

BENCHMARKING GENERATIVE AI MODELS FOR DEEP LEARNING TEST INPUT GENERATION











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Ministero dell'Università e della Ricerca







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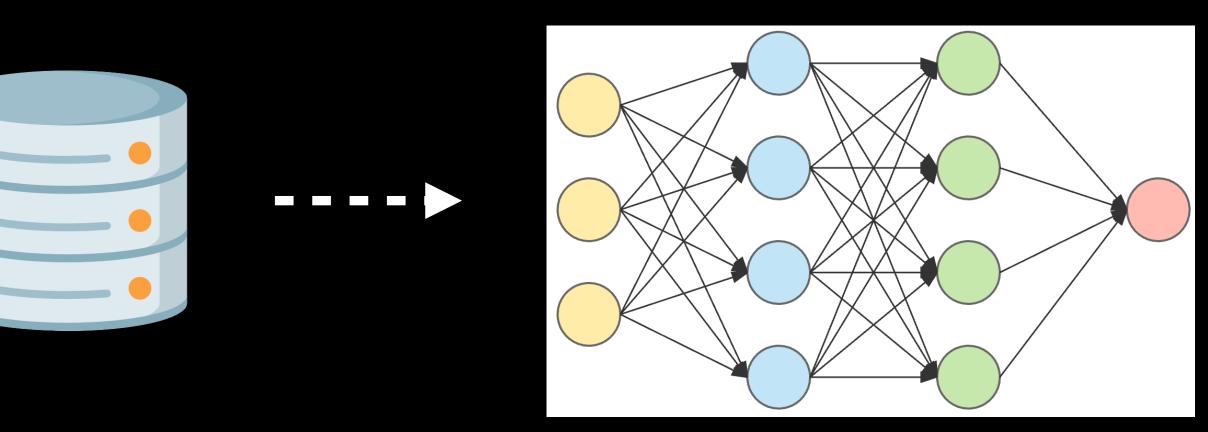








DNN ASSESSMENT







Problem: What is the performance of a DNN for inputs beyond its original dataset?

