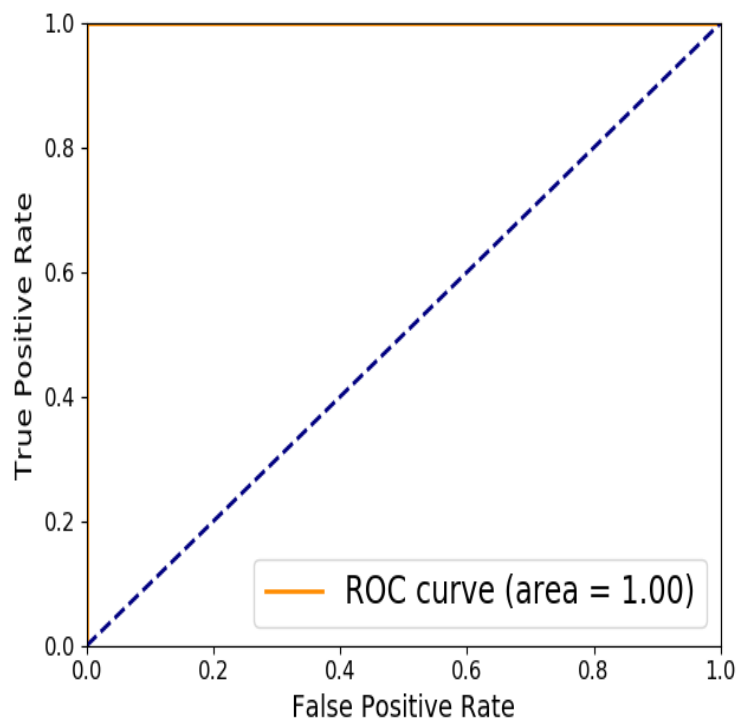
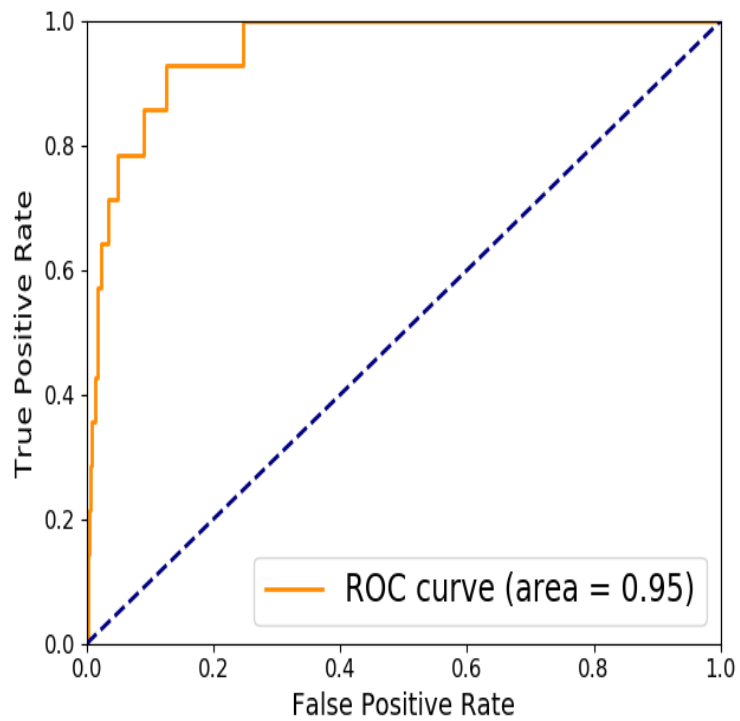


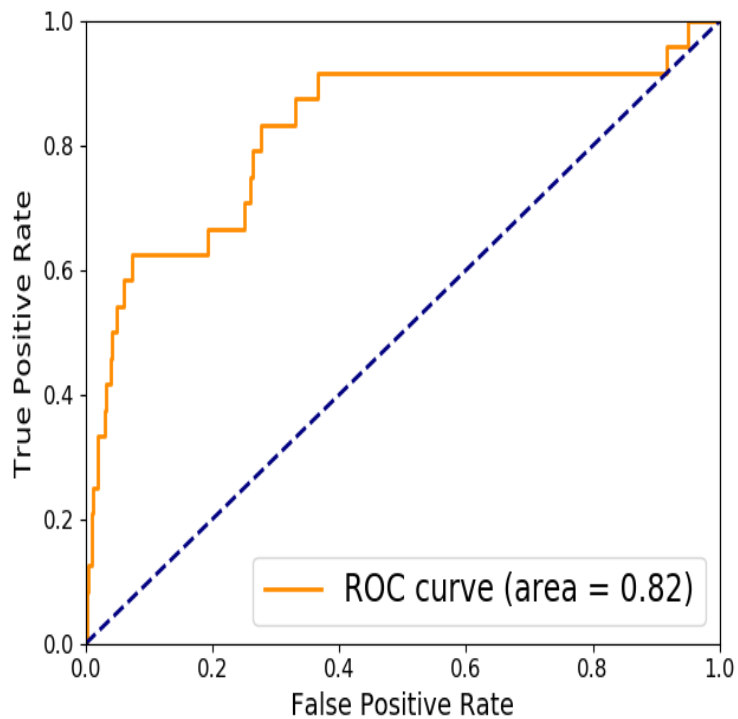
Diffusion ROC from fGOnly_tOverlap STRING10_combined.png



