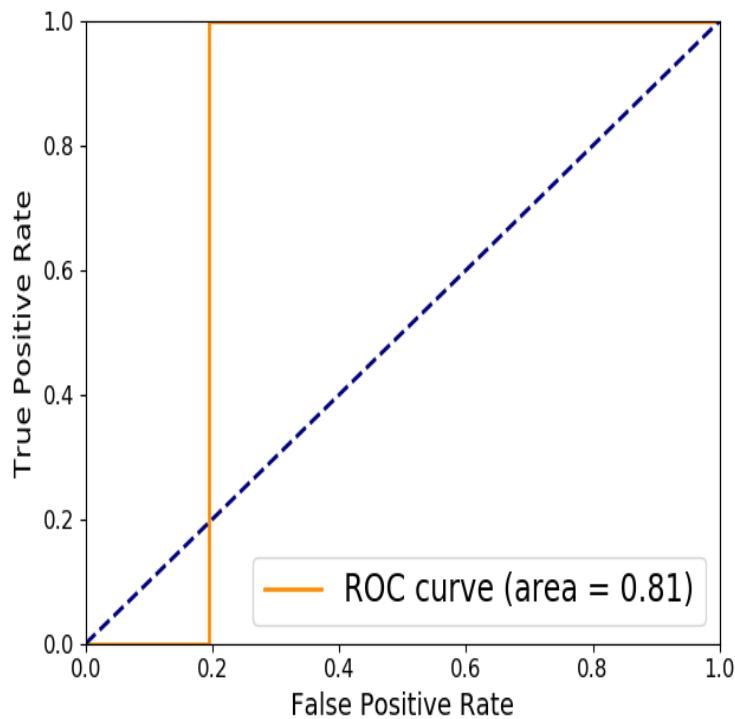
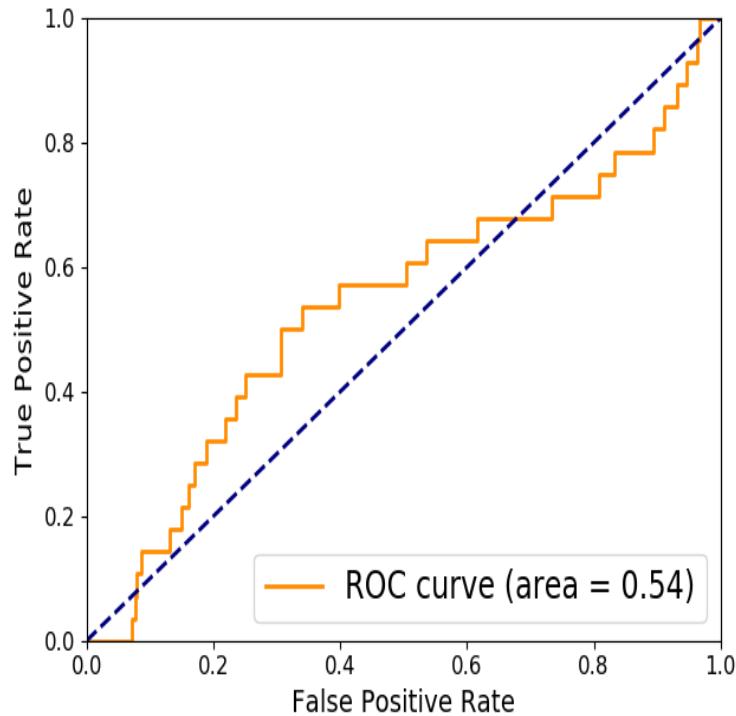


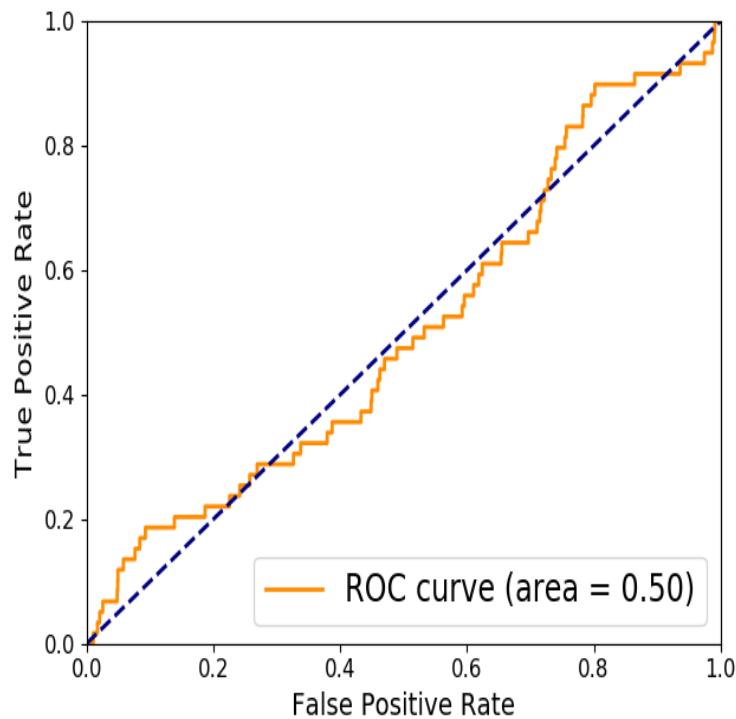
Diffusion ROC from fGSonly_tOverlap STRING10_combined.png



