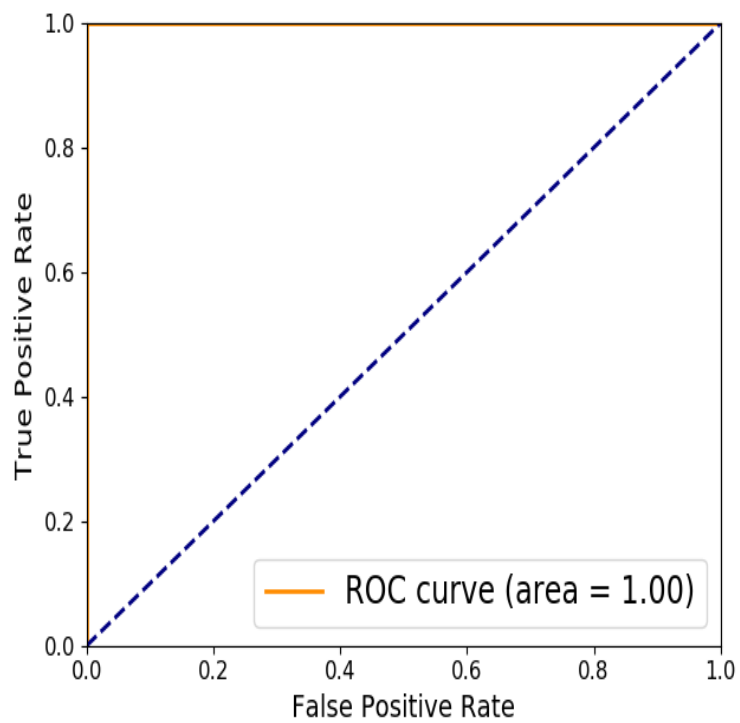
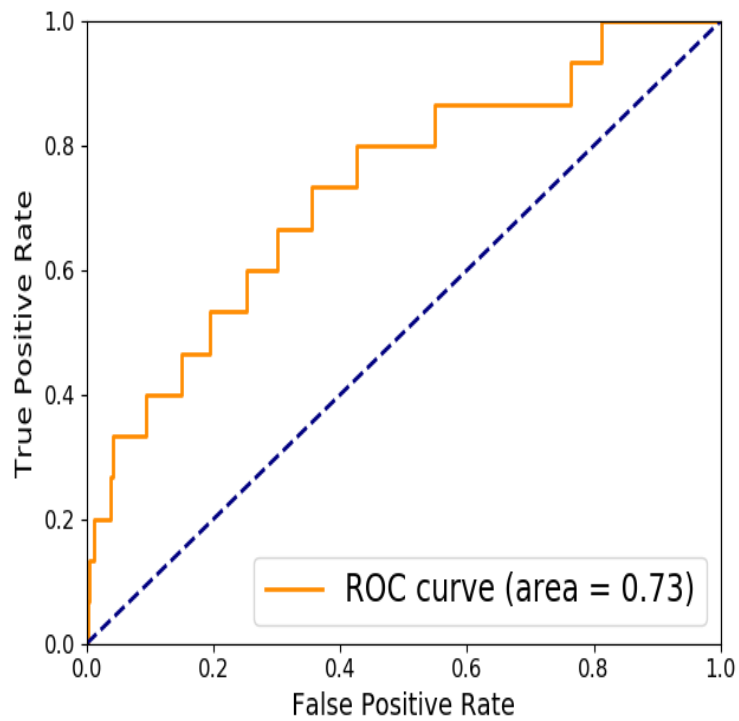


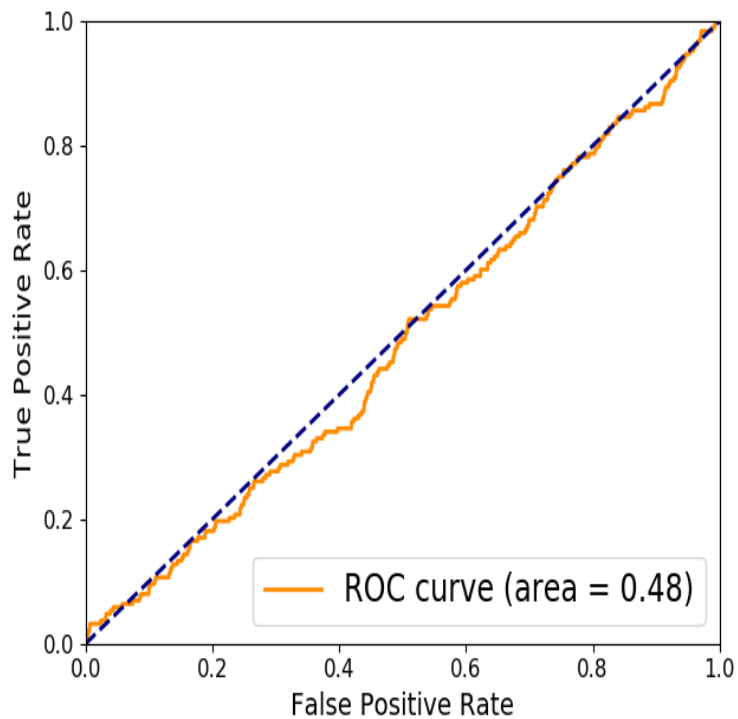
## Diffusion ROC from fGOnly\_tOverlap STRING10\_combined.png



