

The Effect of Data Poisoning on Counterfactual Explanations – Appendix

No Author Given

No Institute Given

1 Proofs

1.1 Proof of Theorem 1

Proof. Sketch: For any $\vec{x}_{\text{orig}}, h(\vec{x}_{\text{orig}}) = y_{\text{orig}}$, assume uniqueness of the solution \vec{x}' – i.e. the closest sample to \vec{x}_{orig} on the decision boundary:

$$\begin{aligned} & \arg \min_{\vec{x}' \in \mathbb{R}^d} \|\vec{x}' - \vec{x}_{\text{orig}}\|_p \text{ s.t.} \\ & \exists i \neq j : (\vec{x}_i, y_i), (\vec{x}_j, y_j) \in \mathcal{D}, y_i \neq y_j, \text{ with } \|\vec{x}' - \vec{x}_i\|_p = \|\vec{x}' - \vec{x}_j\|_p \end{aligned} \quad (1)$$

where we (w.l.o.g.) assume the use of the p-norm as the distance function in the 1-nearest neighbor classifier.

Adding $(\vec{x}', y_{\text{orig}})$ to the training data \mathcal{D} implies that \vec{x}' is no longer the solution to Eq. (1). Therefore, the new closest sample on the decision boundary must have a larger distance to \vec{x}_{orig} than \vec{x}' , otherwise it would have been \vec{x}' before! \square

1.2 Proof of Theorem 2

Proof. Sketch: From the triangle-inequality and $\lambda > \|\vec{x}_i - \vec{x}_j\|_2$ it follows that:

$$\|\vec{x}_i - \vec{x}_j\|_2 + \delta'_j \geq \underbrace{\delta_i + \lambda}_{\delta'_i} \Leftrightarrow \delta'_j \geq \delta_i + \lambda - \|\vec{x}_i - \vec{x}_j\|_2 \quad (2)$$

Because of $\delta_j > \delta_i$, we know that $\delta_j = \alpha \delta_i$ for some $\alpha > 1$. This allows us to rewrite Eq. (2):

$$\delta'_j \geq \underbrace{\frac{\delta_j}{\alpha}}_{\delta_i} + \lambda - \|\vec{x}_i - \vec{x}_j\|_2 \quad (3)$$

The desired results follows from choosing $\lambda \geq 2\alpha\delta_j + \|\vec{x}_i - \vec{x}_j\|_2$ yields:

$$\begin{aligned} \delta'_j & \geq \frac{\delta_j}{\alpha} + \lambda - \|\vec{x}_i - \vec{x}_j\|_2 \\ & \geq \frac{\delta_j}{\alpha} + 2\alpha\delta_j + \|\vec{x}_i - \vec{x}_j\|_2 - \|\vec{x}_i - \vec{x}_j\|_2 \\ & = \delta_j \end{aligned} \quad (4)$$

\square

2 Experiments

2.1 Details on the Classifiers

- **RandomForest:** 10 decision tree classifiers each with a maximum depth of 7.
- **DNN:** 3-layer neural network with ReLU activation functions.

2.2 Local Poisoning Attack

Classifier	Data set	Nearest \uparrow	DiCE \uparrow	FACE \uparrow	Proto \uparrow
DNN	Diabetes	1.59	1.42	1.24	1.89

Table 1: Difference in the cost of recourse: no vs. local poisoning. Positive numbers denote an increase in the cost of recourse. We report the median (over all folds) rounded to two decimal places.