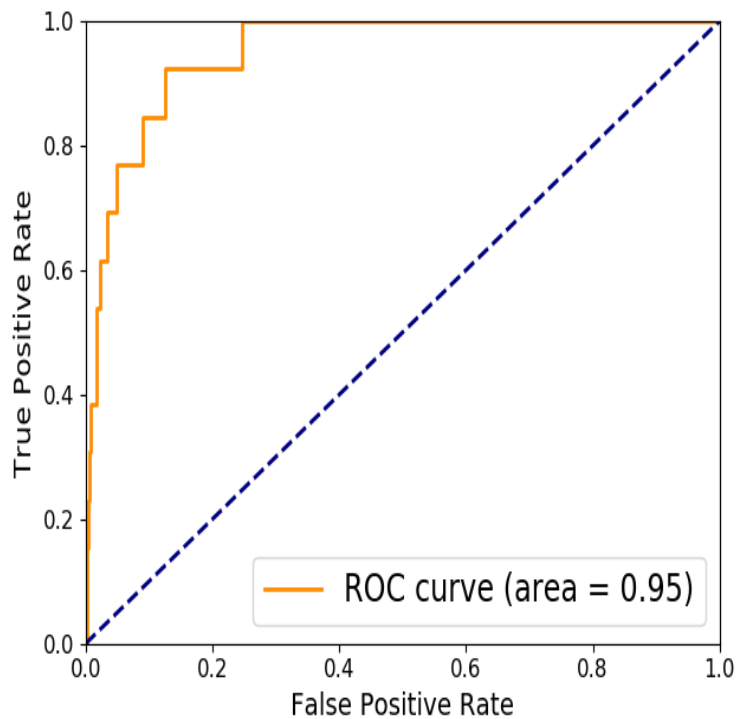
Diffusion ROC from fTestOnly_tGS STRING10_combined.png



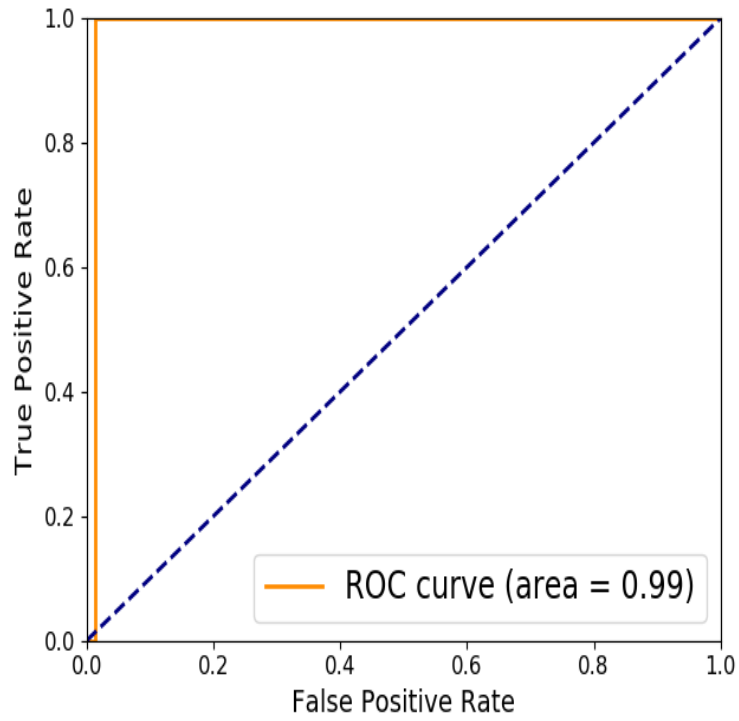
Diffusion ROC from fGOnly_tTestOnly STRING10_combined.png



