
Submitted by:

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Solution:

1. True. If we use reconstruction model for outlier detection.
2. True. Distances between any two vertices are of one length in a fully connected graph.
3. Data samples that are far away from each other are outliers, the samples that are close to each other are less likely to be outliers.
4. Sparse interactions / Parameter sharing / Translational equivalent / less computational expensive.
5. Translational equivalent means the model can produce the same output no matter where the input is placed, and it is achieved by parameter sharing (convolution).
Translational invariance means the model can produce the same output when the input is translated and it is achieved by pooling operation.
6. Dropout layer can be seen as a regularization on the model itself to force the model to be more robust to noise, and to learn more generalizable features.
- 7.

Solution:

- 2.A.** Numerator is the number of samples o' "near" the sample o , denominator is the number of samples in D , thus LHS is the ratio of samples from D that is close to o .

Therefore this ratio is less than $\pi \Leftrightarrow$ few samples from D is close to $o \Leftrightarrow o$ is a distance-based outlier.

- 2.B.**

$$\begin{aligned} \frac{\|o' | dist(o, o') \leq r\|}{\|D\|} &\leq \pi \\ \Leftrightarrow \|o' | dist(o, o') \leq r\| &\leq \|D\|\pi \\ \Leftrightarrow \|o' | dist(o, o') \leq r\| &< \lceil \pi \|D\| \rceil \\ \Leftrightarrow \|o' | dist(o, o') > r\| &> \lceil \pi \|D\| \rceil \\ \Leftrightarrow \|o' | dist(o, o') > r\| &> k \end{aligned}$$

Therefore if $dist(o, o_k) > r \forall o_k$, then $\|o' | dist(o, o') > r\| > k$, thus o is an outlier. ■

- 2.C.** Denote $f(o) = \frac{\|o' | dist(o, o') \leq r\|}{\|D\|}$, then

$f(-4.5) = 0.3 > \pi$, -4.5 is not an outlier.

$f(-4) = 0.3 > \pi$, -4 is not an outlier.

$f(-3) = 0.3 > \pi$, -3 is not an outlier.

$f(-2.5) = 0.3 > \pi$, -2.5 is not an outlier.

$f(3) = 0.4 > \pi$, 3 is not an outlier.

$f(3.5) = 0.4 > \pi$, 3.5 is not an outlier.

$f(4) = 0.4 > \pi$, 4 is not an outlier.

$f(4.5) = 0.4 > \pi$, 4.5 is not an outlier.

$f(5) = 0.4 > \pi$, 5 is not an outlier.

$f(0) = 0.0 \leq \pi$, 0 is an outlier.

Solution:

- 3.A.** Distance-based approach defines the outlier samples with the property that there are not enough neighbouring data samples around. Density-based approach defines the outlier samples with the property that the data sample density is significant lower than its neighbours.

3.B.
$$lrd_k(o) = \frac{\|N_k(o)\|}{\sum_{o' \in N_k(o)} reachdist_k(o' \leftarrow o)}$$

3.B.i. We have
$$\frac{\sum_{o' \in N_k(o)} reachdist_k(o' \leftarrow o)}{\|N_k(o)\|} = \frac{\sum_{o' \in N_k(o)} \max\{dist_k(o'), dist(o', o)\}}{k}$$

Which is the average reachability distance from o to $o' \in N_k(o)$.

- 3.B.ii.** Intuitively, when the density of o is small, in other words, if $o \notin dist_k(o')$, then $reachdist_k(o', o)$ will be larger and the average reachability distance from o to $o' \in N_k(o)$ will increase and $lrd_k(o)$ will decrease. Therefore, if the sample o is closer to a cluster, which can be interpreted as $o \in dist_k(o')$ for most $o' \in dist_k(o)$, then $lrd_k(o)$ will increase.

3.C.
$$LOF_k(o) = \frac{\sum_{o' \in N_k(o)} \frac{lrd_k(o')}{lrd_k(o)}}{\|N_k(o)\|}$$

By calculating the division, we can compare the density of o and o' , $LOF_k(o)$ is then defined as the average density ratio of the density of o 's k nearest neighbours and the density of o .

MY_{LOF_k}

Solution:

4.A. a: local maximal; b: saddle point; c: cliff; d: local minimal

4.B. We have $\frac{\partial O}{\partial I} = O(1 - O)$

4.B.i.

$$\begin{aligned}\delta_k &= \frac{\partial L_k}{\partial I_k} \\ &= -(T - O_k) \frac{\partial O_k}{\partial I_k} \\ &= O_k(1 - O_k)(O_k - T) \blacksquare\end{aligned}$$

4.B.ii.

$$\begin{aligned}\delta_i &= \frac{\partial L_i}{\partial I_i} \\ &= \frac{\partial(\frac{1}{m} \sum_j^m w_{ij} L_j)}{\partial I_i} \\ &= \frac{\partial O_i}{\partial I_i} \sum_j^m w_{ij} \delta_j \\ &= O_k(1 - O_k) \sum_j^m w_{ij} \delta_j \blacksquare\end{aligned}$$

