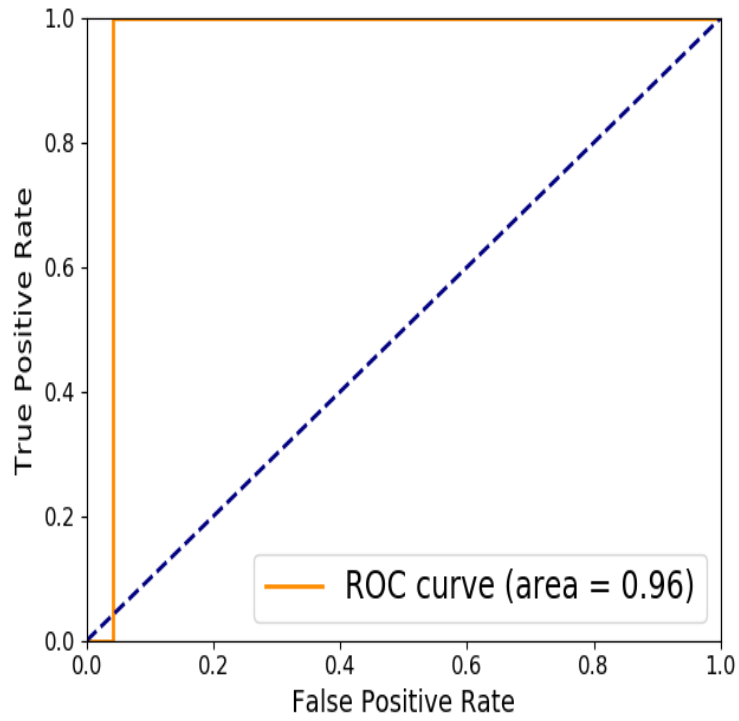
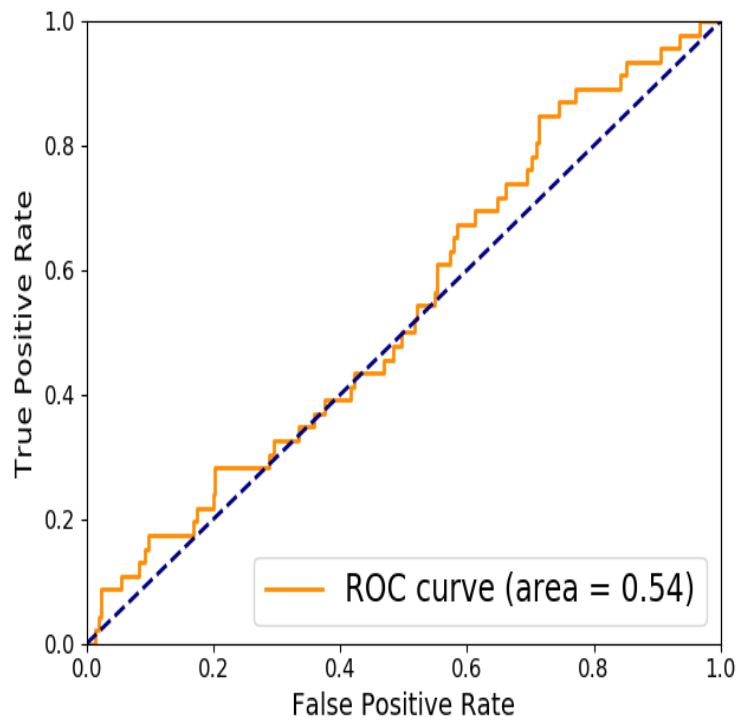


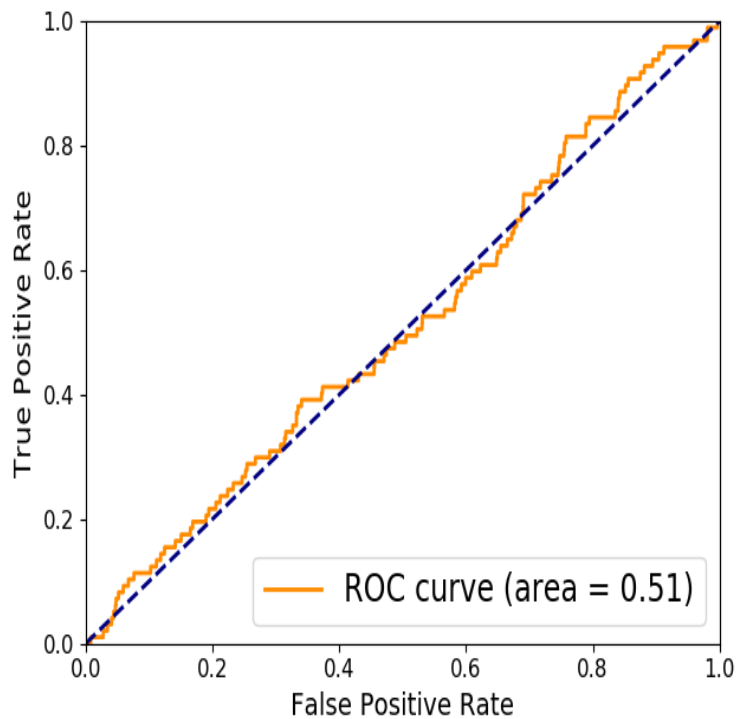
# Diffusion ROC from fGOnly\_tOverlap STRING10\_combined.png



