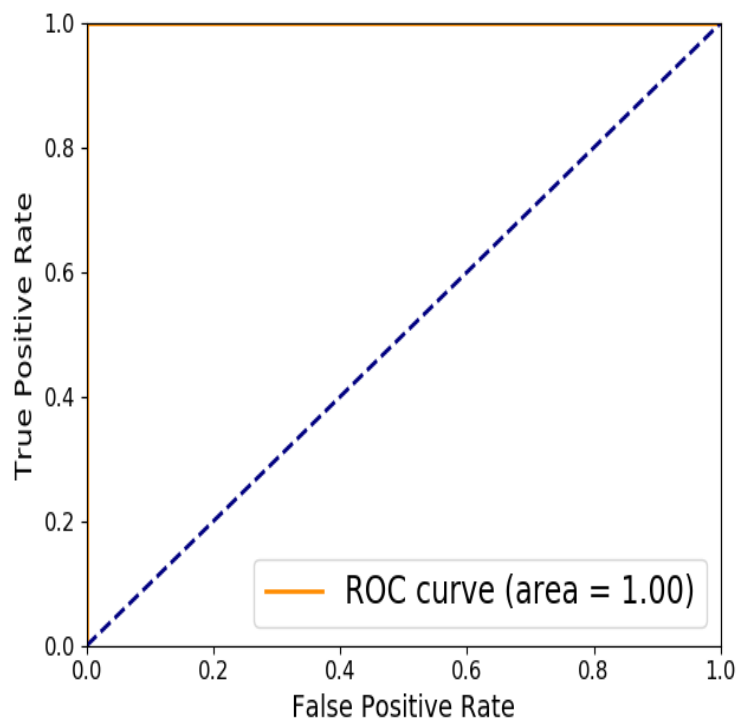
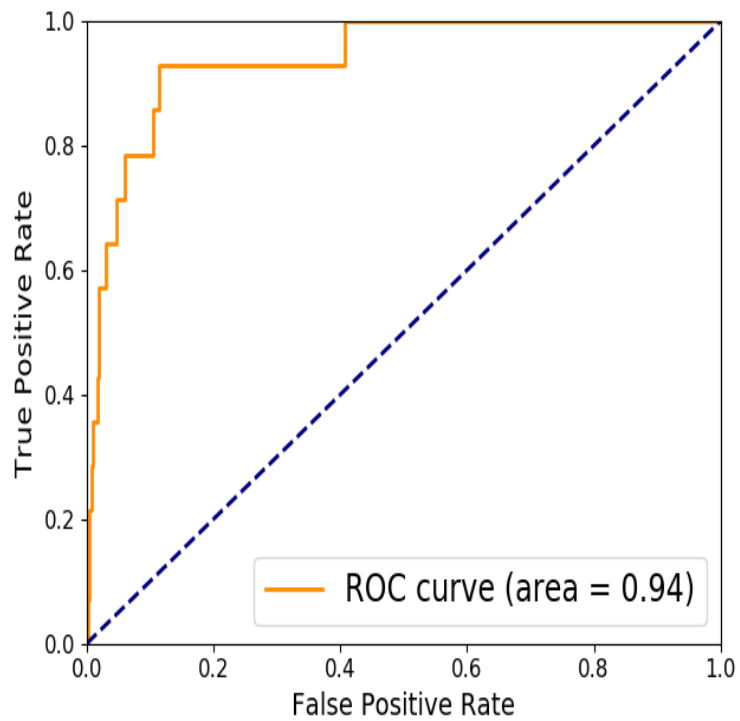


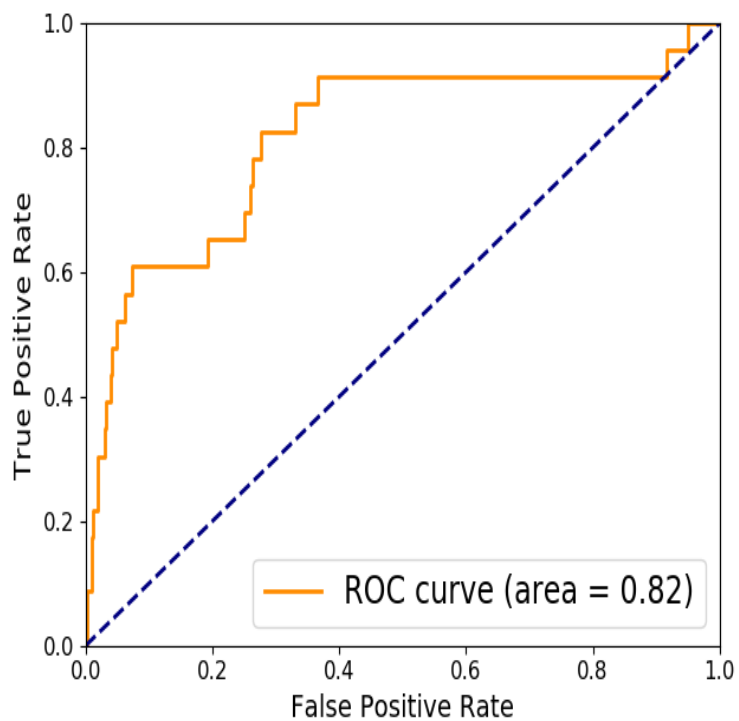
## Diffusion ROC from fGOnly\_tOverlap STRING10\_combined.png



