

# Support Vector Inductive Logic Programming



Review of an article

Oksana Korol

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# Main points and motivation

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Reviewed article:

## **“Support Vector Inductive Logic Programming”**

By Stephen Muggleton, Huma Lodhi,  
Ata Amini, and Michael J. E. Sternberg

# Main points and motivation

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- ▶ Goal: prediction of toxicity
- ▶ Intersection of SVM and ILP
- ▶ SVM provides for dimensionality independence
- ▶ ILP kernel captures relational information

# Chemistry



## • Molecular structure of toxic chemicals

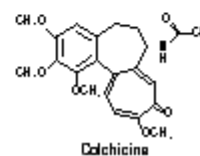
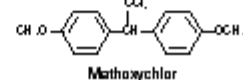
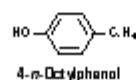
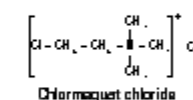
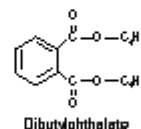
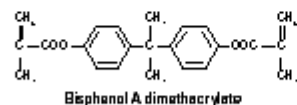
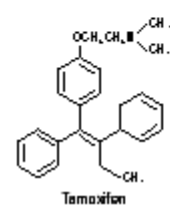
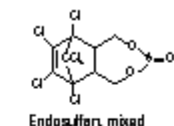
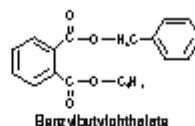
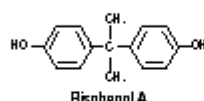
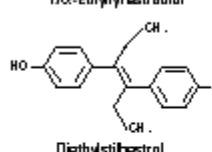
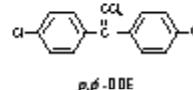
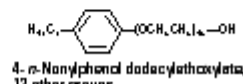
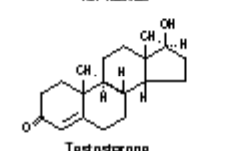
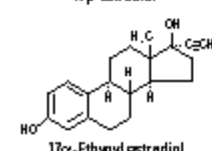
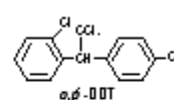
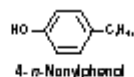
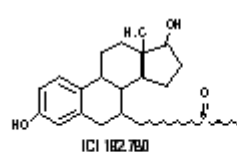
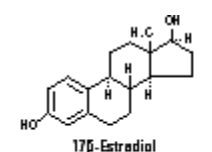


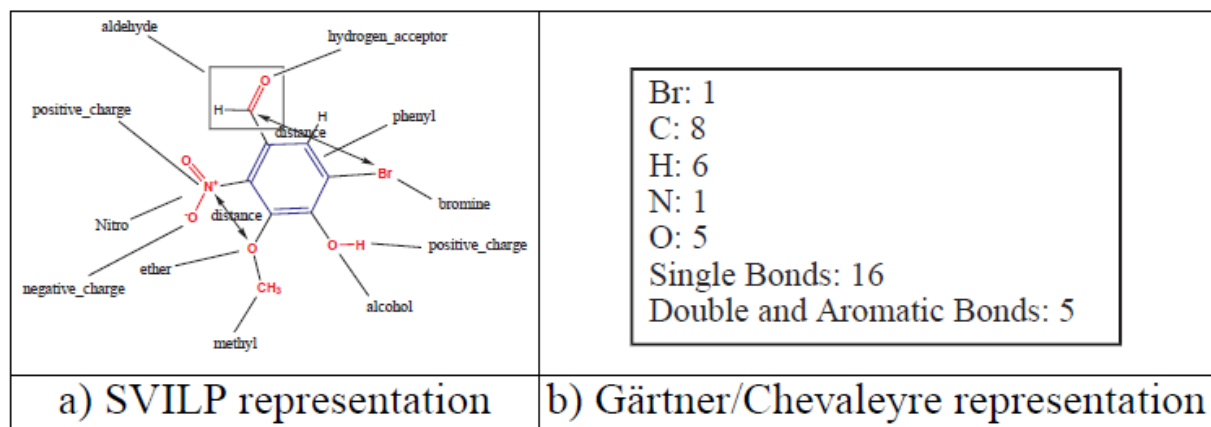
Table 1. COMPACT parameters for 30 chemicals.

