

# Web genre classification with methods for structured output prediction<sup>1</sup>

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## Abstract

This is the supplementary information for the paper. It contains all of the results obtained with the experimental evaluation using single tree models.

*Keywords:* web genre classification; hierarchy construction; hierarchical multi-label classification

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## 1. Complete results using single tree models

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Table 2: The performance of the different machine learning tasks applied on the different features using single tree models. The hierarchy construction method are abbreviated as follows: balanced  $k$ -means clustering (BkM), predictive clustering tree (PCT), clustering with complete linkage (CL), clustering with single linkage (SL), random (RND) and manual (MAN). The evaluation measure is *accuracy* where larger value means better result.

	HMC										HSC										MLC	SC					
	CL	SL	B2M	B3M	B4M	PCT	RND	MAN	CL	SL	B2M	B3M	B4M	PCT	RND	MAN	CL	SL	B2M	B3M			B4M	PCT	RND	MAN	
<i>20-genes</i>	0.105	0.11	0.089	0.087	0.084	0.088	0.081	0.013	0.013	0.013	0.013	0.013	0.013	0.012	0.014	0.013	0.013	0.013	0.013	0.013	0.013	0.012	0.014	0.013	0.092	0.078	
BOWsrf	0.097	0.092	0.081	0.083	0.082	0.086	0.096	0.011	0.011	0.011	0.012	0.012	0.012	0.012	0.012	0.01	0.011	0.011	0.011	0.012	0.012	0.012	0.012	0.01	0.102	0.083	
TFIDFSrf	0.101	0.115	0.107	0.109	0.105	0.114	0.092	0.014	0.014	0.014	0.014	0.014	0.014	0.014	0.015	0.014	0.014	0.014	0.014	0.014	0.014	0.014	0.015	0.014	0.092	0.102	
Struc	0.119	0.117	0.092	0.102	0.096	0.102	0.093	0.04	0.04	0.036	0.037	0.04	0.04	0.04	0.036	0.033	0.04	0.04	0.036	0.037	0.04	0.04	0.036	0.033	0.092	0.104	
Pres	0.097	0.091	0.091	0.092	0.092	0.1	0.091	0.094	0.074	0.066	0.066	0.066	0.065	0.065	0.065	0.063	0.074	0.074	0.066	0.066	0.065	0.065	0.065	0.063	0.089	0.087	
Context	0.105	0.109	0.109	0.113	0.115	0.11	0.109	0.109	0.013	0.012	0.012	0.012	0.012	0.011	0.013	0.011	0.013	0.012	0.012	0.012	0.012	0.011	0.013	0.011	0.091	0.103	
PV_R	0.125	0.114	0.1	0.1	0.103	0.105	0.095	0.011	0.01	0.01	0.01	0.01	0.01	0.01	0.012	0.01	0.011	0.01	0.01	0.01	0.01	0.01	0.012	0.01	0.091	0.097	
PV_C	0.093	0.096	0.106	0.103	0.126	0.113	0.097	0.121	0.013	0.012	0.012	0.012	0.011	0.011	0.013	0.012	0.013	0.012	0.012	0.012	0.011	0.011	0.013	0.012	0.091	0.105	
PV_GLR	0.11	0.109	0.128	0.093	0.12	0.093	0.136	0.012	0.012	0.011	0.011	0.011	0.011	0.011	0.012	0.011	0.012	0.011	0.011	0.011	0.011	0.011	0.012	0.011	0.091	0.102	
PV_GLC	0.089	0.105	0.095	0.094	0.091	0.091	0.092	0.014	0.014	0.013	0.013	0.013	0.013	0.013	0.014	0.014	0.014	0.014	0.013	0.013	0.013	0.013	0.014	0.014	0.091	0.086	
Ngrams	SPIRIT																										
BOWsrf	0.1	0.101	0.1	0.102	0.103	0.103	0.101	0.103	0.012	0.011	0.014	0.012	0.012	0.012	0.016	0.014	0.012	0.011	0.014	0.012	0.012	0.012	0.012	0.016	0.014	0.109	0.109
TFIDFSrf	0.1	0.105	0.107	0.104	0.103	0.103	0.101	0.105	0.011	0.01	0.011	0.011	0.011	0.011	0.012	0.011	0.011	0.01	0.011	0.011	0.011	0.011	0.012	0.011	0.109	0.114	
Struc	0.104	0.106	0.106	0.097	0.104	0.104	0.109	0.105	0.012	0.011	0.012	0.012	0.012	0.012	0.013	0.011	0.012	0.011	0.012	0.012	0.012	0.011	0.013	0.011	0.114	0.119	
Pres	0.103	0.118	0.109	0.109	0.107	0.107	0.103	0.107	0.047	0.048	0.045	0.045	0.045	0.048	0.044	0.052	0.047	0.048	0.045	0.045	0.047	0.048	0.044	0.052	0.144	0.11	
Context	0.113	0.113	0.13	0.107	0.109	0.109	0.11	0.106	0.082	0.082	0.083	0.084	0.082	0.083	0.085	0.088	0.082	0.082	0.083	0.084	0.082	0.083	0.085	0.088	0.129	0.136	
PV_R	0.119	0.119	0.112	0.108	0.11	0.116	0.11	0.108	0.011	0.01	0.012	0.011	0.011	0.011	0.013	0.012	0.011	0.01	0.012	0.011	0.01	0.011	0.013	0.012	0.108	0.126	
PV_C	0.118	0.117	0.117	0.115	0.111	0.111	0.114	0.113	0.011	0.01	0.012	0.011	0.011	0.011	0.013	0.012	0.011	0.01	0.012	0.011	0.01	0.011	0.013	0.012	0.117	0.128	
PV_GLR	0.126	0.117	0.125	0.128	0.122	0.122	0.125	0.118	0.011	0.01	0.012	0.011	0.011	0.011	0.013	0.012	0.011	0.01	0.012	0.011	0.01	0.011	0.013	0.012	0.136	0.13	
PV_GLC	0.123	0.124	0.116	0.115	0.115	0.116	0.122	0.115	0.011	0.01	0.013	0.011	0.011	0.011	0.013	0.012	0.011	0.01	0.013	0.011	0.01	0.011	0.013	0.012	0.129	0.129	
Ngrams	0.117	0.115	0.116	0.119	0.122	0.108	0.115	0.118	0.016	0.014	0.017	0.014	0.014	0.015	0.015	0.013	0.016	0.014	0.014	0.014	0.014	0.015	0.015	0.013	0.113	0.12	

Table 3: The performance of the different machine learning tasks applied on the different features using single tree models. The hierarchy construction method are abbreviated as follows: balanced  $k$ -means clustering (BkM), predictive clustering tree (PCT), clustering with complete linkage (CL), clustering with single linkage (SL), random (RND) and manual (MAN). The evaluation measure is *precision* where larger value means better result.