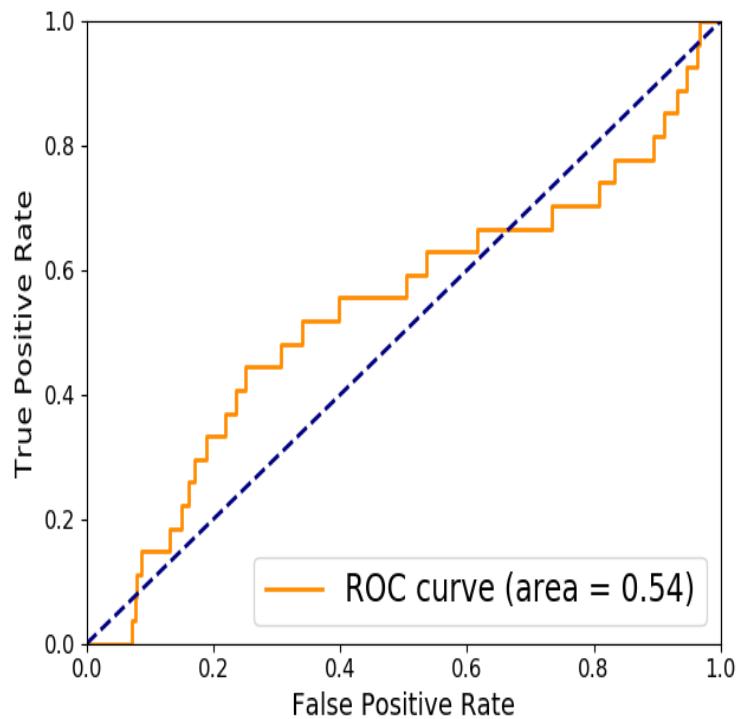
Diffusion ROC from fTestOnly_tGS STRING10_combined.png



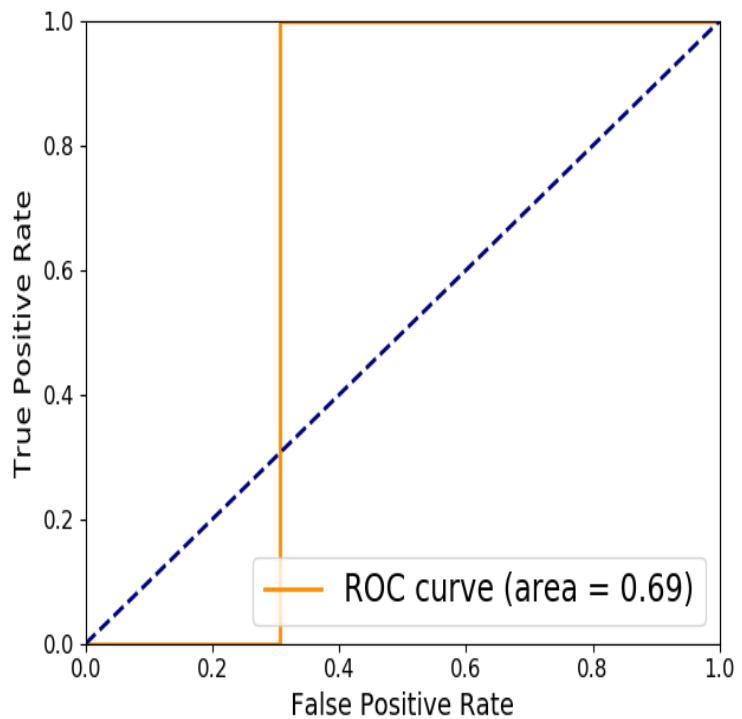
Diffusion ROC from fGSonly_tTestOnly STRING10_combined.png



