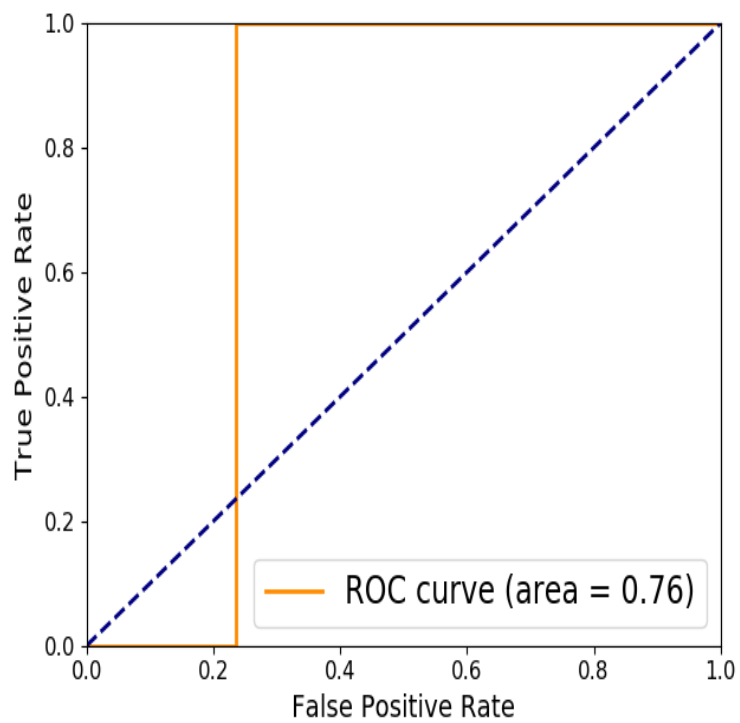
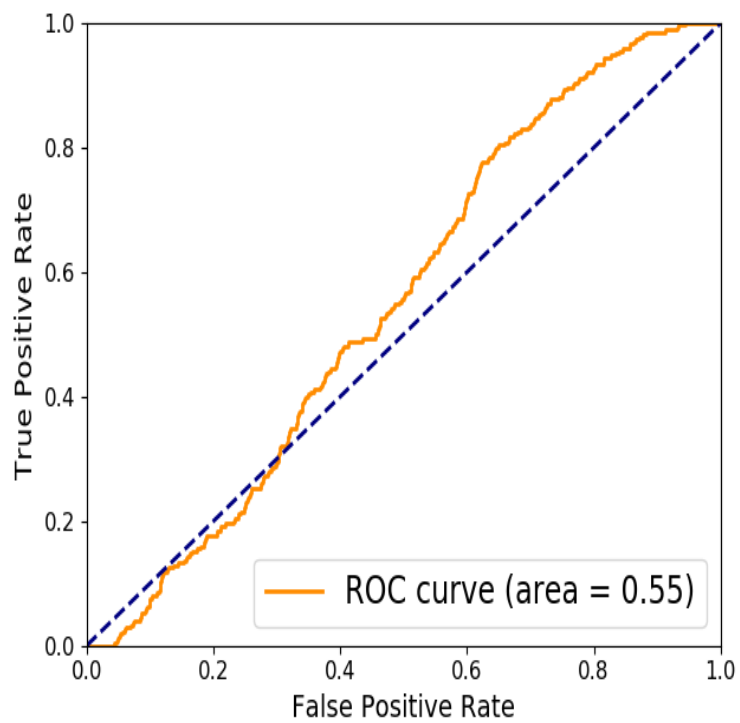


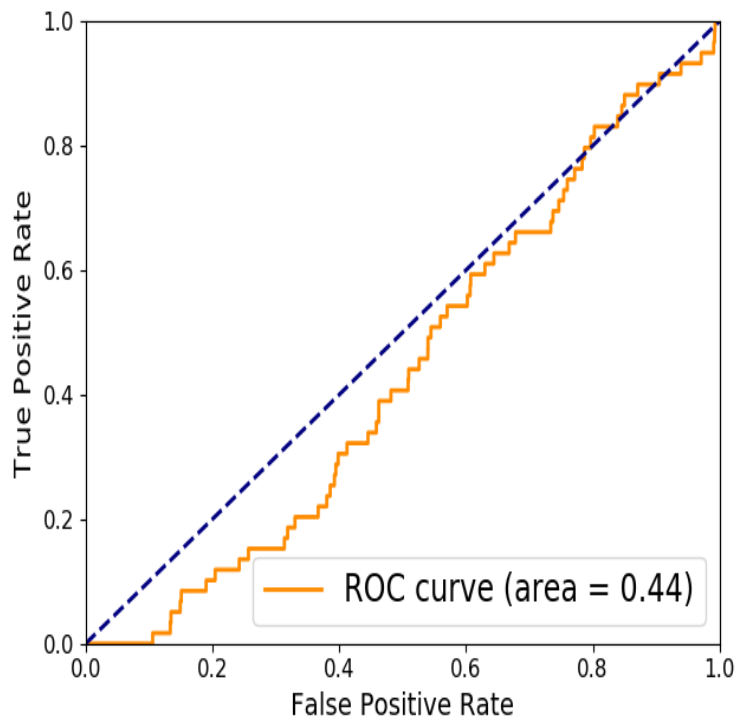
Diffusion ROC from fGOnly_tOverlap STRING10_combined.png