	HMC										HSC					MLC		SC	
	CL	SL	B2M	B3M	B4M	PCT	RND	MAN	CL	SL	B2M	B3M	B4M	PCT	RND	MAN			
<i>20-genes</i>																			
BOWStf	0.243	0.256	0.314	0.323	0.311	0.323	0.294	0.302	0.886	0.892	0.889	0.888	0.894	0.895	0.88	0.89	0.086	0.341	
TFIDFStf	0.186	0.218	0.306	0.339	0.341	0.335	0.244	0.312	0.902	0.901	0.91	0.893	0.905	0.908	0.897	0.917	0.147	0.326	
Struc	0.159	0.16	0.224	0.221	0.242	0.213	0.177	0.197	0.876	0.878	0.878	0.885	0.886	0.882	0.873	0.885	0.086	0.176	
Pres	0.188	0.165	0.185	0.23	0.205	0.23	0.187	0.183	0.666	0.656	0.691	0.678	0.661	0.653	0.666	0.7	0.086	0.204	
Context	0.212	0.194	0.273	0.266	0.265	0.25	0.237	0.264	0.398	0.401	0.394	0.401	0.403	0.403	0.412	0.395	0.12	0.183	
PV_R	0.17	0.166	0.199	0.196	0.199	0.196	0.157	0.2	0.886	0.891	0.896	0.894	0.897	0.899	0.884	0.897	0.093	0.188	
PV_C	0.184	0.142	0.194	0.188	0.2	0.211	0.181	0.2	0.902	0.908	0.912	0.91	0.912	0.912	0.899	0.91	0.093	0.23	
PV_GLR	0.114	0.119	0.181	0.185	0.164	0.182	0.181	0.174	0.887	0.891	0.891	0.896	0.893	0.897	0.879	0.89	0.093	0.175	
PV_GLC	0.165	0.174	0.184	0.158	0.19	0.158	0.219	0.183	0.898	0.904	0.911	0.911	0.909	0.91	0.9	0.906	0.093	0.201	
Ngrams	0.194	0.258	0.28	0.272	0.248	0.236	0.212	0.259	0.881	0.882	0.894	0.884	0.886	0.885	0.868	0.878	0.093	0.309	
<i>SPRIT</i>																			
BOWStf	0.631	0.622	0.591	0.616	0.612	0.594	0.587	0.614	0.938	0.948	0.926	0.943	0.946	0.948	0.914	0.932	0.611	0.51	
TFIDFStf	0.628	0.607	0.574	0.602	0.6	0.6	0.591	0.601	0.948	0.946	0.946	0.943	0.94	0.946	0.94	0.943	0.611	0.48	
Struc	0.577	0.569	0.581	0.61	0.579	0.579	0.56	0.577	0.944	0.944	0.946	0.942	0.942	0.943	0.943	0.946	0.574	0.468	
Pres	0.505	0.556	0.549	0.549	0.549	0.547	0.538	0.549	0.772	0.777	0.784	0.779	0.781	0.765	0.791	0.752	0.525	0.464	
Context	0.538	0.537	0.485	0.552	0.543	0.545	0.537	0.545	0.635	0.636	0.628	0.627	0.633	0.631	0.621	0.6	0.455	0.414	
PV_R	0.477	0.501	0.537	0.558	0.55	0.535	0.543	0.56	0.944	0.948	0.934	0.94	0.946	0.942	0.931	0.932	0.555	0.411	
PV_C	0.533	0.532	0.532	0.541	0.536	0.535	0.512	0.532	0.943	0.949	0.947	0.949	0.951	0.949	0.935	0.938	0.548	0.418	
PV_GLR	0.528	0.535	0.529	0.524	0.535	0.535	0.529	0.557	0.945	0.953	0.941	0.948	0.951	0.951	0.936	0.937	0.416	0.395	
PV_GLC	0.486	0.5	0.536	0.536	0.536	0.534	0.518	0.536	0.941	0.946	0.937	0.944	0.947	0.946	0.932	0.94	0.46	0.419	
Ngrams	0.512	0.53	0.544	0.534	0.512	0.495	0.553	0.542	0.913	0.924	0.904	0.925	0.921	0.922	0.918	0.933	0.549	0.444	

Table 4: The performance of the different machine learning tasks applied on the different features using single tree models. The hierarchy construction method are abbreviated as follows: balanced  $k$ -means clustering (BkM), predictive clustering tree (PCT), clustering with complete linkage (CL), clustering with single linkage (SL), random (RND) and manual (MAN). The evaluation measure is *recall* where larger value means better result.

	HMC										HSC					MLC	SC
	CL	SL	B2M	B3M	B4M	PCT	RND	MAN	CL	SL	B2M	B3M	B4M	PCT	RND		
<i>20-genes</i>	0.29	0.298	0.281	0.283	0.276	0.29	0.258	0.269	0.91	0.917	0.92	0.919	0.919	0.921	0.905	0.914	
BOWSrf	0.256	0.27	0.271	0.295	0.306	0.299	0.311	0.343	0.906	0.903	0.951	0.899	0.907	0.908	0.9	0.954	
TFIDFSrf	0.221	0.23	0.264	0.254	0.277	0.269	0.232	0.215	0.91	0.918	0.916	0.932	0.924	0.927	0.916	0.93	
Struc	0.233	0.199	0.167	0.232	0.189	0.234	0.254	0.184	0.739	0.725	0.756	0.728	0.733	0.72	0.72	0.734	
Pres	0.187	0.152	0.242	0.246	0.231	0.277	0.202	0.255	0.521	0.521	0.447	0.462	0.459	0.458	0.468	0.443	
Context	0.179	0.185	0.221	0.238	0.242	0.221	0.222	0.222	0.911	0.921	0.926	0.925	0.927	0.931	0.91	0.924	
PV_R	0.262	0.208	0.181	0.198	0.207	0.221	0.179	0.204	0.919	0.931	0.936	0.935	0.936	0.939	0.921	0.937	
PV_C	0.137	0.142	0.165	0.161	0.229	0.198	0.164	0.221	0.913	0.925	0.92	0.927	0.925	0.933	0.909	0.924	
PV_GLR	0.213	0.223	0.275	0.148	0.264	0.148	0.222	0.296	0.924	0.935	0.933	0.939	0.937	0.941	0.923	0.933	
PV_GLC	0.195	0.281	0.27	0.243	0.249	0.23	0.214	0.267	0.912	0.915	0.923	0.919	0.916	0.921	0.896	0.92	
Ngrams																	
<b>SPIRIT</b>																	
BOWSrf	0.577	0.624	0.534	0.585	0.587	0.579	0.529	0.593	0.936	0.949	0.953	0.943	0.946	0.95	0.946	0.94	
TFIDFSrf	0.576	0.629	0.551	0.593	0.582	0.58	0.53	0.599	0.958	0.95	0.956	0.957	0.954	0.962	0.949	0.958	
Struc	0.536	0.539	0.561	0.534	0.536	0.536	0.534	0.552	0.945	0.951	0.944	0.952	0.95	0.951	0.939	0.947	
Pres	0.428	0.584	0.524	0.524	0.501	0.505	0.431	0.501	0.821	0.79	0.824	0.804	0.827	0.799	0.814	0.769	
Context	0.497	0.497	0.555	0.439	0.445	0.443	0.433	0.409	0.675	0.684	0.654	0.673	0.682	0.668	0.654	0.637	
PV_R	0.526	0.522	0.543	0.524	0.507	0.532	0.538	0.529	0.945	0.953	0.956	0.958	0.951	0.948	0.958	0.953	
PV_C	0.511	0.524	0.555	0.554	0.487	0.492	0.556	0.509	0.962	0.963	0.962	0.962	0.966	0.963	0.954	0.957	
PV_GLR	0.528	0.468	0.566	0.566	0.577	0.577	0.566	0.53	0.966	0.955	0.963	0.947	0.955	0.951	0.96	0.952	
PV_GLC	0.51	0.53	0.552	0.549	0.549	0.551	0.573	0.549	0.955	0.958	0.949	0.958	0.955	0.957	0.95	0.954	
Ngrams	0.484	0.475	0.586	0.566	0.583	0.456	0.593	0.574	0.929	0.941	0.914	0.933	0.94	0.934	0.946	0.949	