Chemical	Area/ Depth <sup>2</sup>	ΔE (eV)	COMPACT radius*	CYP1 interaction	Molecular collision diameter	CYP2E <sup>b</sup> activation	Overall COMPACT prediction
1. Scopolamine	1.6	15.4	8.6	—	7.9	—	—
2. Codeine	1.4	14.1	7.9	—	8.1	—	—
3. 1,2-Dihydro-2,2,4-trimethylquinoline	1.8	13.1	7.0	—	6.9	—	—
4. Nitromethane	1.4	14.8	8.3	—	4.7	+	+
5. Tetrahydrofuran	1.6	20.9	13.0	—	5.1	—	—
6. t-Butylhydroquinone	1.8	14.3	7.7	—	6.7	—	—
7. Ethylbenzene	2.1	16.7	9.2	—	6.0	—	—
8. Chloroprene	3.7	13.5	5.7	±	5.3	+	+
9. Cobalt sulfate	NP	NP	NP	NP	NP	NP	NP
10. D & C Yellow No. 11	2.0	13.0	6.8	—	7.8	—	—
11. Isobutyraldehyde	1.5	17.5	10.2	—	5.3	—	—
12. Molybdenum trioxide	NP	NP	NP	NP	NP	NP	NP
13. 1-Chloro-2-propanol	1.5	16.4	9.3	—	5.3	—	—
14. Diethanolamine	2.9	20.3	11.9	—	5.8	—	—
15. Phenolphthalein	2.0	13.9	7.3	—	8.1	—	—
16. Pyridine	4.3	16.2	7.6	—	5.2	—	—
17. Xylenesulfonic acid	3.7	13.9	6.0	±	6.7	—	±
18. Furfuryl alcohol	2.6	15.3	7.8	—	5.4	±	±
19. Primidone	1.7	15.9	9.8	—	7.3	—	—
20. Ethylene glycol monobutyl ether	3.3	20.5	11.9	—	6.4	—	—
21. Gallium arsenide	NP	NP	NP	NP	NP	NP	NP
22. Isobutene	2.3	17.9	10.0	—	5.1	—	—
23. Methylugenol	1.8	14.8	8.0	—	7.1	—	—
24. Oxymetholone	2.7	13.9	6.7	—	8.5	—	—
25. Anthraquinone	9.4	13.5	4.3	+	7.0	—	+
26. Emodin	6.6	12.3	3.1	+	7.5	—	+
27. Citral	1.5	14.2	7.9	—	6.8	—	—
28. Sodium nitrite	NP	NP	NP	NP	NP	NP	NP
29. Cinnamaldehyde	6.9	13.3	3.9	+	6.2	+	+
30. Vanadium pentoxide	NP	NP	NP	NP	NP	NP	NP

Abbreviations: NP, not predicted; +, positive; —, negative. \*COMPACT radius,  $\sqrt{\Delta E - 9.5} + \{a/a^2 - 7.8\}^E$ . Positive CYP1 if COMPACT radius < 5.5; ± if radius is between 5.5 and 6.5. <sup>b</sup>CYP2E activation, molecular collision diameter < 6.5 and ΔE < 15.0. <sup>c</sup>Overall COMPACT prediction is the summation of CYP1 interaction and CYP2E activation.

Images from Environmental Health Perspectives

# Chemistry



**Fig. 1.** Molecule represented using a) SVILP representation which employs a kernel based on domain-expert informed chemical background knowledge indicated by the annotations on the figure and b) Gärtner/Chevaleyre bag-of-atoms uses Multi-Instance (MI) kernel based on frequency of occurrences of atoms and atom pairs.

# Inductive Logic Programming

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- ▶ Introduced by Stephen Muggleton in 1992

Inductive Logic Programming (ILP)  
=  
Machine Learning  $\wedge$  Logic Programming  
=  
Learning with Logic

# Inductive Logic Programming

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- ▶ Induction – reasoning from specific to general
- ▶ Logic programs are the set of Horn clauses that follow the rules of the first order logic:

$mother(X, Y) \leftarrow parent(X, Y) \wedge female(X) .$

$female(Jane).$

$female(Ann).$

$male(Jack).$

$male(John).$

$parent(Jane, Ann).$

$parent(Jack, Ann).$

- ▶ Questions: Is Jane a mother of Ann?

Who is a mother of Ann?



# Inductive Logic Programming

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ILP is represented in logic programs which are used to derive a solution to a problem by inducing a hypothesis based on a set of positive and negative examples.

