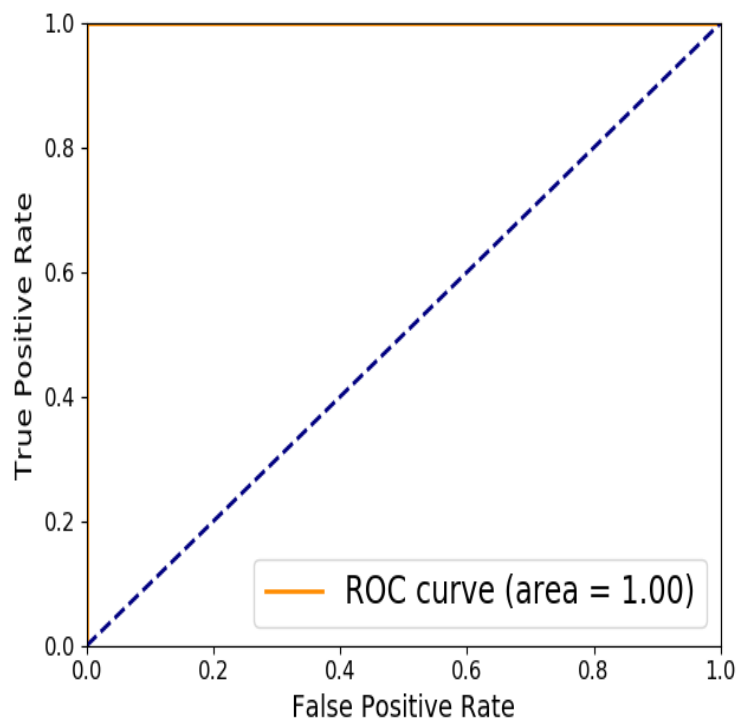
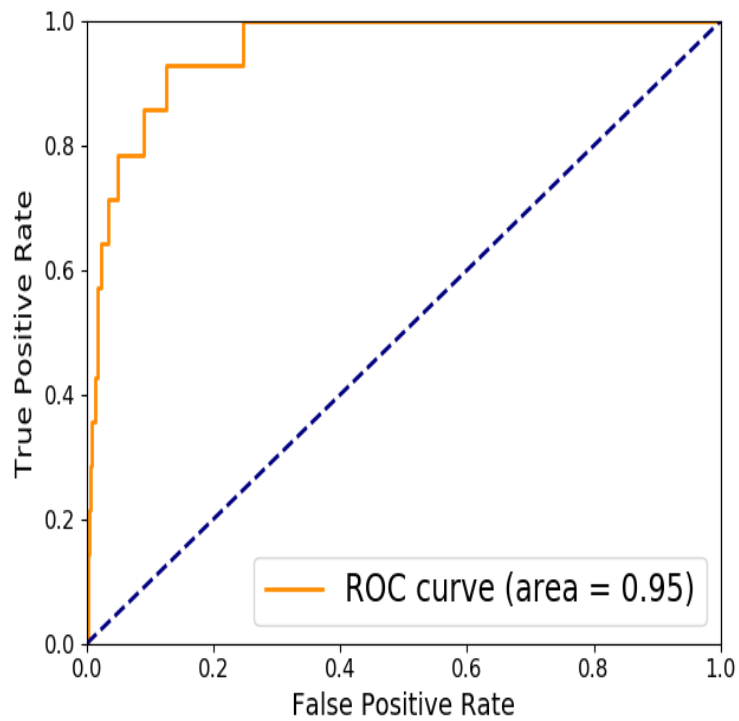


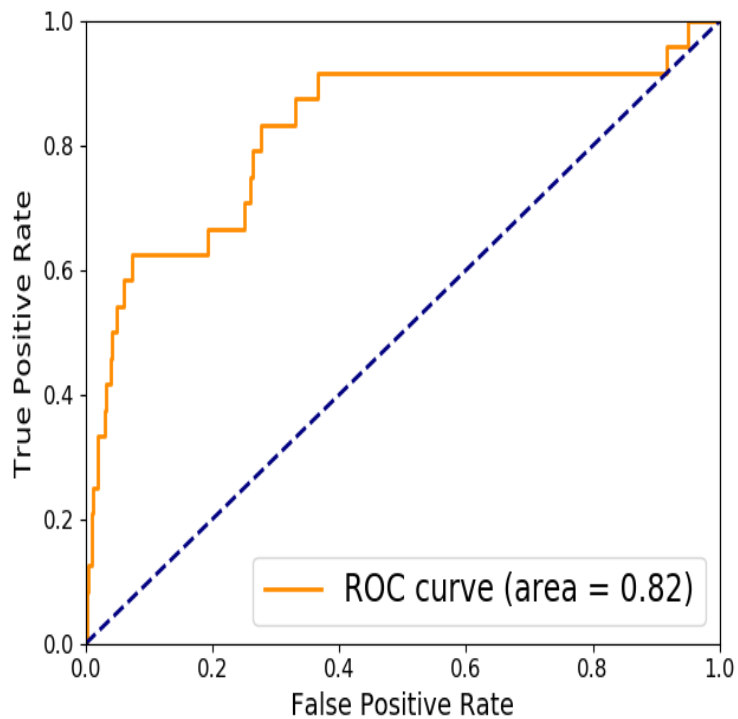
## Diffusion ROC from fGOnly\_tOverlap STRING10\_combined.png



