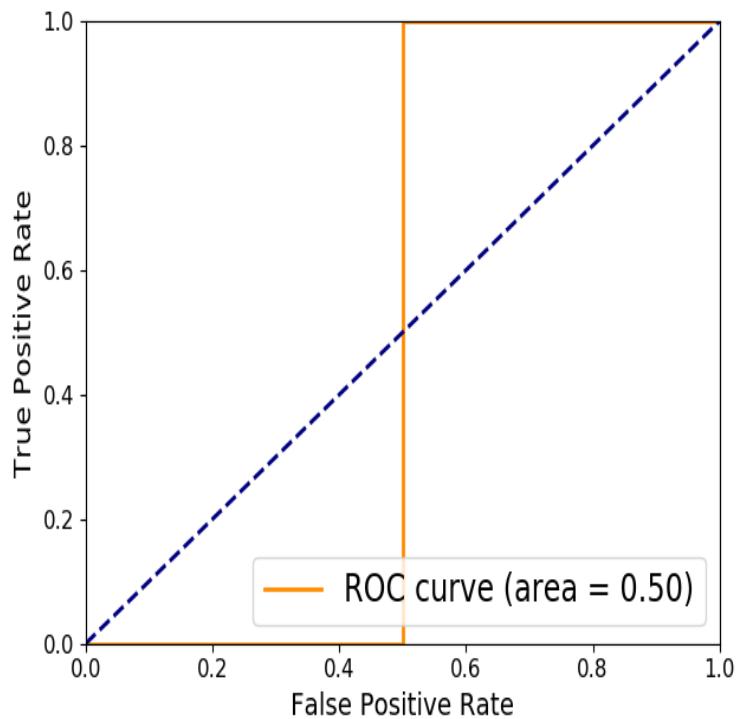
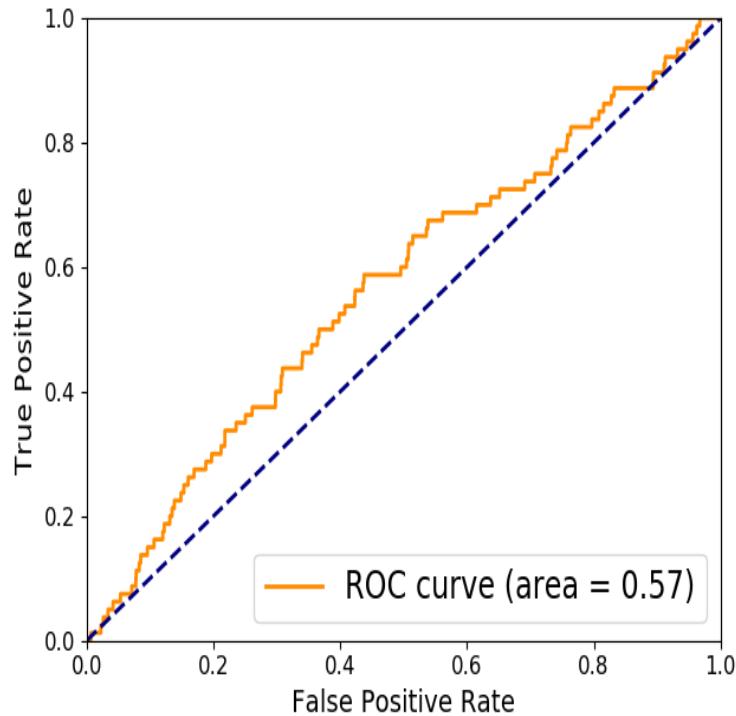


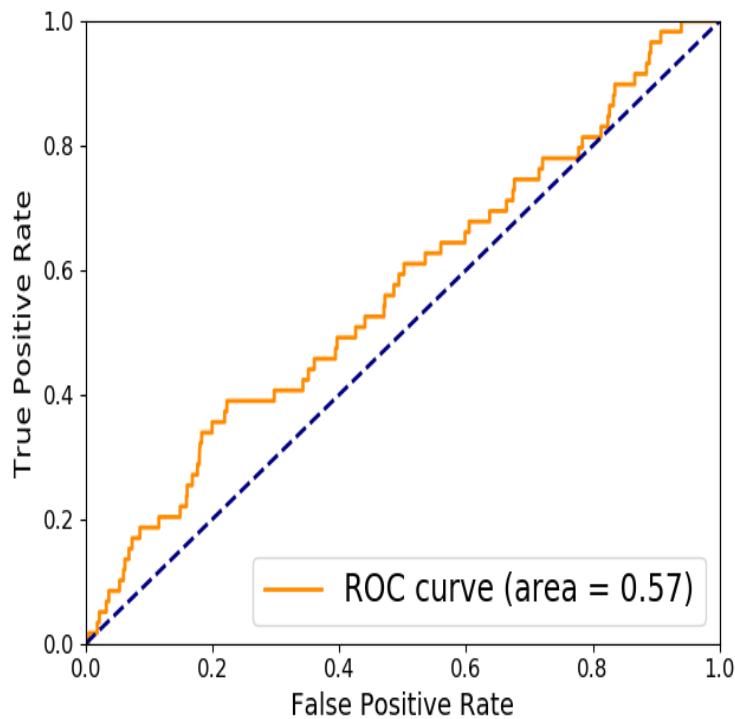
Diffusion ROC from fGSonly_tOverlap STRING10_combined.png