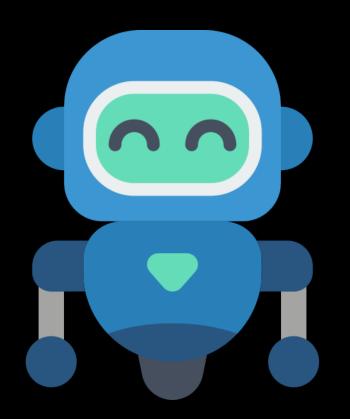
ACC = 0.95

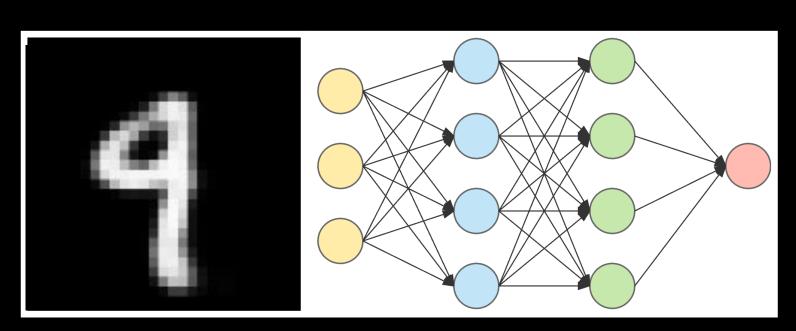
DNN UNDER TEST

PERFORMANCE METRIC



AUTOMATED TEST INPUT GENERATION FOR DNNS





TEST GENERATOR

Target Label

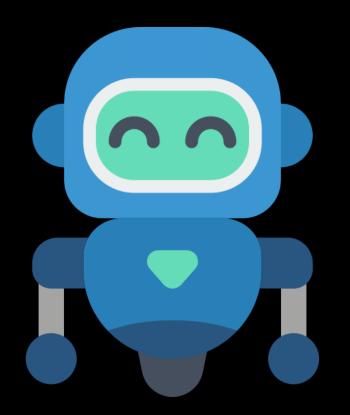
9

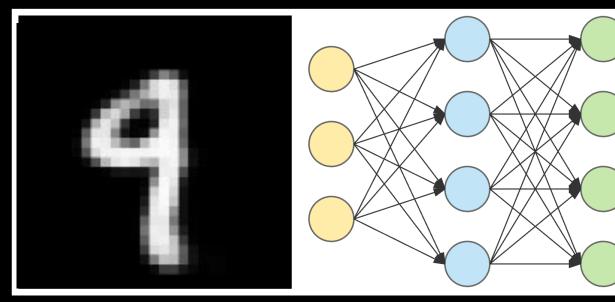
Predicted Label

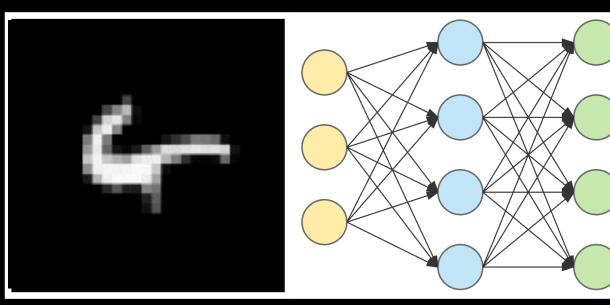




AUTOMATED TEST INPUT GENERATION FOR DNNS

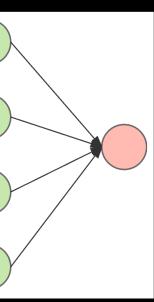






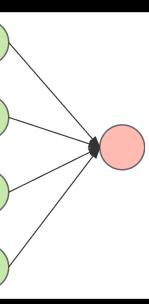
TEST GENERATOR











Predicted Label



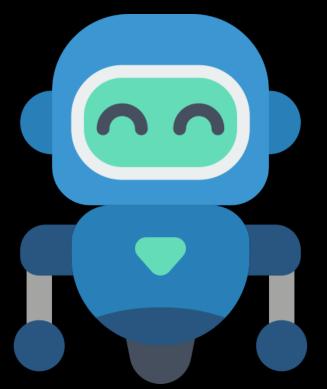
Problem #1:

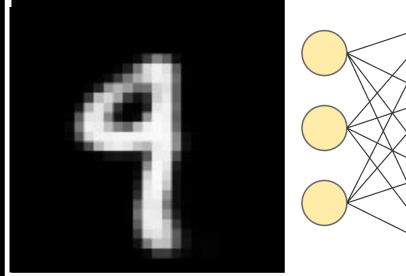
invalid inputs, not recognisable by domain experts in the input domain

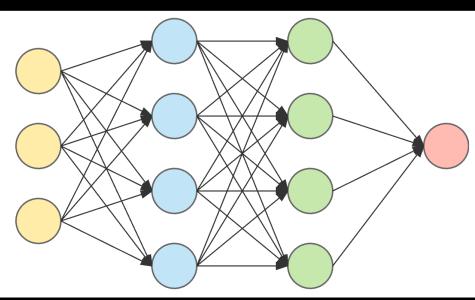




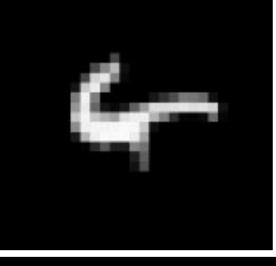
AUTOMATED TEST INPUT GENERATION FOR DNNS

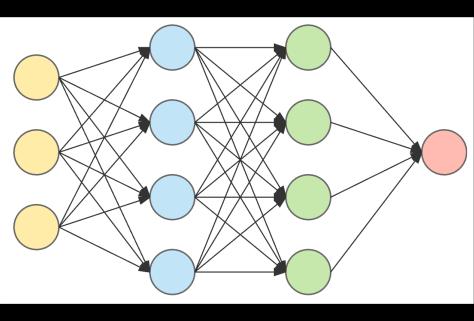




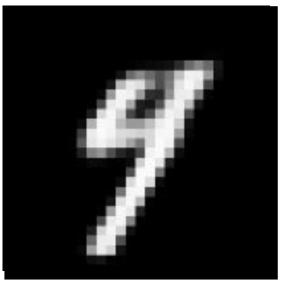


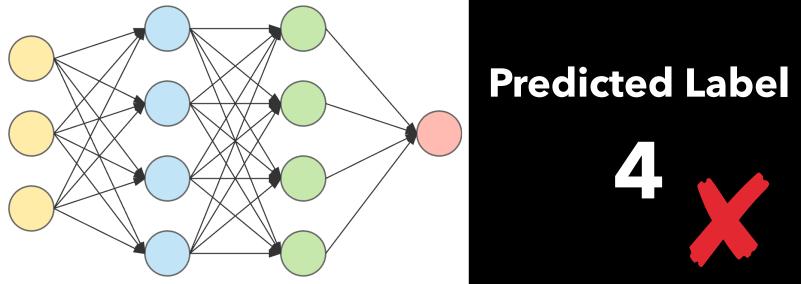
TEST GENERATOR















Predicted Label



Problem #1:

invalid inputs, not recognisable by domain experts in the input domain

Problem #2:

