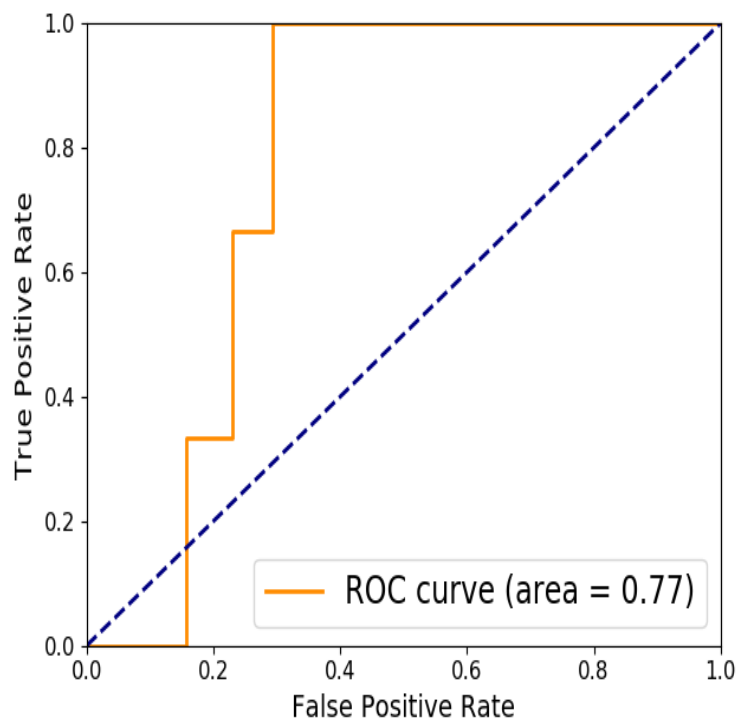
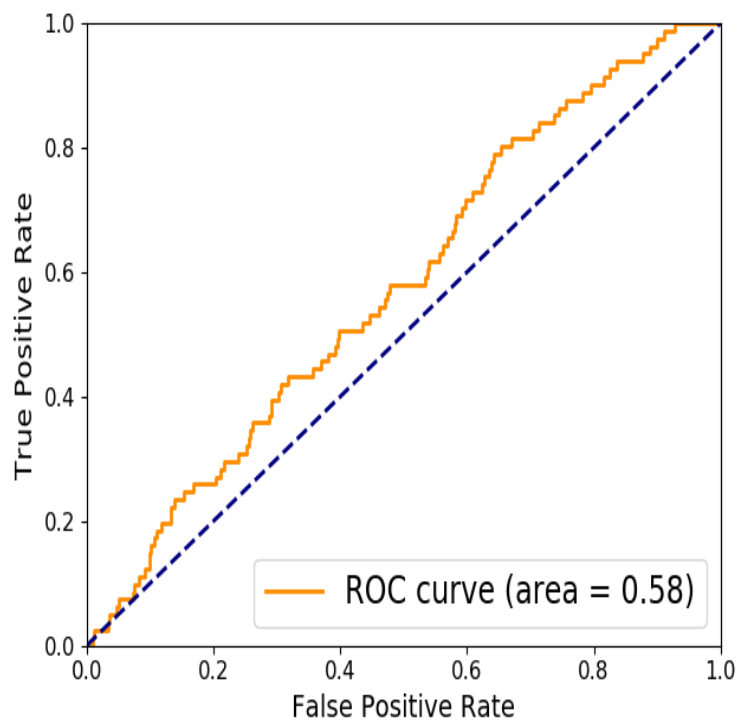


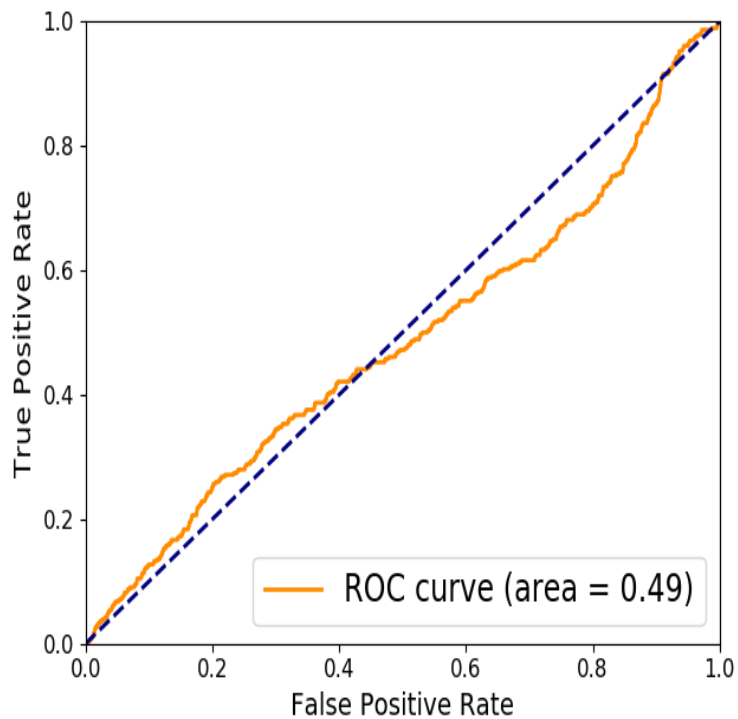
## Diffusion ROC from fGOnly\_tOverlap STRING10\_combined.png



