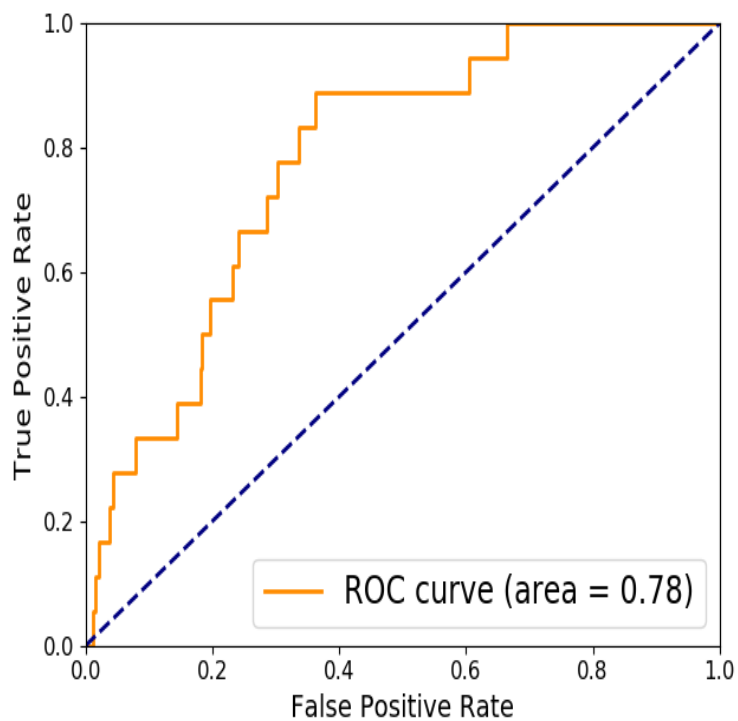
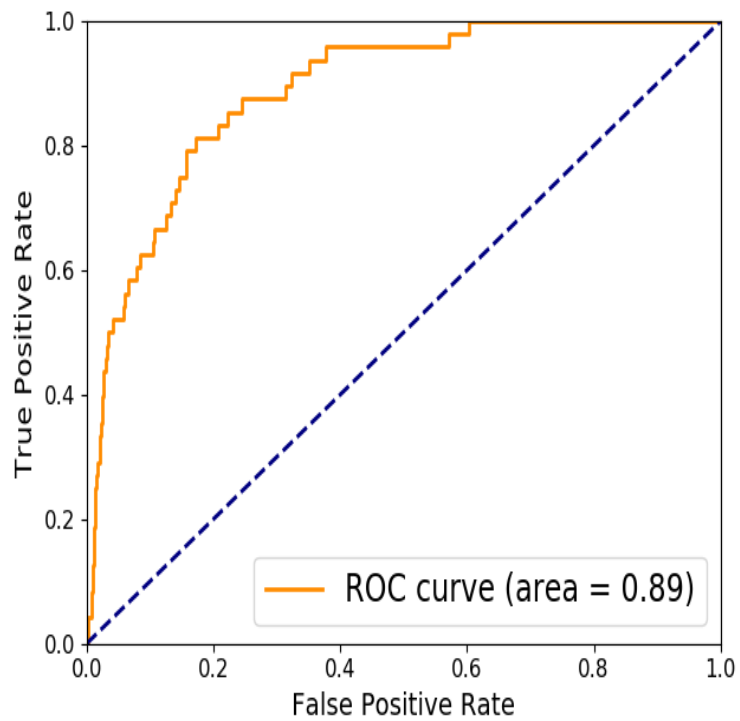


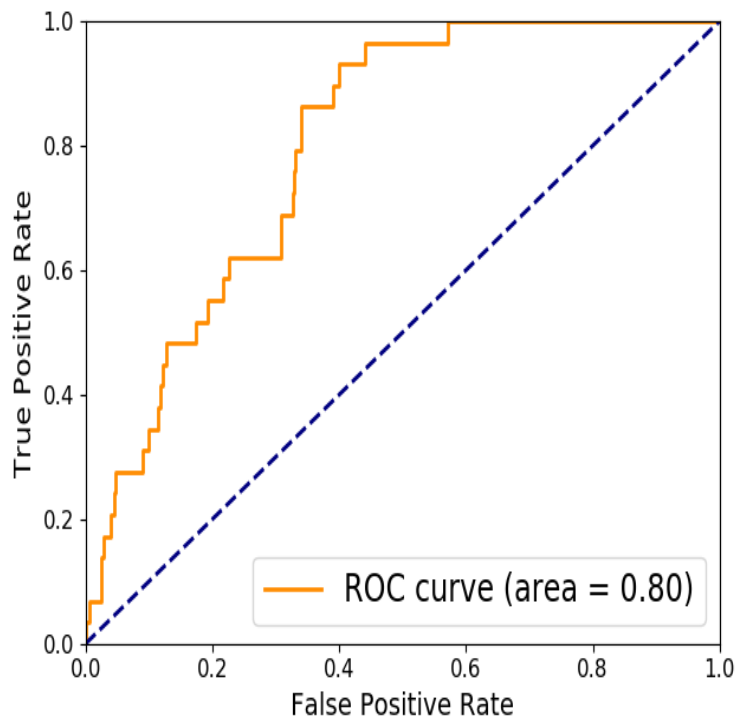
## Diffusion ROC from fGOnly\_tOverlap STRING10\_combined.png