original label is not preserved

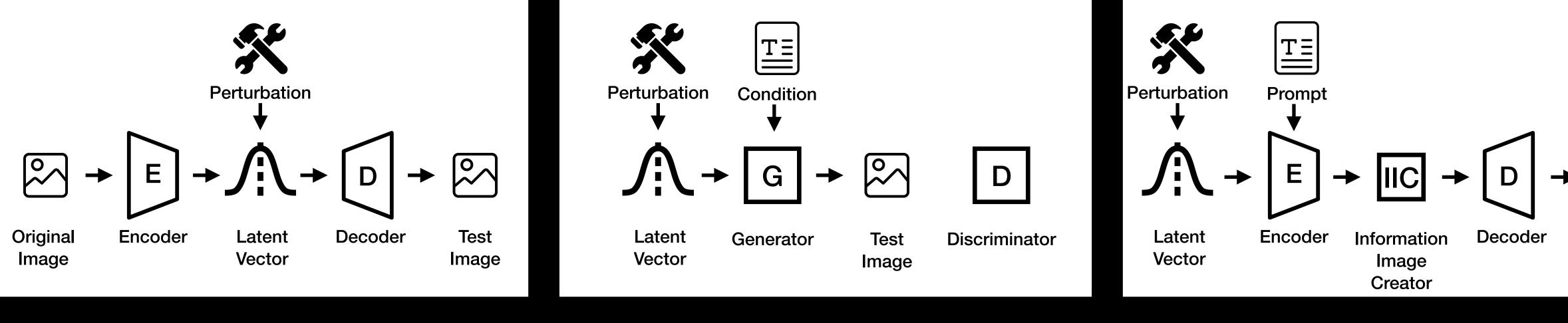






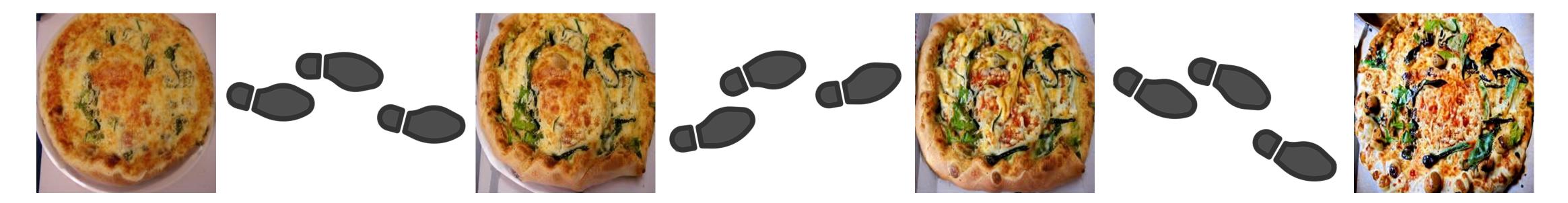






Variational AutoEncoder (VAE)

RANDOM WALK IN THE LATENT SPACE

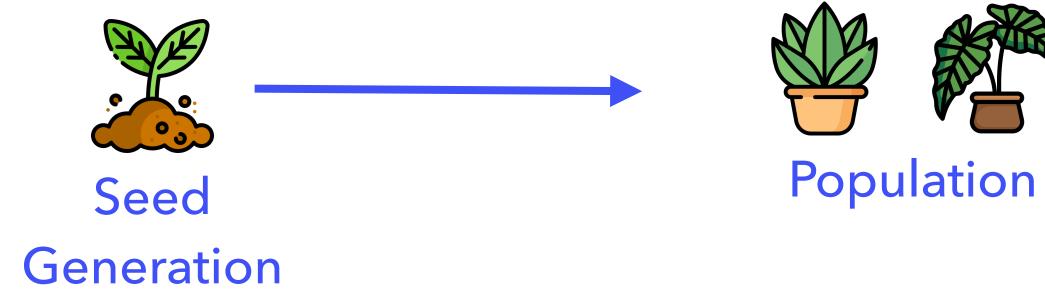


Generative Adversarial Network (GAN)

Diffusion Model (DM)



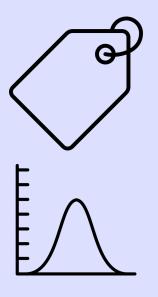








VAE: Image fed to the Encoder



GAN: Sample from target distribution + target label



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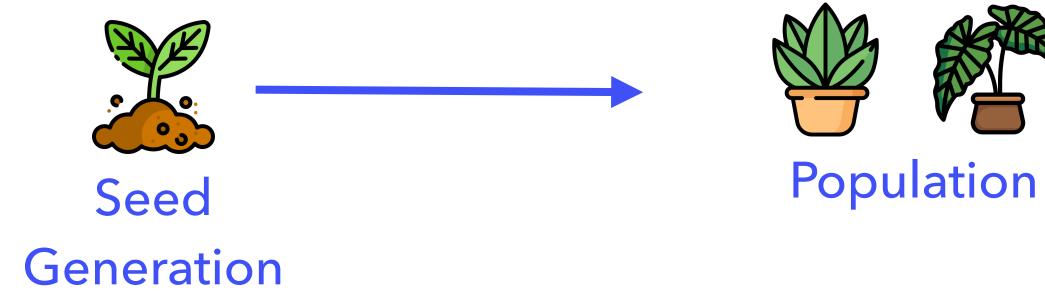
DM: Sample from target distribution + prompt