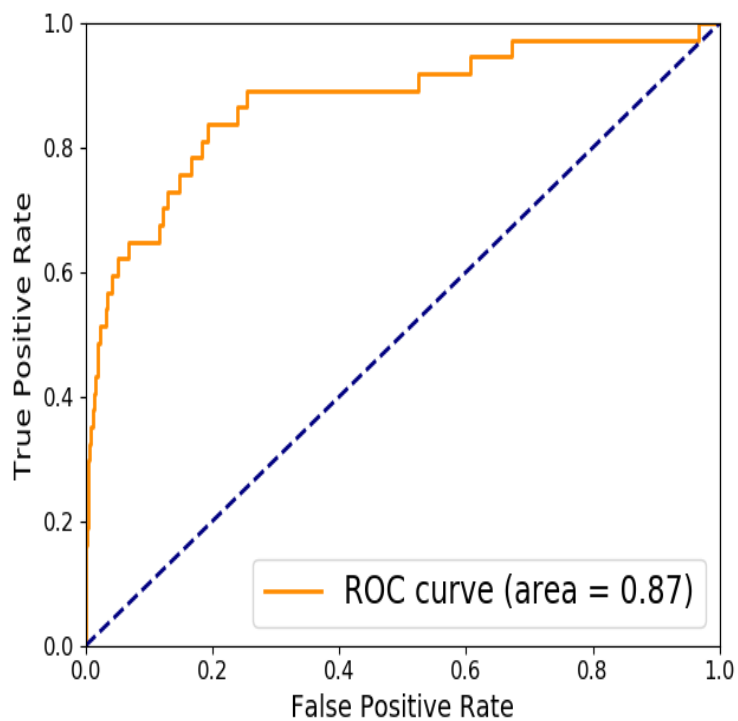
Diffusion ROC from fTestOnly_tGOnly STRING10_combined.png



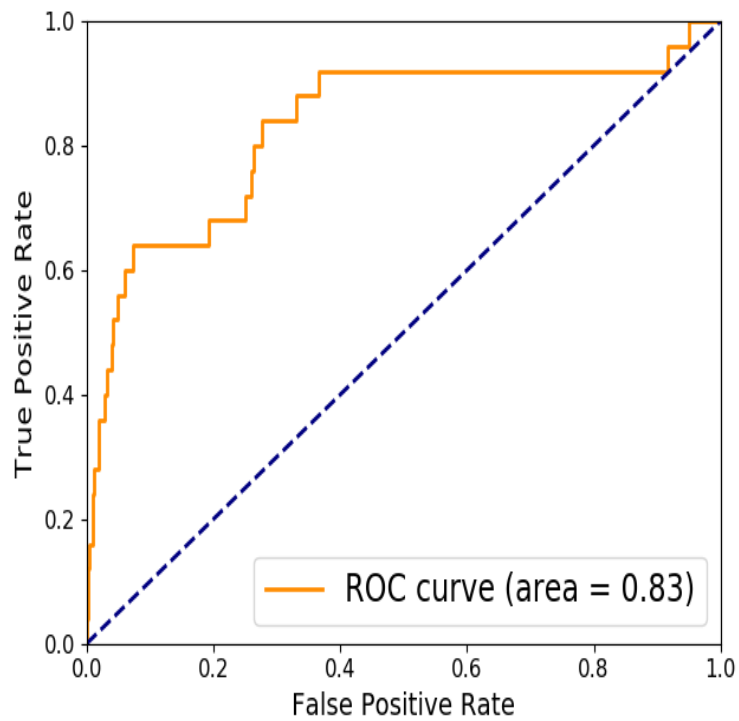
Diffusion ROC from fTestOnly_tOverlap STRING10_combined.png



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	5.808	5.745	4.4121273	4.011
				00069795		
				(invalid)		
Test_only	GS_only	0.952	4.522	5.772	3.8741309	3.631251
				96172138	31180195	
				(invalid)	34 (inva	
					lid)	
GS_only	Overlap	1.000	1.2651087	1.6757087	2.517	1.972
			61887014	429405007		
			(invalid)	(invalid)		
GS_only	Test	0.832	5.659	6.223	4.0742588	3.426592
				08605841	89706479	
				(invalid)	1 (inval	
					id)	
Test_only	Overlap	0.987	1.6859370	1.6870855	2.3078861	2.004043
			625636396	429412166	7726413	73226435
			(invalid)	(invalid)	(invalid)	96 (inva
						lid)
Test_only	GS	0.954	5.620	5.682	3.4346244	3.603515
				43710806	06238194	
				(invalid)	16 (inva	
					lid)	
Overlap	Onlies	0.873	7.143	8.502	2.3090680	2.633
				92245713		
				(invalid)		

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

(end of file)