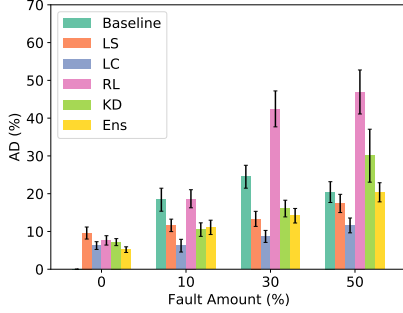
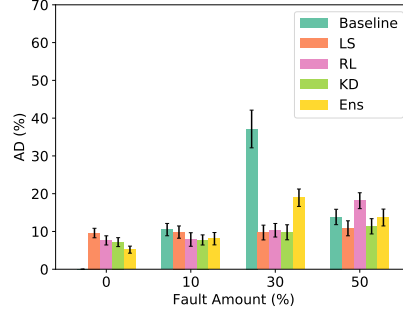


Accuracy Delta by Model, Fault Type, Dataset

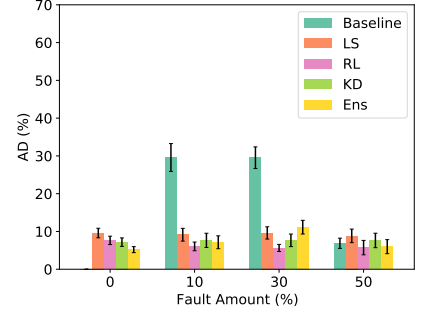
I. CIFAR-10



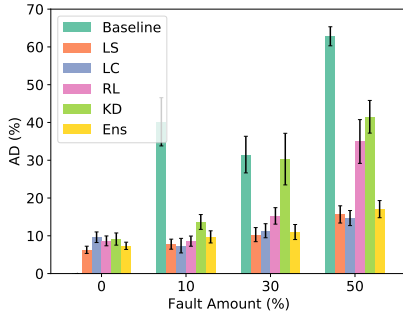
(a) CIFAR-10, ConvNet, Mislabelling



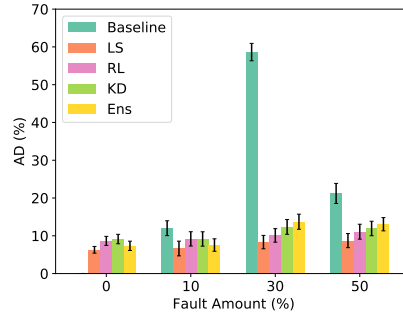
(b) CIFAR-10, ConvNet, Removal



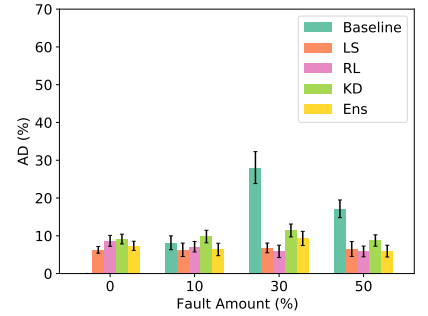
(c) CIFAR-10, ConvNet, Repetition



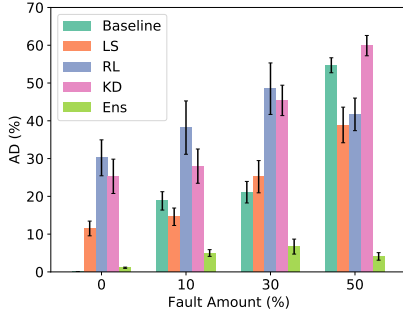
(d) CIFAR-10, DeconvNet, Mislabelling



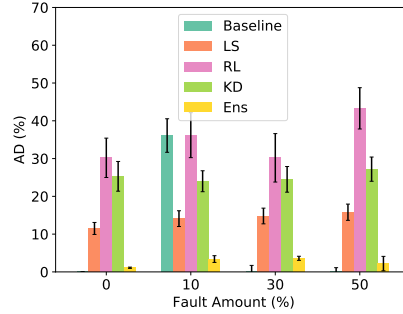
(e) CIFAR-10, DeconvNet, Removal



