

Background Knowledge Injection for Interpretable Sequence Classification

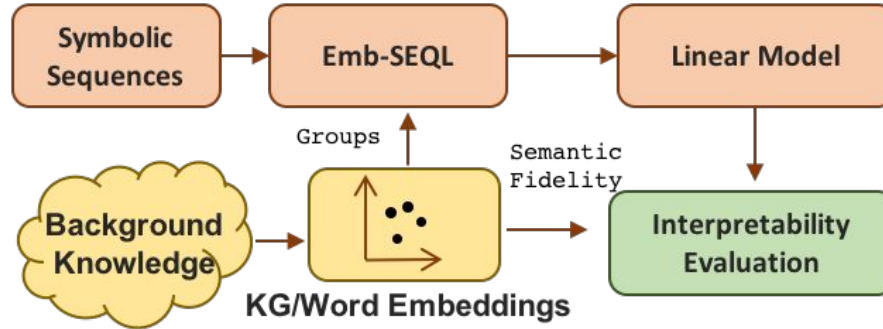
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Contributions



Groups:

Regex-like expansion of traditional k -mers

Emb-SEQL:

Injection of Background Knowledge into Sequence Learning Algorithm

Semantic Fidelity:

Metric to quantify interpretability

Symbolic Sequence Classification

Sequence	Class
ABBCBAABCBAABCBAABBBBCBABBABBCB	+1
CCBBABACAABBABBAABBBCBBAABA	-1
ACBBCACCCBAABCBAABCCABCAABCCA	+1
BABACCBABCTABCBAABBCAABCBBBCA	?

$\Sigma = \{A, B, C\}$

k-mers: Consecutive sequences *k* of symbols

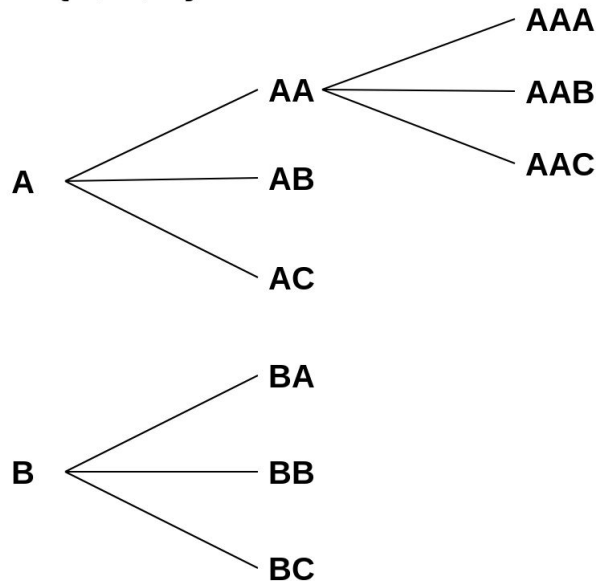
2-mer: AB

5-mer: BCTCB

SEQL - Sequence Learner

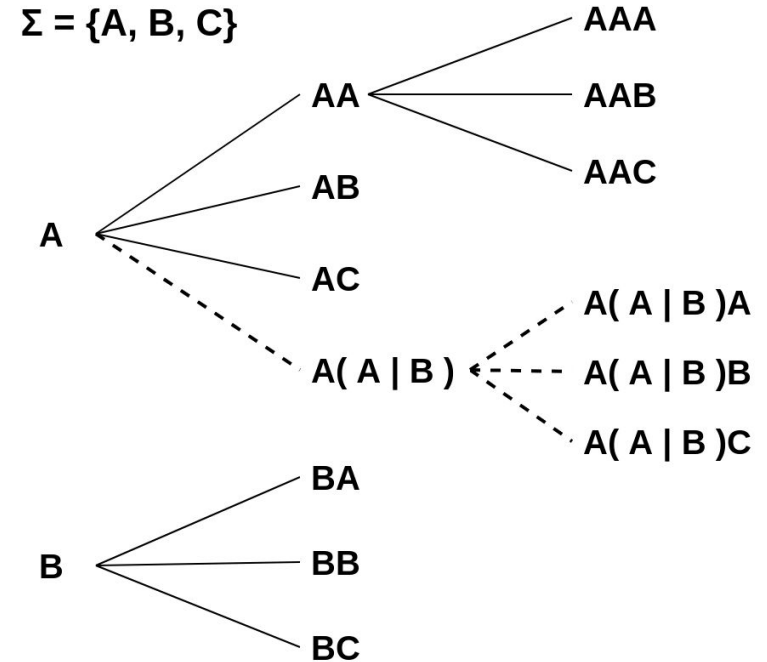
- Integrated approach
- Learns sparse k -mer based linear models
- Feature space of all possible k -mers
- Gauss-Southwell coordinate descent
- Iteratively add best k -mer to model
- Exploits **structure in feature space**