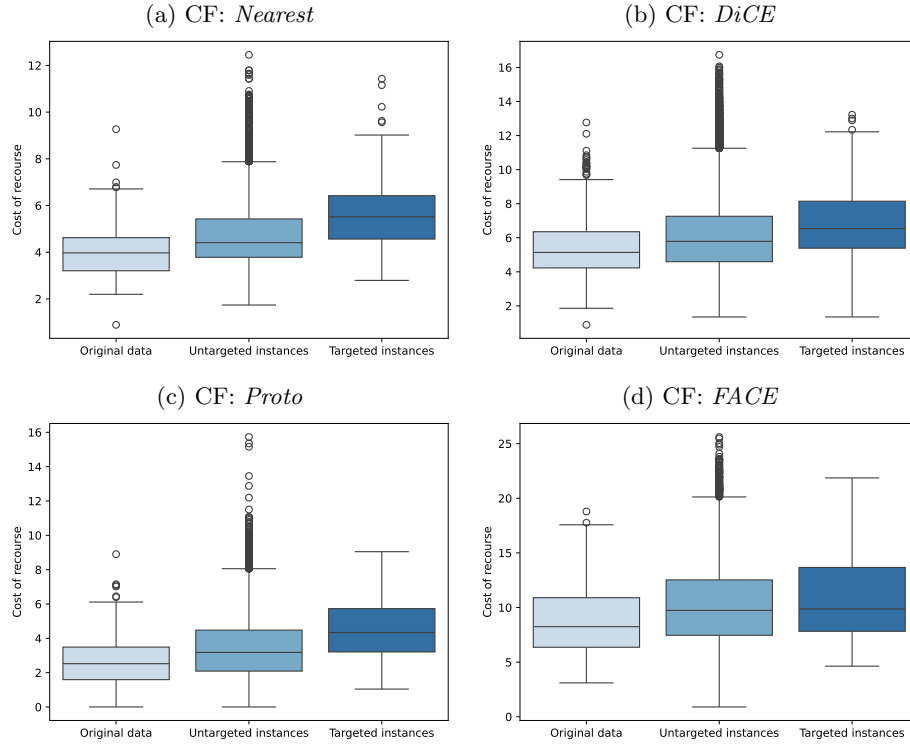
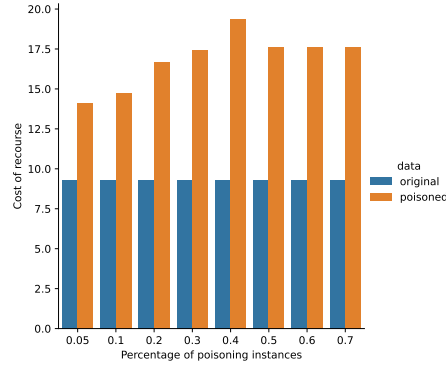
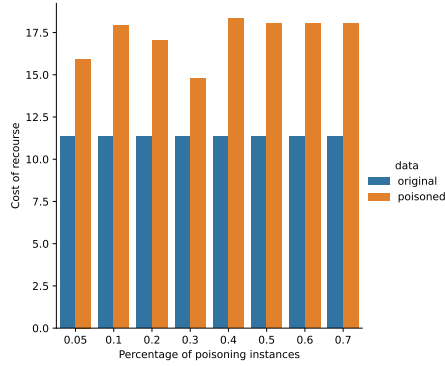
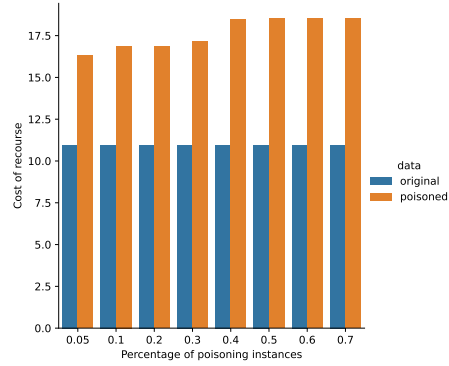
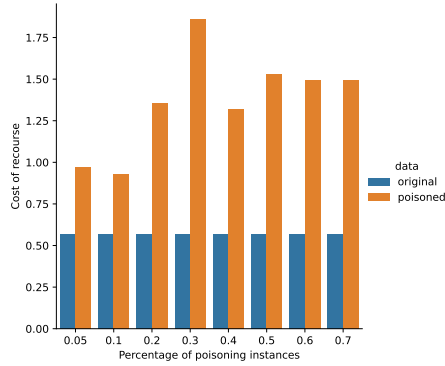
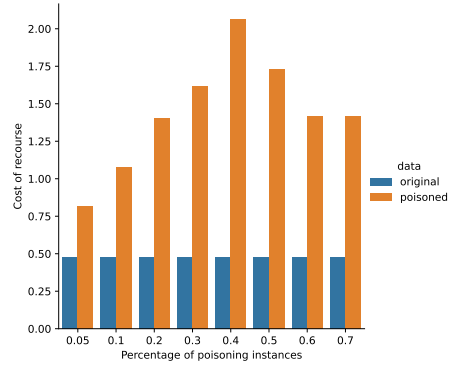
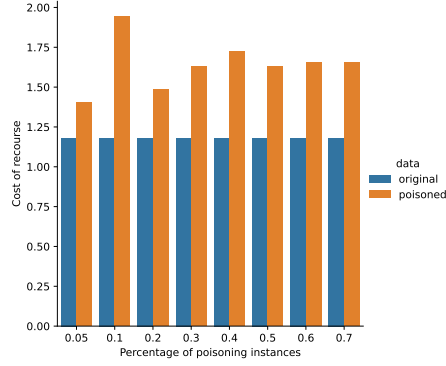
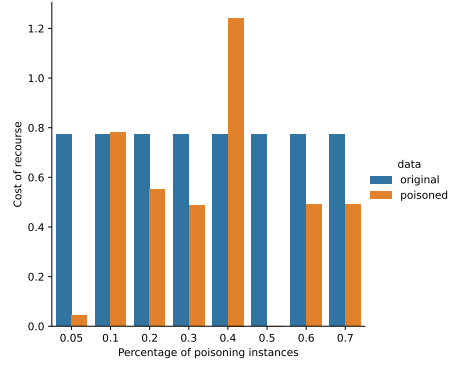
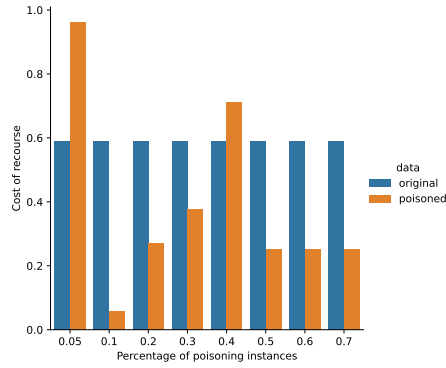
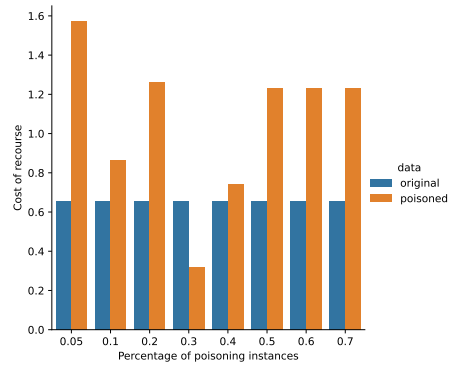
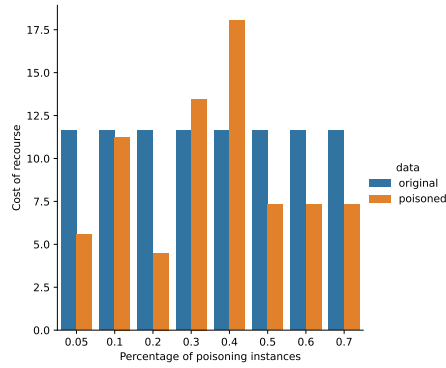
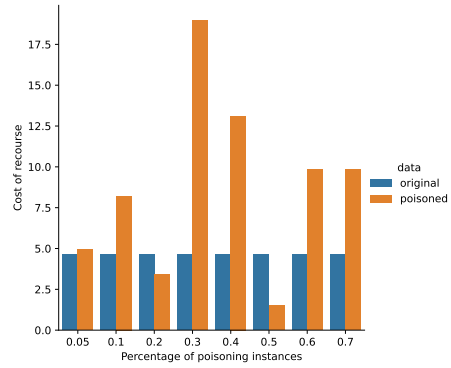
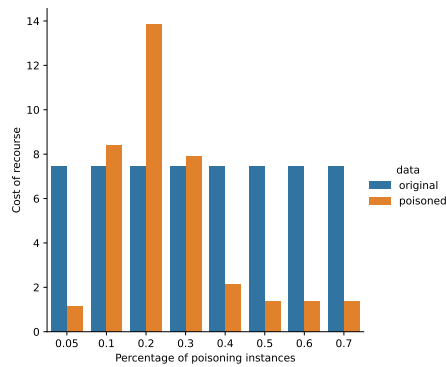
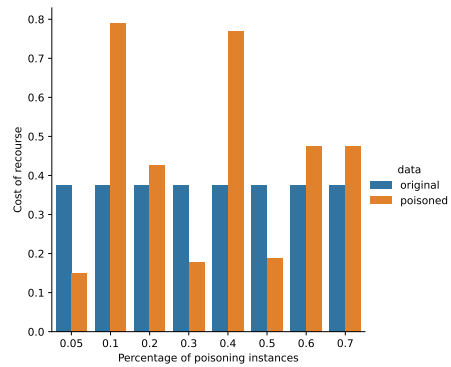
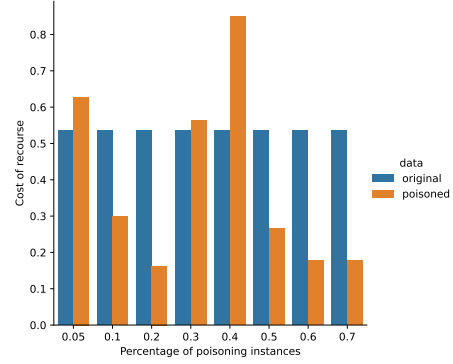
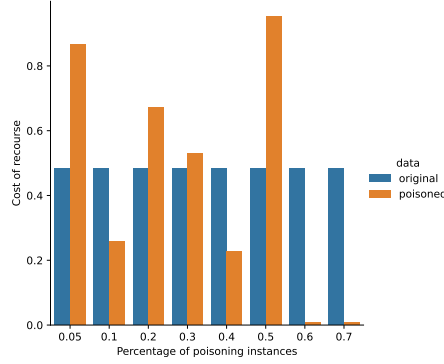
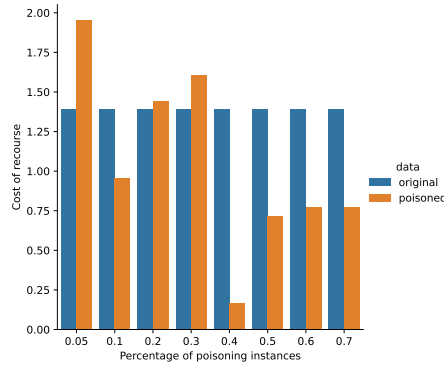
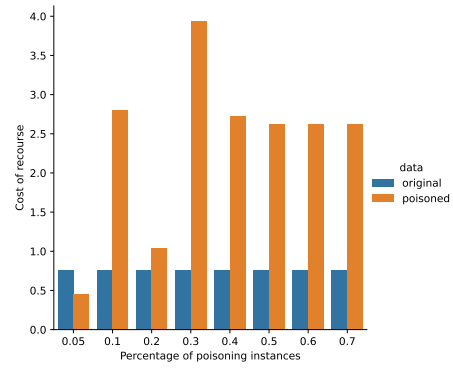
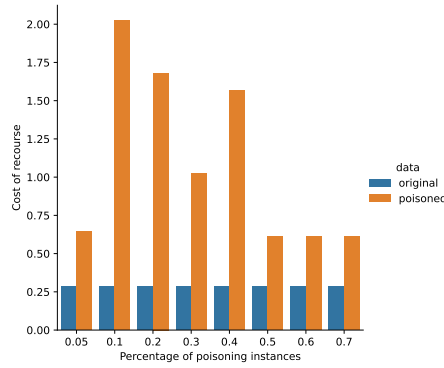
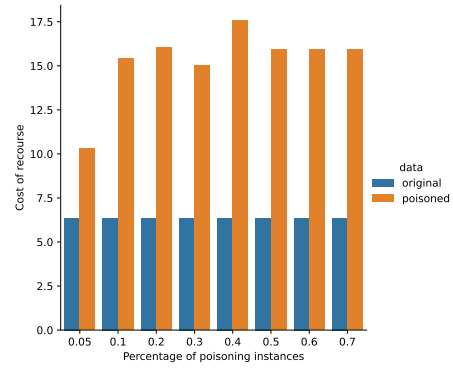
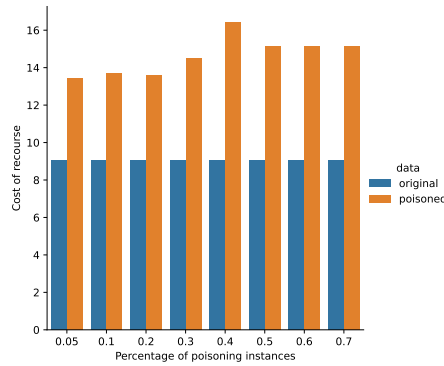
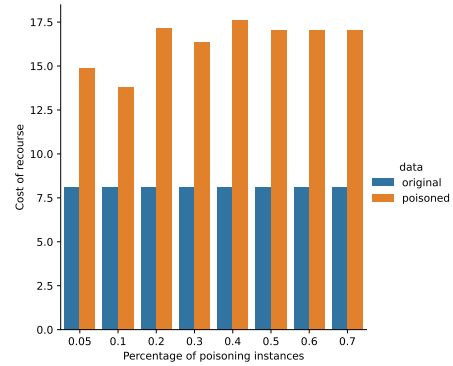
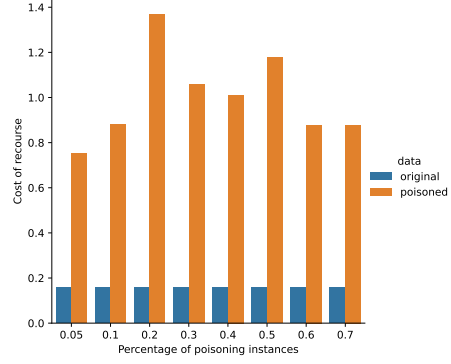
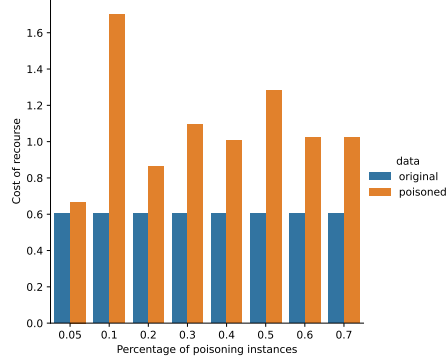
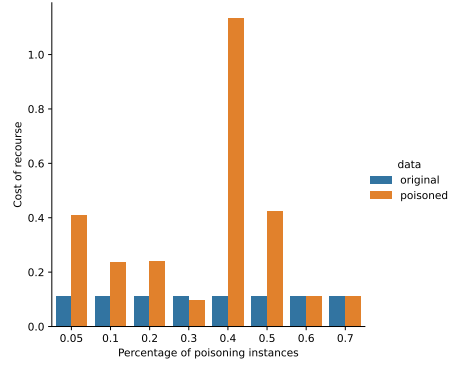
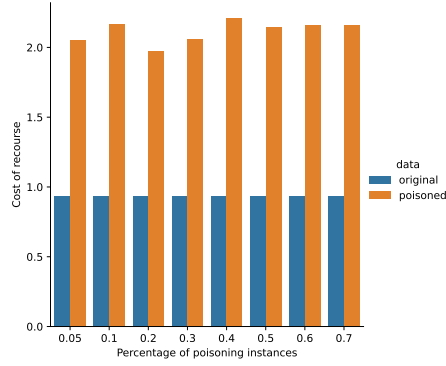
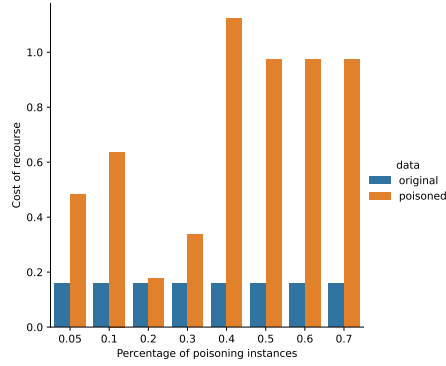
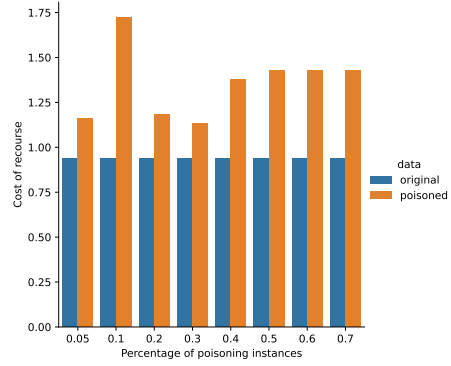
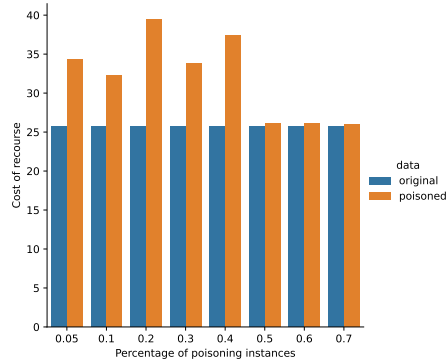
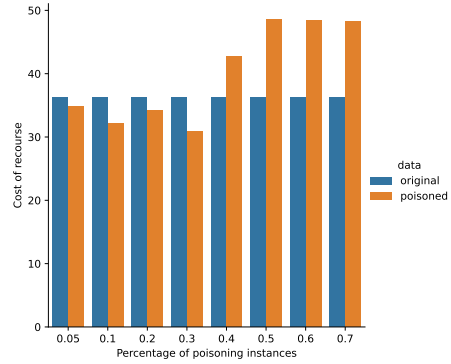