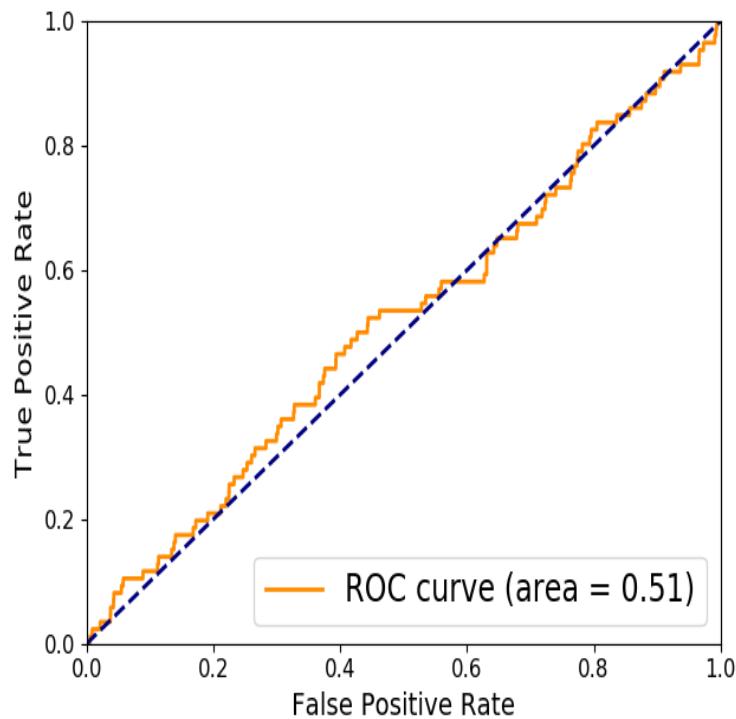
Diffusion ROC from fTestOnly_tGSonly STRING10_combined.png



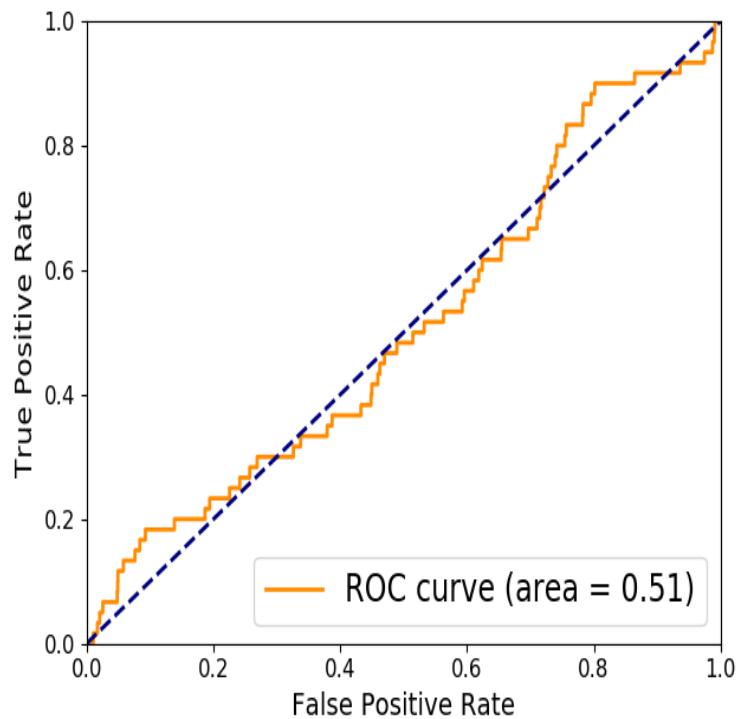
Diffusion ROC from fTestOnly_tOverlap STRING10_combined.png



Diffusion ROC from fOverlap_tOnlies STRING10_combined.png



Diffusion ROC from fGSonly_tTest STRING10_combined.png



<i>Diff_FROM</i>	<i>Diff_TO</i>	<i>AUC</i>	<i>Zt_wdegre</i>	<i>Zt</i>	<i>Zf_wdegre</i>	<i>Zf</i>
/	/	e	/	e	/	/
GS_only Test_only 0.502 -0.012 0.028 0.071 0.712						
Test_only GS_only 0.537 2.0406742 0.564 2.002 1.863						
/	/	62808161	/	/	/	/
/	/	(invalid)	/	/	/	/
GS_only Overlap 0.806 1.679 1.0423704 1.2637593 1.559969						
/	/	/	154164364	77048576	18976440	/
/	/	/	(invalid)	(invalid)	6 (invalid)	/
/	/	/	/	/	id)	/
GS_only Test 0.507 0.390 0.127 0.385 0.689						
Test_only Overlap 0.693 0.9341320 0.5083131 0.8019180 0.692748						
/	/	091701606	201711619	548295594	27248605	/
/	/	/	(invalid)	(invalid)	(invalid)	44 (invalid)
/	/	/	/	/	lid)	/
Test_only GS 0.542 1.926 0.887 1.931 1.972						
Overlap Onlies 0.515 0.863 0.4529314 0.9015424 0.956						
/	/	/	859080995	969200674	/	/
/	/	/	6	(invalid)	/	/
/	/	/	(invalid)	/	/	/

**	#Mapped	#Total	Not mapped genes	/	/
GS 27 27 / N/A					
Test_only 59 62 C19orf45;WDR52;GRXCR2 N/A					
Overlap 1 1 / MAP9					

(end of file)