$\Sigma = \{A, B, C\}$

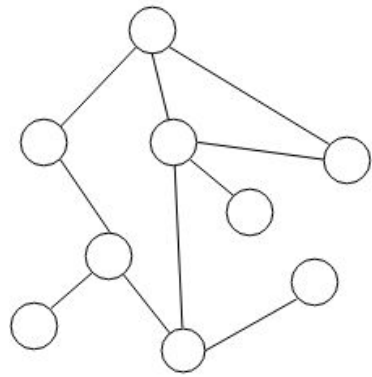


SEQL - Sequence Learner with *Groups*

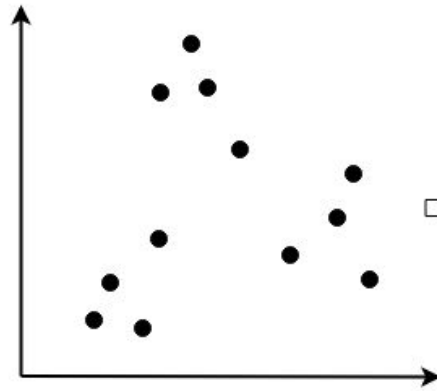
- Exploit **structure in symbol space**
- Use *groups* to gain more flexibility
- *Groups* are built by combining basic symbols with OR
- Groups predefined by user or automatically generated



