

Additional Table 2. Average AUROC across terms and average F_{max} , Precision and Recall across genes of HTD and TPR ensemble variants using RANKS and SVMs as base learner and the UA integrated network. Results are estimated through 5-fold cross-validation. For each sub-ontology and each metric best results are highlighted in bold. Results significantly better than all the others methods according to the Wilcoxon Rank Sum test ($\alpha = 10^{-6}$) are underlined.

Subontology	Method	AUROC	F_{max}	Precision	Recall
Organ	RANKS	0.8493	0.3106	0.2407	0.4377
	SVM	0.7128	0.1668	0.1688	0.1648
	HTD-RANKS	0.8446	0.3411	0.2717	0.4583
	HTD-SVM	0.8328	0.3155	0.2370	0.4718
	TPR-T-RANKS	0.8646	0.3575	0.2933	0.4586
	TPR-T-SVM	0.8240	0.3155	0.2370	0.4718
	TPR-D-RANKS	0.8408	<u>0.3773</u>	<u>0.3231</u>	0.4538
	TPR-D-SVM	0.7883	0.3209	0.2800	0.3782
	TPR-TF-RANKS	0.8420	0.3547	0.2880	0.4615
	TPR-TF-SVM	0.7060	0.2359	0.2074	0.2736
Inheritance	TPR-W-RANKS	0.8377	0.3620	0.3025	0.4515
	TPR-W-SVM	0.8238	0.3171	0.2343	0.4916
	RANKS	0.8715	0.5986	0.4709	0.8215
	SVM	0.7637	0.1668	0.1688	0.1648
	HTD-RANKS	0.8846	0.6034	0.4709	0.8396
	HTD-SVM	0.7991	0.5429	0.3885	0.9008
	TPR-T-RANKS	0.8708	0.6066	0.4795	0.8276
	TPR-T-SVM	0.7909	0.5444	0.3892	0.9063
	TPR-D-RANKS	0.8659	0.6064	0.4793	0.8278
	TPR-D-SVM	0.7919	0.5441	0.3889	0.9063
Onset	TPR-TF-RANKS	0.8675	0.6029	0.4757	0.8228
	TPR-TF-SVM	0.7546	0.4480	0.2972	0.9099
	TPR-W-RANKS	0.8636	<u>0.6070</u>	0.4769	0.8355
	TPR-W-SVM	0.7900	0.5448	0.3897	0.9059
	RANKS	0.8159	0.4494	0.3821	0.5455
	SVM	0.7025	0.3978	0.2717	0.7421
	HTD-RANKS	0.8253	0.4466	0.3563	0.5981
	HTD-SVM	0.7781	0.4492	0.3374	0.6719
	TPR-T-RANKS	0.8249	0.4631	0.3791	0.6021
	TPR-T-SVM	0.7511	0.4517	0.3385	0.6813