Table 5: The performance of the different machine learning tasks applied on the different features using single tree models. The hierarchy construction method are abbreviated as follows: balanced  $k$ -means clustering (BKM), predictive clustering tree (PCT), clustering with complete linkage (CL), clustering with single linkage (SL), random (RND) and manual (MAN). The evaluation measure is  $F_1$  score where larger value means better result.

	HMC										HSC					MLC		SC	
	CL	SL	B2M	B3M	B4M	PCT	RND	MAN	CL	SL	B2M	B3M	B4M	PCT	RND	MAN	MAN		
<i>20-genes</i>																			
BOWStf	0.264	0.276	0.296	0.302	0.292	0.306	0.275	0.285	0.897	0.904	0.904	0.904	0.907	0.908	0.892	0.902	0.072	0.367	
TFIDFStf	0.216	0.241	0.287	0.315	0.322	0.316	0.274	0.326	0.904	0.902	0.93	0.896	0.906	0.908	0.898	0.935	0.123	0.357	
Struc	0.185	0.189	0.243	0.237	0.258	0.238	0.201	0.206	0.893	0.898	0.897	0.908	0.904	0.904	0.894	0.907	0.072	0.201	
Pres	0.208	0.18	0.175	0.231	0.197	0.232	0.215	0.184	0.701	0.689	0.722	0.702	0.695	0.685	0.692	0.717	0.072	0.235	
Context	0.199	0.171	0.257	0.256	0.247	0.263	0.218	0.26	0.451	0.453	0.419	0.43	0.429	0.429	0.438	0.417	0.105	0.186	
PV_R	0.175	0.175	0.209	0.215	0.218	0.208	0.184	0.21	0.898	0.906	0.911	0.909	0.912	0.915	0.897	0.911	0.078	0.215	
PV_C	0.216	0.169	0.187	0.193	0.203	0.216	0.18	0.202	0.91	0.919	0.924	0.922	0.924	0.925	0.91	0.923	0.078	0.26	
PV_GLR	0.125	0.129	0.173	0.172	0.191	0.19	0.172	0.195	0.9	0.908	0.905	0.911	0.909	0.915	0.894	0.907	0.078	0.199	
PV_GLC	0.186	0.195	0.221	0.153	0.221	0.153	0.22	0.226	0.911	0.919	0.922	0.925	0.922	0.925	0.911	0.92	0.078	0.225	
Ngrams	0.194	0.269	0.275	0.257	0.249	0.233	0.213	0.263	0.896	0.898	0.908	0.901	0.901	0.903	0.882	0.899	0.078	0.34	
SPIRIT																			
BOWStf	0.603	0.623	0.561	0.6	0.599	0.587	0.557	0.603	0.937	0.949	0.939	0.943	0.946	0.949	0.93	0.936	0.606	0.527	
TFIDFStf	0.601	0.617	0.562	0.598	0.591	0.59	0.559	0.6	0.953	0.948	0.951	0.95	0.947	0.954	0.944	0.951	0.606	0.502	
Struc	0.556	0.554	0.57	0.57	0.557	0.557	0.547	0.564	0.944	0.948	0.945	0.947	0.946	0.947	0.941	0.946	0.567	0.484	
Pres	0.464	0.57	0.536	0.536	0.524	0.525	0.478	0.524	0.796	0.784	0.803	0.791	0.803	0.782	0.802	0.76	0.601	0.441	
Context	0.517	0.516	0.517	0.489	0.489	0.489	0.48	0.467	0.654	0.659	0.641	0.649	0.657	0.649	0.637	0.618	0.489	0.488	
PV_R	0.501	0.512	0.54	0.541	0.528	0.533	0.541	0.544	0.945	0.95	0.945	0.949	0.948	0.945	0.944	0.942	0.501	0.438	
PV_C	0.522	0.528	0.543	0.547	0.51	0.513	0.533	0.52	0.952	0.956	0.954	0.956	0.959	0.956	0.944	0.947	0.533	0.434	
PV_GLR	0.528	0.499	0.547	0.544	0.556	0.556	0.547	0.543	0.955	0.954	0.951	0.948	0.953	0.951	0.948	0.944	0.484	0.42	
PV_GLC	0.498	0.514	0.544	0.543	0.543	0.542	0.544	0.543	0.948	0.952	0.943	0.951	0.951	0.952	0.941	0.947	0.502	0.442	
Ngrams	0.497	0.501	0.564	0.55	0.545	0.475	0.572	0.557	0.921	0.933	0.909	0.929	0.93	0.928	0.932	0.941	0.507	0.466	

Table 6: The performance of the different machine learning tasks applied on the different features using single tree models. The hierarchy construction method are abbreviated as follows: balanced  $k$ -means clustering (BkM), predictive clustering tree (PCT), clustering with complete linkage (CL), clustering with single linkage (SL), random (RND) and manual (MAN). The evaluation measure is *subset accuracy* where larger value means better result.