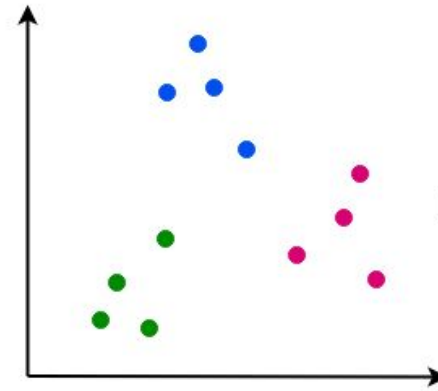
Automatic Group Generation



**Knowledge Graph
with Symbols**



**Embedding Space
of Symbols**



**Clustering
of Symbols**



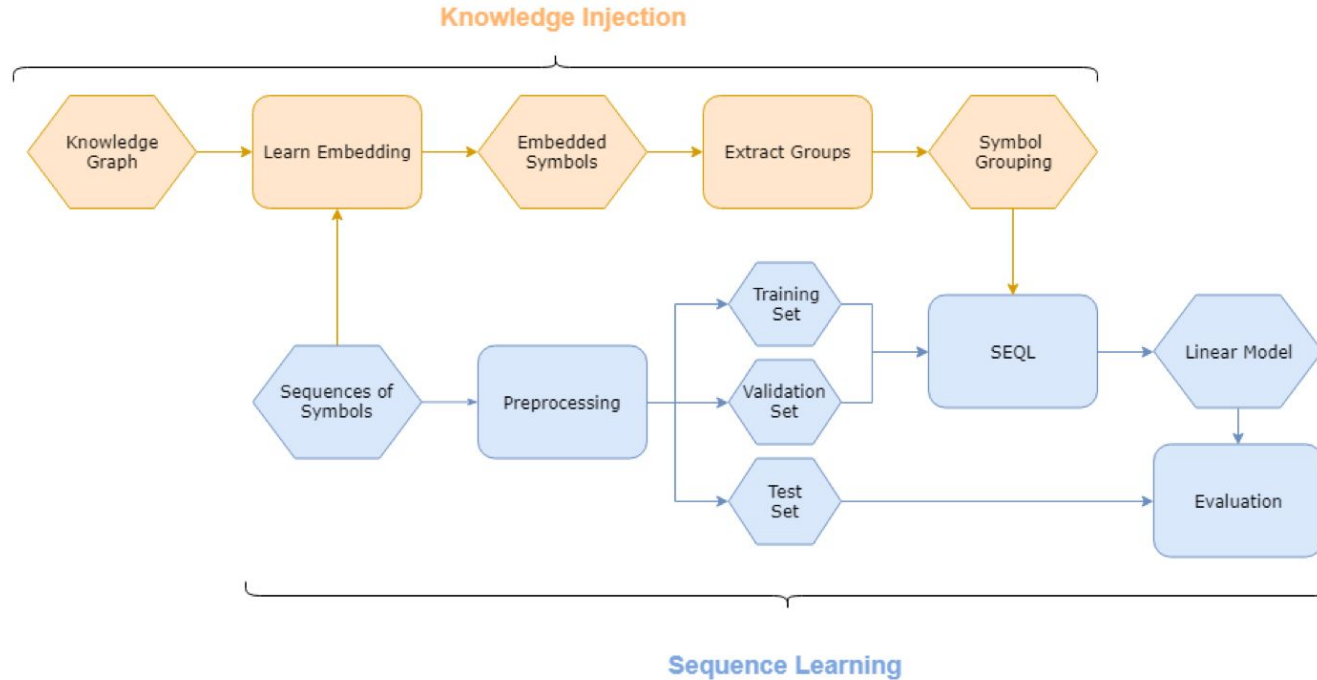
Group I: A B C D

Group II: E F G H

Group III: I J K L

Extracted Groups

Emb-SEQL Pipeline



Interpretability

- Interpretability is crucial in some problems
- Accuracy-interpretability trade-off
- Measuring interpretability of models is an open question

Semantic Fidelity intuition:

- **Positive** features should be “close” to target class
- **Negative** features should be “close” to non-target class

Functional grounded protocol as proxy measurement

Semantic Fidelity

k-mer - Target Class Distance

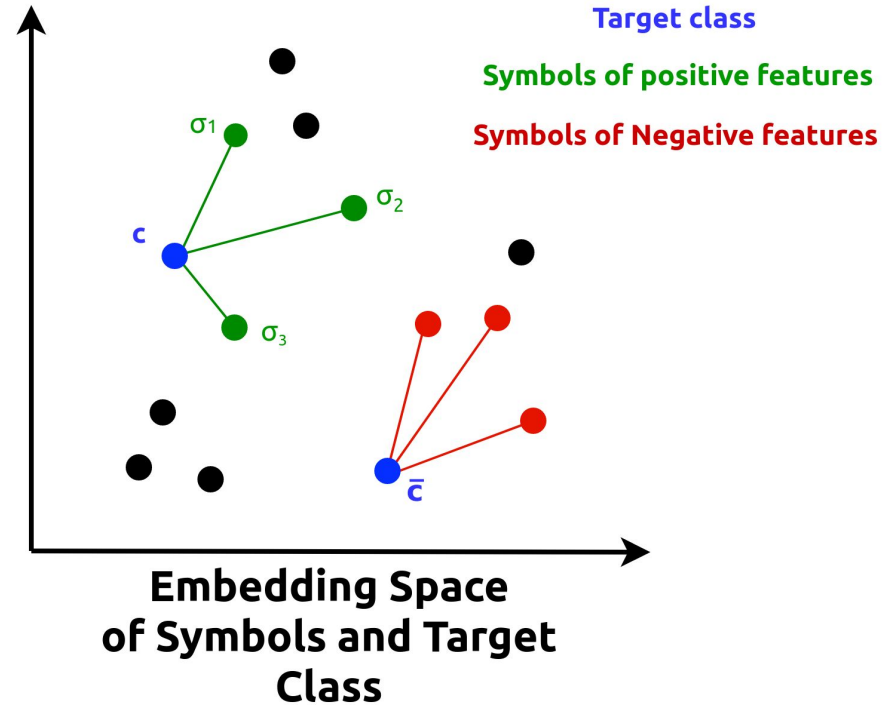
$$d_c(\phi, c) = \frac{1}{n_\phi} \sum_{\sigma_j \in \phi} \|\mathbf{E}_{\sigma_j} - \mathbf{e}_c\|$$

Weighted Target Class Distance

$$h(\phi) = |w| \begin{cases} d(\phi, c) & \text{if } w \geq 0 \\ d(\phi, \bar{c}) & \text{otherwise} \end{cases}$$