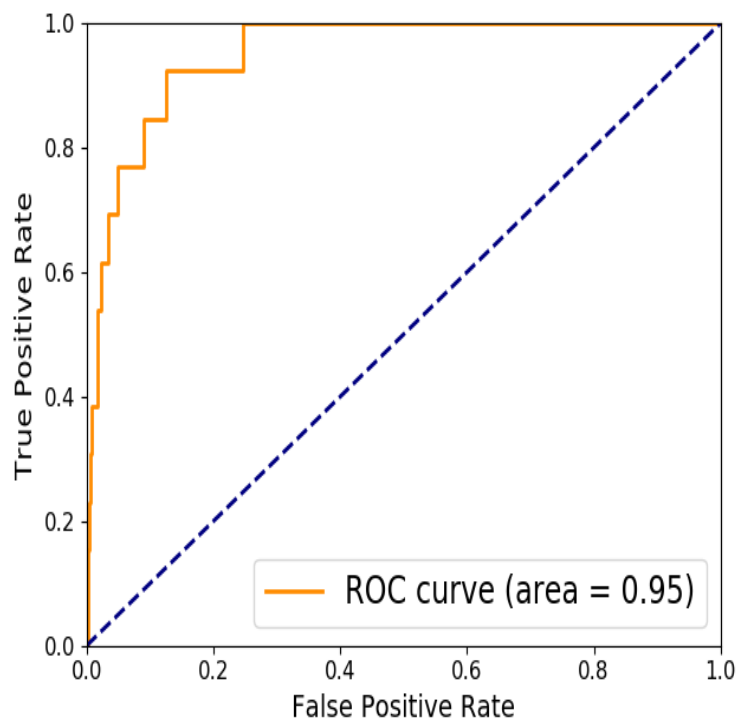
# Diffusion ROC from fTestOnly\_tGS STRING10\_combined.png



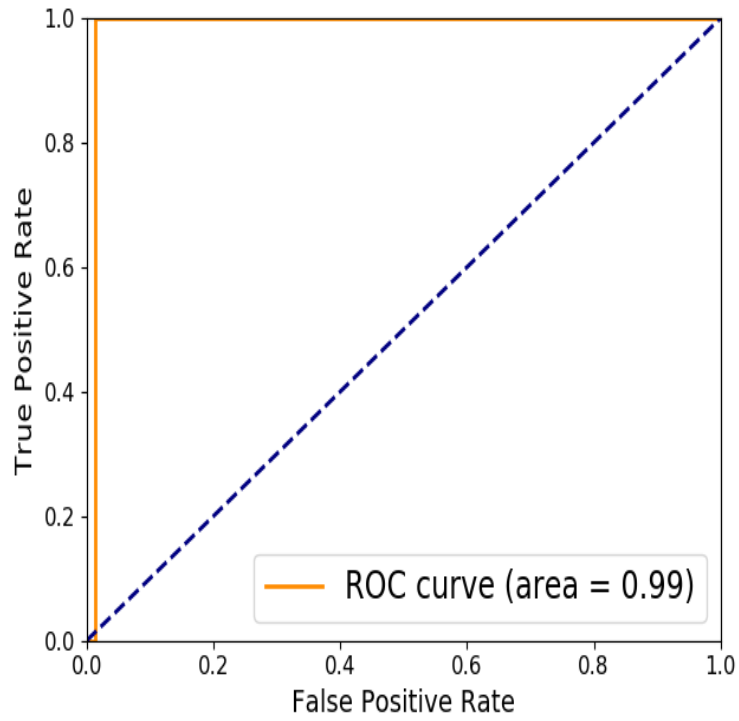
# Diffusion ROC from fGOnly\_tTestOnly STRING10\_combined.png