20-genres	HMC										HSC										MLC	SC					
	CL	SL	B2M	B3M	B4M	PCT	RND	MAN	CL	SL	B2M	B3M	B4M	PCT	RND	MAN	CL	SL	B2M	B3M			B4M	PCT	RND	MAN	
BOWSrf	0.101	0.11	0.176	0.179	0.171	0.182	0.157	0.163	0.766	0.775	0.77	0.774	0.776	0.785	0.76	0.775	0.766	0.775	0.77	0.774	0.776	0.785	0.76	0.775	0.04	0.173	
TFIDFSrf	0.075	0.086	0.176	0.198	0.186	0.194	0.091	0.156	0.811	0.807	0.815	0.785	0.809	0.812	0.793	0.829	0.811	0.807	0.815	0.785	0.809	0.812	0.793	0.829	0.069	0.141	
Struc	0.047	0.048	0.07	0.079	0.082	0.068	0.046	0.067	0.763	0.764	0.747	0.765	0.764	0.762	0.756	0.761	0.763	0.764	0.747	0.765	0.764	0.762	0.756	0.761	0.04	0.066	
Pres	0.069	0.043	0.093	0.104	0.101	0.104	0.061	0.074	0.445	0.437	0.469	0.465	0.434	0.423	0.474	0.493	0.445	0.437	0.469	0.465	0.434	0.423	0.474	0.493	0.04	0.071	
Context	0.107	0.103	0.153	0.148	0.149	0.122	0.129	0.142	0.181	0.18	0.211	0.21	0.216	0.214	0.224	0.209	0.181	0.18	0.211	0.21	0.216	0.214	0.224	0.209	0.069	0.073	
PV_R	0.075	0.061	0.055	0.056	0.049	0.055	0.023	0.055	0.78	0.79	0.795	0.788	0.794	0.797	0.78	0.794	0.78	0.79	0.795	0.788	0.794	0.797	0.78	0.794	0.044	0.075	
PV_C	0.062	0.044	0.099	0.09	0.087	0.105	0.077	0.1	0.804	0.812	0.82	0.816	0.82	0.819	0.799	0.817	0.804	0.812	0.82	0.816	0.82	0.819	0.799	0.817	0.044	0.089	
PV_GLR	0.045	0.038	0.072	0.083	0.034	0.062	0.086	0.039	0.784	0.79	0.785	0.795	0.796	0.799	0.774	0.79	0.784	0.79	0.785	0.795	0.796	0.799	0.774	0.79	0.044	0.063	
PV_GLC	0.055	0.055	0.045	0.077	0.045	0.077	0.098	0.021	0.799	0.8	0.808	0.807	0.806	0.805	0.793	0.799	0.799	0.8	0.808	0.807	0.806	0.805	0.793	0.799	0.044	0.073	
Ngrams	0.102	0.113	0.167	0.168	0.137	0.141	0.108	0.141	0.762	0.765	0.779	0.768	0.77	0.771	0.754	0.761	0.762	0.765	0.779	0.768	0.77	0.771	0.754	0.761	0.044	0.127	
SPIRIT																											
BOWSrf	0.288	0.274	0.264	0.261	0.261	0.262	0.254	0.257	0.84	0.848	0.829	0.848	0.843	0.848	0.803	0.826	0.84	0.848	0.829	0.848	0.843	0.848	0.803	0.826	0.26	0.227	
TFIDFSrf	0.288	0.262	0.235	0.251	0.27	0.268	0.265	0.251	0.867	0.866	0.866	0.86	0.855	0.865	0.855	0.858	0.867	0.866	0.866	0.86	0.855	0.865	0.855	0.858	0.26	0.211	
Struc	0.255	0.252	0.264	0.295	0.264	0.264	0.241	0.27	0.853	0.855	0.853	0.851	0.852	0.854	0.845	0.853	0.853	0.855	0.853	0.851	0.852	0.854	0.845	0.853	0.235	0.177	
Pres	0.251	0.207	0.281	0.281	0.275	0.275	0.243	0.275	0.525	0.511	0.53	0.533	0.518	0.497	0.533	0.471	0.525	0.511	0.53	0.533	0.518	0.497	0.533	0.471	0.189	0.2	
Context	0.19	0.191	0.138	0.25	0.237	0.238	0.24	0.267	0.313	0.316	0.313	0.296	0.315	0.309	0.291	0.292	0.313	0.316	0.313	0.296	0.315	0.309	0.291	0.292	0.096	0.052	
PV_R	0.194	0.204	0.216	0.251	0.255	0.237	0.233	0.248	0.861	0.867	0.848	0.861	0.863	0.86	0.85	0.855	0.861	0.867	0.848	0.861	0.863	0.86	0.85	0.855	0.271	0.14	
PV_C	0.227	0.218	0.203	0.226	0.234	0.237	0.197	0.224	0.855	0.867	0.858	0.86	0.872	0.862	0.842	0.852	0.855	0.867	0.858	0.86	0.872	0.862	0.842	0.852	0.247	0.152	
PV_GLR	0.206	0.213	0.193	0.193	0.197	0.197	0.193	0.235	0.865	0.873	0.856	0.86	0.868	0.869	0.849	0.852	0.865	0.873	0.856	0.86	0.868	0.869	0.849	0.852	0.052	0.126	
PV_GLC	0.182	0.189	0.217	0.217	0.217	0.217	0.197	0.217	0.857	0.864	0.846	0.862	0.868	0.862	0.836	0.847	0.857	0.864	0.846	0.862	0.868	0.862	0.836	0.847	0.098	0.15	
Ngrams	0.223	0.247	0.238	0.241	0.207	0.237	0.243	0.243	0.813	0.829	0.806	0.826	0.826	0.818	0.815	0.835	0.813	0.829	0.806	0.826	0.826	0.818	0.815	0.835	0.268	0.169	

Table 7: The performance of the different machine learning tasks applied on the different features using single tree models. The hierarchy construction method are abbreviated as follows: balanced  $k$ -means clustering (BKM), predictive clustering tree (PCT), clustering with complete linkage (CL), clustering with single linkage (SL), random (RND) and manual (MAN). The evaluation measure is *micro precision* where larger value means better result.

