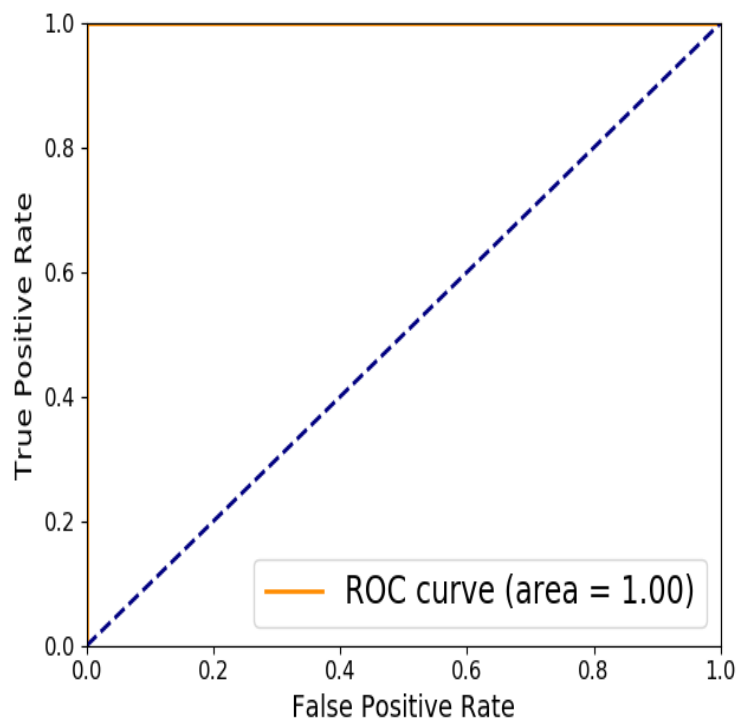
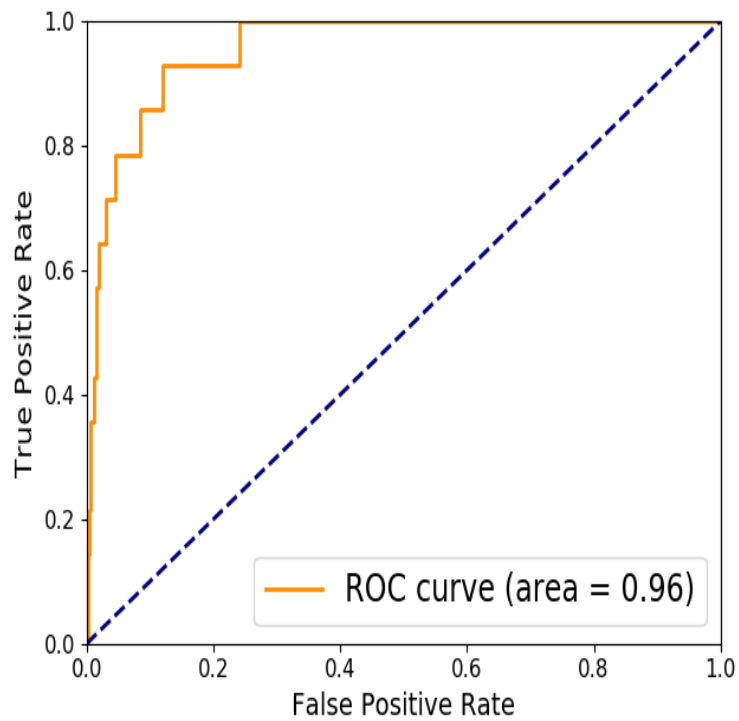


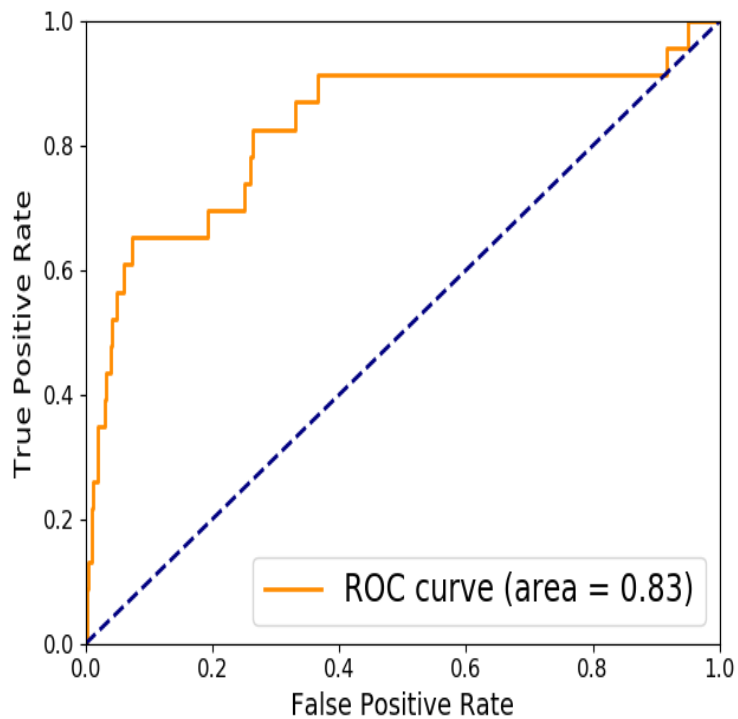
Diffusion ROC from fGOnly_tOverlap STRING10_combined.png