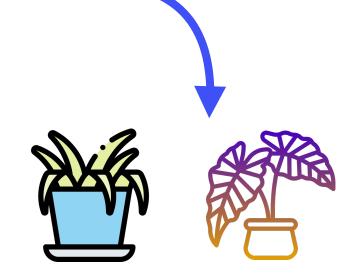












Crossover and Mutation in the Latent Space



Mutation: Random Latent Walk



Crossover: One-**Point Crossover** of Latent Vectors



Constraint: Clamping Vectors to Target Distribution

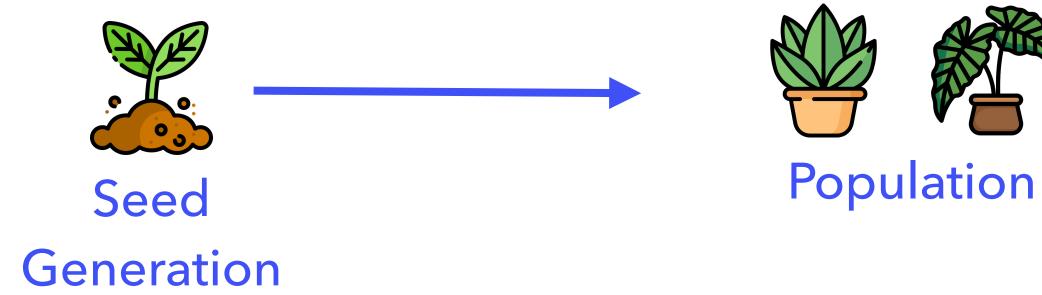






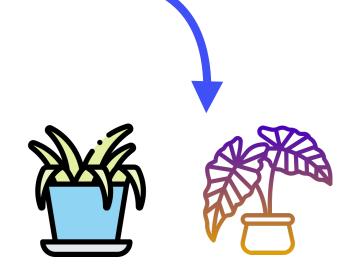




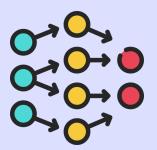








Crossover and Mutation in the Latent Space



Evaluation of generated images on the **DNN under test**

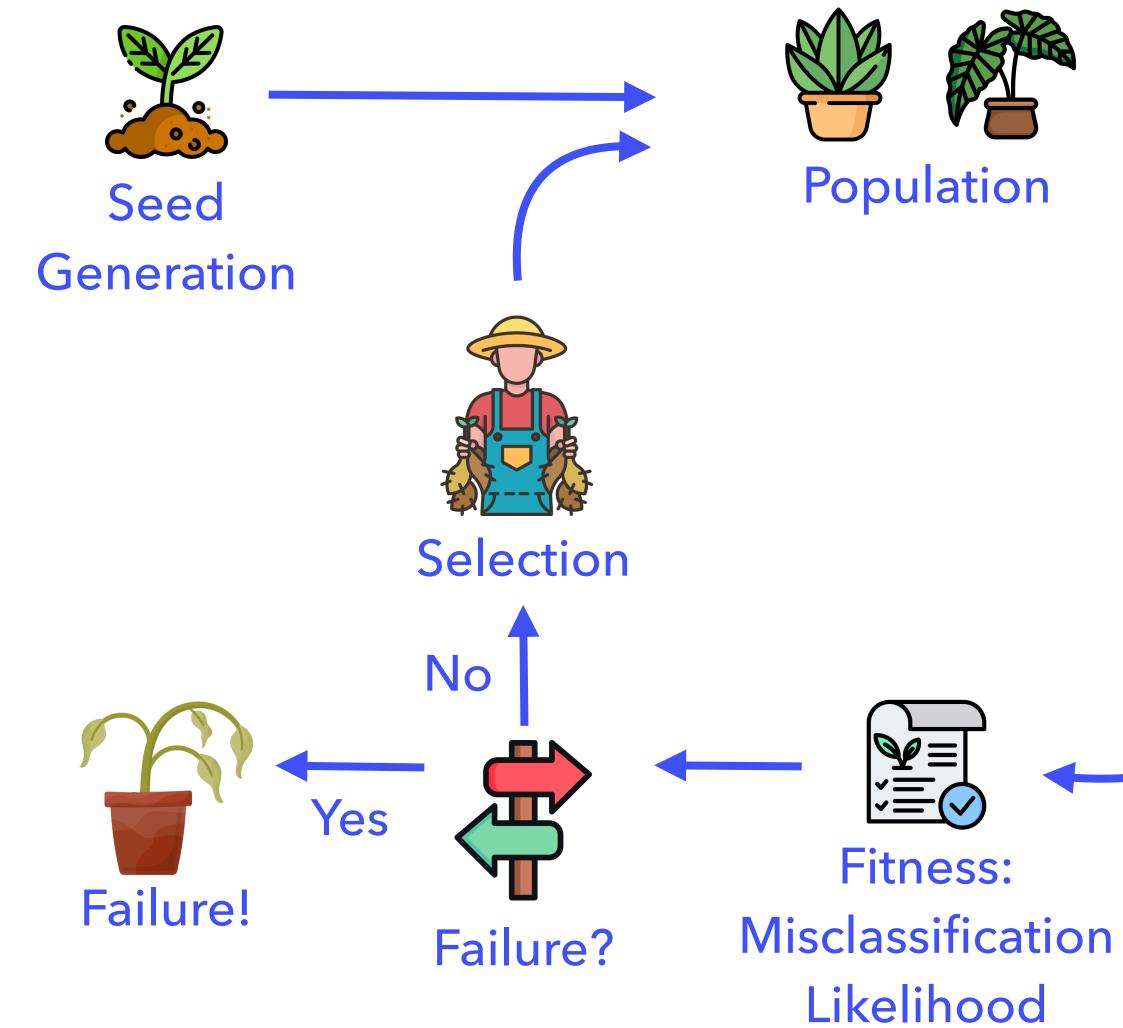


Compares confidence assigned to the target class VS other classes

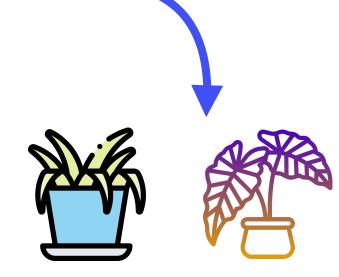