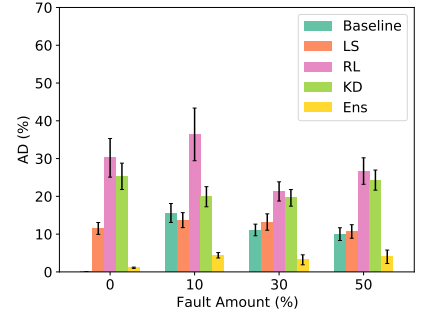
(f) CIFAR-10, DeconvNet, Repetition



(g) CIFAR-10, MobileNet, Mislabelling

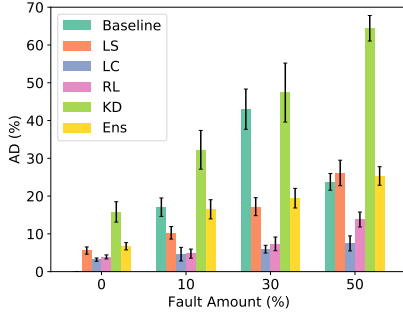


(h) CIFAR-10, MobileNet, Removal

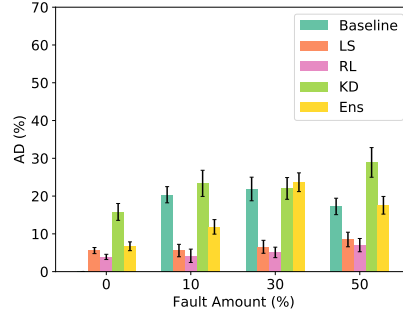


(i) CIFAR-10, MobileNet, Repetition

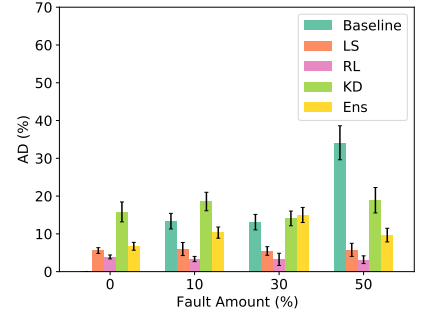
Fig. 1: AD of individual models, compared with models protected with TDFM techniques when trained with faulty CIFAR-10 datasets. The error bars in the results indicate the 95% confidence intervals. Lower values are better.