# Inductive Logic Programming

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- ▶ **Concept learning:** given a background knowledge  $B$  and experimental observations  $E$  (consisting of positive  $E+$  and negative  $E-$  examples) find a hypothesis  $H$  such that:

$$B \wedge H \models E$$

- ▶  $B$ ,  $E$  and  $H$  are each logic programs

# Inductive Logic Programming

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- ▶  $B$ ,  $H$ , and  $E$  should satisfy the following conditions:

**Prior Satisfiability.**  $B \wedge E^- \not\models$

**Posterior Satisfiability.**  $B \wedge H \wedge E^- \not\models$

**Prior Necessity.**  $B \not\models E^+$

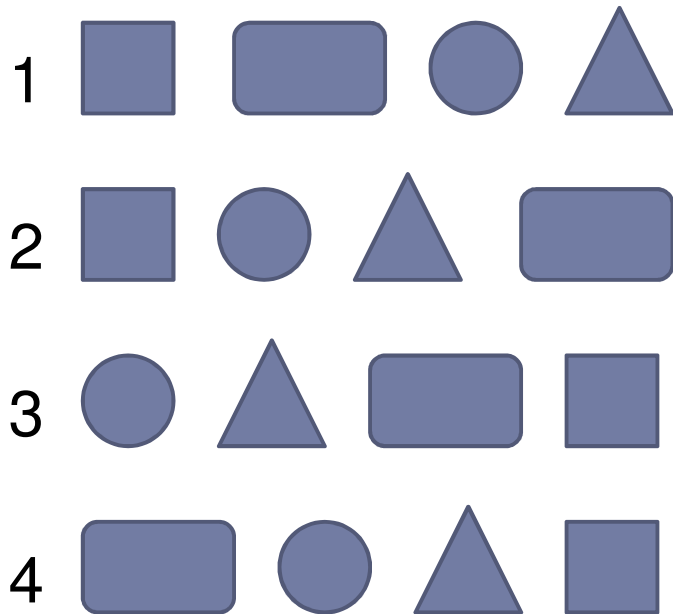
**Posterior Sufficiency.**  $B \wedge H \models E^+$

# Inductive Logic Programming

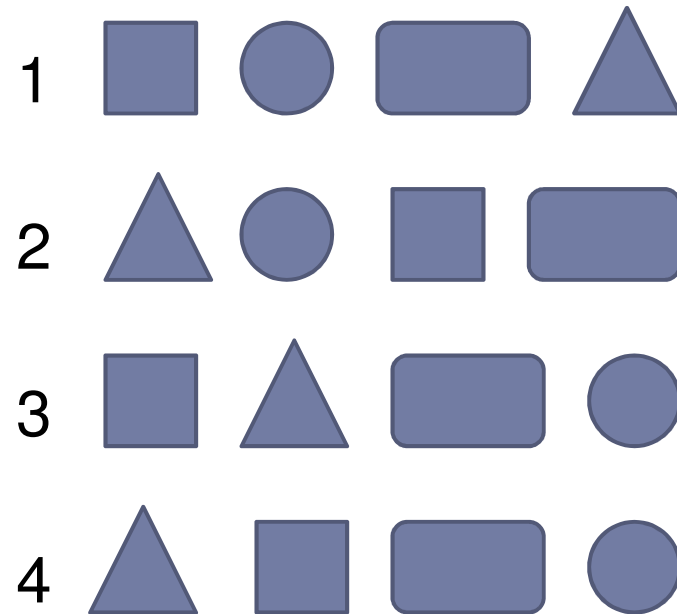


- Classify the following:

