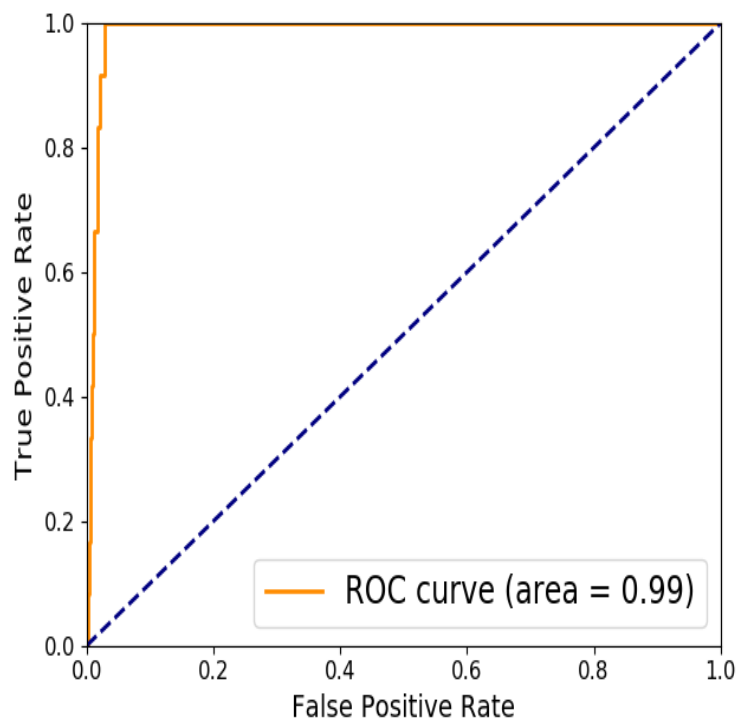
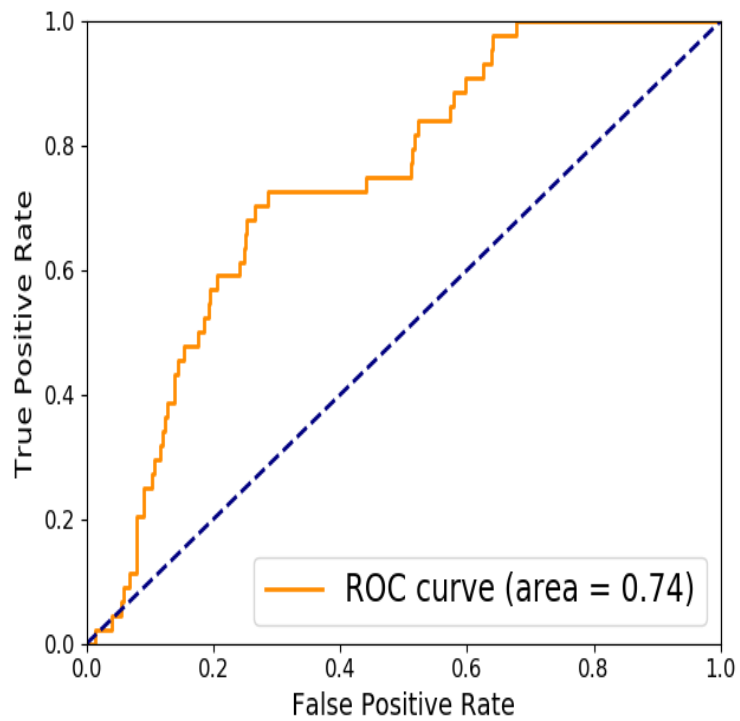


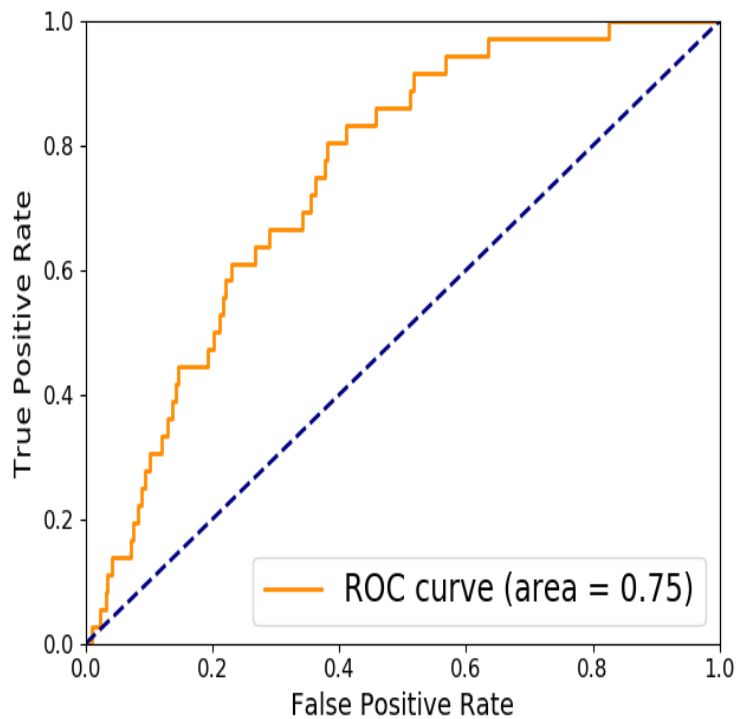
Diffusion ROC from fGOnly_tOverlap STRING10_combined.png



