

HW3 Report

Section 1

Q1:

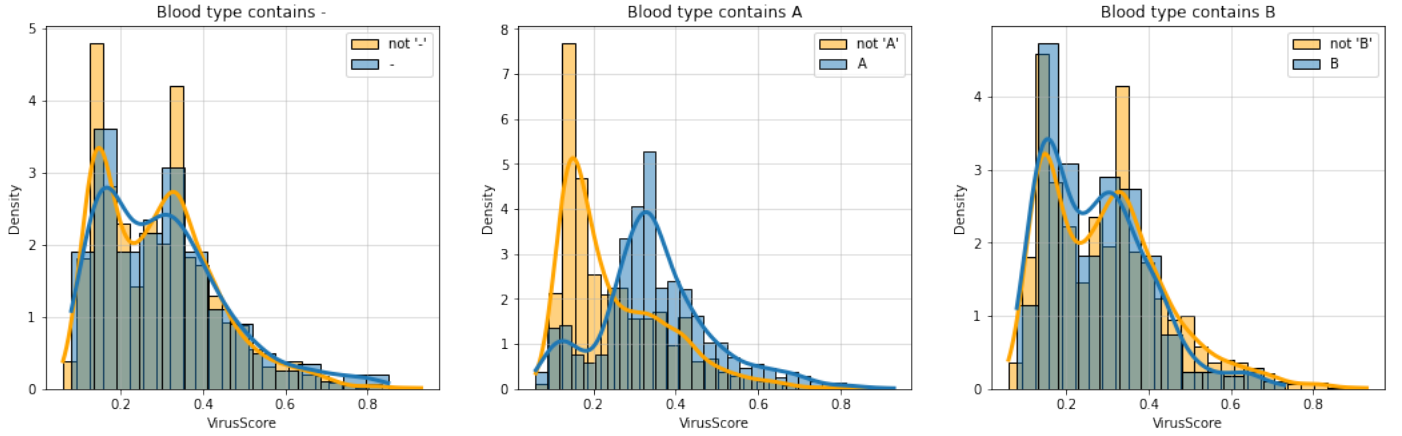


Figure 1: KDE plots of **VirusScore** conditioned different conditions of **blood_type**

Q2:

In figure 1 in the plot of A versus not A, we observe that the groups of patients with and without "A" in their blood types are mostly separable along a boundary that is approximately the **VirusScore** of 0.225.

Therefore, the condition of contains/does not contain A would be most informative for learning **VirusScore**. As it turns out, we decided already in hw1 to create this feature.

Q3:

$$\begin{aligned}
 \frac{\partial}{\partial b} \mathcal{L}(\mathbf{w}, b) &\stackrel{a}{=} \frac{\partial}{\partial b} \frac{1}{m} \sum_{i=1}^m (\mathbf{w}^\top \mathbf{x}_i - b - y_i)^2 \stackrel{b}{=} \frac{1}{m} \frac{\partial}{\partial b} \sum_{i=1}^m (\mathbf{w}^\top \mathbf{x}_i - b - y_i)^2 \stackrel{c}{=} \frac{1}{m} \sum_{i=1}^m \frac{\partial}{\partial b} (\mathbf{w}^\top \mathbf{x}_i - b - y_i)^2 \stackrel{d}{=} \\
 &\stackrel{d}{=} \frac{1}{m} \sum_{i=1}^m 2(\mathbf{w}^\top \mathbf{x}_i - b - y_i) \cdot (-1) \stackrel{e}{=} \frac{2}{m} \sum_{i=1}^m (b + y_i - \mathbf{w}^\top \mathbf{x}_i) \stackrel{f}{=} \frac{2}{m} \left[mb + \sum_{i=1}^m (y_i - \mathbf{w}^\top \mathbf{x}_i) \right] \stackrel{g}{=} \\
 &\stackrel{g}{=} 2b + \frac{2}{m} \sum_{i=1}^m (y_i - \mathbf{w}^\top \mathbf{x}_i) \Rightarrow \frac{\partial}{\partial b} \mathcal{L}(\mathbf{w}, b) = 2b + \frac{2}{m} \sum_{i=1}^m (y_i - \mathbf{w}^\top \mathbf{x}_i)
 \end{aligned}$$

a : Definition of $\mathcal{L}(\mathbf{w}, b)$
 b : $\frac{1}{m}$ is scalar
 c : Derivative of a sum is the sum of derivatives
 d : Derivative of $(\mathbf{w}^\top \mathbf{x}_i - b - y_i)^2$ w.r.t b
 e : 2 is scalar
 f : sum of b
 g : Removing b from the sum