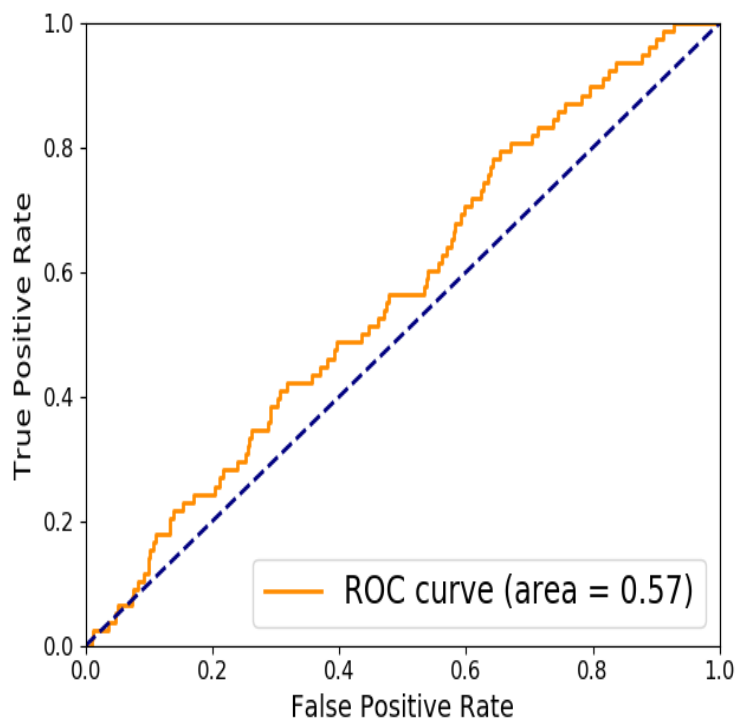
## Diffusion ROC from fTestOnly\_tGS STRING10\_combined.png



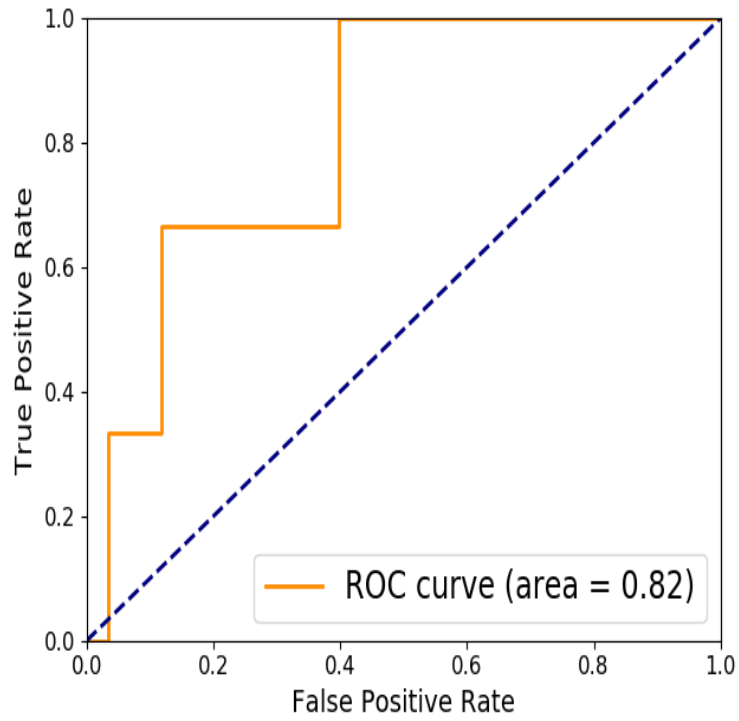
# Diffusion ROC from fGOnly\_tTestOnly STRING10\_combined.png