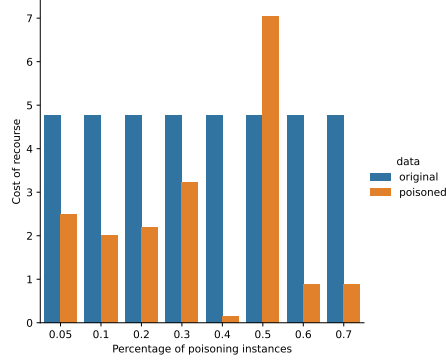
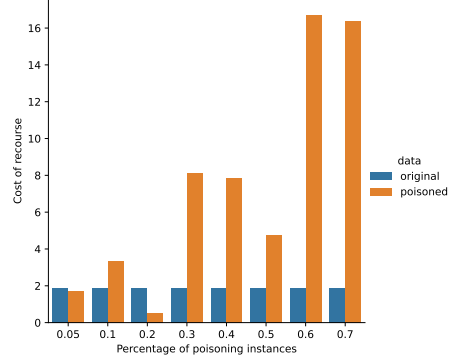
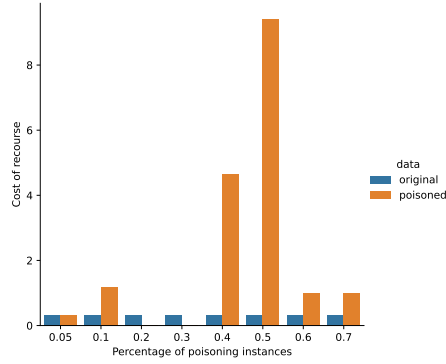
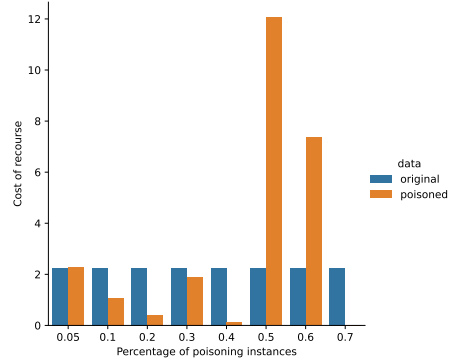
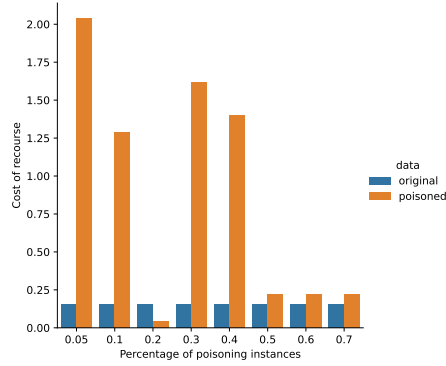
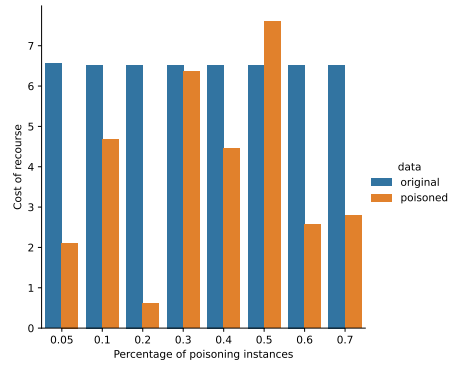
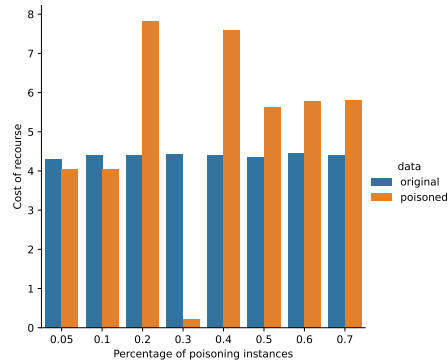
Fig. 1: *Local* data poisoning: Cost of recourse (over all test samples) in the case of the diabetes data set and a DNN classifier. Cost of recourse without any data poisoning, of untargeted instances and targeted instances in a local data poisoning.