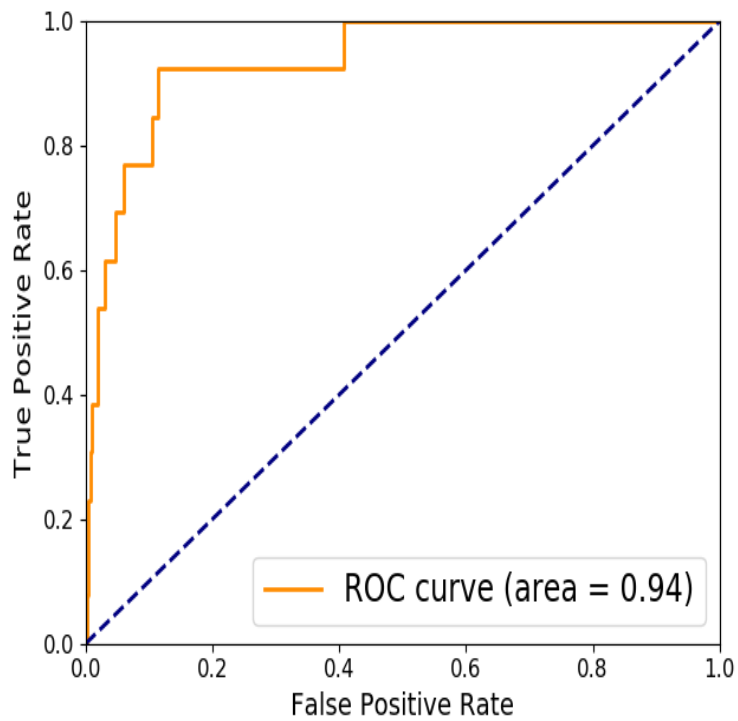
## Diffusion ROC from fTestOnly\_tGS STRING10\_combined.png



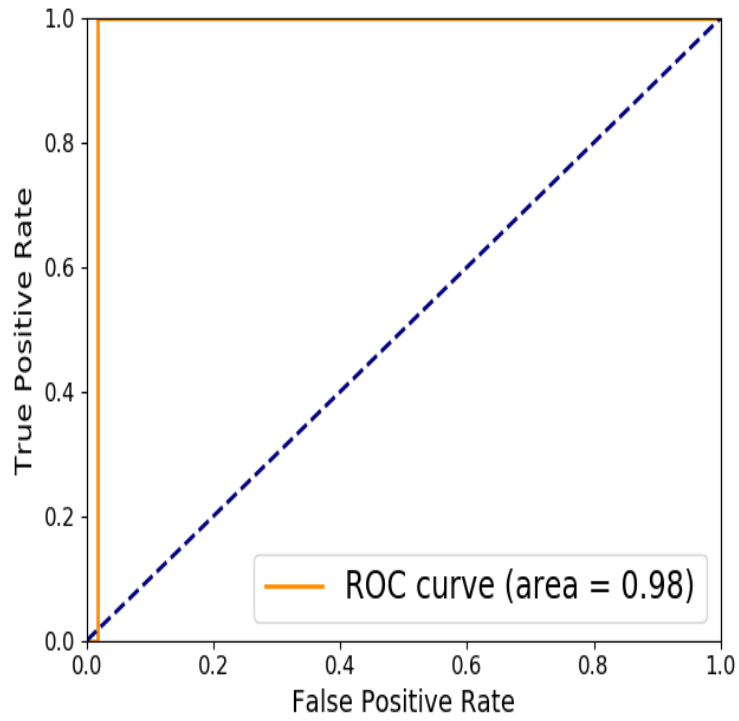
## Diffusion ROC from fGOnly\_tTestOnly STRING10\_combined.png