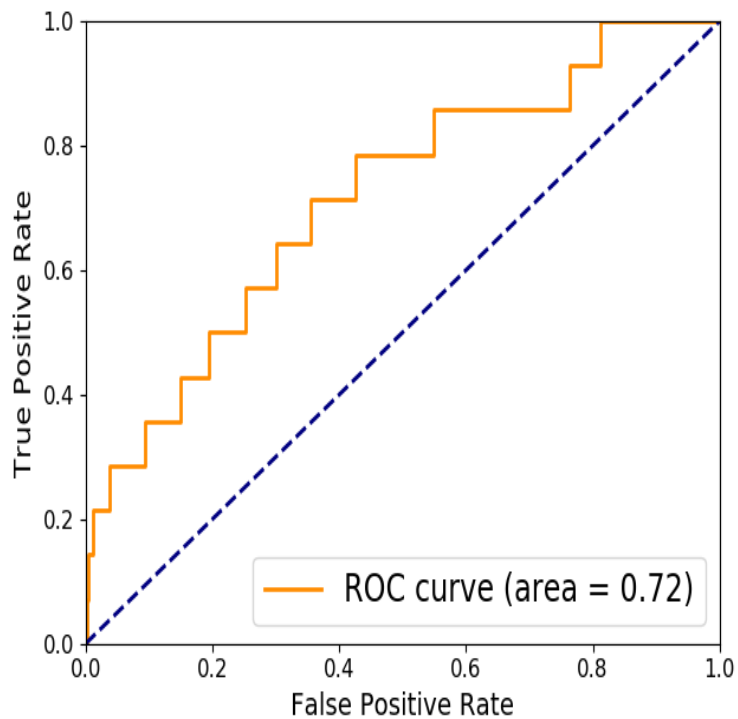
# Diffusion ROC from fTestOnly\_tGS STRING10\_combined.png



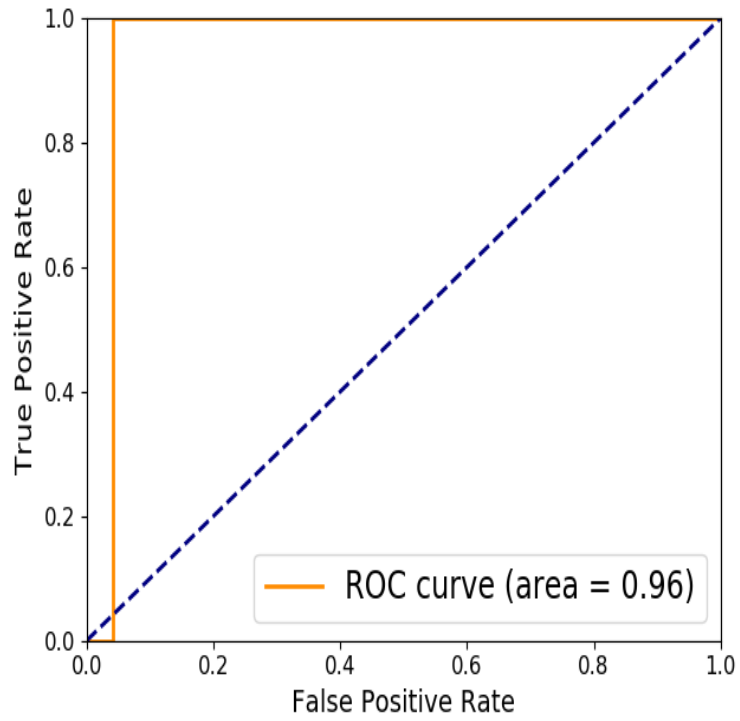
# Diffusion ROC from fGOnly\_tTestOnly STRING10\_combined.png