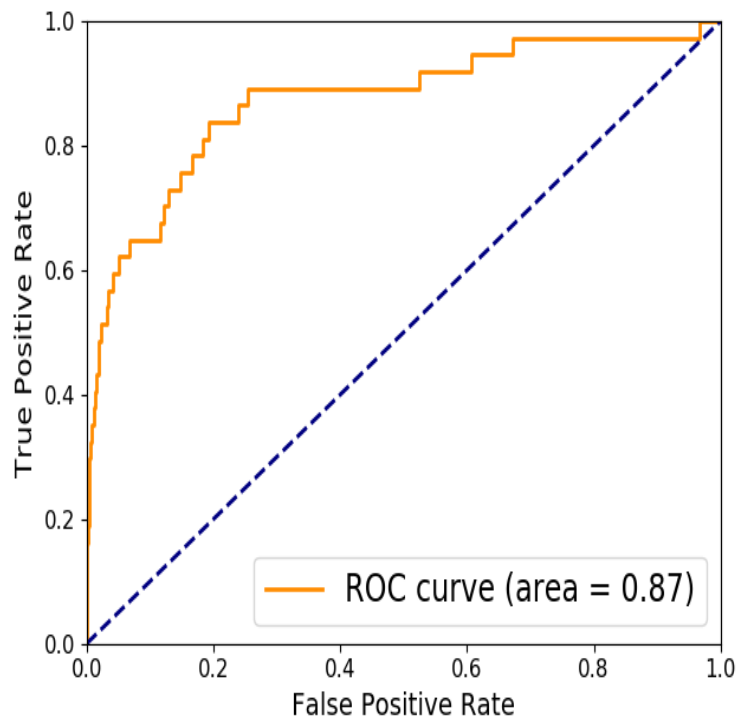
## Diffusion ROC from fTestOnly\_tGOnly STRING10\_combined.png



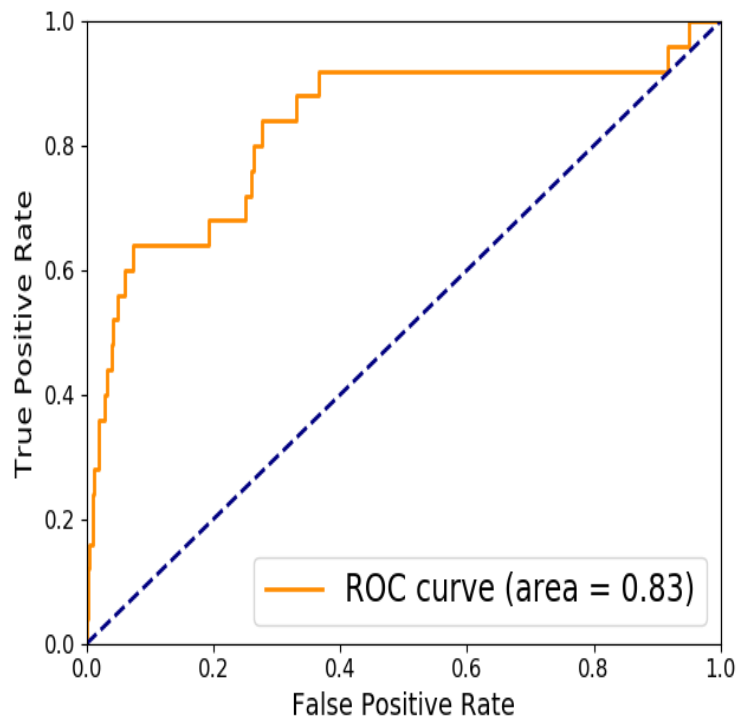
# Diffusion ROC from fTestOnly\_tOverlap STRING10\_combined.png



# 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	0.825	5.811	5.261	4.380	3.303
Test_only	GS_only	0.952	5.773	6.264	3.7742774	3.853
				27986478		
				(invalid)		
GS_only	Overlap	1.000	1.3068489	1.6481980	2.5693476	2.004093
			85115075	47206458	1128933	41634608
			(invalid)	(invalid)	(invalid)	35 (inva
						lid)
GS_only	Test	0.832	5.700	5.767	4.105	3.063
Test_only	Overlap	0.987	1.3999409	1.6909703	1.9166431	2.037782
			726551324	01899999	275188032	71775642
			(invalid)	(invalid)	(invalid)	67 (inva
						lid)
Test_only	GS	0.954	5.841	6.164	3.1934100	4.204
					255835793	
					(invalid)	
Overlap	Onlies	0.873	6.839	8.238	1.9766910	2.749
					184231297	
					(invalid)	

**	#Mapped	#Total	Not mapped genes
GS	13	13	N/A
Test_only	24	24	N/A
Overlap	1	1	ADAM10

(end of file)