27 **2.3 Sub-group Poisoning Attack**(a) \mathcal{D} : Crime – $h(\cdot)$: DNN – CF: $DiCE$ (b) \mathcal{D} : Crime – $h(\cdot)$: RNF – CF: $DiCE$ (c) \mathcal{D} : Crime – $h(\cdot)$: SVC – CF: $DiCE$ (d) \mathcal{D} : Diabetes – $h(\cdot)$: DNN – CF: $DiCE$ (e) \mathcal{D} : Diabetes – $h(\cdot)$: RNF – CF: $DiCE$ 

(a) \mathcal{D} : Diabetes – $h(\cdot)$: SVC – CF: *DiCE*(b) \mathcal{D} : Credit – $h(\cdot)$: DNN – CF: *DiCE*(c) \mathcal{D} : Credit – $h(\cdot)$: RNF – CF: *DiCE*(d) \mathcal{D} : Credit – $h(\cdot)$: SVC – CF: *DiCE*(e) \mathcal{D} : Crime – $h(\cdot)$: DNN – CF: *FACE*(f) \mathcal{D} : Crime – $h(\cdot)$: RNF – CF: *FACE*(g) \mathcal{D} : Crime – $h(\cdot)$: SVC – CF: *FACE*(h) \mathcal{D} : Diabetes – $h(\cdot)$: DNN – CF: *FACE*

(a) \mathcal{D} : Diabetes – $h(\cdot)$: RNF – CF: *FACE* (b) \mathcal{D} : Diabetes – $h(\cdot)$: SVC – CF: *FACE*(c) \mathcal{D} : Credit – $h(\cdot)$: DNN – CF: *FACE*(d) \mathcal{D} : Credit – $h(\cdot)$: RNF – CF: *FACE*(e) \mathcal{D} : Credit – $h(\cdot)$: SVC – CF: *FACE*(f) \mathcal{D} : Crime – $h(\cdot)$: DNN – CF: *Nearest*(g) \mathcal{D} : Crime – $h(\cdot)$: RNF – CF: *Nearest*(h) \mathcal{D} : Crime – $h(\cdot)$: SVC – CF: *Nearest*

(a) \mathcal{D} : Diabetes – $h(\cdot)$: DNN – CF: *Nearest* (b) \mathcal{D} : Diabetes – $h(\cdot)$: RNF – CF: *Nearest*(c) \mathcal{D} : Diabetes – $h(\cdot)$: SVC – CF: *Nearest* (d) \mathcal{D} : Credit – $h(\cdot)$: SVC – CF: *Nearest*(e) \mathcal{D} : Credit – $h(\cdot)$: RNF – CF: *Nearest*(f) \mathcal{D} : Credit – $h(\cdot)$: SVC – CF: *Nearest*(g) \mathcal{D} : Crime – $h(\cdot)$: DNN – CF: *Proto*(h) \mathcal{D} : Crime – $h(\cdot)$: SVC – CF: *Proto*

(a) \mathcal{D} : Diabetes – $h(\cdot)$: DNN – CF: *Proto*(b) \mathcal{D} : Diabetes – $h(\cdot)$: SVC – CF: *Proto*(c) \mathcal{D} : Credit – $h(\cdot)$: DNN – CF: *Proto*(d) \mathcal{D} : Credit – $h(\cdot)$: SVC – CF: *Proto*(e) \mathcal{D} : Credit – $h(\cdot)$: RNF – CF: *Proto*(f) \mathcal{D} : Diabetes – $h(\cdot)$: RNF – CF: *Proto*(g) \mathcal{D} : Crime – $h(\cdot)$: RNF – CF: *Proto*

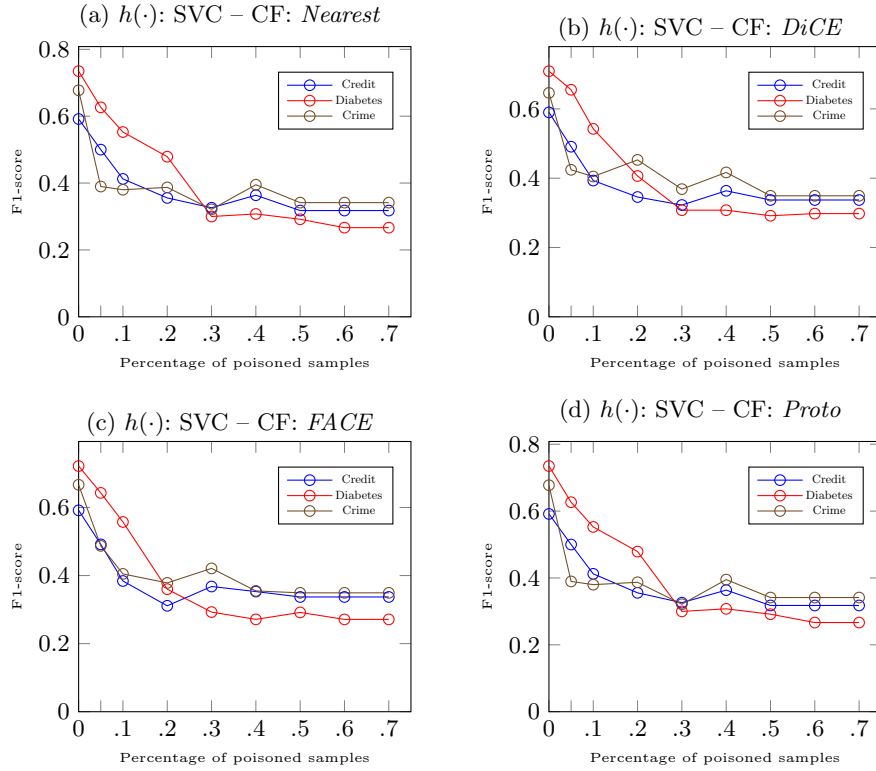


Fig. 7: Sub-group data poisoning attack: Median (over all folds) F1-score of the classifier for different percentages of poisoned samples (0% to 70%).

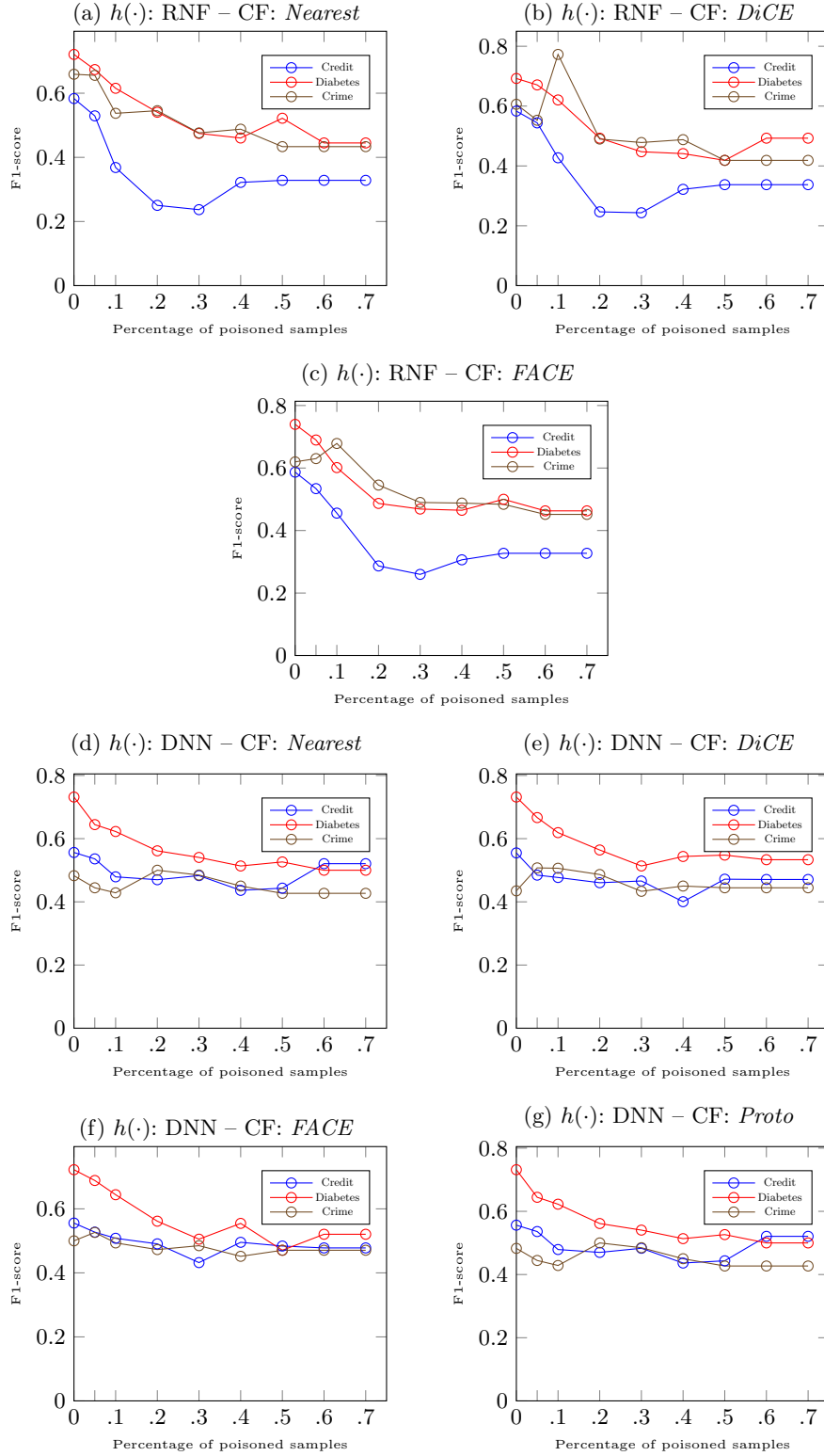


Fig. 8: Sub-group data poisoning attack: Median (over all folds) F1-score of the classifier for different percentages of poisoned instances (0% to 70%).