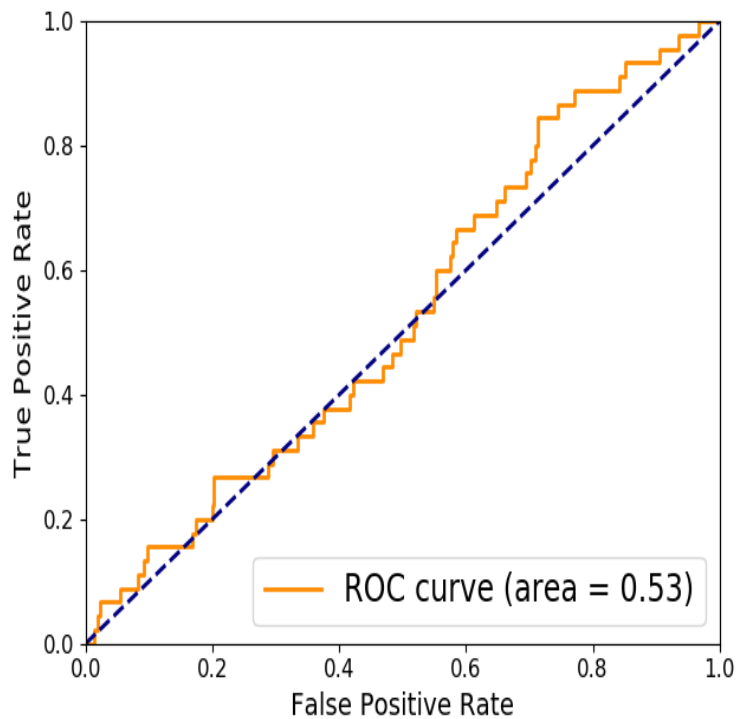
# Diffusion ROC from fTestOnly\_tGS STRING10\_combined.png



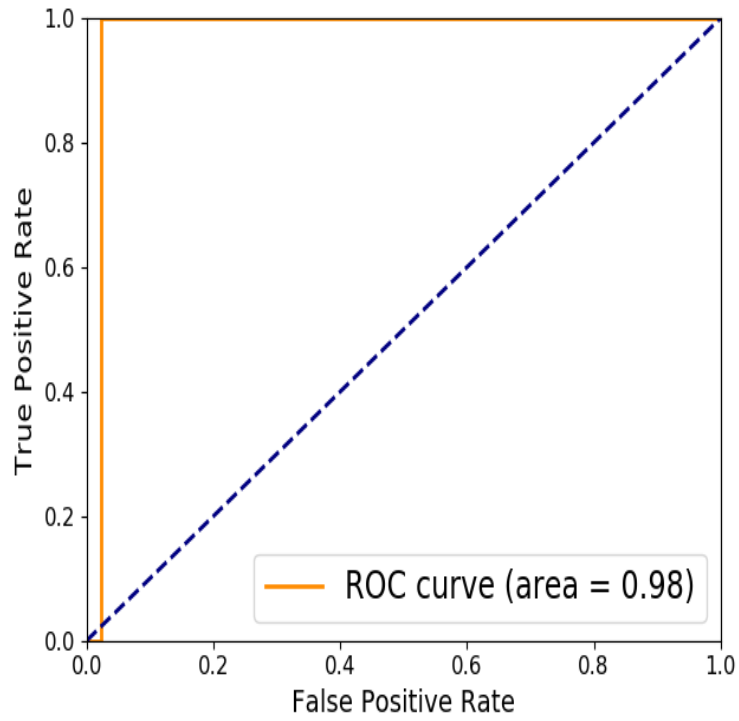
# Diffusion ROC from fGOnly\_tTestOnly STRING10\_combined.png



