Exercise 2

1. Algorithm:

Go through every x in the sample and check whether it belongs to our discrete domain &XX.

If we find such X, then function hz, where Z=X
is the ontcome of our algorithm. Due to realizability
assumption, there can only be one such X in the sample
and, therefore, L_s(hz) will be O.

If we don't find such x, then h is the outcome of our algorithm. Since in this case we don't have any positives in the sample, Ls(h) will be O.

2. If the algorithm described above outputs hz, then, due to the realizability assumption it will mean that our hypothesis hz correctly identifies the only positive in our domain, and hence, Lp, f (hz) with be 0, which is less than & for all &>0.

Since in this case, the population risk will 7 be equal to $P_D(x_3)$, we get the inequality for the size of the sumple m: $(1-\epsilon)^m \in \mathcal{O}$

 $m \geq \lceil \log_{(1-\epsilon)} S \rceil$

the required upper bound on the sample complexity

Exercise 3 Fix some distribution Dover X.

Assuming realizability, Addition Let R* with radius n* he the concentric circle that generates the labels and let f be the Corresponding hypothesis, Let r<r* be assessable to once radius of the concentric circle R, such that the probability mass of the area between R and R* is exactly E. (Let's denote that area R)

For perhas rapositive rexemple Allowalthe sugapher 8 Let A be the algorithm that returns the smallest contangles evolvsing all positive examples in the training set. From realizability assumption we get that A does not mislable any negative examples (if it does, there should be smaller circle enclosing all positive examples, which contradicts definition of A(SI), Hence, A is an ERM.

Since A(S) is the smallest circle enclosing all positive examples, A(S) = R*. IF R contains any positive example, then the boundary of A lies between R* and R. Since the probability mass of that region is E, empirial risk of A(5) in this case will be at most E. The probability of positive example being inside R is 1-E, hence the probability of all positive samples

the be inside R is (1-E) , where m is the Size of the sample. Therefore, the probability that at least

one positive is in R is 1-(1-E)m

We set that P(Lo, p(A(S)) < E) > 1-(1-E) = 1-ème

Rewriting this inequality, we get $m \not\in \frac{\log(1/\delta)}{\varepsilon}$.

Therefore, we have show that \exists algorithm f, s.b. \forall ε , $\delta > 0$, \forall distributions D, \forall labeling fous ε , s.t. D, ε ealizable by \mathcal{H} , given $m \geq \lceil \frac{\log(1/\delta)}{\varepsilon} \rceil$, then $W \cdot P$. And at least $1-\delta$ over $S \cap D^m$, we have L_{D}, ε (A(S)) $< \varepsilon$

tience, It is PAC-learnable and its sample complexity is bounded by $m_{\mathcal{H}}(\xi, \delta) \leq \Gamma \frac{\log(1/\sigma)}{\varepsilon}$

q. e. d.