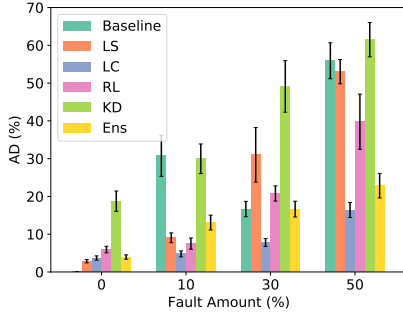
(a) CIFAR-10, ResNet18, Mislabelling



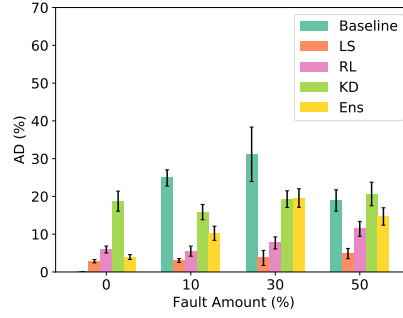
(b) CIFAR-10, ResNet18, Removal



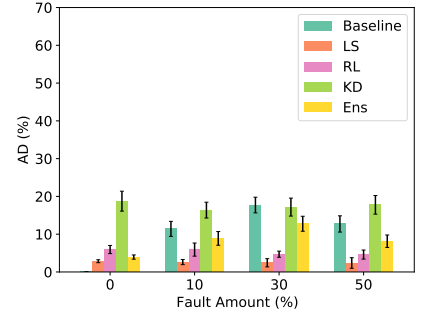
(c) CIFAR-10, ResNet18, Repetition



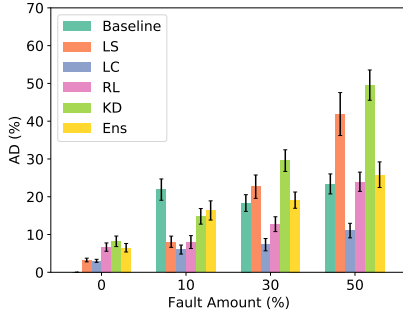
(d) CIFAR-10, ResNet50, Mislabelling



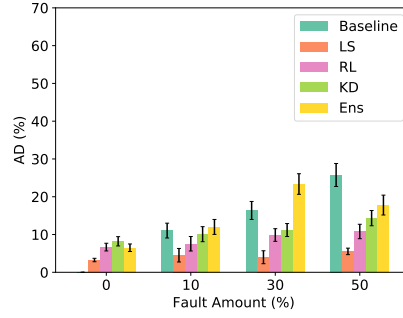
(e) CIFAR-10, ResNet50, Removal



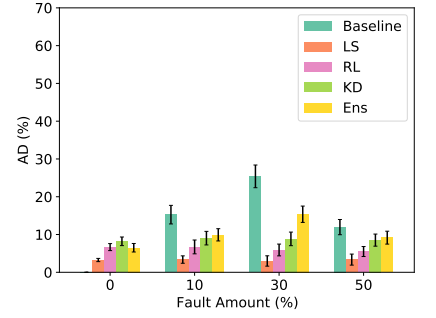
(f) CIFAR-10, ResNet50, Repetition



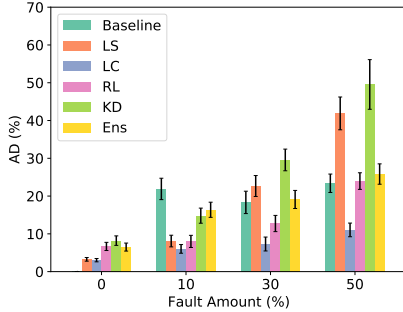
(g) CIFAR-10, VGG11, Mislabelling