Positive examples



Negative examples

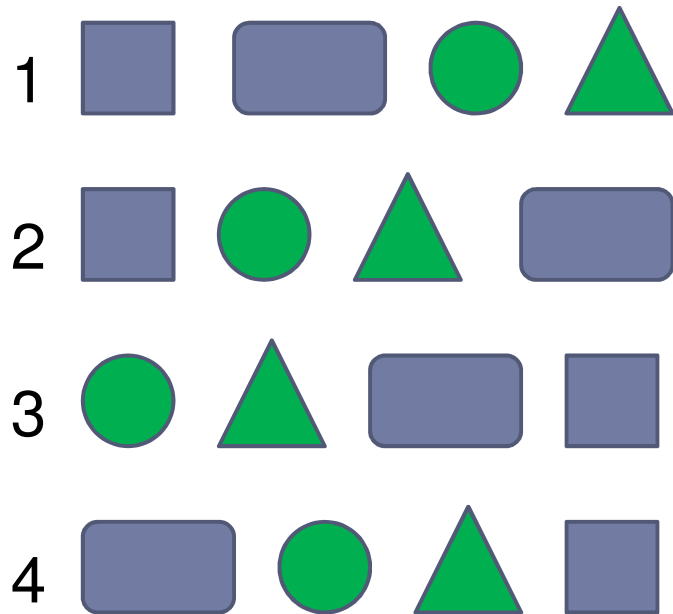


# Inductive Logic Programming

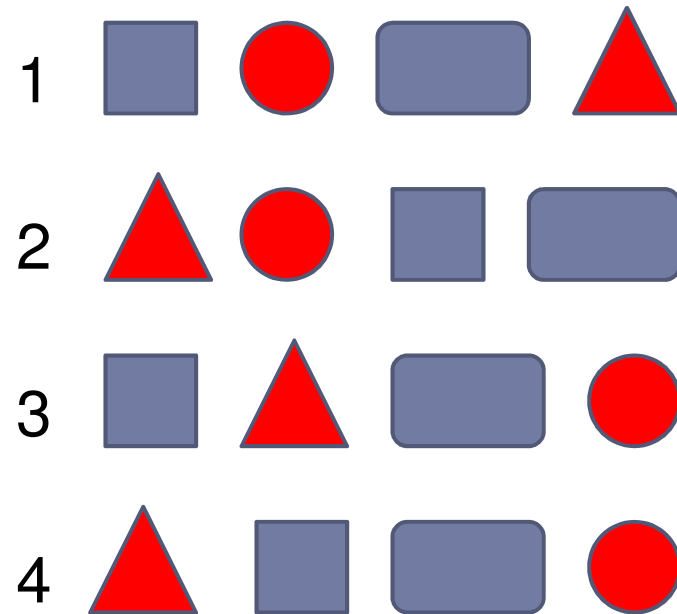


- Classify the following:

Positive examples

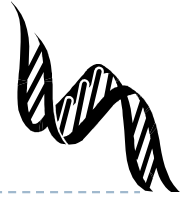


Negative examples



# Inductive Logic Programming

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- ▶  $B$  would specify following rules:
  - ▶  $\text{before}(X, Y) \text{ :- } \langle \text{when position of } X \text{ is less than position of } Y \rangle$
  - ▶  $\text{adjacent}(X, Y) \text{ :- } \langle \text{when there is no other object between } X \text{ and } Y \rangle$
- ▶ Then the resulting theory  $H$  will be:
  - ▶  $\text{positive} \text{ :- } \text{before}(\text{circle}, \text{triangle}), \text{adjacent}(\text{triangle}, \text{circle}).$

# Support Vector Machines

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- ▶ Take any problem and transform it into a high dimensional space, so that it becomes linearly separable, but
- ▶ Calculations to obtain the separability plane can be done in the original input space (*kernel trick*)

# Support Vector Machines

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- ▶ *I.e. SVM learning process consists of 2 stages:*
  1. Map the input data,  $d_1, \dots, d_n \in D$ , into some higher dimensional space  $H$  through a non-linear mapping  $\varphi$  that is given by  $\varphi : D \rightarrow H$ .
  2. Construct a linear function  $f$  in the space

