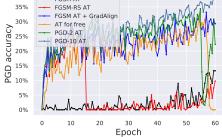
- We thank all reviewers for the encouraging feedback and detailed comments which we'll integrate into the next version.
- 2 R1, R2, R3: FGSM+GradAlign is slower than FGSM.
- 3 We admit that GradAlign leads to a slowdown: e.g. on CIFAR-10 the runtime of FGSM AT is 9.5 min while FGSM
- 4 + GradAlign AT takes 30.9 min on an NVIDIA V100 GPU. However, we present GradAlign as a proof of concept
- 5 motivated by our empirical and theoretical analysis. We hope that the same effect (stability of the gradients under
- 6 random noise) can be achieved in future work with other regularization methods that avoid double backpropagation.
- 7 **R1**: What makes the grad. alignment low by FGSM training? Why a noise-sensitive filter is learned by FGSM training?
- 8 These are very interesting questions, and we believe they are connected to the finding of [25] that SGD for neural
- 9 networks learns models of increasing complexity, e.g., measured in terms of local linearity.
- 10 R2: Running stronger attacks, e.g., AutoAttack or MultiTargeted
- 11 After running AutoAttack, we observe that it proportionally reduces the adversarial accuracy for all methods. E.g., for
- 12 $\varepsilon = 8/255$, FGSM+GradAlign achieves $44.54\pm0.24\%$ adversarial accuracy while FGSM-RS achieves $42.80\pm0.58\%$.
- This is consistent with the results of the AutoAttack paper where they show an average reduction of 2%-3% adversarial
- 14 accuracy compared to most of the evaluations that were originally done with variants of PGD.
- 15 **R2**: Discussion on LLR [26] and CURE [24]. Their exact goal is to make the loss surface smoother.
- 16 Indeed, the goals of LLR/CURE and GradAlign wrt smoothness are similar, however the important difference is that
- we were not looking for a replacement of adv. training, but rather for a *complement* that would prevent catastrophic
- overfitting. Related to this, in Table 7 we also provide the results for FGSM+CURE where we can see that CURE
- 19 also stabilizes FGSM training, but performs worse than FGSM+GradAlign. Finally, GradAlign does not have any
- 20 worst-case motivation unlike CURE (uses the FGSM point) or LLR (uses a point with the worst-case linear violation).
- **R2**: Line 118: How is the alpha step-size tuned for this experiments? (is it 1.25ε ?)
- Yes, we used $\alpha = 1.25\varepsilon$ since it was the recommended choice of Wong et al. [44] on all the datasets they considered.
- **R2**: Performance of FGSM+GradAlign in large ε settings (16/255) on ImageNet against random targeted attacks.
- First, we note that for FGSM+GradAlign on CIFAR-10 we did not encounter cat. overfitting even with $\varepsilon=16/255$. We
- briefly tried targeted AT with $\varepsilon=16/255$ on ImageNet but we did not succeed at training a sufficiently robust model. We
- 26 think that it is likely that more epochs (e.g., LLR [26] used 110 epochs instead of 15 epochs as we did following [44])
- 27 and different hyperparameters are needed, but tuning them on ImageNet was too computationally expensive for us.
- **R3**: Learning curve with robust accuracy and the effect of the regularizer
- on grad. alignment, e.g., on CIFAR-10 with $\varepsilon=14$ and 60 epochs.
- We present this experiment in Fig. A. The only two methods that do not fail at epoch 60 are PGD-10 and FGSM+GradAlign which is also reflected
- by their gradient alignment, i.e. cosine distances (highest for GradAlign).
- **R3**: *I'd vote for early stop since it is not complicated and rather efficient.*
- We'd like to clarify that using early stopping leads to worse PGD accuracy
- (particularly for high ε) and much worse clean accuracy as we comment in lines 307-310. Thus, early stopping alone is not a satisfying solution.
- **R3**: What happens when we use $\varepsilon > 4$ for ImageNet? Do all methods
- including the proposed one fail? We have the results for $\varepsilon=6$ in Table 6 and catastrophic overfitting there
- occurs *only* for the FGSM-RS model. However, the results on ImageNet
- are not fully conclusive since we could not repeat the experiments over
- multiple random seeds due to the computational constraints.
- **R3**: Lemma 1: what if $\eta \sim \mathcal{U}([-\beta, \beta]^d)$? Will β appear in the bound?
- We were interested in $\beta = \varepsilon$ since it was the setting of [44]. Indeed, β will
- appear in the bound, but this will not change the message of Lemma 1.
- **R4**: The empirical analysis is mainly done on CIFAR10 only.
- 47 We'd like to emphasize that we have provided experiments on ImageNet
- to illustrate that GradAlign can be scaled to large datasets. But due to our
- 49 limited computational resources, we could not do replications over random seeds and a thorough comparison to other
- methods (particularly, to PGD-10) for a range of ε as on CIFAR-10 (e.g., Fig. 1 required to train 480 different models).
- **84**: Why does cat. overfitting not occur on ImageNet [for $\varepsilon \in \{2, 4\}$]? How does the gradient alignment evolve?
- 52 The main reason is that these ε values are sufficiently small (we observed the same also on CIFAR/SVHN for $\varepsilon \le 4/255$).
- The gradient alignment decreases gradually over epochs, but without a sharp drop that would indicate cat. overfitting.



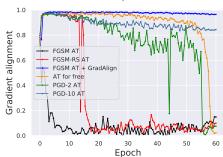


Figure A: Illustration of catastrophic overfitting for various AT methods with $\varepsilon = 14/255$.