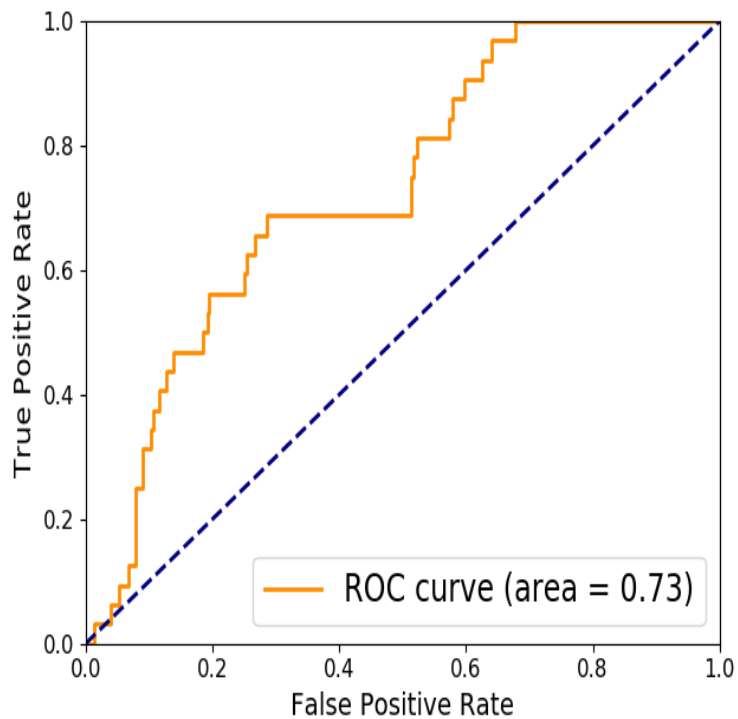
Diffusion ROC from fTestOnly_tGS STRING10_combined.png



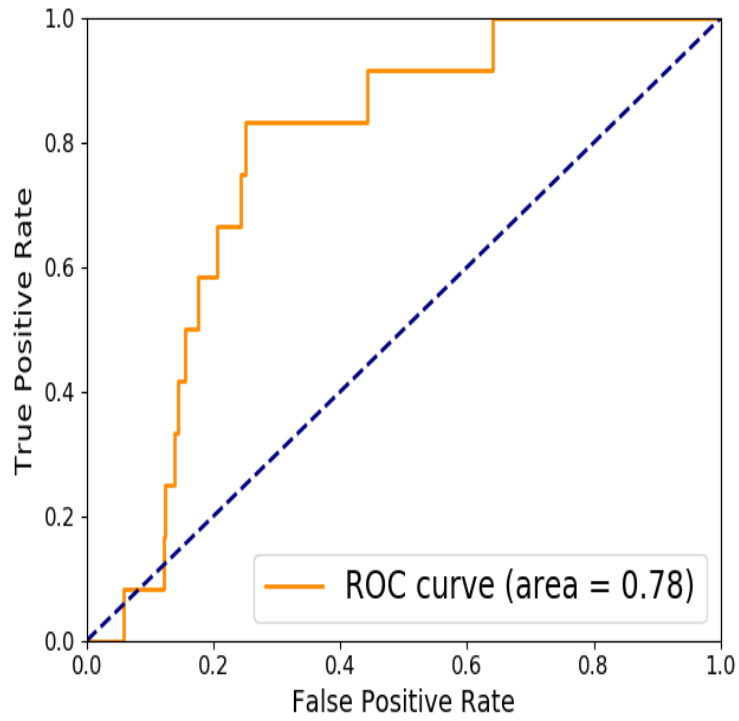
Diffusion ROC from fGOnly_tTestOnly STRING10_combined.png