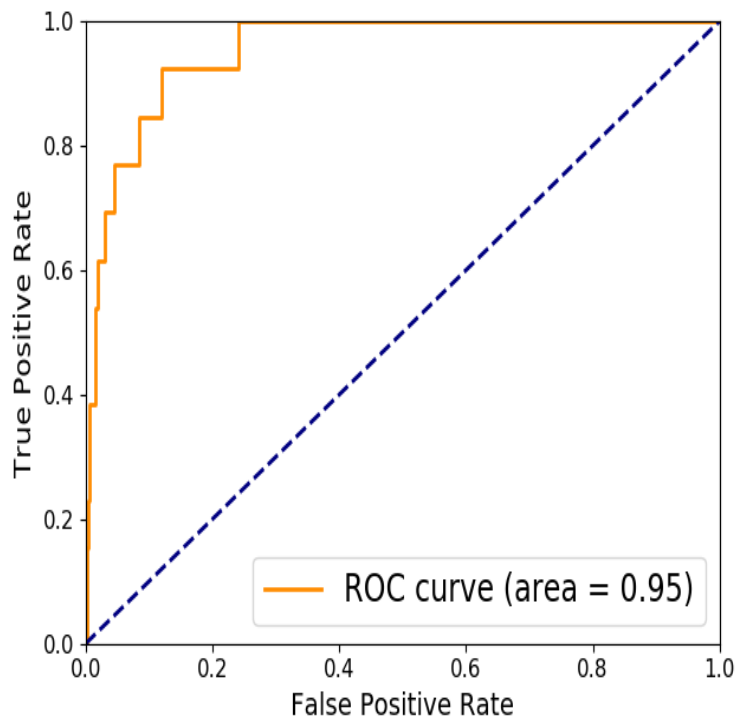
Diffusion ROC from fTestOnly_tGS STRING10_combined.png



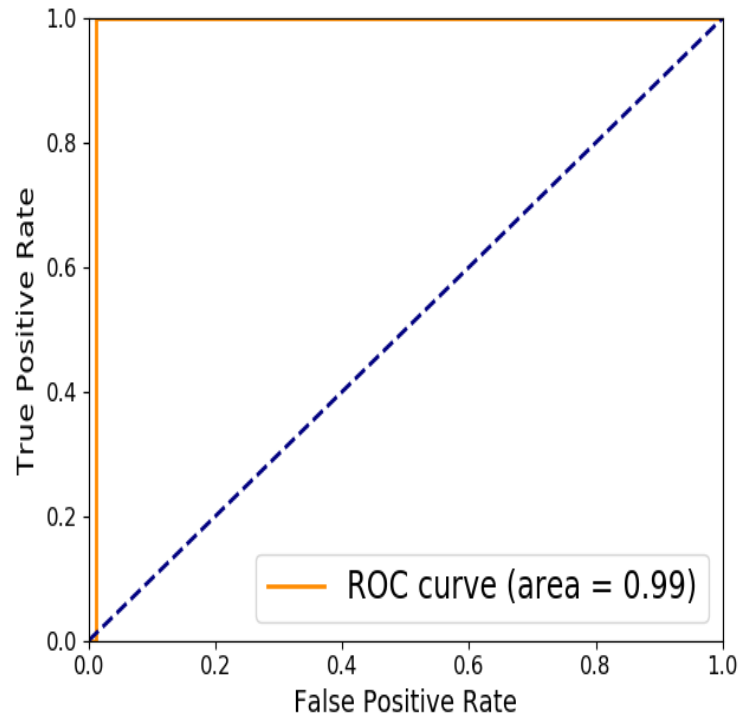
Diffusion ROC from fGOnly_tTestOnly STRING10_combined.png