	HMC										HSC					MLC		SC	
	CL	SL	B2M	B3M	B4M	PCT	RND	MAN	CL	SL	B2M	B3M	B4M	PCT	RND	MAN			
<i>20-genes</i>																			
BOWStf	0.248	0.24	0.315	0.324	0.339	0.319	0.308	0.357	0.899	0.9	0.891	0.894	0.899	0.903	0.896	0.902	0.13	0.41	
TFIDFStf	0.265	0.293	0.358	0.356	0.362	0.34	0.241	0.304	0.938	0.939	0.904	0.928	0.942	0.943	0.933	0.91	0.147	0.384	
Struc	0.226	0.196	0.232	0.219	0.244	0.214	0.209	0.262	0.889	0.885	0.876	0.879	0.885	0.88	0.878	0.877	0.13	0.229	
Pres	0.187	0.174	0.23	0.227	0.229	0.228	0.196	0.238	0.695	0.694	0.722	0.731	0.697	0.697	0.736	0.772	0.13	0.246	
Context	0.224	0.229	0.282	0.281	0.274	0.26	0.257	0.276	0.453	0.453	0.513	0.507	0.514	0.522	0.519	0.537	0.176	0.273	
PV_R	0.193	0.185	0.201	0.197	0.196	0.199	0.204	0.201	0.906	0.906	0.909	0.904	0.907	0.907	0.908	0.912	0.14	0.233	
PV_C	0.184	0.179	0.211	0.22	0.22	0.218	0.231	0.24	0.925	0.92	0.921	0.918	0.922	0.919	0.915	0.918	0.14	0.278	
PV_GLR	0.201	0.196	0.178	0.187	0.168	0.181	0.208	0.172	0.907	0.902	0.906	0.907	0.908	0.906	0.903	0.902	0.14	0.216	
PV_GLC	0.198	0.207	0.184	0.219	0.197	0.219	0.215	0.181	0.912	0.908	0.916	0.91	0.912	0.909	0.911	0.906	$\infty$	0.239	
Ngrams	0.266	0.251	0.282	0.267	0.296	0.275	0.246	0.297	0.889	0.888	0.894	0.889	0.891	0.889	0.893	0.882	0.14	0.359	
SPIRIT																			
BOWStf	0.558	0.546	0.561	0.545	0.543	0.541	0.556	0.539	0.956	0.955	0.927	0.952	0.954	0.954	0.914	0.942	0.515	0.517	
TFIDFStf	0.556	0.528	0.522	0.537	0.543	0.542	0.558	0.531	0.951	0.958	0.95	0.946	0.946	0.949	0.948	0.948	0.515	0.492	
Struc	0.539	0.528	0.528	0.577	0.539	0.538	0.517	0.531	0.955	0.951	0.956	0.945	0.951	0.951	0.954	0.955	0.493	0.47	
Pres	0.558	0.479	0.518	0.518	0.527	0.525	0.563	0.527	0.778	0.795	0.791	0.805	0.779	0.784	0.808	0.775	0.414	0.515	
Context	0.496	0.495	0.433	0.53	0.518	0.52	0.515	0.54	0.631	0.633	0.637	0.625	0.63	0.63	0.624	0.612	0.434	0.418	
PV_R	0.47	0.471	0.5	0.518	0.51	0.486	0.51	0.521	0.961	0.958	0.94	0.947	0.958	0.956	0.93	0.944	0.522	0.439	
PV_C	0.473	0.481	0.479	0.489	0.505	0.504	0.492	0.499	0.943	0.951	0.941	0.948	0.951	0.95	0.934	0.942	0.478	0.429	
PV_GLR	0.445	0.479	0.452	0.442	0.462	0.462	0.452	0.476	0.94	0.961	0.938	0.958	0.957	0.962	0.933	0.945	0.416	0.422	
PV_GLC	0.455	0.452	0.486	0.487	0.487	0.487	0.465	0.487	0.951	0.954	0.944	0.948	0.956	0.954	0.937	0.945	0.435	0.431	
Ngrams	0.477	0.487	0.485	0.474	0.464	0.525	0.49	0.479	0.931	0.934	0.934	0.94	0.932	0.935	0.922	0.939	0.498	0.465	



Table 8: The performance of the different machine learning tasks applied on the different features using single tree models. The hierarchy construction method are abbreviated as follows: balanced  $k$ -means clustering (BkM), predictive clustering tree (PCT), clustering with complete linkage (CL), clustering with single linkage (SL), random (RND) and manual (MAN). The evaluation measure is *micro recall* where larger value means better result.

	HMC										HSC				MLC	SC		
	CL	SL	B2M	B3M	B4M	PCT	RND	MAN	CL	SL	B2M	B3M	B4M	PCT			RND	MAN
<i>20-genes</i>	0.276	0.294	0.272	0.275	0.274	0.277	0.25	0.267	0.9	0.911	0.91	0.909	0.911	0.914	0.892	0.906	0.064	0.383
BOWSrf	0.25	0.267	0.266	0.288	0.297	0.292	0.294	0.338	0.9	0.899	0.946	0.891	0.901	0.902	0.891	0.949	0.109	0.383
TFIDFSrf	0.21	0.228	0.256	0.247	0.268	0.264	0.221	0.204	0.899	0.911	0.906	0.924	0.916	0.921	0.903	0.921	0.064	0.223
Struc	0.233	0.198	0.159	0.216	0.181	0.22	0.247	0.176	0.724	0.71	0.75	0.715	0.718	0.704	0.718	0.724	0.064	0.265
Pres	0.183	0.149	0.235	0.239	0.227	0.269	0.192	0.25	0.511	0.512	0.434	0.449	0.442	0.445	0.457	0.433	0.089	0.178
Context	0.177	0.183	0.21	0.224	0.23	0.21	0.213	0.21	0.901	0.914	0.921	0.918	0.922	0.928	0.898	0.918	0.069	0.234
PV_R	0.251	0.196	0.18	0.195	0.209	0.217	0.176	0.194	0.91	0.926	0.931	0.928	0.933	0.935	0.909	0.931	0.069	0.278
PV_C	0.13	0.141	0.161	0.158	0.221	0.194	0.158	0.213	0.905	0.921	0.914	0.921	0.923	0.931	0.897	0.918	0.069	0.215
PV_GLR	0.209	0.223	0.264	0.149	0.258	0.149	0.22	0.292	0.911	0.926	0.925	0.93	0.931	0.935	0.909	0.925	0.069	0.24
PV_GLC	0.185	0.283	0.267	0.234	0.253	0.222	0.203	0.267	0.905	0.91	0.916	0.913	0.91	0.916	0.893	0.913	0.069	0.366
Ngrams	SPIRIT																	
BOWSrf	0.533	0.586	0.5	0.547	0.55	0.547	0.496	0.551	0.932	0.943	0.95	0.938	0.939	0.942	0.945	0.934	0.568	0.518
TFIDFSrf	0.531	0.592	0.513	0.558	0.551	0.548	0.498	0.559	0.954	0.949	0.952	0.953	0.952	0.959	0.944	0.954	0.568	0.502
Struc	0.502	0.504	0.534	0.506	0.505	0.505	0.499	0.526	0.941	0.949	0.938	0.948	0.946	0.948	0.932	0.943	0.513	0.477
Pres	0.399	0.549	0.487	0.487	0.465	0.468	0.389	0.465	0.813	0.775	0.811	0.789	0.811	0.785	0.797	0.755	0.677	0.398
Context	0.444	0.443	0.502	0.386	0.391	0.39	0.383	0.358	0.643	0.651	0.618	0.638	0.649	0.633	0.622	0.606	0.48	0.546
PV_R	0.498	0.5	0.507	0.501	0.478	0.509	0.503	0.503	0.943	0.951	0.953	0.957	0.949	0.946	0.959	0.952	0.422	0.448
PV_C	0.472	0.488	0.514	0.521	0.454	0.46	0.523	0.476	0.958	0.959	0.954	0.954	0.961	0.959	0.95	0.953	0.501	0.43
PV_GLR	0.486	0.432	0.52	0.523	0.528	0.528	0.52	0.481	0.965	0.951	0.959	0.941	0.95	0.947	0.956	0.951	0.528	0.426
PV_GLC	0.474	0.495	0.522	0.519	0.519	0.522	0.54	0.519	0.953	0.955	0.944	0.955	0.953	0.954	0.946	0.948	0.513	0.454
Ngrams	0.451	0.441	0.562	0.544	0.56	0.408	0.568	0.55	0.93	0.943	0.917	0.933	0.94	0.933	0.943	0.945	0.425	0.465



Table 10: The performance of the different machine learning tasks applied on the different features using single tree models. The hierarchy construction method are abbreviated as follows: balanced  $k$ -means clustering (BkM), predictive clustering tree (PCT), clustering with complete linkage (CL), clustering with single linkage (SL), random (RND) and manual (MAN). The evaluation measure is *macro precision* where larger value means better result.