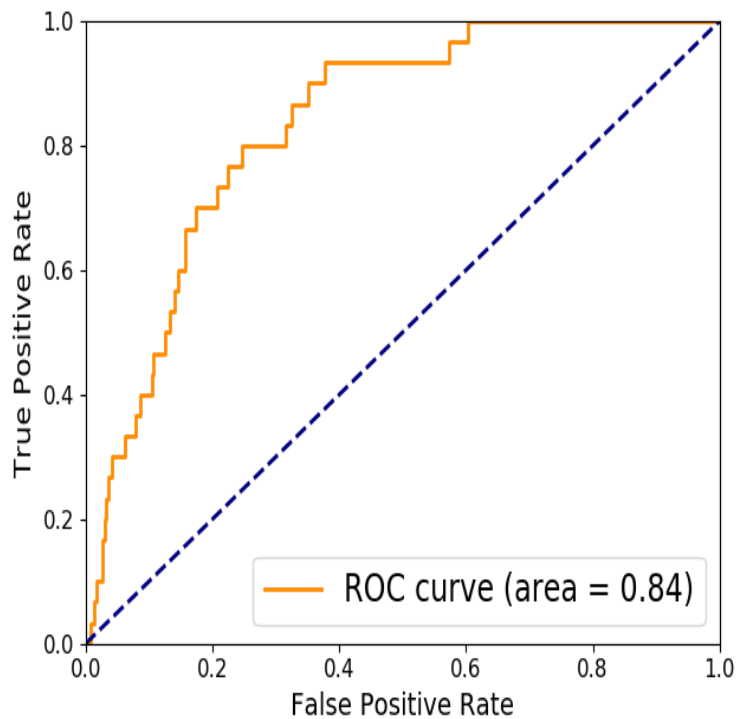
## Diffusion ROC from fTestOnly\_tGS STRING10\_combined.png



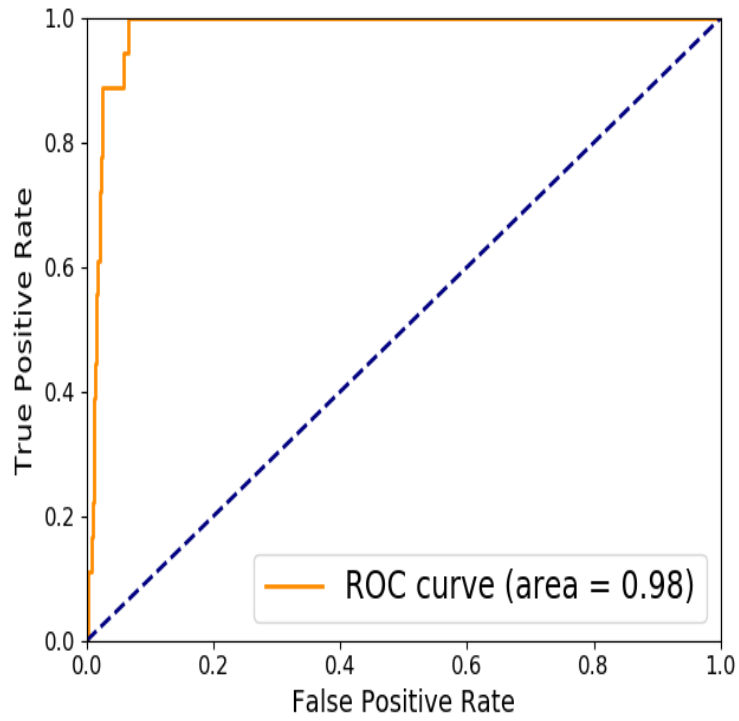
# Diffusion ROC from fGOnly\_tTestOnly STRING10\_combined.png