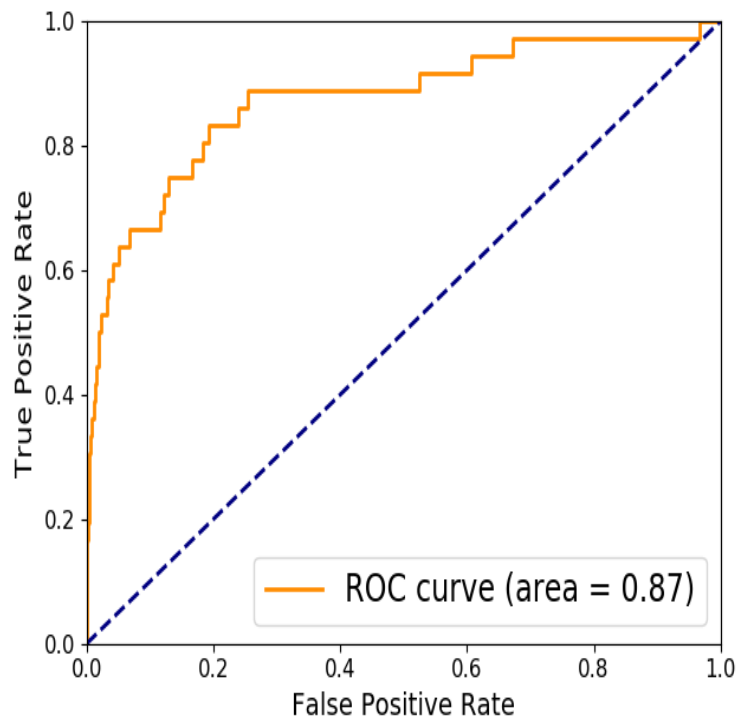
Diffusion ROC from fTestOnly_tGOnly STRING10_combined.png



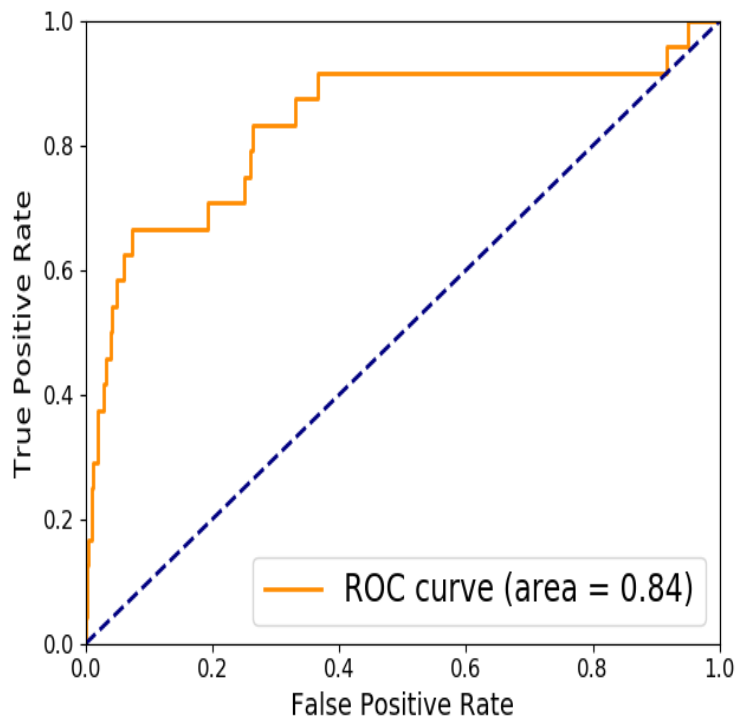
Diffusion ROC from fTestOnly_tOverlap STRING10_combined.png



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	0.829	5.494	6.489	4.2212415	4.298
				83225865		
				(invalid)		
Test_only	GS_only	0.955	6.417	6.208	3.6126220	4.023
				4469314		
				(invalid)		
GS_only	Overlap	1.000	1.3232708	1.7714785	2.369	1.934370
			077856592	122840488		24150662
			(invalid)	(invalid)		84 (inva
						lid)
GS_only	Test	0.836	6.711	5.9533821	3.6399124	3.984
				24512128	528259885	
				(invalid)	(invalid)	
Test_only	Overlap	0.989	1.5254653	1.6579969	2.2553313	1.902727
			572040724	575215068	26729682	36303021
			(invalid)	(invalid)	(invalid)	(invalid)
)
Test_only	GS	0.957	5.720	6.418	4.880	4.004
Overlap	Onlies	0.874	7.907	6.983	1.9435597	2.296
				6152941		
				(invalid)		

**	#Mapped	#Total	Not mapped genes
GS	13	13	N/A
Test_only	23	24	HLA -DRB1
Overlap	1	1	ADAM10

(end of file)