20-genes	HMC										HSC										MLC	SC														
	CL	SL	B2M	B3M	B4M	PCT	RND	MAN	CL	SL	B2M	B3M	B4M	PCT	RND	MAN	CL	SL	B2M	B3M			B4M	PCT	RND	MAN										
	BOWsrf	0.176	0.191	0.229	0.279	0.289	0.269	0.205	0.311	0.916	0.917	0.908	0.914	0.915	0.92	0.917	0.919	0.013	0.439	0.176			0.191	0.229	0.279	0.289	0.269	0.205	0.311	0.916	0.917	0.908	0.914	0.915	0.92	0.917
TFIDFSrf	0.202	0.259	0.309	0.31	0.341	0.318	0.196	0.379	0.947	0.948	0.919	0.944	0.95	0.953	0.948	0.921	0.054	0.419	0.202	0.259	0.309	0.31	0.341	0.318	0.196	0.379	0.947	0.948	0.919	0.944	0.95	0.953	0.948	0.921	0.054	0.419
Struc	0.109	0.124	0.173	0.16	0.184	0.169	0.066	0.164	0.898	0.894	0.885	0.887	0.893	0.886	0.888	0.885	0.013	0.224	0.109	0.124	0.173	0.16	0.184	0.169	0.066	0.164	0.898	0.894	0.885	0.887	0.893	0.886	0.888	0.885	0.013	0.224
Pres	0.113	0.076	0.196	0.215	0.214	0.217	0.183	0.131	0.726	0.722	0.736	0.737	0.724	0.726	0.747	0.79	0.013	0.278	0.113	0.076	0.196	0.215	0.214	0.217	0.183	0.131	0.726	0.722	0.736	0.737	0.724	0.726	0.747	0.79	0.013	0.278
Context	0.198	0.061	0.319	0.324	0.324	0.33	0.196	0.383	0.509	0.51	0.537	0.536	0.533	0.547	0.548	0.549	0.062	0.374	0.198	0.061	0.319	0.324	0.324	0.33	0.196	0.383	0.509	0.51	0.537	0.536	0.533	0.547	0.548	0.549	0.062	0.374
PV_R	0.114	0.085	0.067	0.097	0.071	0.067	0.135	0.067	0.914	0.912	0.918	0.912	0.915	0.913	0.913	0.918	0.014	0.235	0.114	0.085	0.067	0.097	0.071	0.067	0.135	0.067	0.914	0.912	0.918	0.912	0.915	0.913	0.913	0.918	0.014	0.235
PV_C	0.151	0.127	0.191	0.227	0.164	0.24	0.125	0.179	0.93	0.925	0.926	0.923	0.927	0.923	0.923	0.922	0.014	0.29	0.151	0.127	0.191	0.227	0.164	0.24	0.125	0.179	0.93	0.925	0.926	0.923	0.927	0.923	0.923	0.922	0.014	0.29
PV_GLR	0.078	0.076	0.046	0.043	0.091	0.093	0.046	0.063	0.914	0.907	0.913	0.911	0.915	0.91	0.907	0.906	0.014	0.207	0.078	0.076	0.046	0.043	0.091	0.093	0.046	0.063	0.914	0.907	0.913	0.911	0.915	0.91	0.907	0.906	0.014	0.207
PV_GLC	0.134	0.121	0.089	0.085	0.083	0.085	0.157	0.084	0.92	0.913	0.922	0.914	0.919	0.913	0.916	0.913	0.014	0.243	0.134	0.121	0.089	0.085	0.083	0.085	0.157	0.084	0.92	0.913	0.922	0.914	0.919	0.913	0.916	0.913	0.014	0.243
Ngrams	0.168	0.18	0.281	0.224	0.302	0.257	0.188	0.282	0.904	0.902	0.907	0.904	0.907	0.903	0.905	0.903	0.014	0.4	0.168	0.18	0.281	0.224	0.302	0.257	0.188	0.282	0.904	0.902	0.907	0.904	0.907	0.903	0.905	0.903	0.014	0.4
SPIRIT																																				
BOWsrf	0.17	0.167	0.163	0.167	0.168	0.187	0.179	0.254	0.958	0.958	0.942	0.956	0.958	0.955	0.94	0.955	0.135	0.267	0.17	0.167	0.163	0.167	0.168	0.187	0.179	0.254	0.958	0.958	0.942	0.956	0.958	0.955	0.94	0.955	0.135	0.267
TFIDFSrf	0.17	0.197	0.169	0.205	0.191	0.189	0.177	0.272	0.925	0.928	0.924	0.921	0.923	0.924	0.926	0.924	0.135	0.253	0.17	0.197	0.169	0.205	0.191	0.189	0.177	0.272	0.925	0.928	0.924	0.921	0.923	0.924	0.926	0.924	0.135	0.253
Struc	0.138	0.164	0.139	0.15	0.14	0.14	0.185	0.153	0.938	0.936	0.938	0.931	0.934	0.934	0.943	0.957	0.117	0.202	0.138	0.164	0.139	0.15	0.14	0.14	0.185	0.153	0.938	0.936	0.938	0.931	0.934	0.934	0.943	0.957	0.117	0.202
Pres	0.167	0.12	0.15	0.15	0.131	0.131	0.109	0.131	0.754	0.787	0.755	0.764	0.779	0.756	0.711	0.72	0.109	0.162	0.167	0.12	0.15	0.15	0.131	0.131	0.109	0.131	0.754	0.787	0.755	0.764	0.779	0.756	0.711	0.72	0.109	0.162
Context	0.065	0.065	0.09	0.065	0.08	0.08	0.076	0.054	0.619	0.62	0.616	0.614	0.623	0.61	0.606	0.611	0.059	0.157	0.065	0.065	0.09	0.065	0.08	0.08	0.076	0.054	0.619	0.62	0.616	0.614	0.623	0.61	0.606	0.611	0.059	0.157
PV_R	0.156	0.137	0.152	0.178	0.144	0.145	0.188	0.181	0.943	0.942	0.929	0.931	0.939	0.94	0.923	0.854	0.123	0.191	0.156	0.137	0.152	0.178	0.144	0.145	0.188	0.181	0.943	0.942	0.929	0.931	0.939	0.94	0.923	0.854	0.123	0.191
PV_C	0.115	0.129	0.123	0.128	0.133	0.166	0.132	0.132	0.862	0.866	0.862	0.865	0.866	0.864	0.864	0.863	0.117	0.2	0.115	0.129	0.123	0.128	0.133	0.166	0.132	0.132	0.862	0.866	0.862	0.865	0.866	0.864	0.864	0.863	0.117	0.2
PV_GLR	0.106	0.117	0.113	0.109	0.116	0.116	0.113	0.112	0.953	0.967	0.952	0.965	0.965	0.968	0.948	0.954	0.059	0.181	0.106	0.117	0.113	0.109	0.116	0.116	0.113	0.112	0.953	0.967	0.952	0.965	0.965	0.968	0.948	0.954	0.059	0.181
PV_GLC	0.138	0.117	0.129	0.129	0.129	0.164	0.12	0.129	0.855	0.856	0.853	0.854	0.857	0.856	0.846	0.859	0.122	0.192	0.138	0.117	0.129	0.129	0.129	0.164	0.12	0.129	0.855	0.856	0.853	0.854	0.857	0.856	0.846	0.859	0.122	0.192
Ngrams	0.139	0.124	0.127	0.125	0.162	0.16	0.129	0.15	0.952	0.953	0.954	0.961	0.954	0.952	0.95	0.959	0.111	0.249	0.139	0.124	0.127	0.125	0.162	0.16	0.129	0.15	0.952	0.953	0.954	0.961	0.954	0.952	0.95	0.959	0.111	0.249



