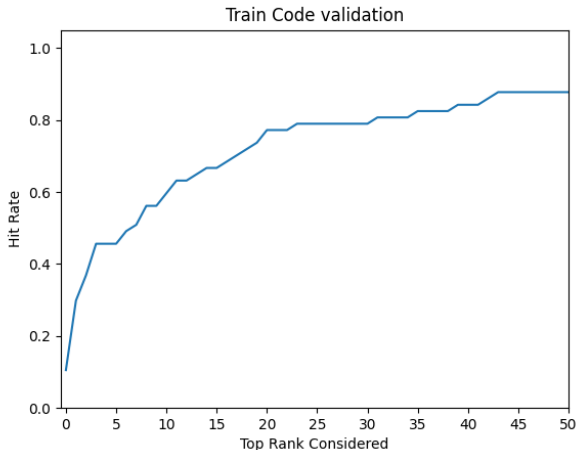
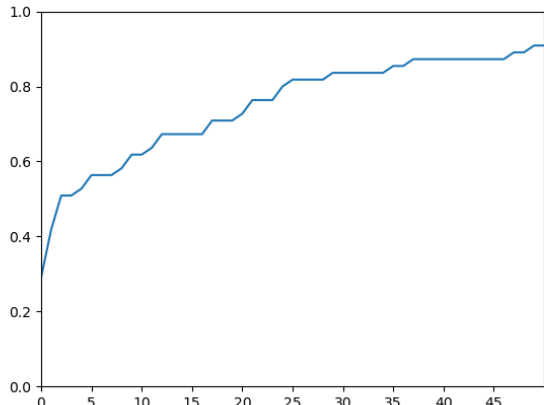


1. Train/Test feature를 그대로 하고 학습하고 평가	
 <p>Train Code validation</p> <p>Hit Rate</p> <p>Top Rank Considered</p>	 <p>Hit Rate</p>
새로 학습해서 그린 hit rate 그래프	기존에 주어진 모델로 그린 hit rate 그래프
학습시간:	
2. Residue 추가한 Train/Test	
학습시간: 약 1시간 * 50 epoch = 50시간	
3. calpep	
capri 기준	
f(nat) thresholds of 0.2, 0.5 and 0.8 now distinguish between high (f(nat)>0.8), medium (f(nat)>0.5), acceptable (f(nat)>0.2) and incorrect models (f(nat)<0.2), or:	
* fnat: incorrect < 0.2 < acceptable < 0.5 < medium < 0.8 < high	
and similarly L-rms and I-rms:	
* lrms: incorrect > 4.0 > acceptable > 2.0 > medium > 1.0 > high	
* irms: incorrect > 2.0 > acceptable > 1.0 > medium > 0.5 > high	
in other words: a high-quality protein-peptide model has f(nat) above 0.8 and either L-rms below 1.0 or I-rms below 0.5.	
>> incorrect = fnat < 0.2 and lrmsd > 4.0 or irmsd > 2.0 를 기준으로 삼았습니다	
>> 231 개	
/mnt/rv1/alhome/jhhwang/PepPro/pepdecoys/ppbenchoutputs-230114/results.txt	