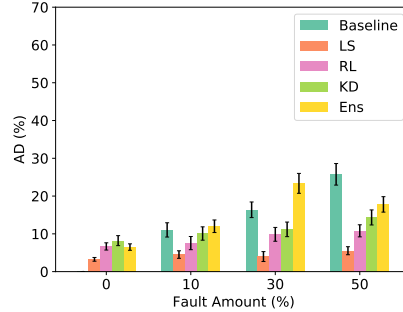
(h) CIFAR-10, VGG11, Removal



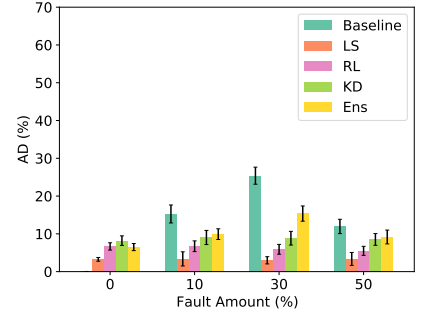
(i) CIFAR-10, VGG11, Repetition



(j) CIFAR-10, VGG16, Mislabelling



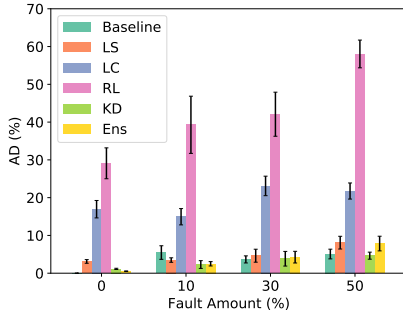
(k) CIFAR-10, VGG16, Removal



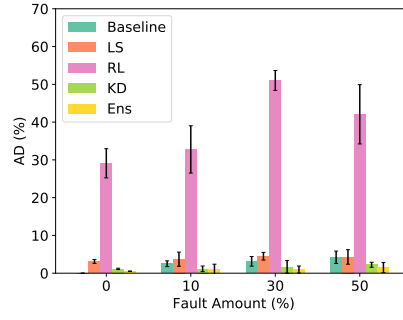
(l) CIFAR-10, VGG16, Repetition

Fig. 2: AD of individual models, compared with models protected with TDFM techniques when trained with faulty CIFAR-10 datasets. The error bars in the results indicate the 95% confidence intervals. Lower values are better.

