A Mishmash of Methods for Mitigating the Model Mismatch Mess

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Image database to study model mismatch

- 9000 JPEG images from each of 9 Flickr users.
- All images share the same quantization table (qf = 85).
- 8 different camera models from 5 manufacturers.
- Stego images embedded with nsF5 with payload 0.05 bpp.
- 7850 dimensional *CF** features used for steganalysis.

The reality of model mismatch

		Testing actor								
		1	2	3	4	5	6	7	8	9
actor	1	0.00	0.01	0.13	0.23	0.01	0.10	0.05	0.04	0.08
	2	0.02	0.00	0.16	0.10	0.02	0.10	0.04	0.04	0.06
	3	0.36	0.33	0.02	0.38	0.24	0.23	0.36	0.29	0.31
	4	0.03	0.01	0.18	0.00	0.01	0.09	0.03	0.04	0.06
ing	5	0.04	0.01	0.11	0.07	0.00	0.09	0.03	0.02	0.06
Traini	6	0.06	0.02	0.16	0.04	0.03	0.06	0.07	0.03	0.07
	7	0.01	0.00	0.21	0.10	0.00	0.11	0.01	0.02	0.06
	8	0.05	0.03	0.12	0.09	0.01	0.09	0.04	0.01	0.08
	9	0.03	0.01	0.19	0.17	0.01	0.11	0.08	0.04	0.05

Testing error of Fisher Linear Discriminant classifiers

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Image: Image:

- Traditional error measures do not quantify the stability of the detector.
- Mean error rate $\mu_1 = \frac{1}{k} \sum_i P_E^i$.
- Root mean square error rate $\mu_2 = \sqrt{\frac{1}{k} \sum_i (P_E^i)^2}$.
- Maximum error rate $\mu_{\infty} = \max_i P_{\rm E}^i$.

Summary of the baseline

ma	atched cas	ses	mismatched cases			
μ_1	μ_2	μ_{∞}	μ_1	μ_2	μ_∞	
0.0204	0.0315	0.0663	0.0981	0.1369	0.3887	

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Mismatch due to shift



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Mean distances of covers

	Actor's whose centroid distance is measured									
		1	2	3	4	5	6	7	8	9
from other actor's centroid	1	0	0.59	3.93	12.58	0.66	1.90	2.82	0.72	0.79
	2	1.40	0	7.27	3.67	1.35	0.38	1.07	0.48	1.02
	3	12.84	12.46	0	19.02	10.56	9.25	13.92	12.00	12.58
	4	5.01	0.50	8.77	0	0.70	0.37	2.01	1.97	1.06
	5	0.49	1.09	2.54	5.39	0	2.17	2.29	2.56	2.00
	6	2.18	0.81	5.09	1.72	0.51	0	1.04	1.09	2.08
	7	0.26	0.35	7.53	2.92	0.53	0.76	0	0.49	0.10
	8	1.65	1.05	2.00	5.09	1.28	1.33	1.86	0	3.69
:	9	0.26	0.61	4.82	3.31	0.44	0.95	1.26	1.30	0

Actor's whose centroid distance is measured...

Correlation with errors: ho=0.91 au=0.51

Centering the cover means helps

	matched cases			mismatched cases		
	μ_1 μ_2 μ_{∞}		μ_1	μ_2	μ_∞	
Baseline	0.0204	0.0315	0.0663	0.0981	0.1369	0.3887
Subtracted mean				0.0691	0.0838	0.2332

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Mismatch due to angle and speed



$$ho = -0.52, \ au = -0.44 \ \cos lpha_{i,j} = rac{w_i \cdot w_j}{\|w_i\| \|w_j\|}$$

$$\rho = -0.10, \ \tau = -0.09$$
$$r_{i,j} = \frac{\left| \|w_i\| - \|w_j\| \right|}{\sqrt{\|w_i\| \|w_j\|}}$$

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Calibrating for the different angle

- Re-embedding of a small payload and estimating the direction of the movement
- Training an ensemble of detectors, each on a different cover source
- weighting their vote based on the alignment measured by

• angle
$$\cos \alpha_i = \frac{w_i \cdot \delta}{\|w_i\| \|\delta\|}$$

Experimental results

	matched cases			mismatched cases			
	μ_1	μ_2	μ_∞	μ_1	μ_2	μ_∞	
Baseline	0.0204	0.0315	0.0663	0.0981	0.1369	0.3887	
Centered				0.0691	0.0838	0.2332	
Ensemble voting							
Equal weight				0.0439	0.0627	0.1366	
Weight by $\cos \alpha_i$				0.0433	0.0593	0.1160	
Ensemble voting,	centered	1					
Equal weight				0.0391	0.0479	0.0776	
Weight by $\cos \alpha_i$				0.0372	0.0468	0.0806	

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Mismatch due to false certainty (1)

matched case

mismatched case



Decision areas of Binary Support Vector Machines

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Abstaining classifiers

- "I don't know" is
 - honest
 - potentially advantageous in pooled steganalysis.
- Implemented as a pair of 1-class Support Vector Machines (1-SVMs).
- 1-SVMs and 2-SVMs have hyper-parameters found by cross-validation.

Mismatch due to false certainty (1)



Decision areas of abstaining SVMs

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Measuring error of abstaining classifiers

The value of abstaining classifiers is in the pooled steganalysis.Introduce deflection coefficient

$$d = \frac{(1 - P_{\text{FP}} - P_{\text{FN}})\sqrt{1 - P_{\text{DK}}}}{\sqrt{P_{\text{FP}}(1 - P_{\text{FP}})} + \sqrt{P_{\text{FN}}(1 - P_{\text{FN}})}},$$

where $P_{\rm DK}$ is the probability of "don't know".

Experimental results

	mismatched cases							
	роо	pooled error rates						
	$P_{\rm FP}$	$P_{\rm FN}$	$P_{\rm DK}$	— <i>u</i> ∞				
2-SVMs	0.165	0.059	0.000	0.465				
1-SVMs	0.028	0.010	0.539	1.017				

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Conclusion

- $\, \bullet \,$ Uncorrected mismatched $\rightarrow \,$ error rates 5× matched case.
- $\bullet\,$ Centering and matching direction $\rightarrow\,$ error rates 1.5-2 $\times\,$ matched case.
- Possibility of answering "don't know".
- This is a mishmash of methods: no definitive answer.
- Research tends to be driven by the metrics:

are we chasing the right ones?