2.4 Global Poisoning Attack

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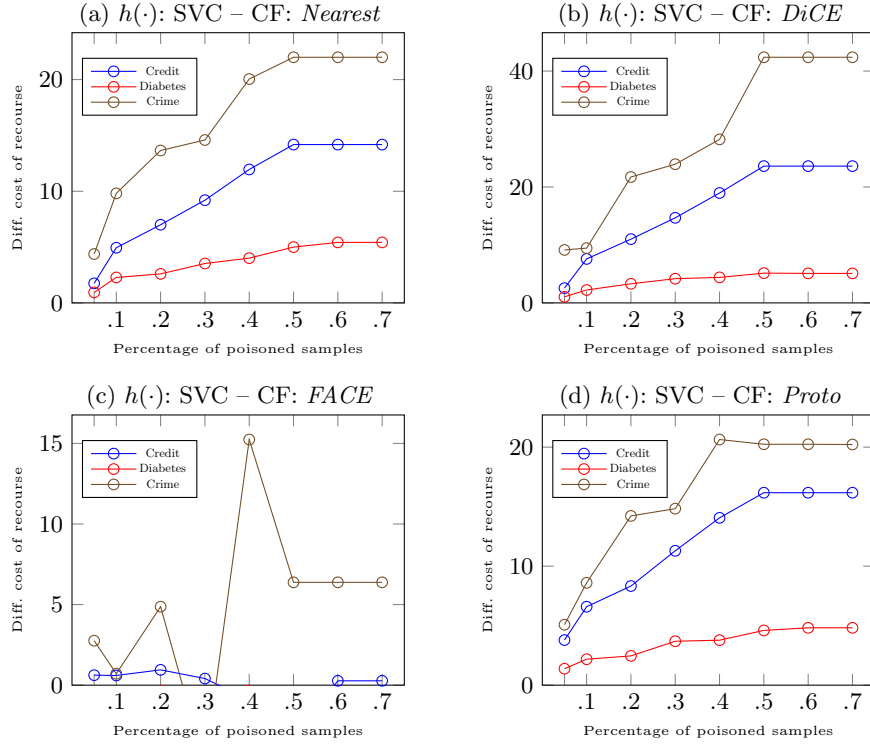


Fig. 9: Global data poisoning attack: Median (over all folds) difference in the cost of recourse vs. percentage of poisoned instances (5% to 70%).

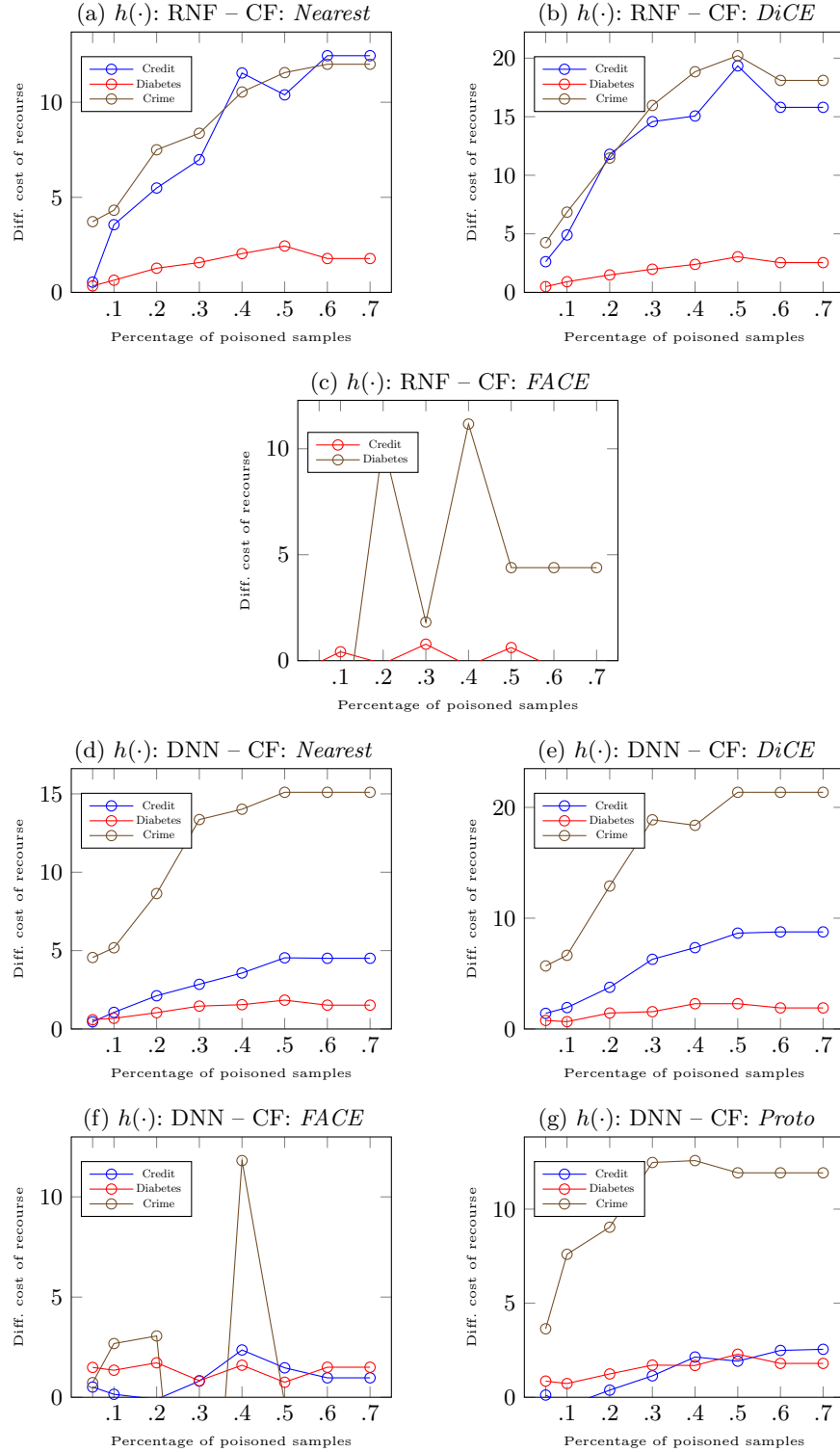


Fig. 10: Global data poisoning attack: Median (over all folds) difference in the cost of recourse vs. percentage of poisoned instances (5% to 70%).