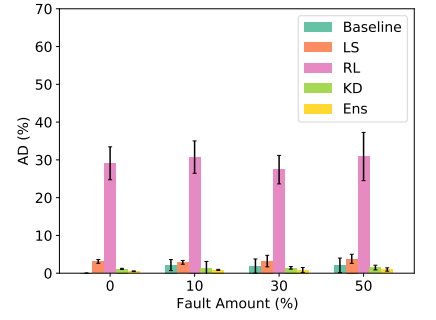
II. GTSRB



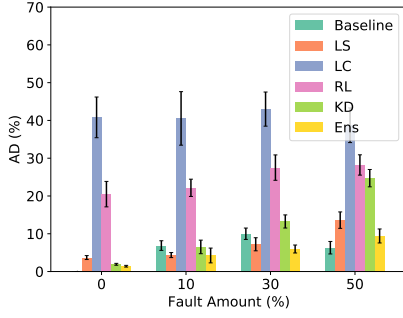
(a) GTSRB, ConvNet, Mislabelling



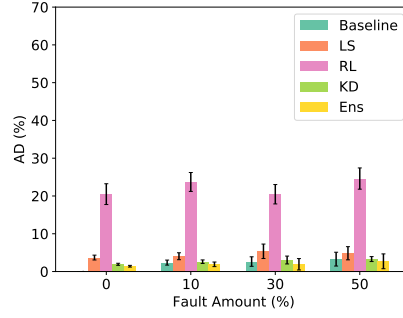
(b) GTSRB, ConvNet, Removal



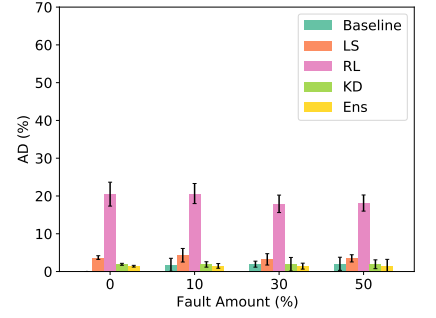
(c) GTSRB, ConvNet, Repetition



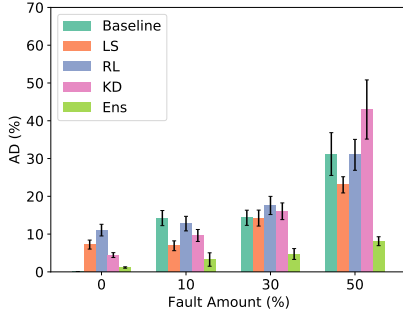
(d) GTSRB, DeconvNet, Mislabelling



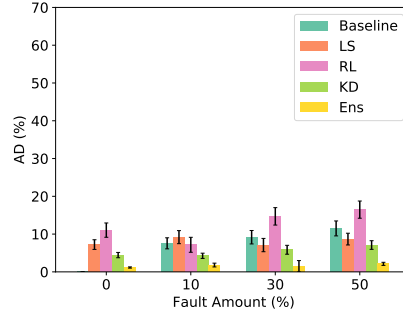
(e) GTSRB, DeconvNet, Removal



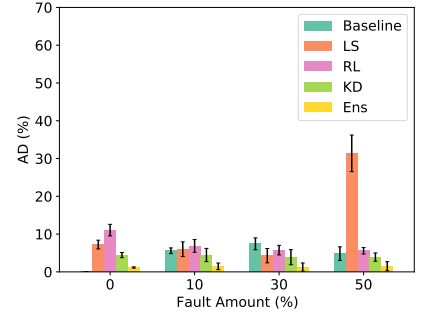
(f) GTSRB, DeconvNet, Repetition



(g) GTSRB, MobileNet, Mislabelling

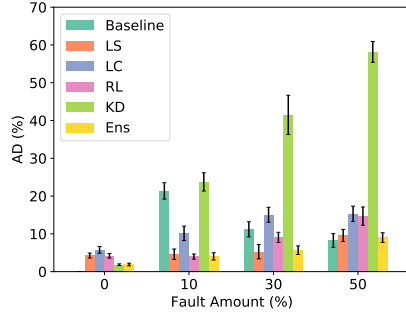


(h) GTSRB, MobileNet, Removal

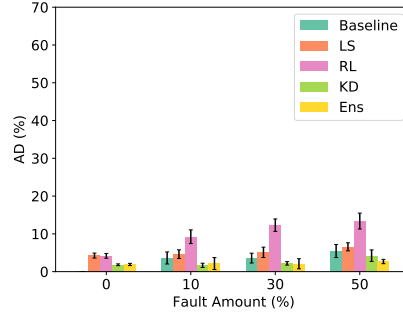


(i) GTSRB, MobileNet, Repetition

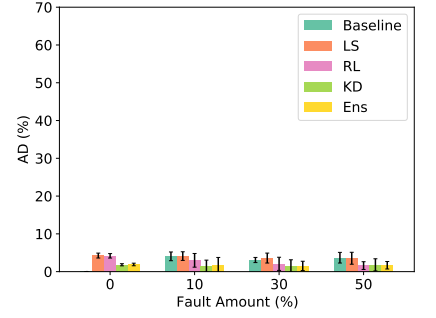
Fig. 3: AD of individual models, compared with models protected with TDFM techniques when trained with faulty GTSRB datasets. The error bars in the results indicate the 95% confidence intervals. Lower values are better.