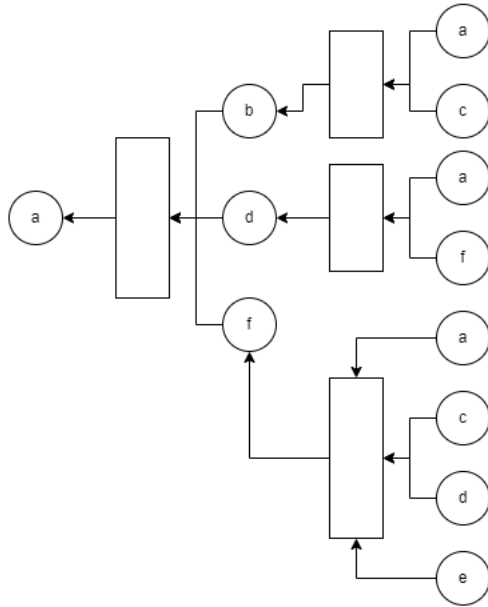
4.B.iii.

$$\begin{aligned}\delta'_k &= \frac{\partial L'_k}{\partial I_k} \\ &= \frac{\partial(-T \log O_k - (1 - T) \log(1 - O_k))}{\partial I_k} \\ &= \frac{\partial O_k}{\partial I_k} \frac{T - O_k}{O_k(1 - O_k)} \\ &= O_k(1 - O_k) \frac{T - O_k}{O_k(1 - O_k)} = T - O_k \blacksquare\end{aligned}$$

4.B.iv. When output unit is saturated, we have $\text{MSE} = O_k(1 - O_k)(O_k - T) \approx 0$ as well, where as for Cross-Entropy loss does not have this issue.

Solution:

- 5.A. Conv with K_1 : $[[0, 2, 1, 0], [1, 1, 0, 2], [2, 0, 2, 1], [1, 2, 0, 1]]$.
 Conv with K_2 : $[[2, 1, 1, 2], [1, 2, 1, 2], [1, 2, 1, 2], [3, 1, 1, 1]]$.
- 5.B. Avg Pooling on feature map with K_1 : $[[1.0, 0.75], [1.25, 1.0]]$.
 Avg Pooling on feature map with K_2 : $[[1.5, 1.75], [1.5, 1.75]]$.
- 5.C. The output shape is $E \times E \times L$, where $E = \lfloor \frac{N-K}{S} \rfloor + 1$.
- 5.D. $h_1 = Wh_0 + Ux_1 = [2, 2]$, $h_2 = Wh_1 + Ux_2 = [5, 3]$, $\hat{y}_2 = Vh_2 = [8, 5]$
- 5.E. Message passing tree for Node a .

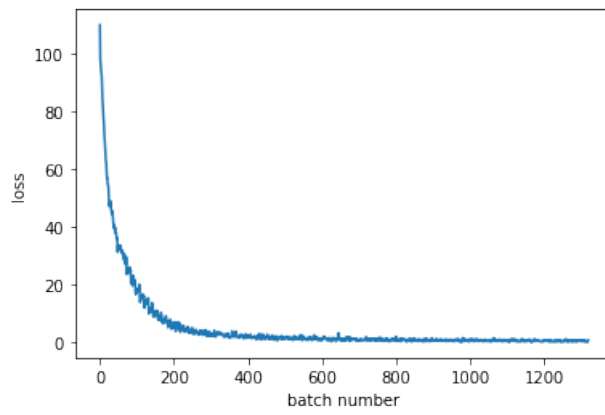


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| Solution:
Holer

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Solution:



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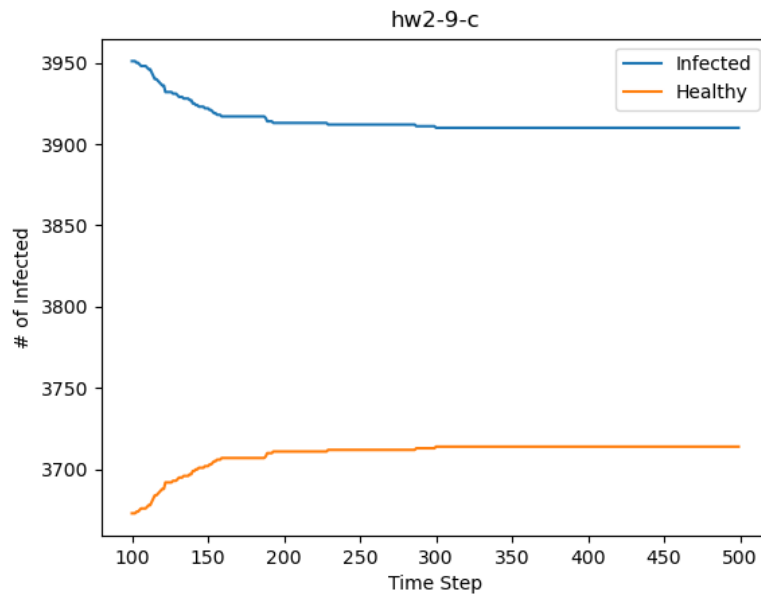
| Solution:
Holer

Solution:

9.A. $1 - (1 - \beta)^n$

9.B. 38.6013

9.C. Treating infected probability ≥ 0.6 as infected, getting an epidemic.



9.D. Treating infected probability ≥ 0.6 as infected, getting no epidemic.