Table 12: The performance of the different machine learning tasks applied on the different features using single tree models. The hierarchy construction method are abbreviated as follows: balanced  $k$ -means clustering (BkM), predictive clustering tree (PCT), clustering with complete linkage (CL), clustering with single linkage (SL), random (RND) and manual (MAN). The evaluation measure is *macro F<sub>1</sub>* where larger value means better result.

	HMC										HSC					MLC	SC
	CL	SL	B2M	B3M	B4M	PCT	RND	MAN	CL	SL	B2M	B3M	B4M	PCT	RND		
<i>20-genes</i>	0.177	0.176	0.194	0.192	0.202	0.206	0.17	0.209	0.897	0.905	0.9	0.904	0.904	0.909	0.895	0.904	
BOWSrf	0.189	0.203	0.223	0.232	0.239	0.237	0.187	0.271	0.918	0.918	0.924	0.912	0.92	0.921	0.914	0.926	
TFIDFSrf	0.104	0.111	0.126	0.11	0.142	0.125	0.077	0.12	0.885	0.89	0.885	0.897	0.893	0.893	0.883	0.891	
Struc	0.118	0.063	0.109	0.138	0.115	0.149	0.126	0.12	0.725	0.711	0.738	0.722	0.712	0.711	0.732	0.734	
Pres	0.101	0.051	0.189	0.189	0.174	0.207	0.118	0.193	0.493	0.495	0.486	0.483	0.484	0.488	0.493	0.486	
Context	0.086	0.084	0.078	0.093	0.091	0.08	0.12	0.078	0.886	0.895	0.904	0.904	0.902	0.907	0.89	0.905	
PV_R	0.134	0.117	0.129	0.139	0.125	0.166	0.104	0.134	0.906	0.914	0.918	0.92	0.92	0.92	0.907	0.919	
PV_C	0.057	0.066	0.055	0.052	0.098	0.084	0.048	0.079	0.89	0.898	0.898	0.905	0.902	0.907	0.888	0.9	
PV_GLR	0.093	0.111	0.106	0.065	0.099	0.065	0.104	0.094	0.9	0.908	0.912	0.915	0.913	0.914	0.903	0.909	
PV_GLC	0.116	0.156	0.203	0.16	0.209	0.177	0.112	0.22	0.902	0.904	0.905	0.906	0.906	0.907	0.899	0.905	
Ngrams																	
<b>SPIRIT</b>																	
BOWSrf	0.165	0.171	0.159	0.173	0.165	0.178	0.165	0.189	0.851	0.854	0.847	0.847	0.852	0.852	0.841	0.821	
TFIDFSrf	0.165	0.195	0.164	0.192	0.182	0.18	0.164	0.197	0.847	0.848	0.846	0.843	0.845	0.848	0.84	0.845	
Struc	0.148	0.156	0.152	0.155	0.148	0.147	0.171	0.154	0.852	0.853	0.85	0.849	0.851	0.852	0.838	0.852	
Pres	0.137	0.142	0.152	0.152	0.142	0.142	0.099	0.142	0.577	0.568	0.568	0.562	0.574	0.557	0.573	0.541	
Context	0.081	0.081	0.101	0.069	0.077	0.077	0.073	0.059	0.377	0.379	0.374	0.383	0.374	0.367	0.398	0.385	
PV_R	0.153	0.15	0.147	0.157	0.142	0.156	0.149	0.156	0.846	0.848	0.84	0.842	0.845	0.842	0.829	0.777	
PV_C	0.126	0.136	0.137	0.142	0.134	0.144	0.133	0.137	0.803	0.806	0.804	0.805	0.806	0.805	0.789	0.787	
PV_GLR	0.121	0.118	0.131	0.131	0.132	0.132	0.131	0.124	0.839	0.84	0.834	0.832	0.836	0.836	0.817	0.782	
PV_GLC	0.131	0.127	0.147	0.146	0.146	0.152	0.141	0.146	0.792	0.794	0.785	0.788	0.794	0.791	0.78	0.772	
Ngrams	0.138	0.127	0.149	0.148	0.166	0.134	0.151	0.155	0.847	0.851	0.846	0.851	0.851	0.85	0.844	0.854	



Table 14: The performance of the different machine learning tasks applied on the different features using single tree models. The hierarchy construction method are abbreviated as follows: balanced  $k$ -means clustering (BkM), predictive clustering tree (PCT), clustering with complete linkage (CL), clustering with single linkage (SL), random (RND) and manual (MAN). The evaluation measure is coverage where smaller value means better result.