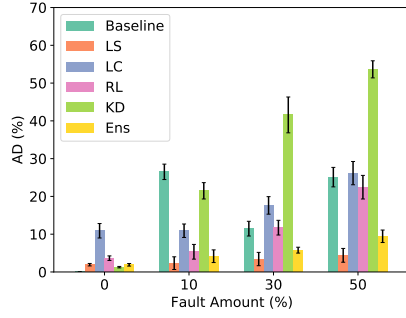
(a) GTSRB, ResNet18, Mislabelling



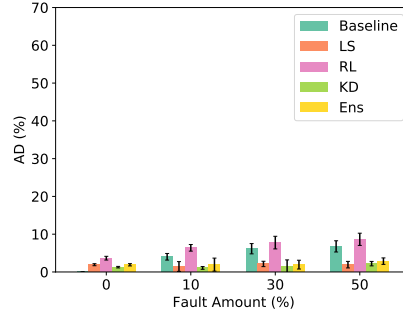
(b) GTSRB, ResNet18, Removal



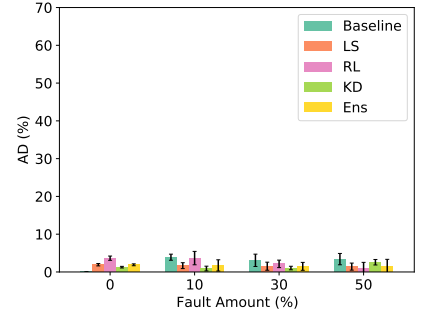
(c) GTSRB, ResNet18, Repetition



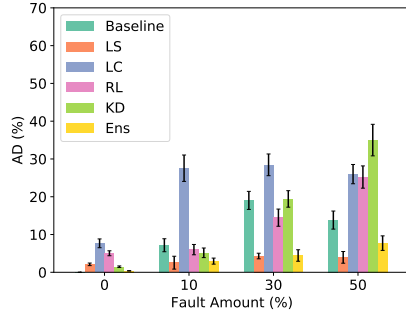
(d) GTSRB, ResNet50, Mislabelling



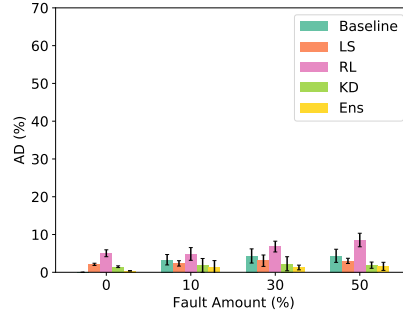
(e) GTSRB, ResNet50, Removal



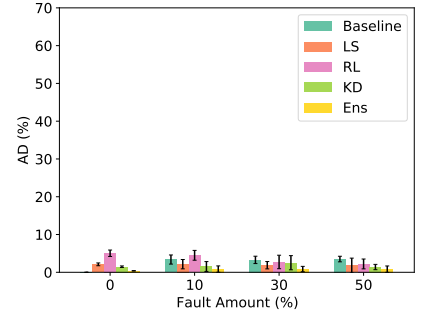
(f) GTSRB, ResNet50, Repetition



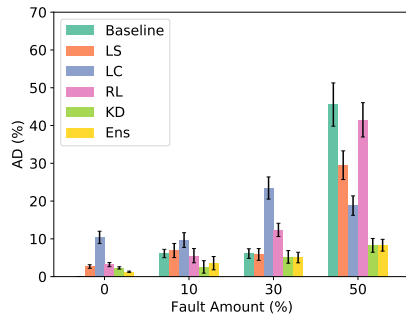
(g) GTSRB, VGG11, Mislabelling



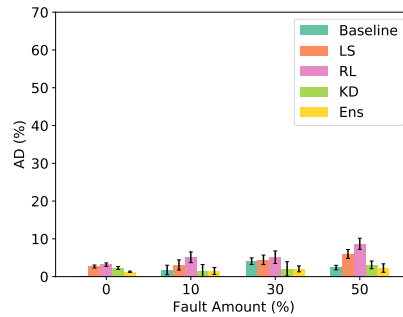
(h) GTSRB, VGG11, Removal



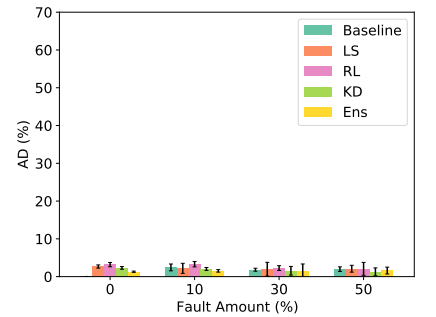
(i) GTSRB, VGG11, Repetition



(j) GTSRB, VGG16, Mislabelling



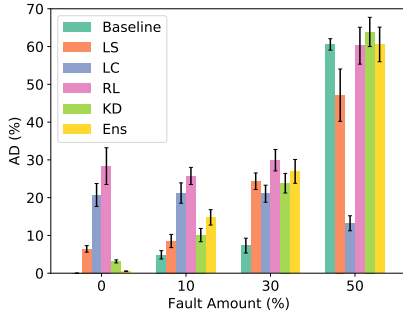
(k) GTSRB, VGG16, Removal



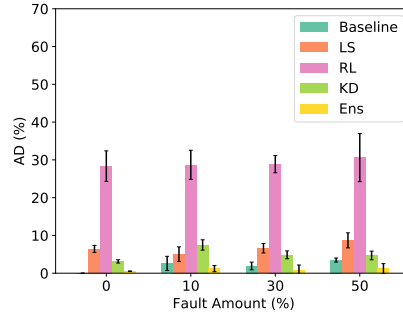
(l) GTSRB, VGG16, Repetition

Fig. 4: AD of individual models, compared with models protected with TDFM techniques when trained with faulty GTSRB datasets. The error bars in the results indicate the 95% confidence intervals. Lower values are better.