Crossover and Mutation in the Latent Space

Termination Conditions





Search budget exhausted









EVALUATION BENCHMARK

DATASETS







Imagenet

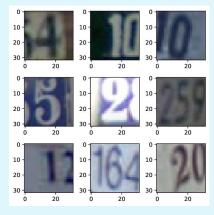








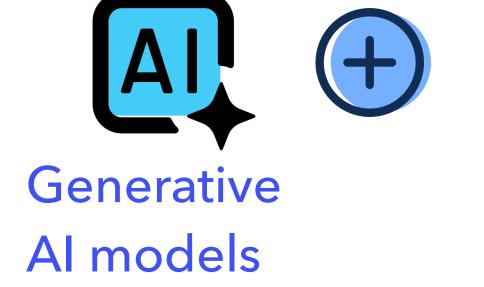
CIFAR-10

















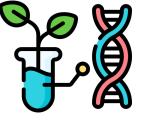
METRICS



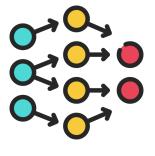
Effectiveness: # Valid labelpreserving misclassification -inducing inputs



Efficiency: # Iterations







DNN

Under Test

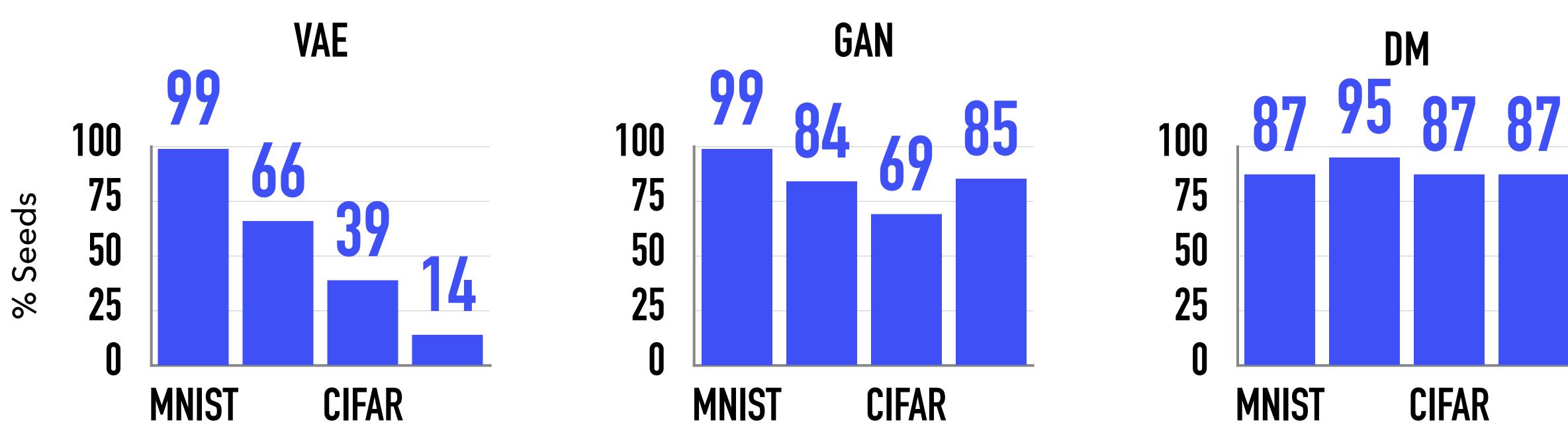
Genetic Algorithm

SETUP

Fixed search budget



GENFRA



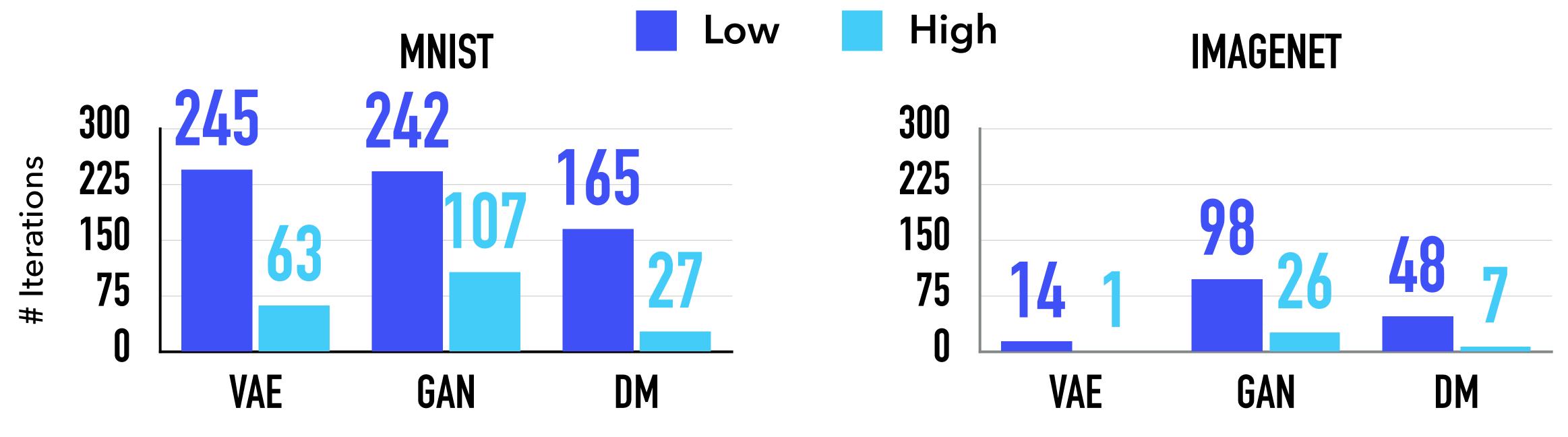
VAE performance declined as dataset complexity increased