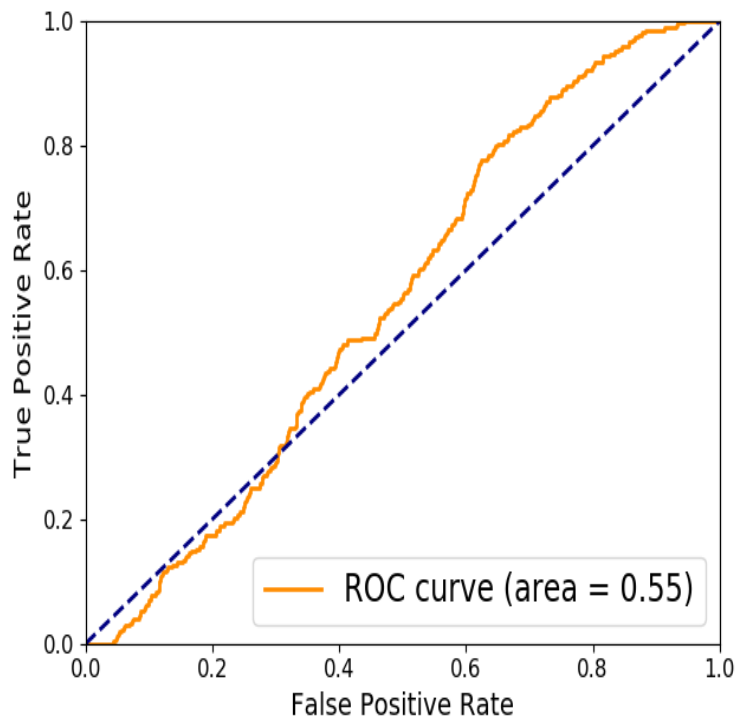
Diffusion ROC from fTestOnly_tGS STRING10_combined.png



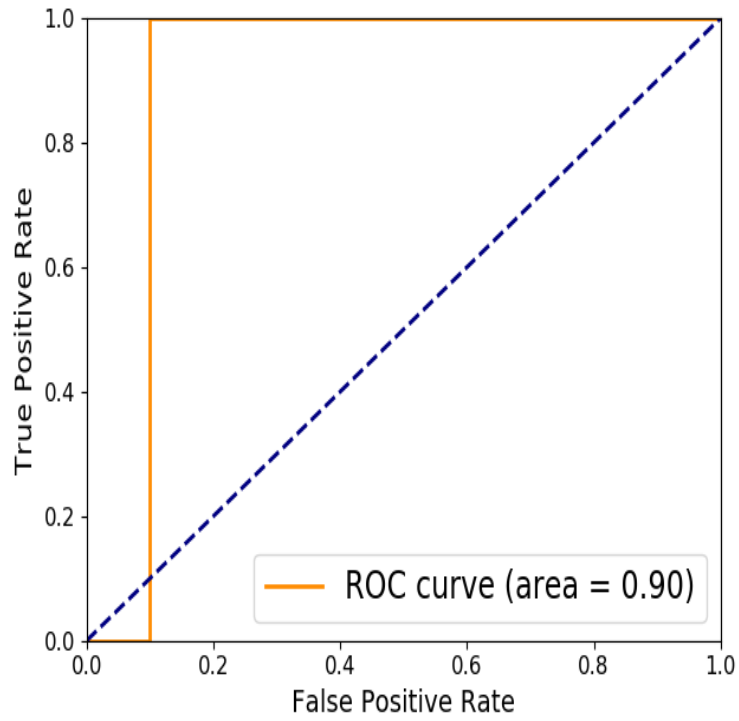
Diffusion ROC from fGOnly_tTestOnly STRING10_combined.png