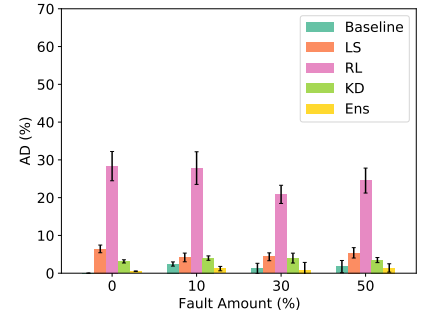
III. PNEUMONIA



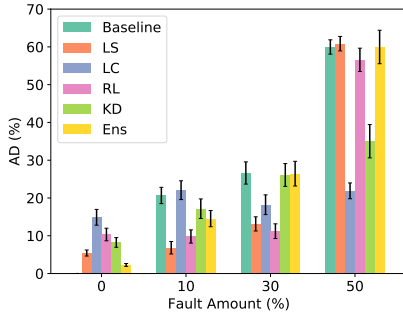
(a) Pneumonia, ConvNet, Mislabelling



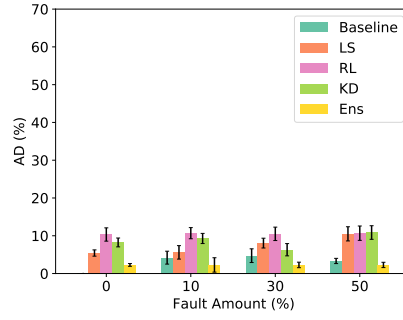
(b) Pneumonia, ConvNet, Removal



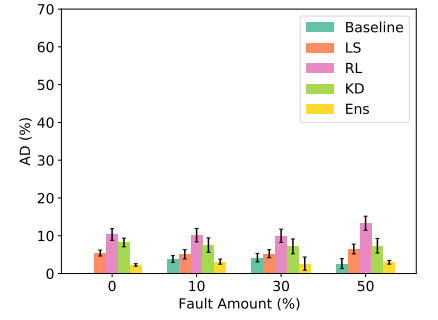
(c) Pneumonia, ConvNet, Repetition



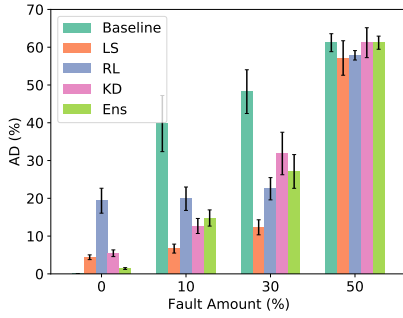
(d) Pneumonia, DeconvNet, Mislabelling



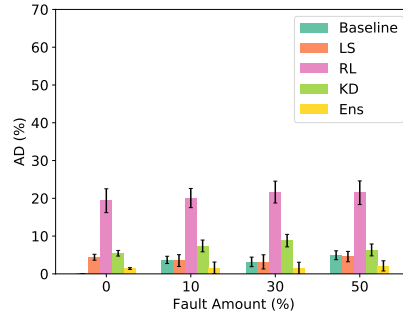
(e) Pneumonia, DeconvNet, Removal



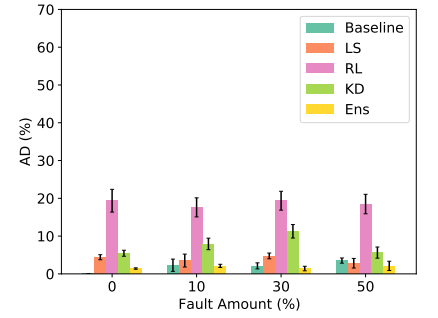
(f) Pneumonia, DeconvNet, Repetition



(g) Pneumonia, MobileNet, Mislabelling

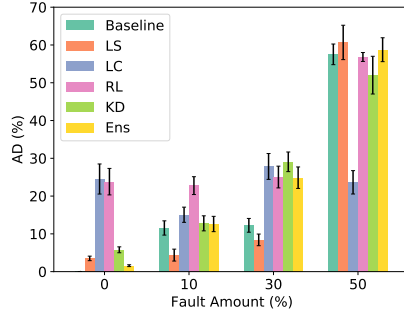


(h) Pneumonia, MobileNet, Removal

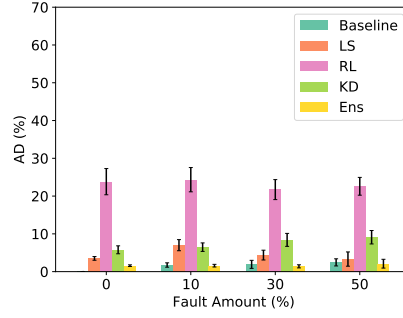


(i) Pneumonia, MobileNet, Repetition

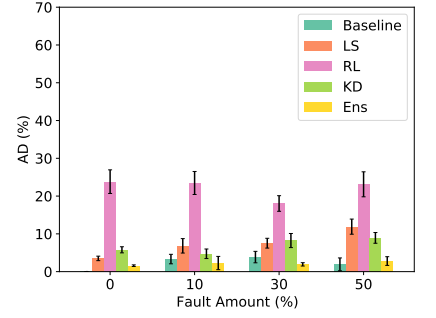
Fig. 5: AD of individual models, compared with models protected with TDFM techniques when trained with faulty Pneumonia datasets. The error bars in the results indicate the 95% confidence intervals. Lower values are better.