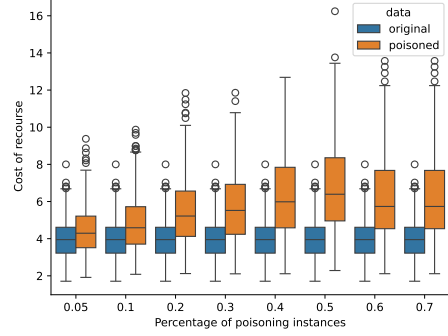
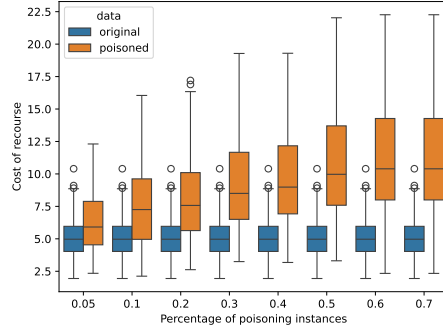
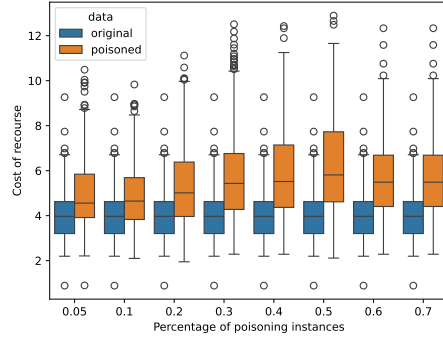
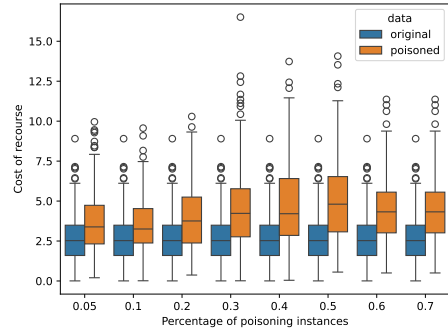
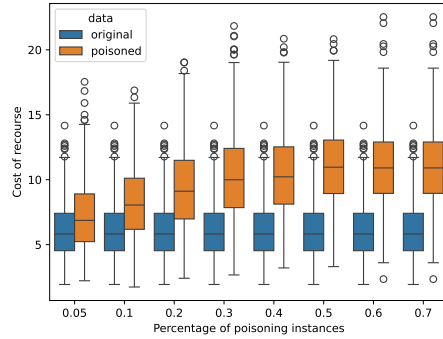
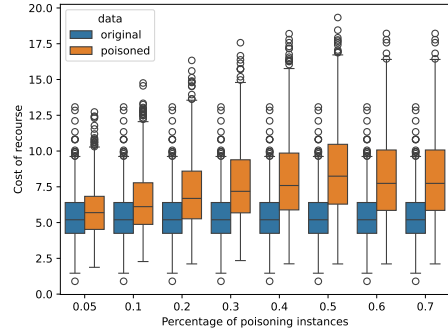
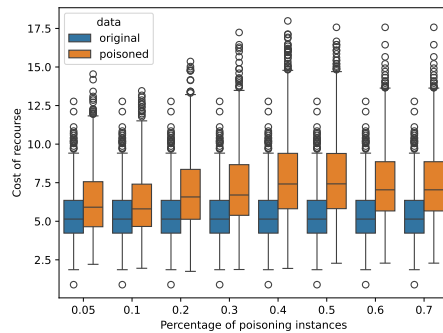
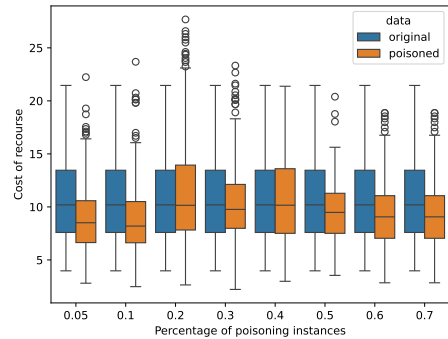
(a) \mathcal{D} : Diabetes – $h(\cdot)$: SVC – CF: *Nearest* (b) \mathcal{D} : Diabetes – $h(\cdot)$: RNF – CF: *Nearest*(c) \mathcal{D} : Diabetes – $h(\cdot)$: DNN – CF: *Nearest*(d) \mathcal{D} : Diabetes – $h(\cdot)$: DNN – CF: *Proto*(e) \mathcal{D} : Diabetes – $h(\cdot)$: SVC – CF: *DiCE*(f) \mathcal{D} : Diabetes – $h(\cdot)$: RNF – CF: *DiCE*(g) \mathcal{D} : Diabetes – $h(\cdot)$: DNN – CF: *DiCE*(h) \mathcal{D} : Diabetes – $h(\cdot)$: SVC – CF: *FACE*

Fig. 11: Cost of recourse (over all folds) of original data vs. poisoned data – (5% to 70% of poisoned instances).

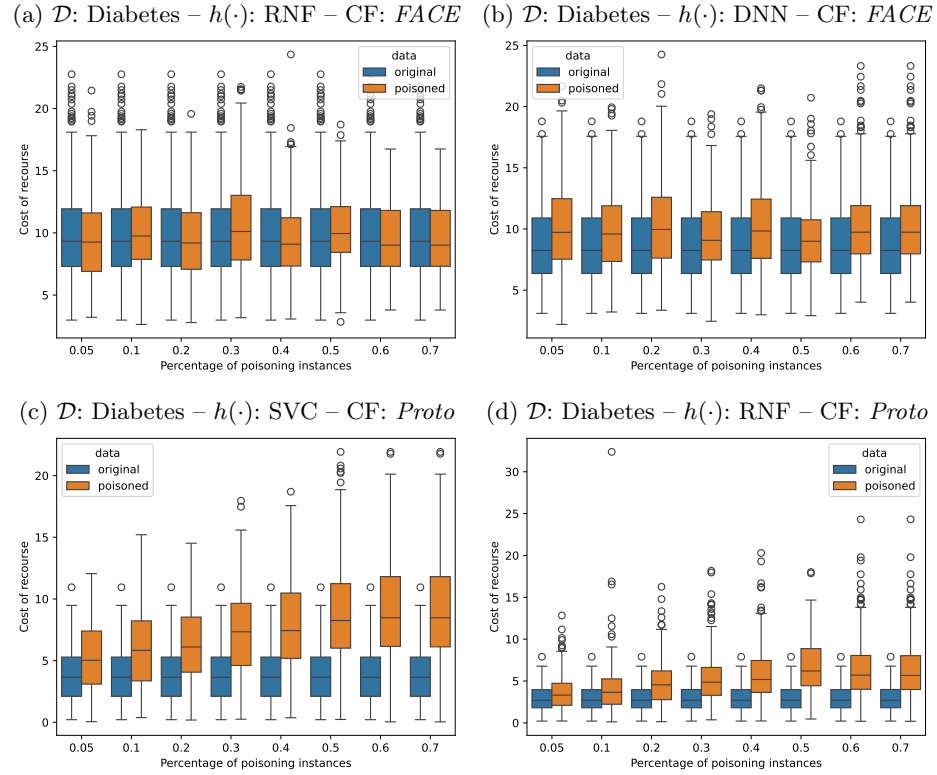


Fig. 12: Cost of recourse (over all folds) of original data vs. poisoned data – (5% to 70% of poisoned instances).

