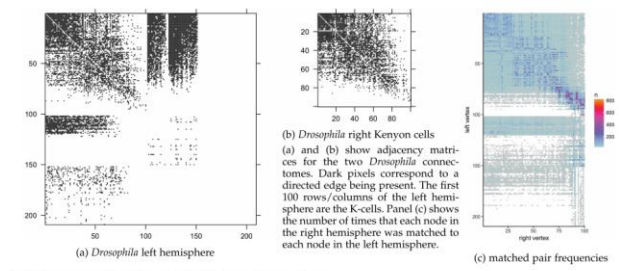


subgraph2vec																																																			
Article	Data	Data	Comments or Questions																																																
Narayanan A , Chandramohan M , Chen L , et al. subgraph2vec: Learning Distributed Representations of Rooted Sub-graphs from Large Graphs[J]. 2016.	<p>Aim:</p> <p>Learning latent representations of rooted subgraphs from large graphs.</p> <p>Contributions:</p> <ul style="list-style-type: none">● Propose subgraph2vec.● Develop a modified version of the skipgram language model.● Discuss how subgraph2vec’s representation learning technique would help to build the deep learning variant of WL kernel. <p>Applications:</p> <ul style="list-style-type: none">● graph classification and clustering tasks● code clone detection● malware detection	<p>Key results:</p> <ul style="list-style-type: none">● Benchmark Datasets <p>Table 3: Average Accuracy (\pm std dev.) for subgraph2vec and state-of-the-art graph kernels on benchmark graph classification datasets</p> <table><tr><th>Dataset</th><th>MUTAG</th><th>PTC</th><th>PROTEINS</th><th>NCI1</th><th>NCI109</th></tr><tr><td>WL [6]</td><td>80.63 \pm 3.07</td><td>56.91 \pm 2.79</td><td>72.92 \pm 0.56</td><td>80.01 \pm 0.50</td><td>80.12 \pm 0.34</td></tr><tr><td>Deep WL_{YV} [7]</td><td>82.95 \pm 1.96</td><td>59.04 \pm 1.09</td><td>73.30 \pm 0.82</td><td>80.31 \pm 0.46</td><td>80.32 \pm 0.33</td></tr><tr><td>subgraph2vec</td><td>87.17 \pm 1.72</td><td>60.11 \pm 1.21</td><td>73.38 \pm 1.09</td><td>78.05 \pm 1.15</td><td>78.39 \pm 1.89</td></tr></table> <p>● Clone Detection</p> <p>Table 5: Clone Detection - Results</p> <table><tr><th>Kernel</th><th>WL [6]</th><th>Deep WL_{YV} [7]</th><th>subgraph2vec</th></tr><tr><td>Pre-training duration</td><td>-</td><td>421.7 s</td><td>409.28 s</td></tr><tr><td>ARI</td><td>0.67</td><td>0.71</td><td>0.88</td></tr></table> <p>● Malware Detection</p> <p>Table 7: Malware Detection - Results</p> <table><tr><th>Classifier</th><th>WL [6]</th><th>Deep WL_{YV} [7]</th><th>subgraph2vec</th></tr><tr><td>Pre-training duration</td><td>-</td><td>2631.17 s</td><td>2219.28 s</td></tr><tr><td>Accuracy</td><td>66.15</td><td>71.03</td><td>74.48</td></tr></table> <p>Background:</p> <ul style="list-style-type: none">● language model <p>Methods:</p> <ul style="list-style-type: none">● Generate rooted subgraphs: WL Relabeling Process● Embeddings of those subgraphs: Radial Skip Gram Model● Approximate the probability distribution: Negative Sampling <p>for each $sg_{cont} \in context_v^{(d)}$ do $J(\Phi) = -\log \Pr (sg_{cont} \Phi(sg_v^{(d)}))$</p>	Dataset	MUTAG	PTC	PROTEINS	NCI1	NCI109	WL [6]	80.63 \pm 3.07	56.91 \pm 2.79	72.92 \pm 0.56	80.01 \pm 0.50	80.12 \pm 0.34	Deep WL _{YV} [7]	82.95 \pm 1.96	59.04 \pm 1.09	73.30 \pm 0.82	80.31 \pm 0.46	80.32 \pm 0.33	subgraph2vec	87.17 \pm 1.72	60.11 \pm 1.21	73.38 \pm 1.09	78.05 \pm 1.15	78.39 \pm 1.89	Kernel	WL [6]	Deep WL _{YV} [7]	subgraph2vec	Pre-training duration	-	421.7 s	409.28 s	ARI	0.67	0.71	0.88	Classifier	WL [6]	Deep WL _{YV} [7]	subgraph2vec	Pre-training duration	-	2631.17 s	2219.28 s	Accuracy	66.15	71.03	74.48	<ul style="list-style-type: none">● When subgraph of degree 0 are considered, subgraph2vec provides node embeddings.● How can we get the initial labels for each node? <input checked="" type="checkbox"/>● What is the Deep WL kernel? <input checked="" type="checkbox"/>
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graph matching			
Article	Data	Data	Comments or Questions