GAN and DMs consistently achieved high accuracy in seed generation, regardless of complexity





FERATIONS TO TRIGGER A FAIL



VAEs need less iterations than other GenAI models for complex datasets, while DMs are the most efficient for simpler datasets

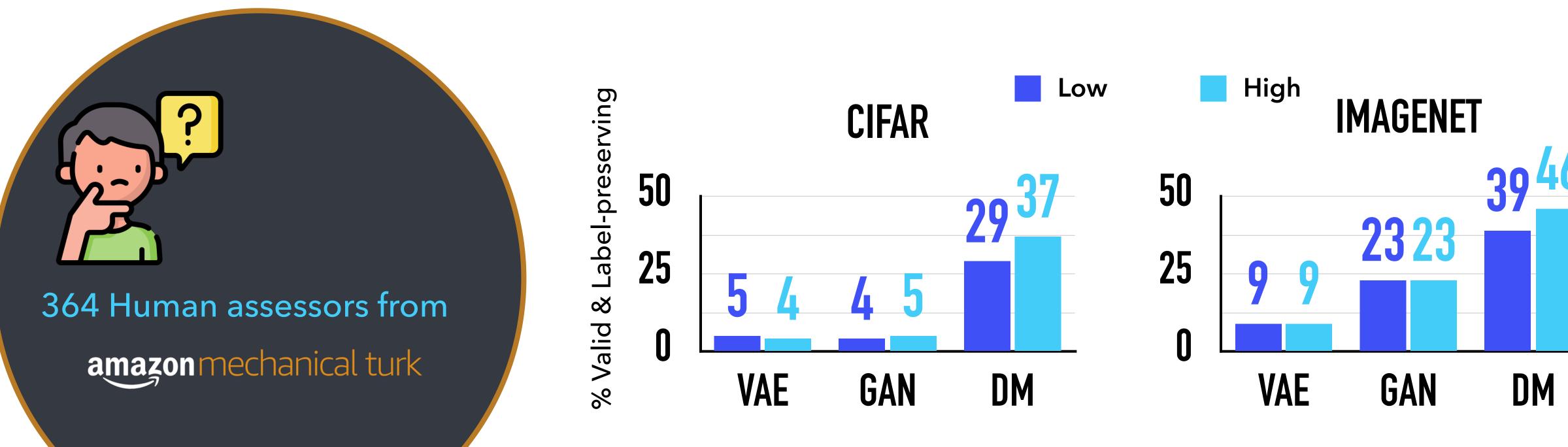
Increased perturbations reduce the number of iterations







VALIDITY AND LABEL PRESERVAT



DMs excel at generating valid misclassification-inducing inputs for complex datasets like CIFAR-10 and ImageNet









KEY INSIGHTS

Diffusion Models excel in complex tasks, but their superior performance comes at a higher cost

Larger perturbation extents speed up test generation without compromising input validity or label preservation

Latent vectors should be carefully constrained and carefully manipulated



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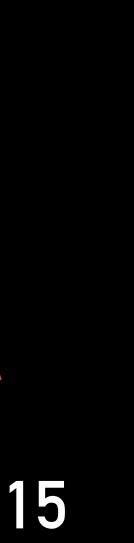
HIGHER FOR DM VS VAE

