

## Explanation

### Functions

The method receives an image  $M$  as input, which in this explanation we will assume to be stretched to fill the  $[0, 1] \times [0, 1]$  domain.

The method works using 3 functions:

$$U, \hat{X}, \hat{Y} : [0, 1] \times [0, 1] \times \mathbb{R}_0^+ \rightarrow \mathbb{R}_0^+$$

The saliency  $U$  starts in  $t=0$  calculated from the image:

$$U(x, y, 0) = \text{Saliency}(M)(x, y) + \varepsilon$$

where  $\varepsilon$  is a relatively small constant that can be added so low information areas of the image are not totally ignored.

For calculating the saliency, different methods can be used, see for instance the ones provided by OpenCV <sup>1</sup>.

The X-osity  $\hat{X}$ , with initial condition in  $t=0$ :

$$\hat{X}(x, y, 0) = x U(x, y, 0)$$

The Y-osity  $\hat{Y}$ , with initial condition in  $t=0$ :

$$\hat{Y}(x, y, 0) = y U(x, y, 0)$$

### Difussion

The saliency is diffused using the heat equation:

$$\frac{\delta U}{\delta t} = \alpha \left( \frac{\delta^2 U}{\delta x^2} + \frac{\delta^2 U}{\delta y^2} \right)$$

with Neumann boundary conditions <sup>2</sup>.

The functions are discretized in a finite grid. A small variation of finite differences is used:

On each time step  $t$ , the amount of saliency  $\Delta U_{i \rightarrow j, t}$  transferred from a cell  $i$  to another cell  $j$  is given by:

$$\Delta U_{i \rightarrow j, t} = \alpha \max(U_{i, t} - U_{j, t}, 0) U_{i, t}$$

So, the saliency of cell  $i$  in the next step is given by:

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<sup>1</sup><https://towardsdatascience.com/opencv-static-saliency-detection-in-a-nutshell-404d4c58fee4>

<sup>2</sup>[https://en.wikipedia.org/wiki/Neumann\\_boundary\\_condition#PDE](https://en.wikipedia.org/wiki/Neumann_boundary_condition#PDE)

$$\begin{aligned}
U_{i,(t+1)} = & U_{i,t} + \\
& \Delta U_{a \rightarrow i,t} - \Delta U_{i \rightarrow a,t} + \\
& \Delta U_{b \rightarrow i,t} - \Delta U_{i \rightarrow b,t} + \\
& \Delta U_{c \rightarrow i,t} - \Delta U_{i \rightarrow c,t} + \\
& \Delta U_{d \rightarrow i,t} - \Delta U_{i \rightarrow d,t}
\end{aligned}$$

where  $a, b, c, d$  are the cells adjacent to  $i$  in the grid.

Smaller values of  $\alpha$  make the simulation more stable and precise, but slower.

Each time an amount of saliency  $\Delta U_{i \rightarrow j,t}$  is transferred from a cell  $i$  to another cell  $j$ , a proportional amount of the X-osity and Y-osity are transferred along with it:

$$\Delta \hat{X}_{i \rightarrow j,t} = \frac{\Delta U_{i \rightarrow j,t}}{U_{i,t}} \hat{X}_{i,t}$$

$$\Delta \hat{Y}_{i \rightarrow j} = \frac{\Delta U_{i \rightarrow j}}{U_i} \hat{Y}_i$$

$\hat{X}_{i,(t+1)}$  and  $\hat{Y}_{i,(t+1)}$  are obtained analogously to  $U_{i,(t+1)}$

Once enough iterations are performed, the saliency is expected to be approximately constant:

$$\lim_{t \rightarrow \infty} U(x, y, t) = C \quad , \text{ with } C \text{ constant.}$$

so the amount of detail of the image is distributed evenly.

To build the final image  $M'$ , the final X-osity and Y-osity should be used to obtain coordinates of the pixels from the original image  $M$  that have to be queried.

$$M'(x, y) = M \left( \frac{\hat{X}(x, y, \infty)}{U(x, y, \infty)}, \frac{\hat{Y}(x, y, \infty)}{U(x, y, \infty)} \right)$$

linear interpolation can be used to obtain the color of positions between the pixels.

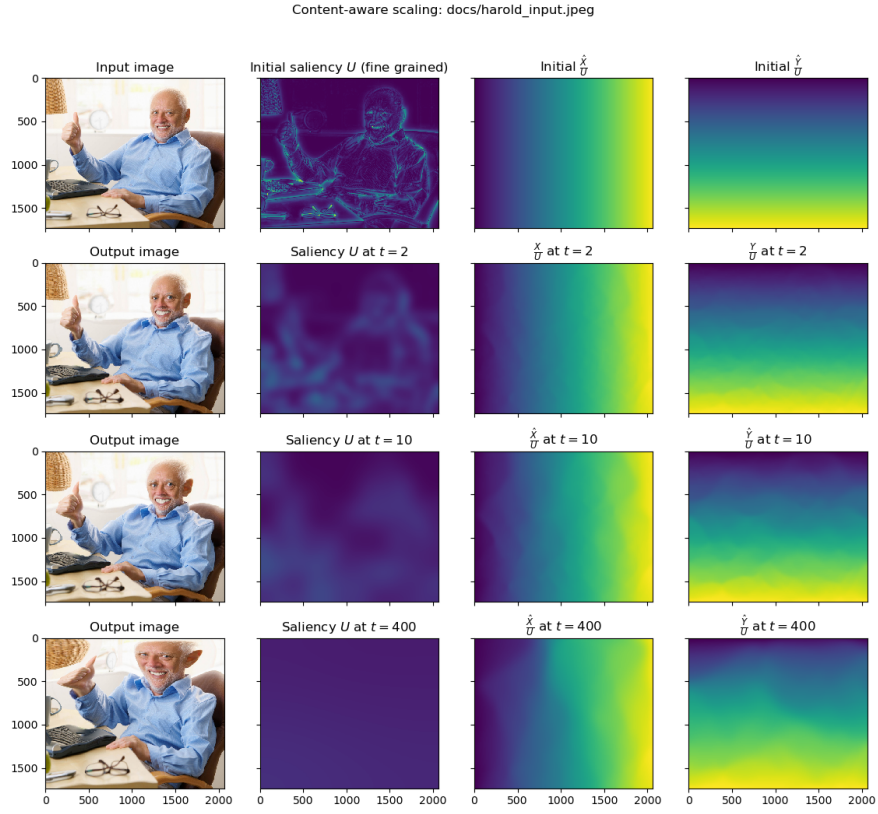


Figure 1: Harold image diffused at different times