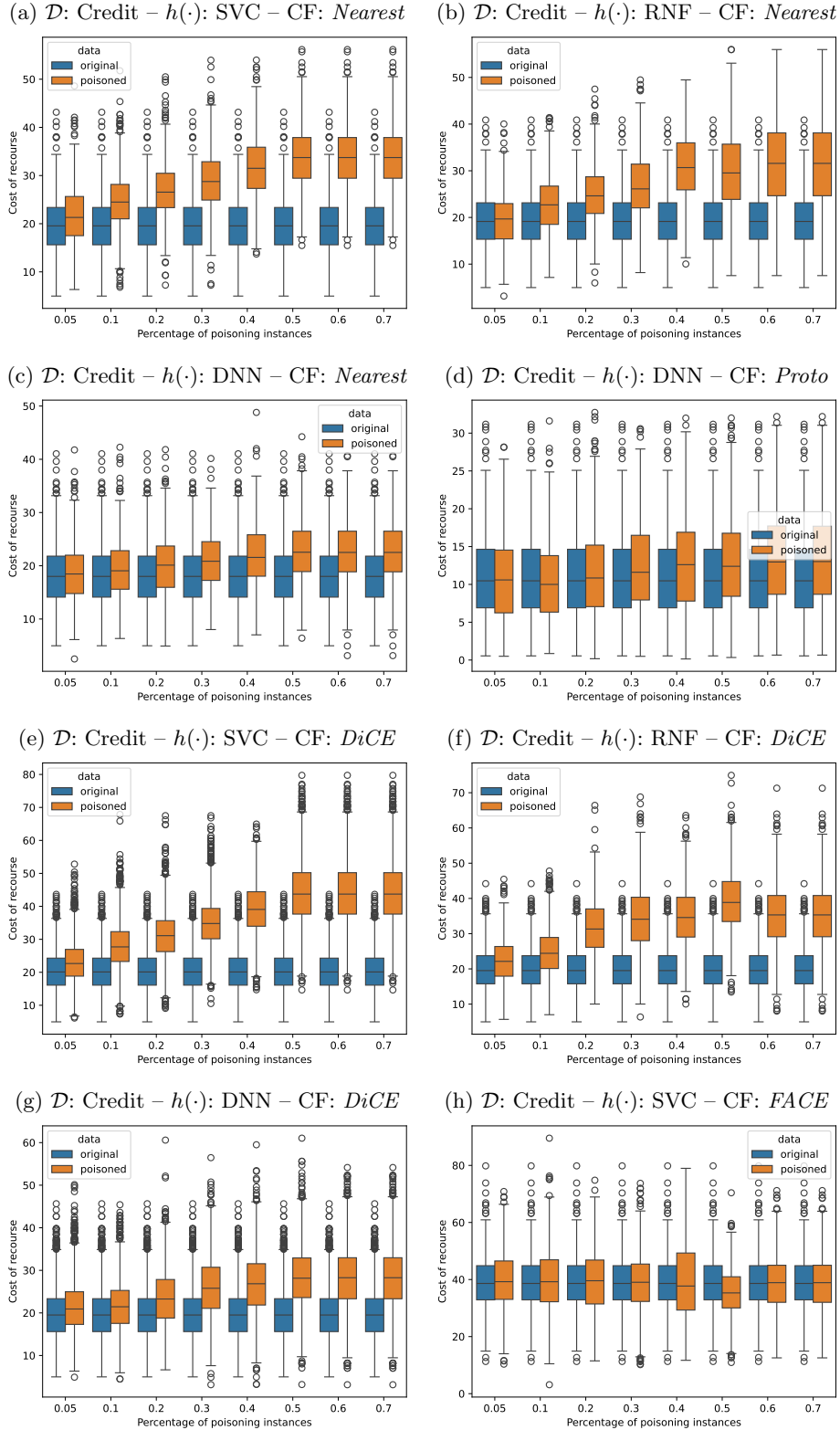


Fig. 13: Cost of recourse (over all folds) of original data vs. poisoned data – (5% to 70% of poisoned instances).

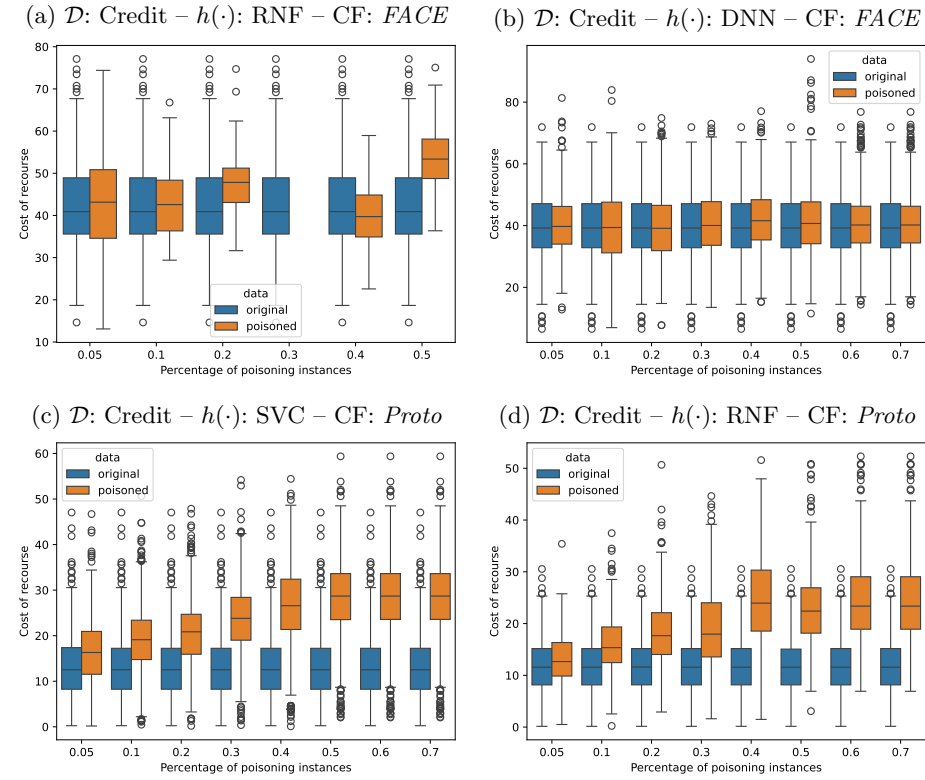


Fig. 14: Cost of recourse (over all folds) of original data vs. poisoned data – (5% to 70% of poisoned instances).

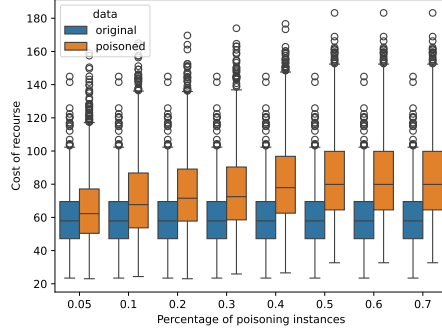
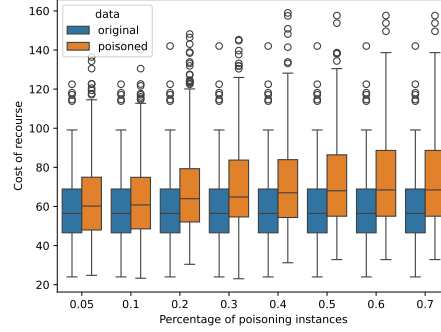
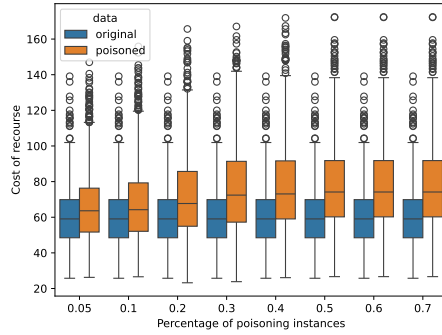
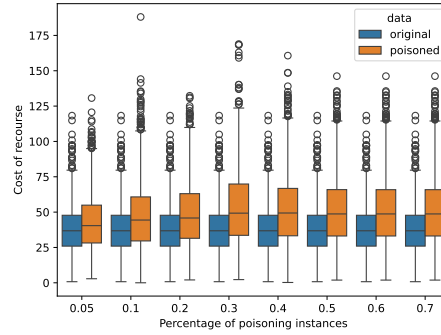
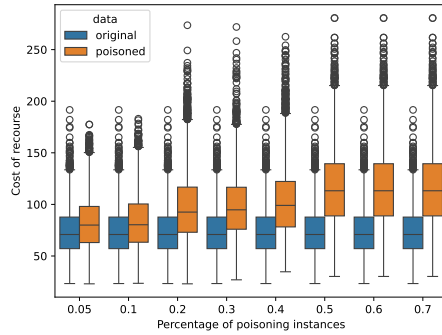
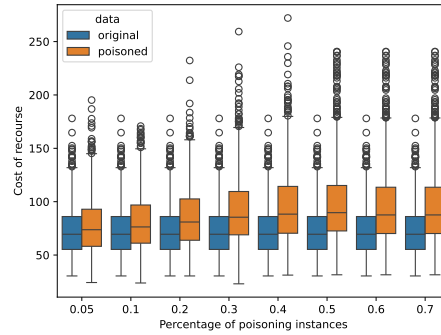
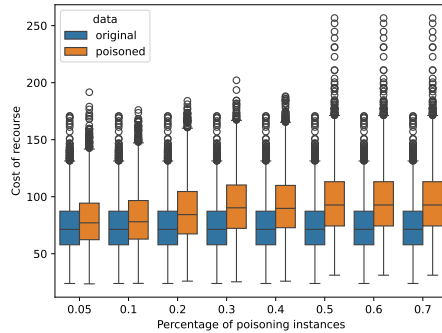
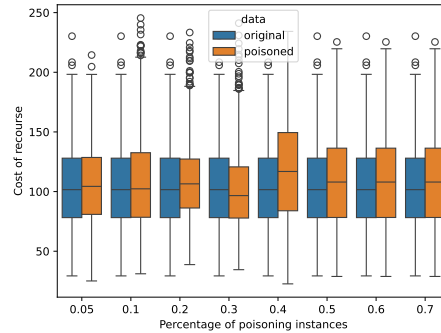
(a) \mathcal{D} : Crime – $h(\cdot)$: SVC – CF: *Nearest*(b) \mathcal{D} : Crime – $h(\cdot)$: RNF – CF: *Nearest*(c) \mathcal{D} : Crime – $h(\cdot)$: DNN – CF: *Nearest*(d) \mathcal{D} : Crime – $h(\cdot)$: DNN – CF: *Proto*(e) \mathcal{D} : Crime – $h(\cdot)$: SVC – CF: *DiCE*(f) \mathcal{D} : Crime – $h(\cdot)$: RNF – CF: *DiCE*(g) \mathcal{D} : Crime – $h(\cdot)$: DNN – CF: *DiCE*(h) \mathcal{D} : Crime – $h(\cdot)$: SVC – CF: *FAE*

Fig. 15: Cost of recourse (over all folds) of original data vs. poisoned data – (5% to 70% of poisoned instances).