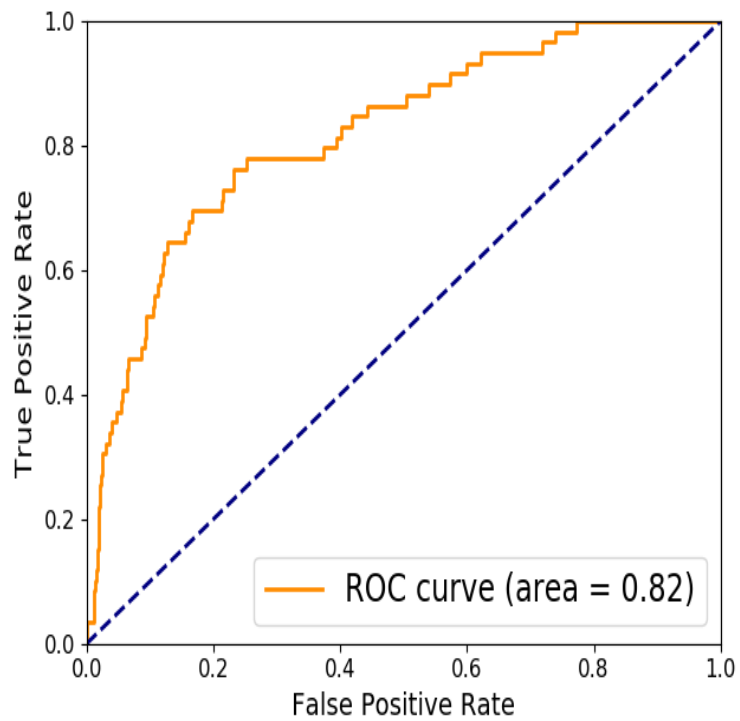
# Diffusion ROC from fTestOnly\_tGOnly STRING10\_combined.png



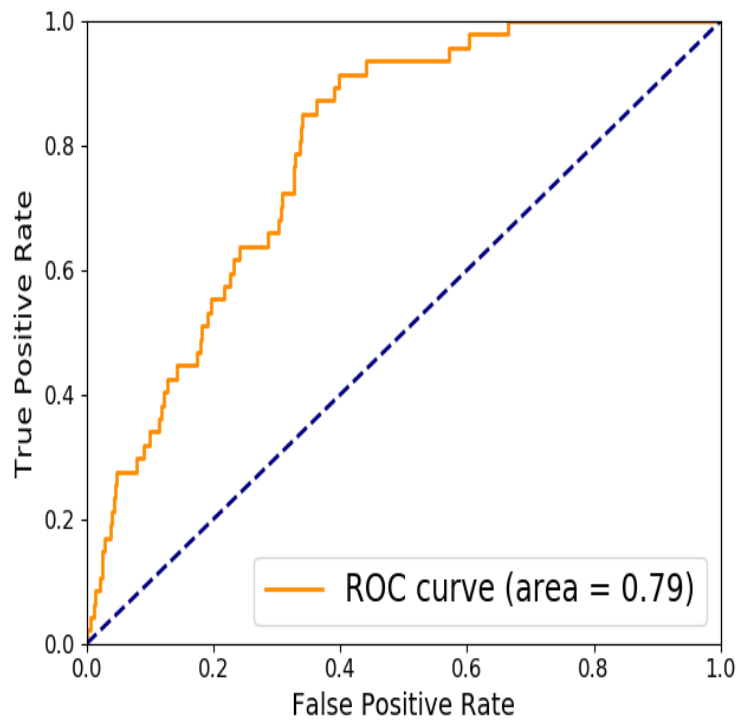
# Diffusion ROC from fTestOnly\_tOverlap STRING10\_combined.png



## Diffusion ROC from fOverlap\_tOnlies STRING10\_combined.png



## Diffusion ROC from fGOnly\_tTest STRING10\_combined.png



Diff_FROM	Diff_TO	AUC	Zt_wdegre	Zt	Zf_wdegre	Zf
		e		e		
GS_only	Test_only	0.801	5.030	5.958	2.3629795	2.686
				360081474		
				(invalid)		
Test_only	GS_only	0.836	6.789	7.123	3.762	3.469
GS_only	Overlap	0.781	2.957	4.655	1.3037320	1.545282
				975074722	09954217	
				(invalid)	23 (inva	
				lid)		
GS_only	Test	0.794	5.647	7.363	1.880	2.210
Test_only	Overlap	0.980	6.119	7.424	4.088	3.759700
				61743522		
				17 (inva		
				lid)		
Test_only	GS	0.890	8.994	9.3820049	6.135	5.278
				50839459		
				(invalid)		
Overlap	Onlies	0.820	8.654	7.779	3.435	3.675

**	#Mapped	#Total	Not mapped genes
GS	30	30	MT-RNR2;MT-RNR1   N/A
Test_only	29	30	MT-RNR2;MT-RNR1;MTATP6P1   N/A
Overlap	18	20	MT-RNR2;MT-RNR1   RPS11;MT-ND2;MT-ND4   ;MT-ATP6;MT-ND4L;MT-RNR1;MT-CYB;MT-CO2   ;MT-CO1;MT-ATP8;MT-CO3;RPS12;MT-RNR2;TMS   B4X;ACTB;FTL;FTH1;MT-ND3;TMSB10;MT-ND1

(end of file)