DMS UP TO 10X MORE

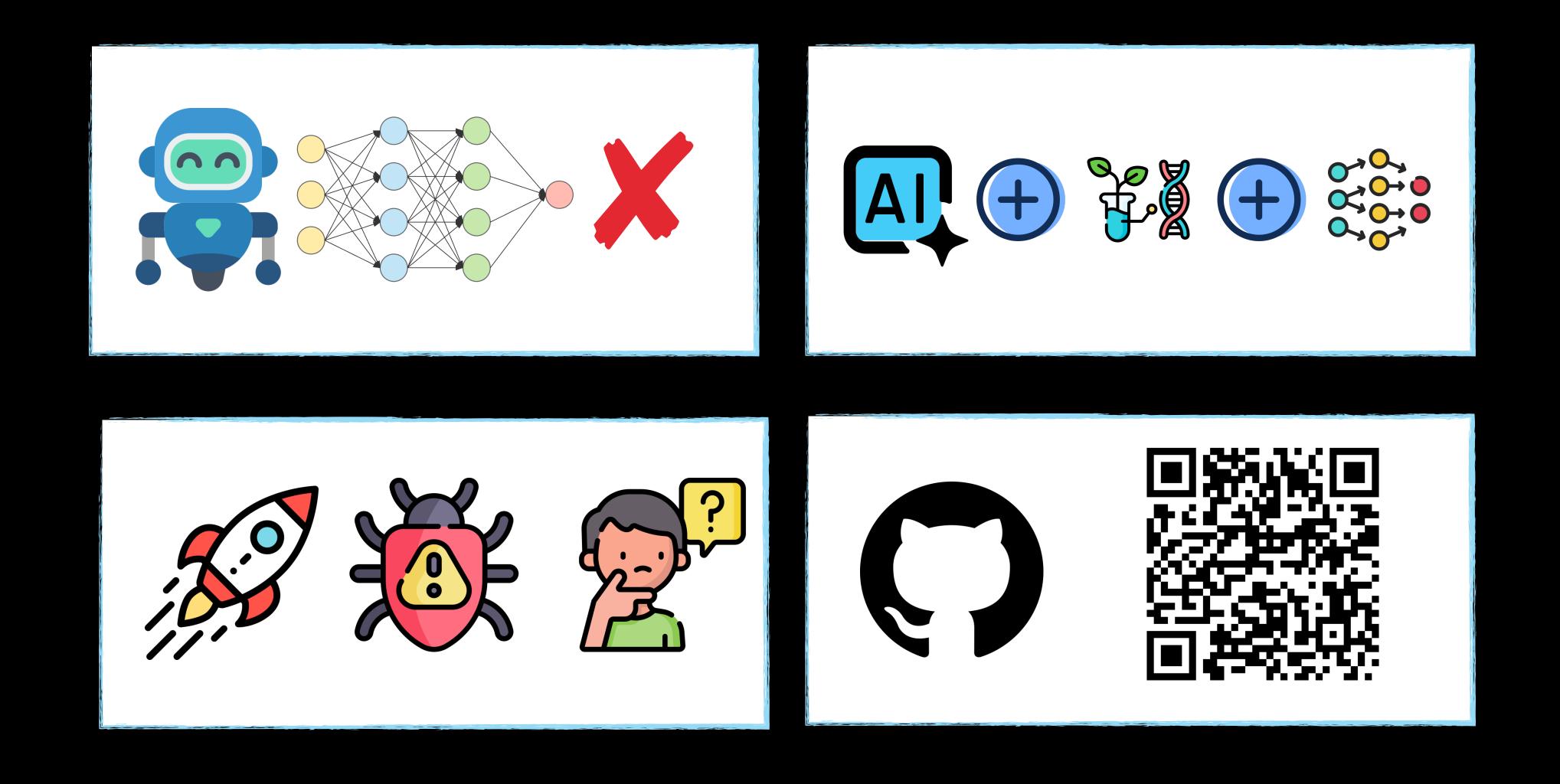
INFERENCE TIME 10X

EFFICIENT FOR IMAGENET AND SVHN AT MOST ONLY 15%

MISCLASSIFICATIONS FOR SVHN ARE VALID AND **PRESERVE THE LABEL**



BENCHMARKING GENERATIVE AI MODELS FOR DEEP LEARNING TEST INPUT GENERATION



Icons from <u>www.flaticon.com</u>





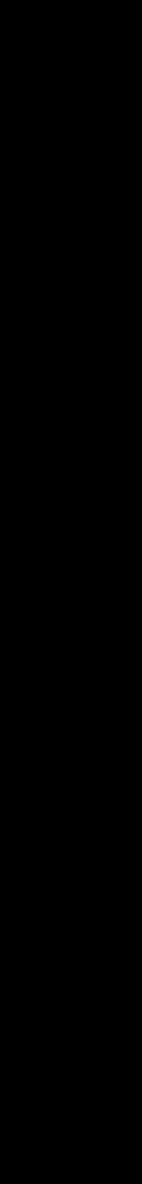




TABLE II CHARACTERISTICS OF THE GENAI MODELS: LATENT VECTOR SIZE, TRAINING TIME UNTIL CONVERGENCE, AVERAGE INFERENCE TIME.

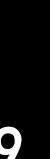
Dataset	Model	LV size	t_{train} (min)	t_{infer} (ms)
	VAE [49]	400	6	0.27
MNIST	GAN [50], [51]	100	9	0.7
	DM [52]	16384	405	960.68
SVHN	VAE [53]	800	93	4.07
	GAN [50], [51]	100	86	1.75
	DM [52]	16384	572	1213.49
	VAE [53]	1024	423	2.51
CIFAR-10	GAN [50], [51]	100	450	1.73
	DM [52]	16384	362	1903.29
	VAE [54]	512	2521	11.92
ImageNet	GAN [55]	128	21600	20.68
	DM [52]	16384	30	1945.77