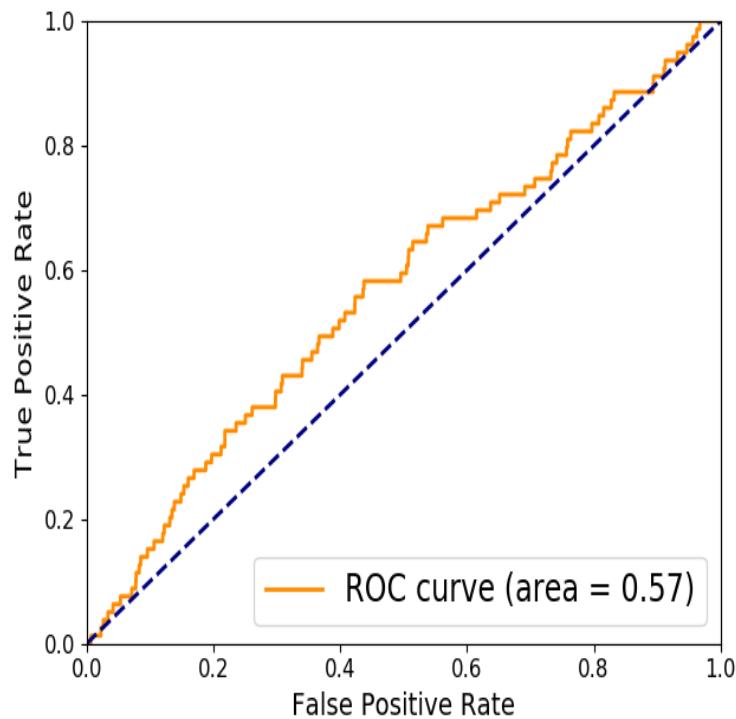
Diffusion ROC from fTestOnly_tGS STRING10_combined.png



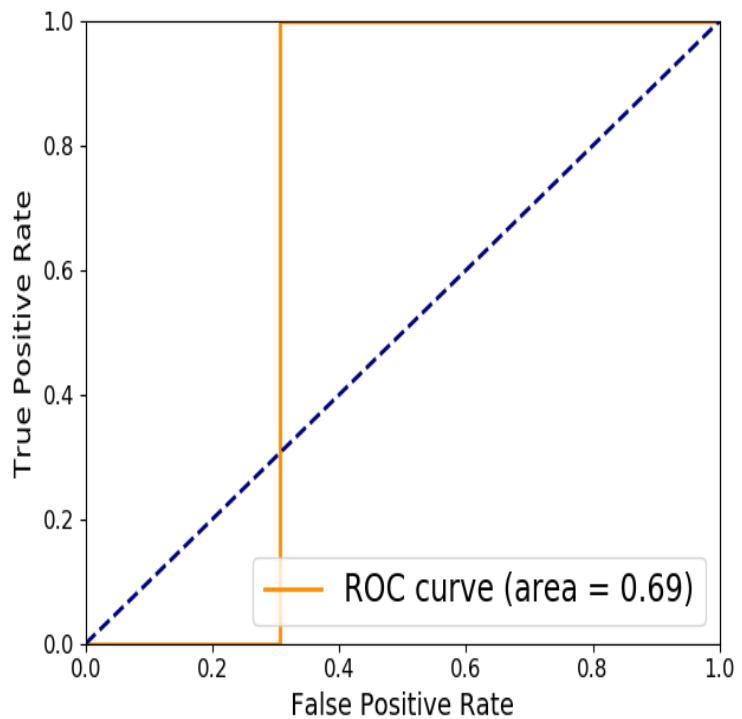
Diffusion ROC from fGSonly_tTestOnly STRING10_combined.png