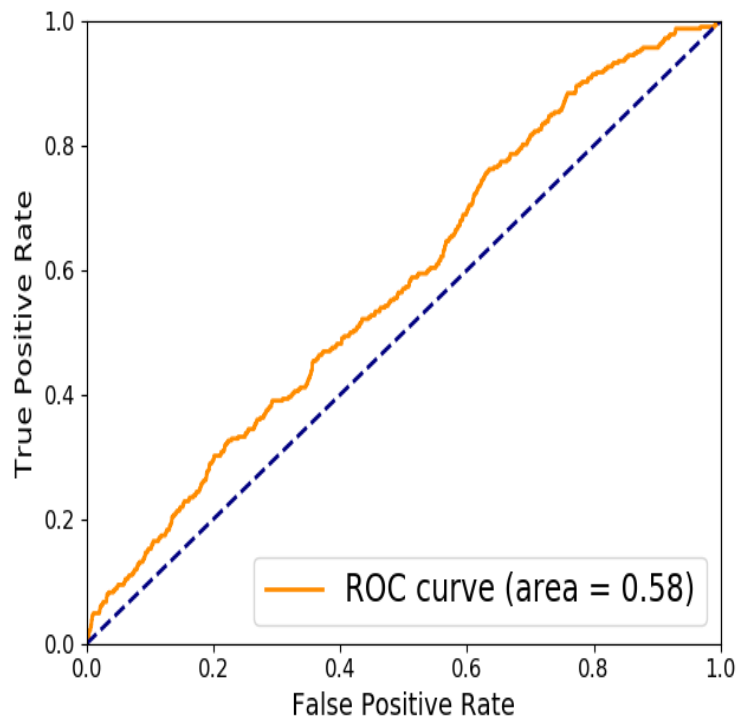
Diffusion ROC from fTestOnly_tGOnly STRING10_combined.png



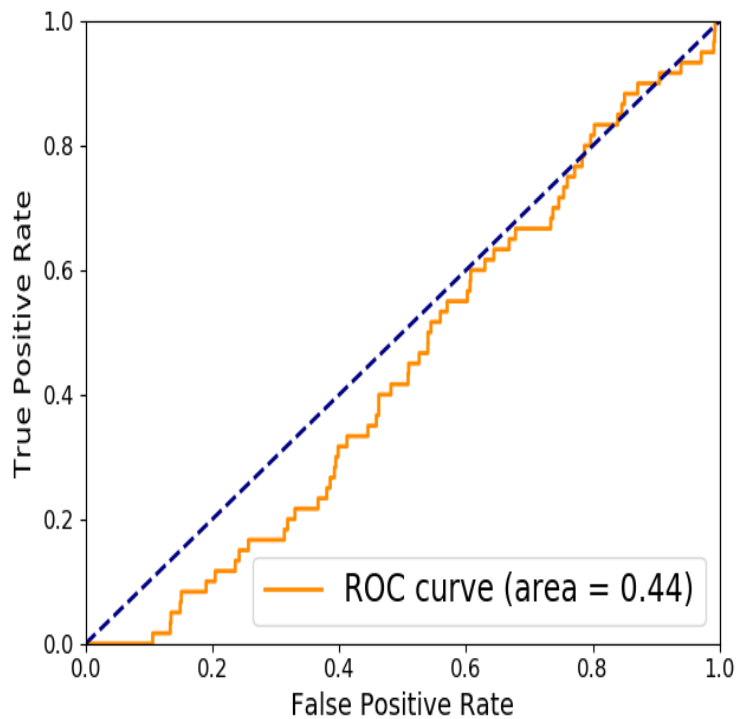
Diffusion ROC from fTestOnly_tOverlap STRING10_combined.png



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	0.438	-1.428	-1.628	-1.524	-1.762
Test_only	GS_only	0.553	-4.010	3.897	-1.355	-2.532
GS_only	Overlap	0.764	1.3449812	0.8370864	2.6065910	1.866835
		100117553	592743662	95291122	72231678	
		(invalid)	(invalid)	(invalid)	09 (inva	
				lid)		
GS_only	Test	0.443	-1.053	-1.485	-1.116	-1.681
Test_only	Overlap	0.901	1.1965233	1.5478333	1.7989227	2.238853
		060623077	393500587	10970993	13175516	
		(invalid)	(invalid)	(invalid)	9 (inval	
				id)		
Test_only	GS	0.555	-4.521	3.115	-1.332	-2.217
Overlap	Onlies	0.578	-0.776	4.714	-0.075	-0.560

**	#Mapped	#Total	Not mapped genes
GS	269	283	RABGAP;UGT2A2;TTL3;MMP5;U2af50-1;MMP6; N/A
			TRIP;U2af50-2;TTL8;ACK1;JMJD7-PLA2G4B;
			MMP4;CDC2L2;RABGAPL
Test_only	59	62	GRXCR2;WDR52;C19orf45 N/A
Overlap	1	1	NEK4

(end of file)