Semantic Fidelity

$$SF = 1 - \frac{1}{2n} \sum_{\phi_i \in \Phi} h(\phi_i)$$



Experiment

Opportunity - Human Composite Activity Recognition (HAR) [\[1\]](#):

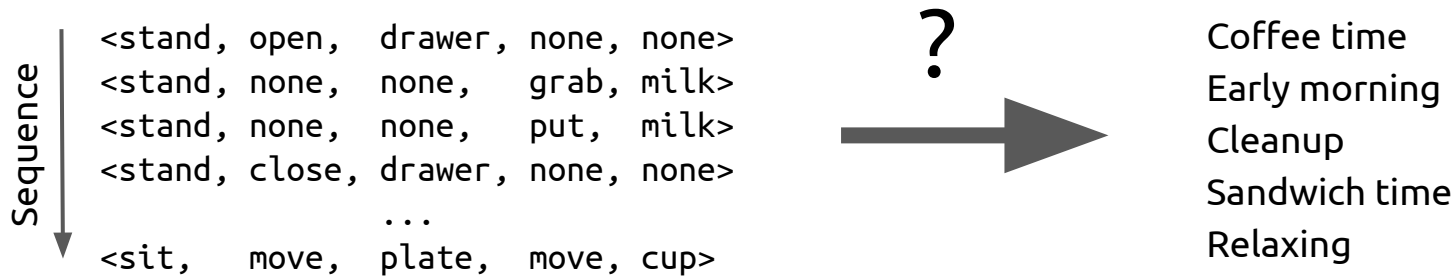
- Predict Composite Activity
- Multiclass classification problem
- Combinations of 5 low level features categories ($|\Sigma| > 1400$ symbols)

PhosphoELM - Protein Classification [\[2\]](#):

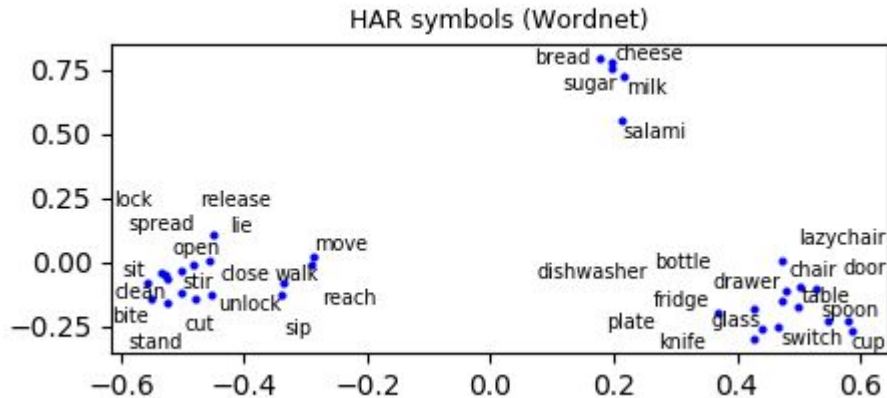
- Binary classification problem
- Predict Kinase group
- Amino acid sequences ($|\Sigma| = 21$ symbols; 438 sequences)

