Feature	AUROC	p-AUROC	p-value
GaDmPmD	0.643	0.547	1.6×10^{-20}
GiGaD	0.558	0.514	2.1×10^{-9}
{Perturbation}	0.740	0.667	2.3×10^{-7}
GeTlD	0.573	0.518	1.8×10^{-5}
{KEGG}	0.613	0.566	5.6×10^{-5}
GaDaGaD	0.633	0.592	1.5×10^{-4}
{BioCarta}	0.548	0.526	0.001
{Reactome}	0.599	0.562	0.002
{Immunologic}	0.703	0.665	0.006
GiGiGaD	0.646	0.621	0.05
{Cancer Module}	0.629	0.612	0.11
GeTeGaD	0.570	0.554	0.14
{TF Target}	0.612	0.596	0.16
{GO Component}	0.560	0.547	0.17
{Positional}	0.529	0.520	0.19
GiGeTlD	0.628	0.616	0.20
{GO Process}	0.626	0.617	0.26
GaD (any disease)	0.683	0.676	0.30
{GO Function}	0.577	0.571	0.31
{Cancer Hood}	0.533	0.532	0.45
GaD (any gene)	0.620	0.620	0.49
GaDlTlD	0.674	0.674	0.49
{Oncogenic}	0.601	0.604	0.60
{miRNA Target}	0.562	0.569	0.72