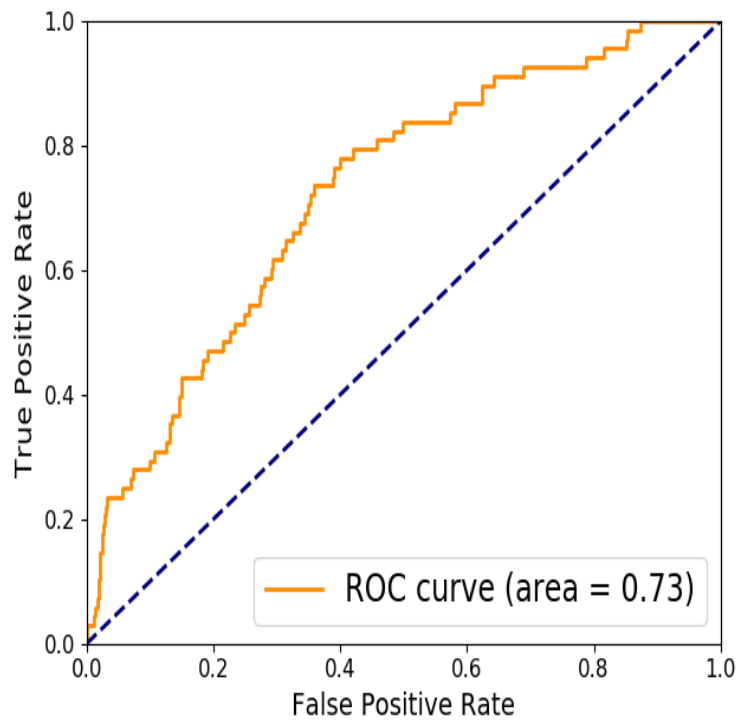
Diffusion ROC from fTestOnly_tGOnly STRING10_combined.png



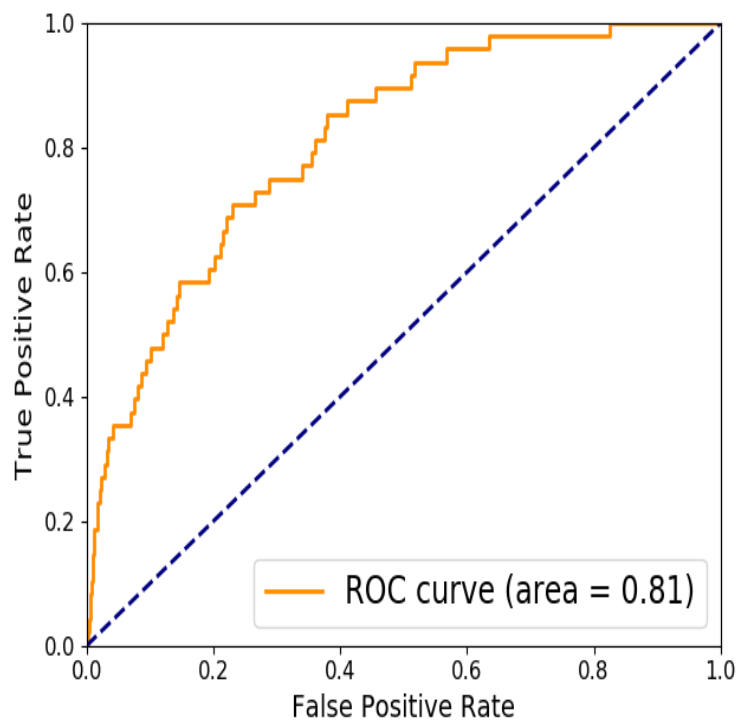
Diffusion ROC from fTestOnly_tOverlap STRING10_combined.png



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	0.753	4.132	4.994	3.6282466	2.398634
				693760234	75571319	
				(invalid)	1 (inval	
				id)		
Test_only	GS_only	0.730	4.899	4.164	1.9696855	2.216
				020808302		
				(invalid)		
GS_only	Overlap	0.988	5.724	5.598	2.3169069	3.050614
				456431943	20664700	
				(invalid)	6 (inval	
				id)		
GS_only	Test	0.812	6.166	7.261	4.6986200	4.231
				467904835		
				(invalid)		
Test_only	Overlap	0.775	2.866	3.462	1.1395935	1.578378
				25180707	28313488	
				(invalid)	76 (inva	
				lid)		
Test_only	GS	0.742	5.095	6.516	1.838	2.478
Overlap	Onlies	0.726	5.523	6.5191807	2.324	1.984
				75970113		
				(invalid)		

**	#Mapped	#Total	Not mapped genes
GS	32	37	MT-RNR2;RP11-331F4.4; N/A
			MTATP6P1;PNLIPRP2;PRS
			S3P2;PRSS3P1
Test_only	36	37	MT-RNR2;MT-RNR1 N/A
Overlap	12	13	MT-RNR2 MT-RNR2;MT-ND3;RPS12
			;MT-ND2;MT-ATP8;MT-
			CYB;MT-ND4;MT-ND1;MT-
			ATP6;MT-CO3;MT-ND4L
			;MT-CO2;MT-CO1

(end of file)