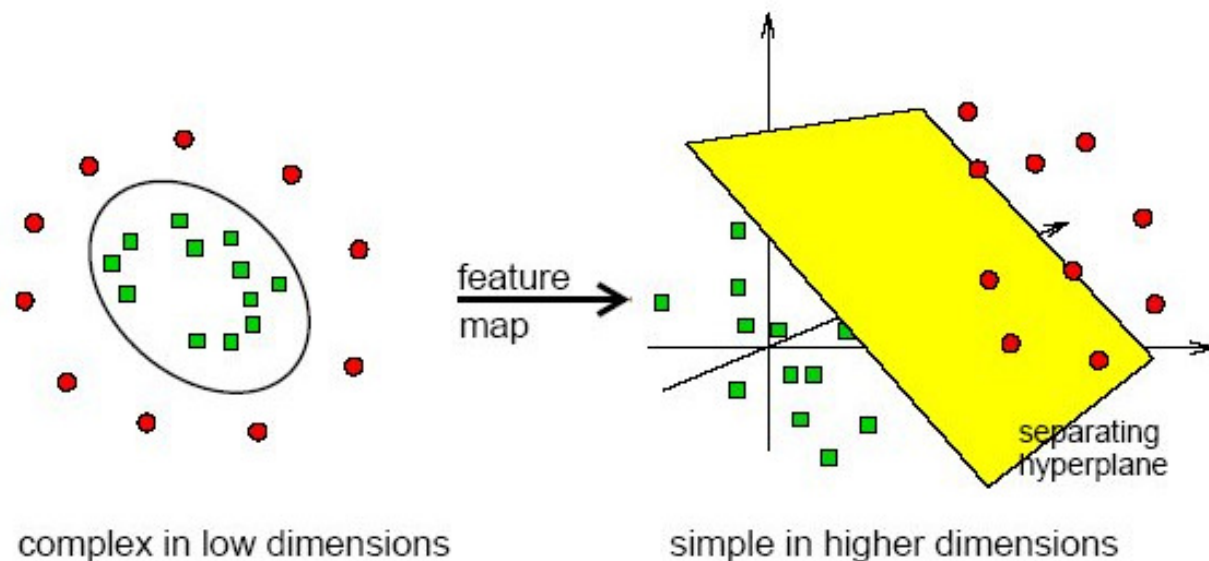


# Support Vector Machines



- ▶ The kernel function may transform the data into a higher dimensional space to make it possible to perform the separation.

Separation may be easier in higher dimensions



Info from <http://www.dtrek.com/svm.htm>



# Propositionalization

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- ▶ Propositionalization – techniques to transform relational (first order logic) representation to propositional (fixed sized feature vectors)
- ▶ Involves construction of structural features from relational background knowledge
- ▶ => Any propositional learner can be applied after propositionalization
- ▶ SVILP is similar in its use of support-vector technology to the domain-dependent bottom-up propositionalisation approach



# Bottom-Up Propositionalization

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- ▶ Discover fragments that occur frequently in the dataset (ex. circle followed by triangle)
- ▶ Bottom-up approach to fragment generation: generate only those fragments that really occur in the examples
- ▶ Algo: depth-first search for fragments for each data point (ex. sequence)



# Support Vector ILP

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- ▶ In short: SVM with ILP as a kernel function
- ▶ Like in ILP, assume background knowledge  $B$ , examples  $E$  and a hypothesis  $H$
- ▶ SVILP bases a kernel on the predictions of the clauses  $h$  in  $H$



# Support Vector ILP

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- ▶ Kernel is built by forming a binary hypothesis-instance association matrix  $M: h_i \times d_j$ , where  $h_i \in H$  and  $d_j \in D$ .

- ▶ For each hypothesis clause  $h$  in  $H$ :

$$h : D \rightarrow \{True, False\}.$$

- ▶ Conversely the  $\tau$  function gives the hypothesised clauses covering any particular instance:

$$\tau(d_i) = \{h : \exists h \in H, (B, h \models d_i)\}$$

- ▶ So the kernel function is as follows:

$$K(d_i, d_j) = f(\tau(d_i) \cap \tau(d_j))$$

# Results

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- ▶ Tested on the new DSSTox dataset (as opposed to Mutagens)
- ▶ Used 5-fold cross validation with mean squared error (MSE) and R-squared evaluation
- ▶ Compared results with well known QSAR software TOPKAT (Toxicity Prediction by Komputer Assisted Technology)
- ▶ Also compared to following techniques: partial least squares (PLS), multi instance kernels (MIK) , an RBF kernel

# Results

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	MSE	R-squared
CHEM	0.811	0.519
PLS	0.671	0.593
MIK	0.838	0.503
SVILP	<b>0.574</b>	<b>0.655</b>

**Fig. 8.** MSE and R-squared for CHEM, PLS, MIK and SVILP.

	Accuracy
ILP (CProgol5.0)	55
CHEM	58
PLS	71
MIK	60
SVILP	<b>73</b>

**Fig. 9.** Accuracy for ILP, CHEM, PLS, MIK and SVILP.

# Conclusion

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- ▶ Accuracy is good, but perhaps not the best way to evaluate the approach
- ▶ No mention of the performance time
- ▶ The kernel works within the standard ILP setting of generalisation with respect to background knowledge (not just atomic generalization)
- ▶ SVILP method shows significant improvement with respect to the other methods
- ▶ Follow up work confirms this (see references)



# References / Further reading

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- ▶ Muggleton, S.H. "**Inverse Entailment and Progol**". In New Generation Computing, 13 (1995) 245-286.
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# Questions

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