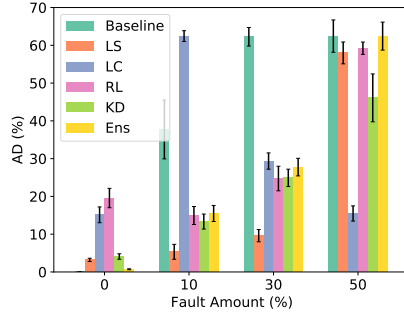
(a) Pneumonia, ResNet18, Mislabelling



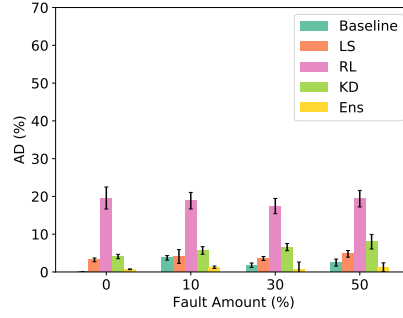
(b) Pneumonia, ResNet18, Removal



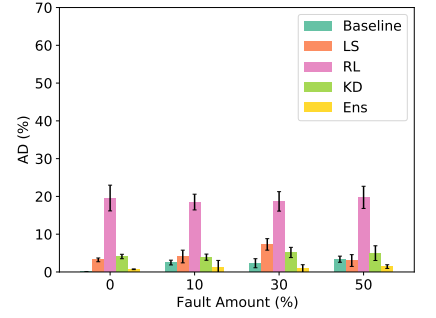
(c) Pneumonia, ResNet18, Repetition



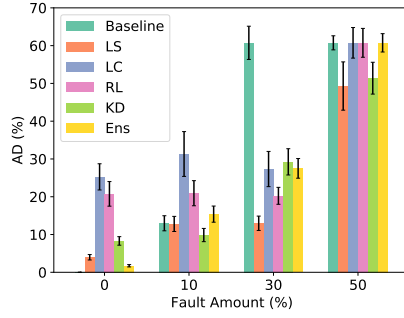
(d) Pneumonia, ResNet50, Mislabelling



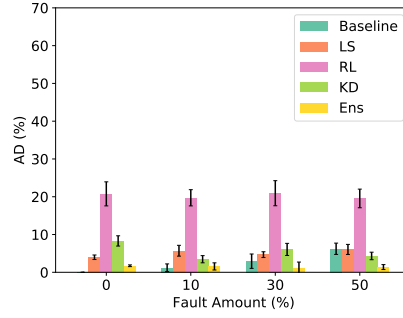
(e) Pneumonia, ResNet50, Removal



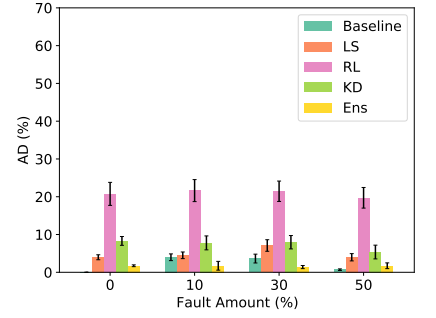
(f) Pneumonia, ResNet50, Repetition



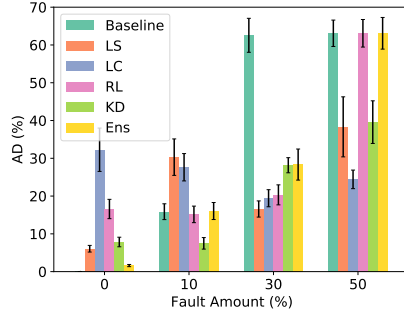
(g) Pneumonia, VGG11, Mislabelling



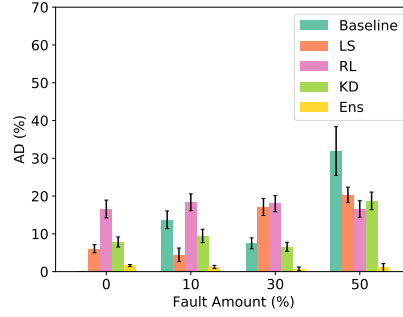
(h) Pneumonia, VGG11, Removal



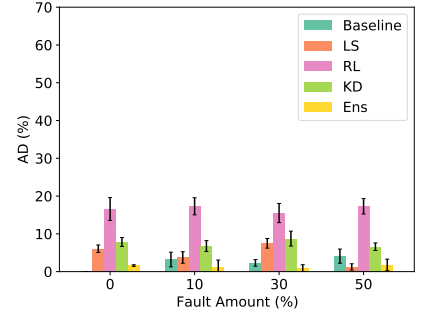
(i) Pneumonia, VGG11, Repetition



(j) Pneumonia, VGG16, Mislabelling



(k) Pneumonia, VGG16, Removal



(l) Pneumonia, VGG16, Repetition

Fig. 6: AD of individual models, compared with models protected with TDFM techniques when trained with faulty Pneumonia datasets. The error bars in the results indicate the 95% confidence intervals. Lower values are better.