Composite Activity Recognition

Predict composite activity based on sequence of low-level activity



Composite Activity Recognition



Atomic symbol embeddings

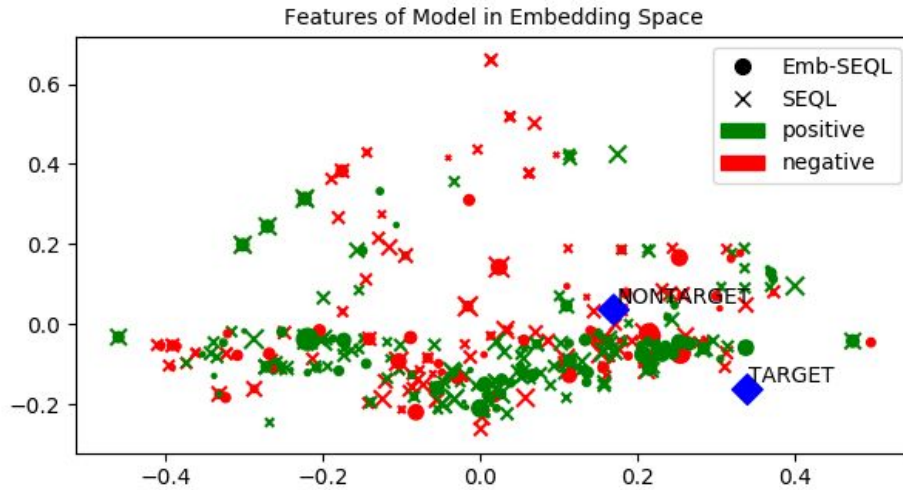
Weight	Feature
1.0	\langle ([sit, release, cheese, move, knife cheese] OR [stand, move, plate, move, plate] OR [stand, move, plate, reach, drawer1] OR [stand, move, plate, open, drawer1] OR [stand, reach, bread, open, drawer3]) \rangle ...
-0.867	\langle ([stand, open, fridge, move, milk] OR [stand, none, none, release, milk] OR [walk, move, plate, none, none] OR [walk, move, plate, release, bread] OR [walk, none, none, move, plate] OR [stand, reach, sugar, reach, spoon]) , [stand, none, none, none, none] \rangle

Sandwich time model

Results - Semantic Fidelity

Dataset	Embeddings	Model	\overline{SF}	std	Class 1	Class 2	Class 3	Class 4	Class 5	
HAR	GloVe	SEQL	0.902	0.028	0.930	0.871	0.925	0.865	0.921	
		Emb-SEQL	0.923	0.025	0.958	0.888	0.931	0.901	0.938	
	ConceptNet	SEQL	0.871	0.033	0.908	0.828	0.895	0.833	0.889	
		Emb-SEQL	0.875	0.025	0.903	0.853	0.887	0.836	0.894	
	YAGO-41	SEQL	0.867	0.029	0.899	0.847	0.893	0.823	0.872	
		Emb-SEQL	0.835	0.043	0.897	0.824	0.861	0.767	0.827	
	WordNet	SEQL	0.894	0.025	0.921	0.879	0.918	0.857	0.895	
		Emb-SEQL	0.936	0.010	0.937	0.945	0.943	0.917	0.939	
	Protein	ChEBI-ChEMBL	SEQL	0.708	-					
			Emb-SEQL	0.719	-					

PCA Visualization



Target: Coffee time
Embedding: WordNet

Results - Classification Quality

Dataset	Model	Embeddings	F1	Accuracy	
HAR	SVM		0.502	0.564	
	LSTM		0.767	0.810	
	SEQL		0.973	0.961	
	Emb-SEQL	ConceptNet		0.965	0.951
		GloVe		0.961	0.945
		WordNet		0.968	0.955
		YAGO-41		0.957	0.941
Protein	SCIS_MA		-	0.948	
	HMM		-	0.918	
	LSTM		0.797	0.796	
	SEQL		0.902	0.903	
	Emb-SEQL	ChEBI-ChEMBL	0.898	0.901	

SCIS_MA and HMM
results from [\[2\]](#)

Achievements & Conclusion

- Introduction of ***Groups***, regex like k-mer symbols
- Generation of *Groups* from background knowledge sources
- **Emb-SEQL**, a method to learn sparse linear models
- **Semantic Fidelity** a way to measure interpretability

- Background knowledge injection improves interpretability measured by Semantic Fidelity without hurting accuracy of learned model

Limitations & Future Work

- High memory demand of Emb-SEQL for large *Groups*
- Clustering method and *Group* size is crucial
- Background knowledge source is needed

- Semantic Fidelity for non-linear models
- Human-based evaluation of Semantic Fidelity

Thank you!

Please email severin.gsponer@insight-centre.org if you have further questions

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References

- [1] D. Roggen, A. Calatroni, M. Rossi, T. Holleczeck, K. Förster, G. Tröster, P. Lukowicz, D. Bannach, G. Pirkl, A. Ferscha, J. Doppler, C. Holzmann, M. Kurz, G. Holl, R. Chavarriaga, H. Sagha, H. Bayati, M. Creatura, and J. d. R. Millàn. Collecting complex activity datasets in highly rich networked sensor environments. In 7th International Conference on Networked Sensing Systems (INSS)
- [2] C. Zhou, B. Cule, and B. Goethals. Pattern based sequence classification. *IEEE Transactions on Knowledge and Data Engineering*, 28(5):1285–1298, 2016.