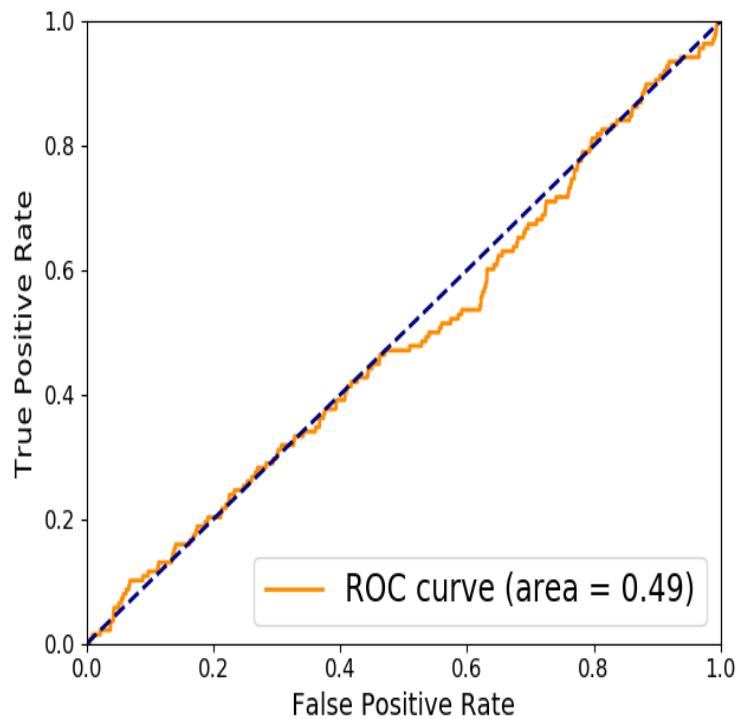
Diffusion ROC from fTestOnly_tGSonly STRING10_combined.png



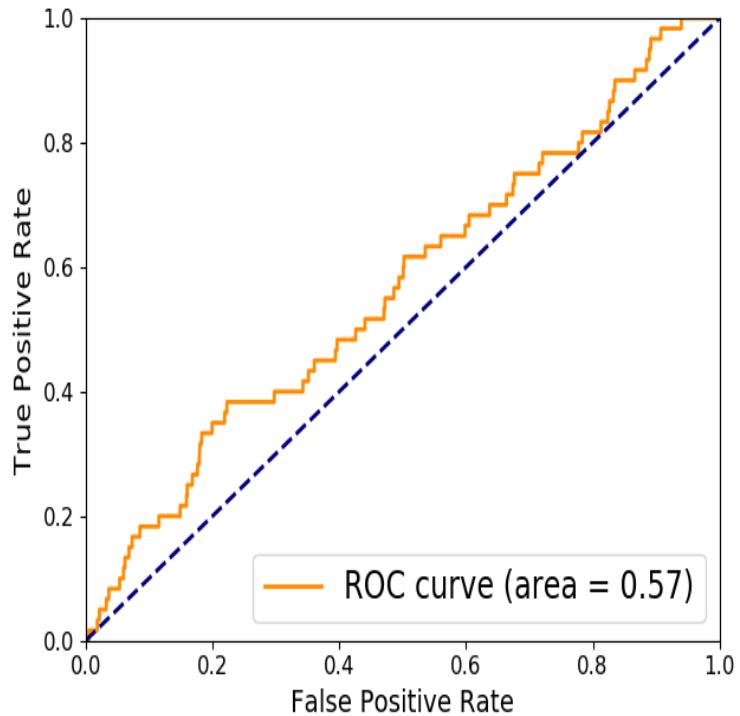
Diffusion ROC from fTestOnly_tOverlap STRING10_combined.png



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 0.567 1.815 1.938 1.460 1.862						
Test_only GS_only 0.568 3.223 2.134 3.624 3.322						
GS_only Overlap 0.499 0.0706205 0.1505870 -0.049026 0.095681						
/	/	/	442851017	778027697	521558618	02353949
/	/	/	6	(invalid)	41	459 (inv)
/	/	/	(invalid)		(invalid)	alid
GS_only Test 0.566 1.959 1.940 1.483 2.090						
Test_only Overlap 0.693 0.8248438 0.6494330 0.8904887 0.747267						
/	/	/	263217647	360193366	099580963	48827714
/	/	/	(invalid)	(invalid)	(invalid)	94 (inva
/	/	/	/	/	/	lid)
Test_only GS 0.569 3.392 2.110 3.522 3.246						
Overlap Onlies 0.489 -0.059 -0.651286 0.096 0.238						
/	/	/	032579569	/	/	/
/	/	/	(invalid)	/	/	/

**	#Mapped	#Total	Not mapped genes	/	/
GS	79	80	<i>IQCF6</i>		N/A
Test_only	59	62	<i>WDR52;C19orf45;GRXCR2</i>		N/A
Overlap	1	1			<i>MAP9</i>

(end of file)