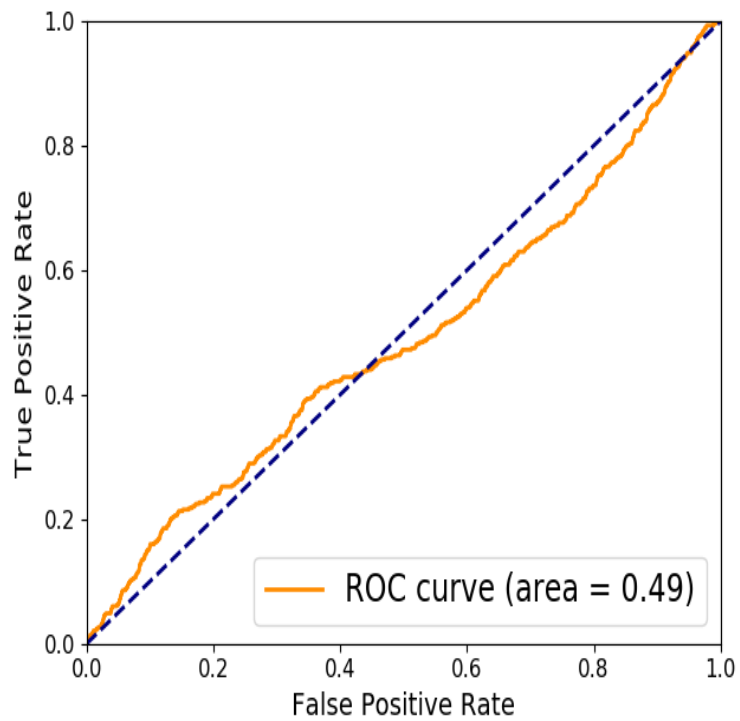
## Diffusion ROC from fTestOnly\_tGOnly STRING10\_combined.png



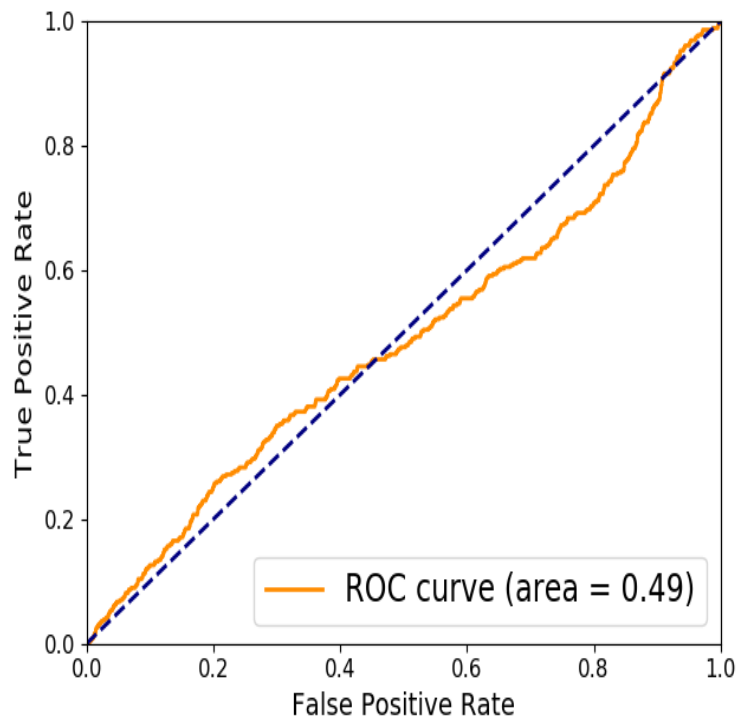
# Diffusion ROC from fTestOnly\_tOverlap STRING10\_combined.png



## 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	0.485	-0.018	-1.198	0.693	0.256709
				77614539		
				73 (inva		
				lid)		
Test_only	GS_only	0.569	0.883	1.9378856	0.325	0.297
				965905857		
				(invalid)		
GS_only	Overlap	0.774	1.792	1.297	2.329	1.860207
					82674318	
					85 (inva	
					lid)	
GS_only	Test	0.488	-0.059	-0.880	0.777	0.234
Test_only	Overlap	0.817	1.7134970	2.076	2.567	2.093
			038521948			
			(invalid)			
Test_only	GS	0.578	1.3663434	2.7121166	0.820	0.787
			672831059	83175867		
			(invalid)	(invalid)		
Overlap	Onlies	0.491	-1.010	-0.528	-0.414	-0.454

**	#Mapped	#Total	Not mapped genes
GS	78	83	RAB29;INTS13;PLPP1;CIPC
			N/A
			;i»¿NPNT
Test_only	354	358	C11orf42;TMEM61;FAM177B
			N/A
			;LY6G5C
Overlap	3	3	EMP3;ZMPSTE24;KIF5B

(end of file)