<p>Sussman Daniel, Park Youngser, Priebe Carey E, et al. Matched Filters for Noisy Induced Subgraph Detection.. 2019,</p>	<p>Aim:</p> <p>Finding the vertex correspondence between two noisy graphs with different number of vertices where the smaller graph is still large.</p> <p>Contributions:</p> <ul style="list-style-type: none"> Proposed a number of padding methods to transform the noisy subgraph detection problem into a graph matching problem. <p>Applications:</p> <ul style="list-style-type: none"> determining whether a particular activity pattern is present in a social network detecting certain shapes in an image discovering motifs in brain networks 	<p>Key results:</p> <ul style="list-style-type: none"> Finding Kenyon Cells in a Drosophila Connectome  <p>Fig. 5. Connectomes and matching analysis of the <i>Drosophila</i> connectomes.</p> <p>Background:</p> <ul style="list-style-type: none"> Subgraph Isomorphism Graph Matching Algorithms Statistical Models <p>Methods:</p> <ul style="list-style-type: none"> Padding Approaches: Naive Padding, Centered Padding, Oracle Padding Graph Matching Approaches: FAQ algorithm 	<ul style="list-style-type: none"> 三种padding方法没看明白。。

Ontology matching																											
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Mohammadi M , Hofman W , Tan Y H . A Comparative Study of Ontology Matching Systems via Inferential Statistics[J]. IEEE Transactions on Knowledge and Data Engineering, 2019, 31(4):615-628.	<p>Aim:</p> <p>This paper examines the alignment systems using statistical inference since averaging is statistically unsafe and inappropriate.</p> <p>Contributions:</p> <ul style="list-style-type: none">It is the first paper considering the statistical inference for comparison of two or more ontology matching systems. <p>Applications:</p> <ul style="list-style-type: none">The statistical methods are then applied to benchmark and multifarm tracks from the ontology matching evaluation initiative (OAEI) 2015 and their results are reported and visualized by critical difference diagrams.	<p>Key results:</p> <ul style="list-style-type: none">Comparison of two systems <div><p>TABLE 1</p><p>The Tests for Comparison of Two Systems over N Datasets</p><table><tr><th>Test</th><th>Presumptions</th><th>Applicability</th></tr><tr><td>Paired t</td><td>Normality of differences</td><td>$N > 30$</td></tr><tr><td>Signed-rank</td><td>symmetry of differences w.r.t median</td><td>$N > 10$</td></tr><tr><td>McNemar</td><td>-</td><td>$N < 10$</td></tr></table><p><i>Applicability is roughly the situation that test can be used and its results are valid and differences refer to the differences in performance scores.</i></p></div> <ul style="list-style-type: none">Comparison of multiple systems <div><p>TABLE 4</p><p>The Tests for Comparison of Multiple Systems over N Datasets</p><table><tr><th>Test</th><th>Presumptions</th><th>Applicability</th></tr><tr><td>ANOVA t</td><td>Sphericity</td><td>$N > 30$</td></tr><tr><td>Friedman</td><td>-</td><td>$N > 10$</td></tr><tr><td>Quade</td><td>-</td><td>$N < 10$</td></tr></table><p><i>Applicability is roughly the situation that test results are valid.</i></p></div>	Test	Presumptions	Applicability	Paired t	Normality of differences	$N > 30$	Signed-rank	symmetry of differences w.r.t median	$N > 10$	McNemar	-	$N < 10$	Test	Presumptions	Applicability	ANOVA t	Sphericity	$N > 30$	Friedman	-	$N > 10$	Quade	-	$N < 10$	<ul style="list-style-type: none">需要补一补概率论的假设检验部分
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