Images 

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## 1. Supplementary Materials

This section provides additional information that we could not include in the paper itself because of space limitations.

### 1.1. Color Normalization in Detail

Let $I_{s} \in \mathbb{R}^{m \times n_{s}}$ be the source image, and let $I_{t} \in \mathbb{R}^{m \times n_{t}}$ be the target image, where $m$ indicates the number of channels, and $n_{s}$ and $n_{t}$ are the number of pixels in the source and target images, respectively. Also, $H \in \mathbb{R}^{r \times n}$ is the stain density map for $r$ stains. The objective is to normalize the colors of the source images into similar color domain of the target, and find the transferred source image denoted as $I_{s}^{\text {norm }}$. Vahadane et al. [2] showed that for both $I_{s}$ and $I_{t}$, we can estimate the normalized source patch $\left(I_{s}^{\text {norm }}\right)$ as:

$$
\begin{gathered}
H_{s}^{\text {norm }}(j,::)=\frac{H_{s}(j,:)}{R M\left(H_{s}(j,:)\right)} R M\left(H_{t}(j,:)\right), j=1, \ldots, r, \\
V_{s}^{\text {norm }}=W_{t} H_{s}^{\text {norm }}, \\
I_{s}^{\text {norm }}=I_{0} \exp \left(-V_{s}^{\text {norm }}\right),
\end{gathered}
$$

where $R M$ denotes the robust maximum of a vector.

### 1.2. Forward Diffusion in Detail

This diffusion process $q$ is a Markov chain and produces latents $x_{1, g_{m}}$ through $x_{T, g_{m}}$ as follows [1]:

$$
\begin{gather*}
q\left(x_{1, g_{m}}, \ldots, x_{T, g_{m}} \mid x_{0, g_{m}}\right)=\prod_{t=1}^{T} q\left(x_{t, g_{m}} \mid x_{t-1}, g_{m}\right),  \tag{1}\\
q\left(x_{t, g_{m}} \mid x_{t-1}, g_{m}\right)=\mathscr{N}\left(x_{t, g_{m}} ;\left(\sqrt{1-\beta_{t}}\right) x_{t-1, g_{m}}, \beta_{t} I\right), \tag{2}
\end{gather*}
$$

and by using reparameterization, latent $x_{t, g_{m}}$ can be derived directly from $x_{0, g_{m}}$ as below:

$$
\begin{equation*}
x_{t, g_{m}}=\sqrt{\alpha_{t}} x_{0, g_{m}}+\sqrt{1-\alpha_{t}} \varepsilon, \tag{3}
\end{equation*}
$$

$0<\beta_{1}, \beta_{2}, \ldots, \beta_{T}<1$ are fixed noise scales for each time step $t$ and $\alpha_{t}:=\prod_{s=1}^{t}\left(1-\beta_{s}\right)$. Also, the distribution of the $\varepsilon$ is as $\varepsilon \sim \mathscr{N}(0, I)$

### 1.3. Model Implementation and Training

We trained our model using PyTorch with 1000 diffusion steps using a workstation with an NVIDIA V100 GPU. The rest of the parameters are available at Table 1 .

| Parameter | Value |
| :--- | :--- |
| Image size | 128 |
| Weighting scheme | P2 |
| Diffusion steps | 1000 |
| Maximum Patches | 100 |
| Channels | 64 |
| Heads | 4 |
| Heads channels | 64 |
| Attention resolution | $32,16,8$ |
| BigGAN upldownsample | yes |
| Num Resblocks | 2 |
| Learning rate | $1 \mathrm{e}-4$ |

Table 1: Hyperparameters for diffusion models used in this paper.

### 1.4. Samples of Real Patches

Table 2 provides the breakdown of the extracted patches based on genomic subtypes and Figure 1 shows samples of real extracted patches after applying the the color normalization module to them.

|  | IDHC | IDHNC | IDHWT | Total |
| :--- | :--- | :--- | :--- | :--- |
| Patches | 12,139 | 16,975 | 4,663 | 33,777 |
| IDH Status | Mutant | Mutant | Wildtype | - |
| 1p19q Status | Codeletion | Retained | - | - |

Table 2: Breakdown of extracted patches per subtype


Figure 1: Selection of patches with size $=128 \times 128$ pixels extracted from the WSI. Each row represents a different genotypes.

### 1.5. Samples of Generated Images by Our Diffusion Probabilistic Model

Figure 2 presents the 10 synthetic images produced by diffusion model, which are significantly close to real images.


Figure 2: Top 10 generated images by our diffusion model.

### 1.6. Survey Visualization

Figure 3 shows a sample section of the web-form that the pathologists completed. Figure 4is the detailed illustration of each expert opinion about each real or synthetic image. Green and yellow colours depict synthetic and real images, respectively. Also, brown presents a synthetic images that classified as real by pathologists and red corresponds to a real images identified as synthetic. Purple shows medium confidence answers.


Figure 3: The web form for subjective study

## References

[1] Jonathan Ho, Ajay Jain, and Pieter Abbeel. Denoising diffusion probabilistic models. Advances in Neural Information Processing Systems, 33:6840-6851, 2020.
[2] Abhishek Vahadane, Tingying Peng, Amit Sethi, Shadi Albarqouni, Lichao Wang, Maximilian Baust, Katja Steiger, Anna Melissa Schlitter, Irene Esposito, and Nassir Navab. Structure-preserving color normalization and sparse stain separation for histological images. IEEE Transactions on Medical Imaging, 35(8):1962-1971, 2016.

| Image Number | Label |
| :---: | :---: |
| 1 | Synthetic |
| 2 | Synthetic |
| 3 | Real |
| 4 | Synthetic |
| 5 | Real |
| 6 | Synthetic |
| 7 | Synthetic |
| 8 | Synthetic |
| 9 | Real |
| 10 | Synthetic |
| 11 | Real |
| 12 | Real |
| 13 | Real |
| 14 | Synthetic |
| 15 | Real |
| 16 | Synthetic |
| 17 | Synthetic |
| 18 | Synthetic |
| 19 | Real |
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| 71 | Real |
| 72 | Synthetic |
| 73 | Real |
| 74 | Real |
| 75 | Synthetic |
| 76 | Real |
| 77 | Synthetic |
| 78 | Synthetic |
| 79 | Synthetic |
| 80 | Synthetic |


| P1 |  |
| :---: | :---: |
| Real | High |
| Real | High |
| Real | High |
| Synthetic | Medium |
| Real | High |
| Synthetic | High |
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| P2 |  |
| :---: | :---: |
| Synthetic | High |
| Real | High |
| Real | High |
| Real | Medium |
| Synthetic | Medium |
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| Real | Medium |
| Synthetic | Medium |
| Synthetic | Medium |
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| Synthetic | Medium |
| Synthetic | Medium |
| Synthetic | Medium |
| Synthetic | Medium |
| Synthetic | Medium |
| Synthetic | High |
| Real | Medium |
| Real | Medium |

Figure 4: Illustration of each pathologist's opinion. Green: synthetic image, yellow: real one. Brown: synthetic images that classified as real by pathologists. Red: real image diagnosed as synthetic image. Purple: medium confidant answers.


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