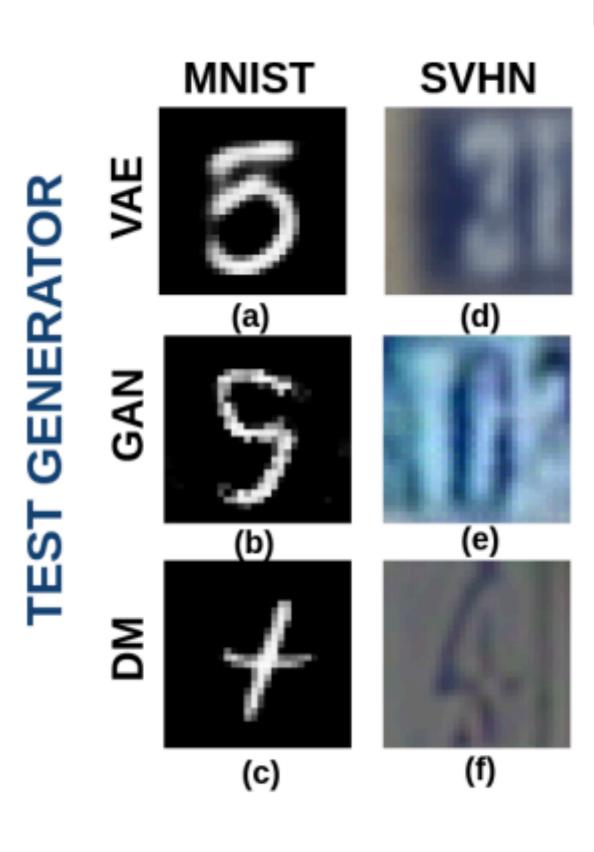


Dataset	Pert. Step (δ_{init})	Model	% Seeds	% Misclass. (#)	# Iterations	% Validity (#)	% Preserved (#)
MNIST		VAE	99	4.04 (4)	245.41	50.00 (2)	100.00 (2)
	Low	GAN	99	8.08 (8)	242.05	75.00 (6)	83.33 (5)
		DM	87	50.57 (44)	164.61	45.45 (20)	30.00 (6)
		VAE	99	100.00 (99)	62.92	73.74 (73)	<u>49.32</u> (36)
	High	GAN	99	96.97 (96)	107.46	<u>69.79</u> (67)	62.69 (42)
		DM	87	100.00 (87)	26.77	40.23 (35)	34.29 (12)
SVHN		VAE	66	50.00 (33)	178.20	51.52 (17)	41.18 (7)
	Low	GAN	84	42.86 (36)	182.22	30.56 (11)	<u>45.45</u> (5)
		DM	95	69.47 (66)	131.04	39.39 (26)	57.69 (15)
		VAE	66	100.00 (66)	27.00	39.39 (26)	30.77 (8
	High	GAN	84	98.81 (83)	39.00	<u>36.14</u> (30)	50.00 (15
	U	DM	95	100.00 (95)	13.23	23.16 (22)	18.18 (4
CIFAR-10		VAE	39	82.05 (32)	118.90	53.13 (17)	29.41 (5
	Low	GAN	69	66.67 (46)	140.32	45.65 (21)	19.05 (4
		DM	87	89.66 (78)	85.63	60.26 (47)	61.70 (29
		VAE	39	100.00 (39)	19.51	30.77 (12)	33.33 (4
	High	GAN	69	100.00 (69)	25.78	31.88 (22)	22.73 (5
	-	DM	87	100.00 (87)	14.18	62.07 (54)	68.52 (37
ImageNet (Teddy Bear)		VAE	14	100.00 (14)	13.57	78.57 (11)	81.82 (9
	Low	GAN	85	100.00 (85)	98.27	74.12 (63)	36.51 (23
		DM	<u>85</u> 87	<u>98.85</u> (86)	48.45	91.86 (79)	49.37 (39
		VAE	14	100.00 (14)	1.36	100.00 (14)	64.29 (9
	High	GAN	<u>85</u>	100.00 (85)	26.38	83.53 (71)	32.39 (23
	-	DM	<u>85</u> 87	100.00 (87)	6.63	<u>94.25</u> (82)	<u>56.10</u> (46
ImageNet (Pizza)		VAE	25	100.00 (25)	12.96	92.00 (23)	91.30 (21
	Low	GAN	99	88.00 (87)	172.88	88.51 (77)	46.75 (36
		DM	73	<u>97.26</u> (71)	83.60	98.59 (70)	92.86 (65
		VAE	25	100.00 (25)	2.60	80.00 (20)	<u>75.00</u> (15
	High	GAN	99	100.00 (99)	47.93	86.87 (86)	51.16 (44
	-	DM	73	100.00 (73)	12.53	100.00 (73)	86.30 (63

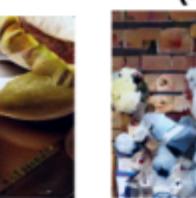
TABLE III

COMPARISON BETWEEN GENAI TIGS ACROSS DIFFERENT DATASETS AND MUTATION EXTENTS IN TERMS OF VIABLE SEEDS, MISCLASSIFICATION-INDUCING INPUTS, NUMBER OF ITERATIONS TO GENERATE FAILURE, INPUT VALIDITY, AND LABEL PRESERVATION. THE BEST RESULTS ARE HIGHLIGHTED IN BOLD, WHILE THE UNDERLINED VALUES ARE NOT STATISTICALLY DIFFERENT FROM THE BEST.





DATASET CIFAR-10 INTB INPI (g) (j) (m)



(n)







Fig. 3. Misclassification-inducing images generated by GenAI TIGs