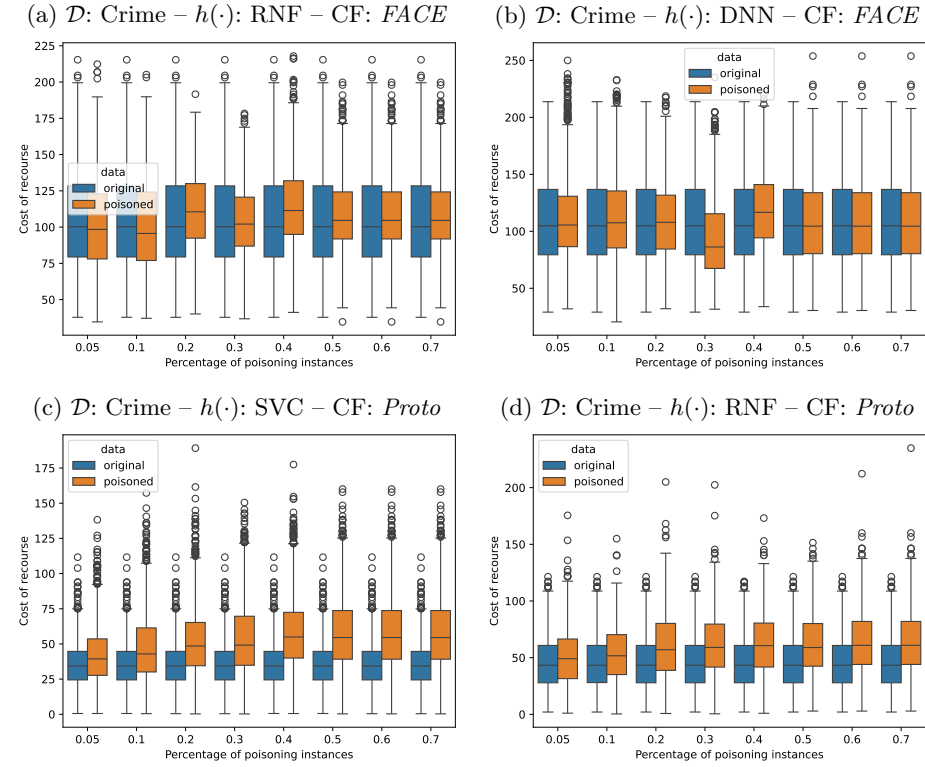


Fig. 16: Cost of recourse (over all folds) of original data vs. poisoned data – (5% to 70% of poisoned instances).

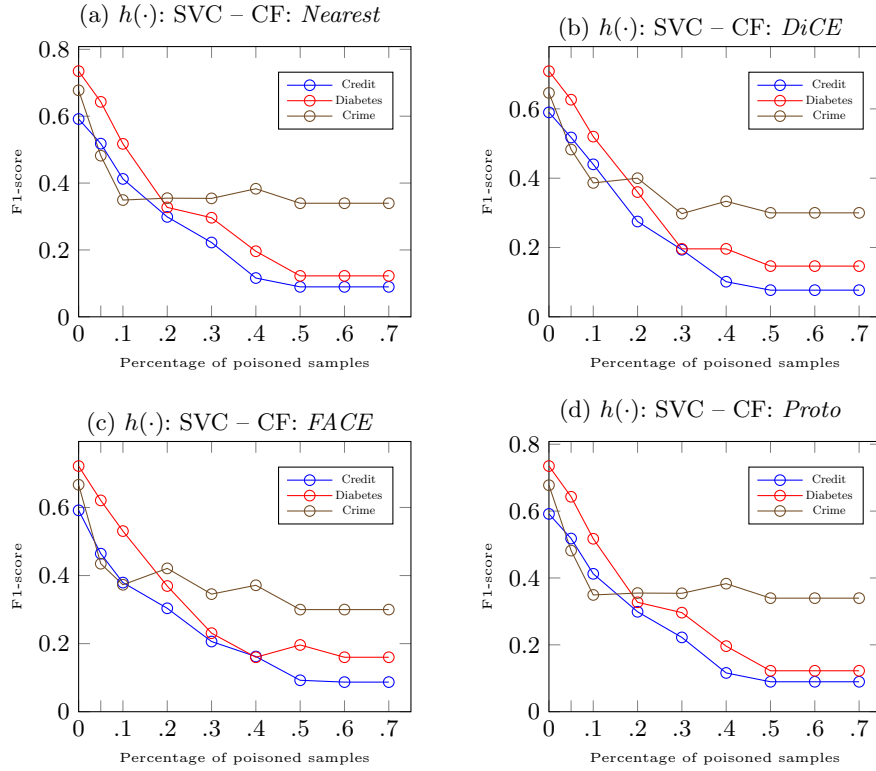


Fig. 17: Global data poisoning attack: Median (over all folds) F1-score of the classifier for different percentages of poisoned samples (0% to 70%).

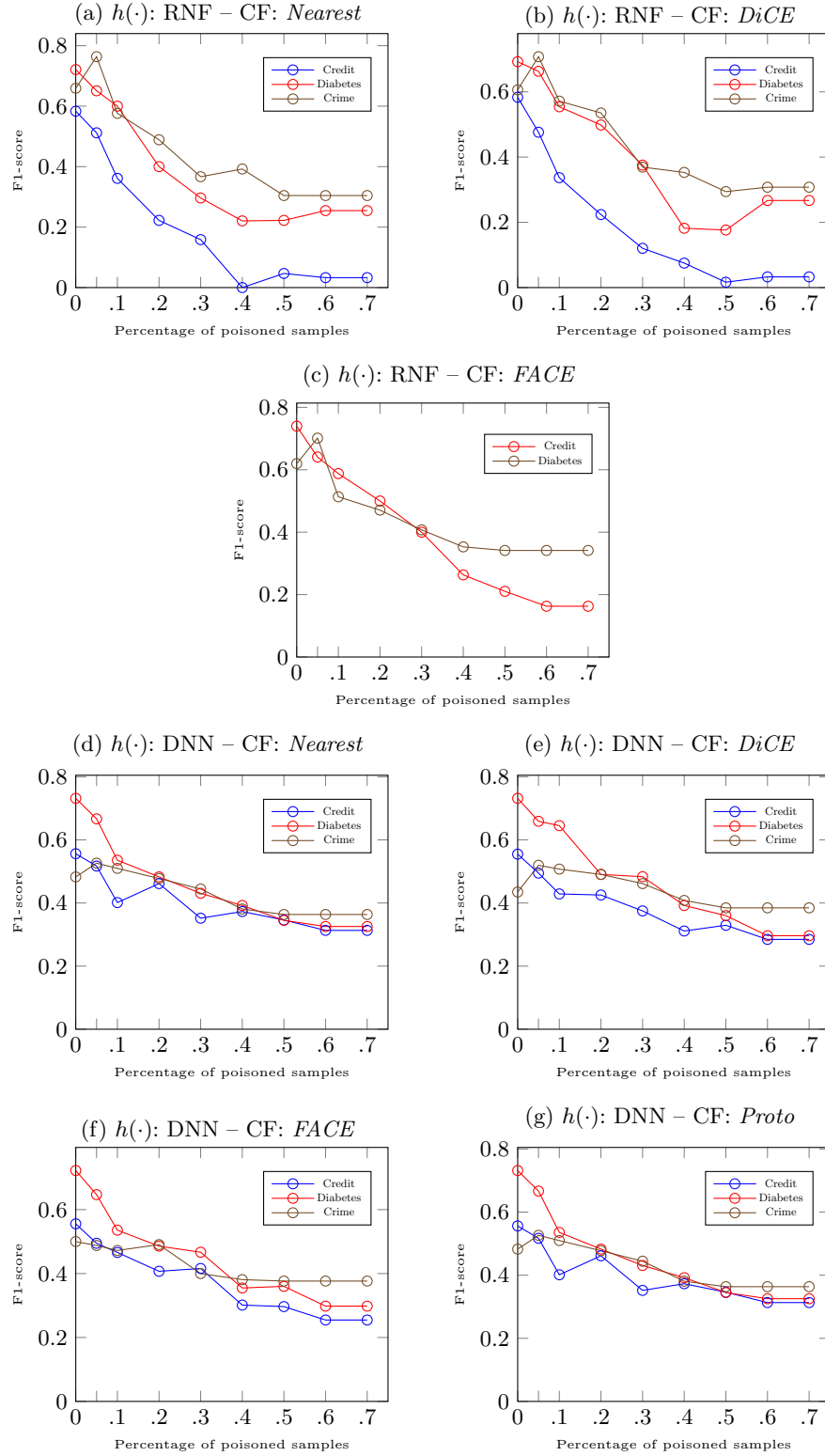


Fig. 18: Global data poisoning attack: Median (over all folds) F1-score of the classifier for different percentages of poisoned samples (0% to 70%).