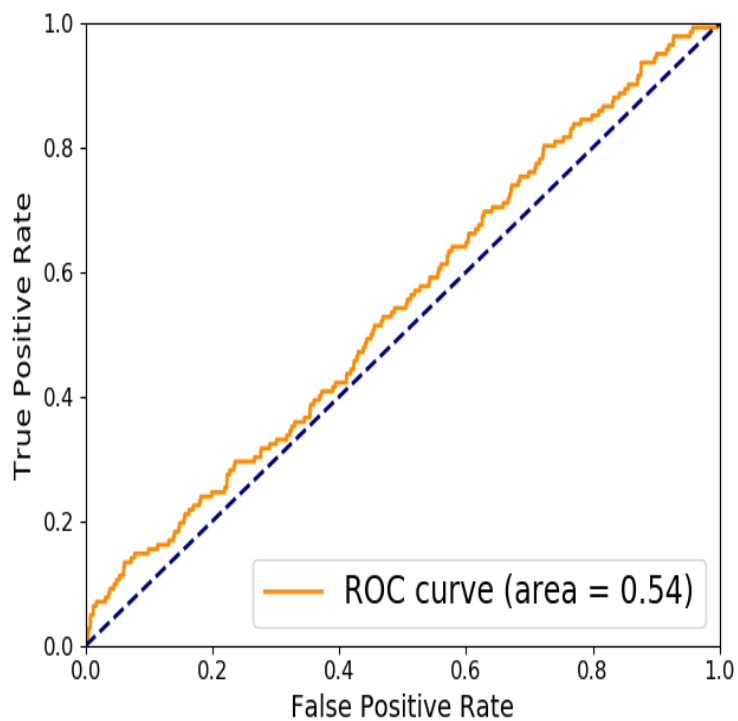
# Diffusion ROC from fTestOnly\_tGOnly STRING10\_combined.png



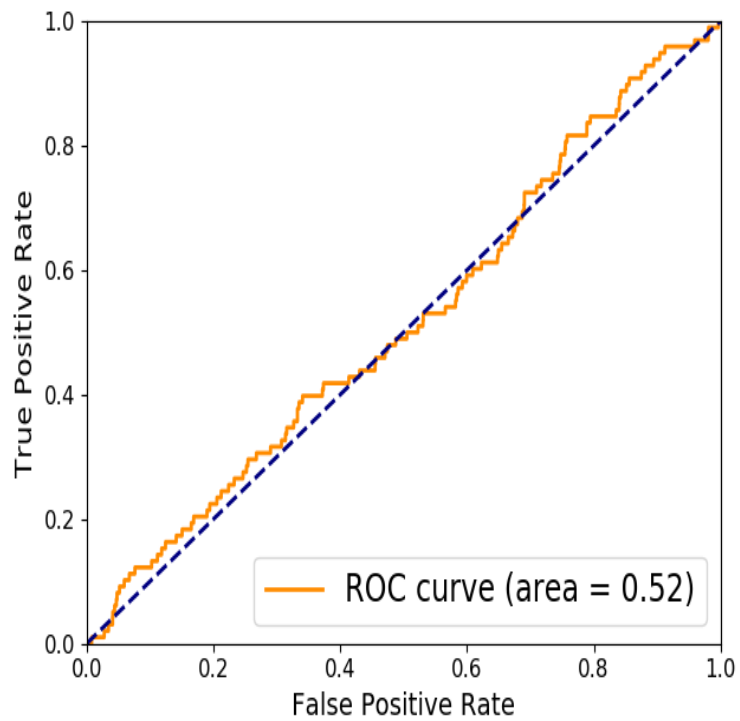
# Diffusion ROC from fTestOnly\_tOverlap STRING10\_combined.png



## 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	0.511	-1.532	0.267	-0.716140	-0.976
				143160088		
				2		
				(invalid)		
Test_only	GS_only	0.530	-0.213	0.765	-1.392	-0.632
GS_only	Overlap	0.959	2.2541142	1.7156541	1.3291460	1.863814
			80362898	77379956	210612935	84608052
			(invalid)	(invalid)	(invalid)	04 (inva
					lid)	
GS_only	Test	0.516	-1.480	0.499	-0.436	-0.460
Test_only	Overlap	0.978	2.6499925	1.8968118	1.9003717	1.828506
			89795067	509254506	417625645	74605585
			(invalid)	(invalid)	(invalid)	4 (inval
					id)	
Test_only	GS	0.539	0.033	1.069	-0.943	-0.350
Overlap	Onlies	0.544	0.265	1.878	0.168	0.365

**	#Mapped	#Total	Not mapped genes
GS	45	49	DGKK;KDM7A;DPH7;MIEF1   N/A
Test_only	97	99	MROH7;CXorf22   N/A
Overlap	1	1	SLC30A8

(end of file)