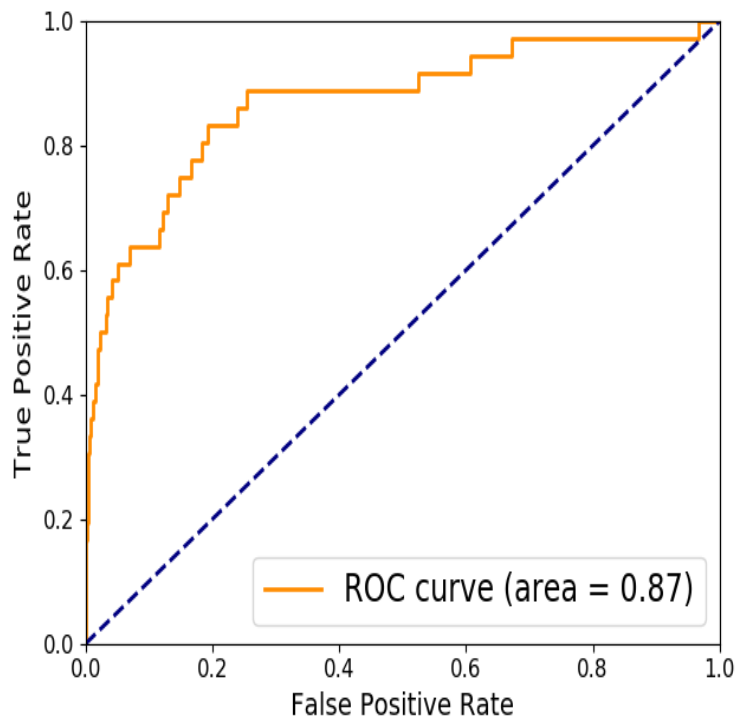
# Diffusion ROC from fTestOnly\_tGOnly STRING10\_combined.png



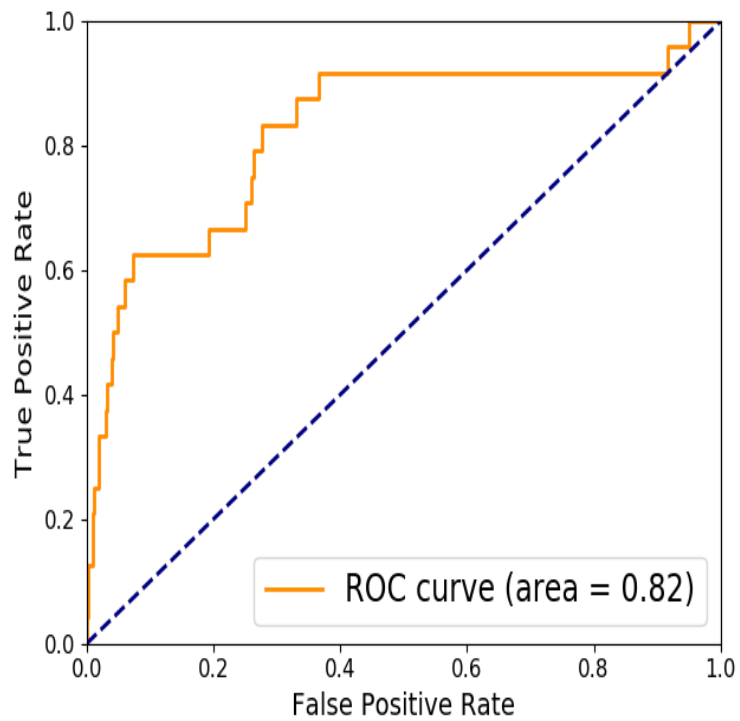
# Diffusion ROC from fTestOnly\_tOverlap STRING10\_combined.png



# 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	0.817	5.294	4.987	4.4448814	3.311
				74270779		
				(invalid)		
Test_only	GS_only	0.936	5.364	5.503	3.8230300	3.535
				79829364		
				(invalid)		
GS_only	Overlap	1.000	1.4398101	1.8319276	2.572	2.204
			151957707	458532034		
			(invalid)	(invalid)		
GS_only	Test	0.825	5.049	5.629	4.922	3.722
Test_only	Overlap	0.982	1.4456659	1.5773526	1.9471834	1.979532
			291635383	664213373	403363417	76406801
			(invalid)	(invalid)	(invalid)	38 (inva
					lid)	
Test_only	GS	0.939	5.124	5.814	3.4171193	3.206
				842004826		
				(invalid)		
Overlap	Onlies	0.870	7.566	7.243	1.9136359	2.564
				177584064		
				(invalid)		

**	#Mapped	#Total	Not mapped genes
GS	13	13	N/A
Test_only	23	23	N/A
Overlap	1	1	ADAM10

(end of file)