	HMC										HSC				MLC	SC		
	CL	SL	B2M	B3M	B4M	PCT	RND	MAN	CL	SL	B2M	B3M	B4M	PCT			RND	MAN
<i>20-genes</i>																		
BOWSrf	6.365	6.532	5.639	5.697	5.615	5.799	5.961	5.626	0.386	0.384	0.39	0.385	0.385	0.385	0.389	0.385	8.829	6.687
TFIDFSrf	6.777	6.617	5.944	5.493	5.334	5.82	6.075	5.696	0.37	0.371	0.365	0.373	0.365	0.363	0.372	0.363	8.55	7.126
Struc	6.917	6.791	6.748	6.865	6.708	6.895	6.899	6.681	0.386	0.383	0.391	0.38	0.386	0.387	0.386	0.382	8.829	8.463
Pres	7.149	7.357	6.94	6.712	6.817	6.552	7.063	6.831	0.802	0.853	0.708	0.827	0.853	0.88	0.784	0.636	8.829	7.258
Context	7.381	7.674	6.858	6.684	6.927	6.526	7.171	6.747	2.774	2.765	2.733	2.814	2.717	2.792	2.683	2.718	8.185	7.321
PV_R	7.024	7.221	7.124	7.122	6.924	7.146	6.988	7.117	0.365	0.365	0.364	0.363	0.365	0.363	0.366	0.36	8.865	9.024
PV_C	7.011	7.156	6.6	6.554	6.562	6.632	6.787	6.329	0.359	0.358	0.363	0.361	0.358	0.36	0.362	0.361	8.865	8.747
PV_GLR	7.831	7.844	7.342	7.376	6.976	7.135	7.607	7.021	0.369	0.368	0.372	0.367	0.37	0.37	0.375	0.371	8.865	9.625
PV_GLC	6.585	6.674	6.546	6.501	6.425	6.499	6.702	6.514	0.366	0.364	0.365	0.364	0.361	0.362	0.366	0.366	8.865	9.214
Ngrams	6.353	6.209	6.053	6.094	5.971	6.174	6.526	5.837	0.395	0.397	0.387	0.39	0.393	0.393	0.395	0.392	8.865	7.076
<b>SPIRIT</b>																		
BOWSrf	2.45	2.427	2.518	2.487	2.448	2.485	2.644	2.462	0.591	0.59	0.591	0.589	0.589	0.589	0.606	0.601	2.577	4.739
TFIDFSrf	2.462	2.644	2.618	2.53	2.56	2.618	2.644	2.533	0.595	0.594	0.599	0.601	0.599	0.594	0.594	0.593	2.577	4.923
Struc	2.427	2.499	2.482	2.353	2.393	2.389	2.611	2.421	0.592	0.589	0.595	0.595	0.594	0.589	0.592	0.587	2.485	4.718
Pres	2.628	2.596	2.603	2.599	2.61	2.586	2.675	2.65	0.828	0.835	0.809	0.839	0.811	0.844	0.789	0.911	2.672	3.315
Context	2.651	2.668	2.647	2.665	2.655	2.64	2.674	2.687	1.504	1.519	1.525	1.508	1.545	1.579	1.513	1.6	2.711	2.915
PV_R	2.936	2.773	2.652	2.609	2.623	2.672	2.668	2.648	0.587	0.585	0.588	0.589	0.587	0.586	0.593	0.588	2.691	4.976
PV_C	2.742	2.875	2.559	2.611	2.586	2.624	2.603	2.62	0.589	0.59	0.591	0.588	0.588	0.591	0.591	0.6	2.643	5.896
PV_GLR	2.833	2.857	2.733	2.732	2.723	2.723	2.723	2.735	0.589	0.585	0.589	0.589	0.585	0.584	0.589	0.588	2.726	5.454
PV_GLC	2.925	2.77	2.495	2.484	2.474	2.511	2.601	2.475	0.581	0.584	0.587	0.588	0.586	0.584	0.59	0.585	2.672	5.855
Ngrams	2.662	2.64	2.583	2.543	2.579	2.597	2.565	2.604	0.612	0.604	0.616	0.609	0.608	0.604	0.606	0.6	2.709	5.811





Table 16: The performance of the different machine learning tasks applied on the different features using single tree models. The hierarchy construction method are abbreviated as follows: balanced  $k$ -means clustering (BkM), predictive clustering tree (PCT), clustering with complete linkage (CL), clustering with single linkage (SL), random (RND) and manual (MAN). The evaluation measure is *average precision* where larger value means better result.

20-genes	HMC										HSC				MLC	SC		
	CL	SL	B2M	B3M	B4M	PCT	RND	MAN	CL	SL	B2M	B3M	B4M	PCT			RND	MAN
	BOWSrf	0.404	0.402	0.444	0.45	0.453	0.446	0.424	0.451	0.986	0.986	0.984	0.986	0.986			0.986	0.984
TFIDFSrf	0.392	0.41	0.454	0.473	0.478	0.459	0.423	0.464	0.99	0.99	0.992	0.989	0.992	0.992	0.989	0.993		
Struc	0.349	0.354	0.378	0.375	0.383	0.367	0.358	0.379	0.985	0.986	0.983	0.986	0.985	0.985	0.985	0.986		
Pres	0.34	0.327	0.362	0.375	0.365	0.378	0.355	0.366	0.881	0.873	0.9	0.881	0.876	0.871	0.886	0.917		
Context	0.344	0.33	0.395	0.398	0.387	0.407	0.364	0.398	0.648	0.65	0.647	0.646	0.648	0.648	0.656	0.648		
PV_R	0.341	0.338	0.345	0.348	0.35	0.343	0.344	0.344	0.993	0.993	0.993	0.993	0.993	0.993	0.992	0.994		
PV_C	0.347	0.335	0.356	0.362	0.363	0.367	0.355	0.377	0.994	0.994	0.992	0.994	0.995	0.994	0.993	0.993		
PV_GLR	0.309	0.31	0.323	0.325	0.331	0.329	0.319	0.329	0.99	0.99	0.989	0.991	0.99	0.99	0.989	0.989		
PV_GLC	0.35	0.351	0.353	0.359	0.361	0.359	0.353	0.352	0.992	0.993	0.992	0.993	0.994	0.993	0.992	0.991		
Ngrams	0.393	0.398	0.421	0.41	0.42	0.413	0.379	0.43	0.982	0.982	0.984	0.983	0.983	0.983	0.981	0.982		
<b>SPIRIT</b>																		
BOWSrf	0.716	0.718	0.707	0.713	0.716	0.715	0.697	0.713	0.994	0.995	0.994	0.994	0.995	0.995	0.99	0.991		
TFIDFSrf	0.715	0.705	0.702	0.71	0.709	0.707	0.697	0.71	0.994	0.994	0.993	0.991	0.992	0.994	0.994	0.994		
Struc	0.705	0.7	0.692	0.71	0.696	0.696	0.681	0.693	0.994	0.995	0.993	0.993	0.994	0.995	0.994	0.996		
Pres	0.679	0.689	0.69	0.69	0.682	0.684	0.678	0.682	0.929	0.927	0.938	0.932	0.932	0.926	0.946	0.914		
Context	0.659	0.658	0.661	0.66	0.661	0.661	0.662	0.662	0.803	0.804	0.802	0.802	0.797	0.797	0.796	0.777		
PV_R	0.658	0.658	0.66	0.68	0.676	0.676	0.656	0.667	0.996	0.997	0.996	0.996	0.997	0.997	0.995	0.997		
PV_C	0.664	0.654	0.674	0.672	0.669	0.669	0.668	0.67	0.996	0.996	0.996	0.996	0.996	0.995	0.995	0.993		
PV_GLR	0.641	0.642	0.653	0.652	0.658	0.658	0.653	0.653	0.995	0.996	0.995	0.995	0.996	0.996	0.996	0.995		
PV_GLC	0.642	0.648	0.675	0.675	0.676	0.674	0.663	0.676	0.997	0.996	0.996	0.996	0.997	0.996	0.996	0.996		
Ngrams	0.66	0.66	0.661	0.659	0.662	0.661	0.663	0.66	0.988	0.99	0.986	0.989	0.989	0.99	0.989	0.99		