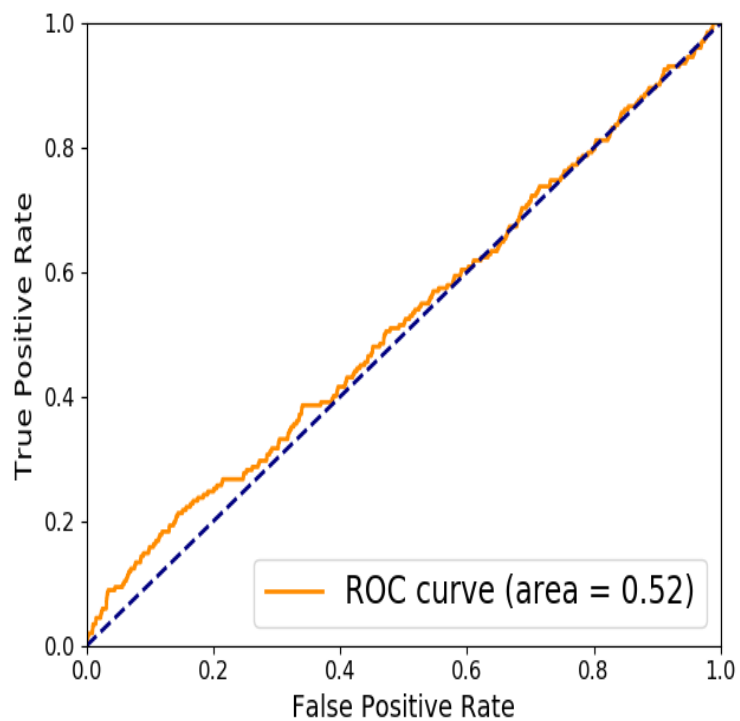
# Diffusion ROC from fTestOnly\_tGOnly STRING10\_combined.png



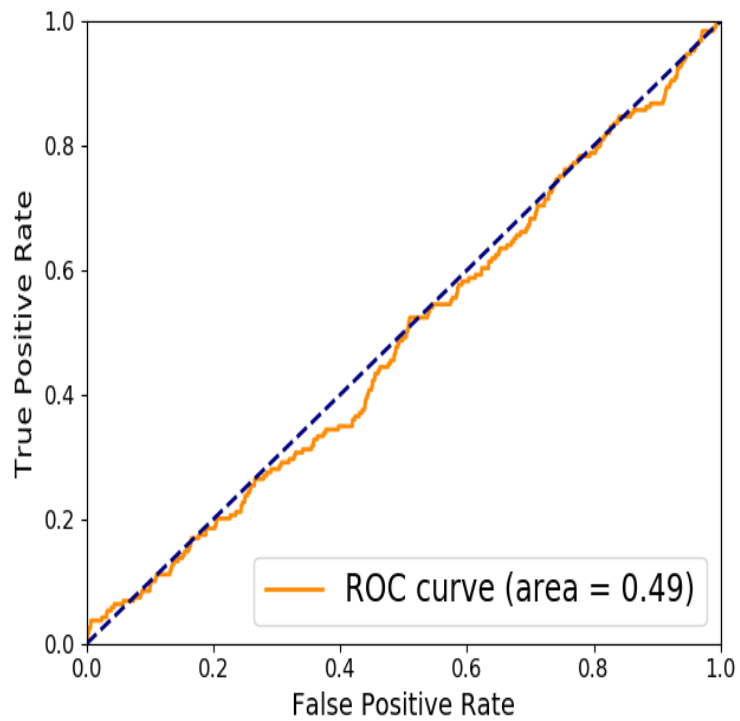
# Diffusion ROC from fTestOnly\_tOverlap STRING10\_combined.png



## 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	0.483	-1.427	-0.816	-1.326	-0.173
Test_only	GS_only	0.718	0.156	2.753	0.108	0.149419
				64711027		
				203 (inv		
				alid)		
GS_only	Overlap	0.999	1.0132336	1.6342291	1.8727077	1.921
			815639735	423434125	626876822	
			(invalid)	(invalid)	(invalid)	
GS_only	Test	0.486	-1.579	-0.576	-1.099	-0.388
Test_only	Overlap	0.959	1.3208668	1.4937335	0.8526299	0.630723
			972606958	720257119	06018727	03139154
			(invalid)	(invalid)	(invalid)	04 (inva
					lid)	
Test_only	GS	0.734	0.147	3.041	0.3375079	-0.035
				072532063		
				4		
				(invalid)		
Overlap	Onlies	0.520	0.090	1.201	9.4862688	0.777
				88026897		
				(invalid)		

**	#Mapped	#Total	Not mapped genes
GS	14	15	CAMK1V   N/A
Test_only	188	199	MICU2;KMT2C;RDH13;LY6G5B;DRC1;OR8G5;KR   N/A
			TAP10.9;KMT2B;43722;KMT2E;PRR3
Overlap	1	1	APOE

(end of file)