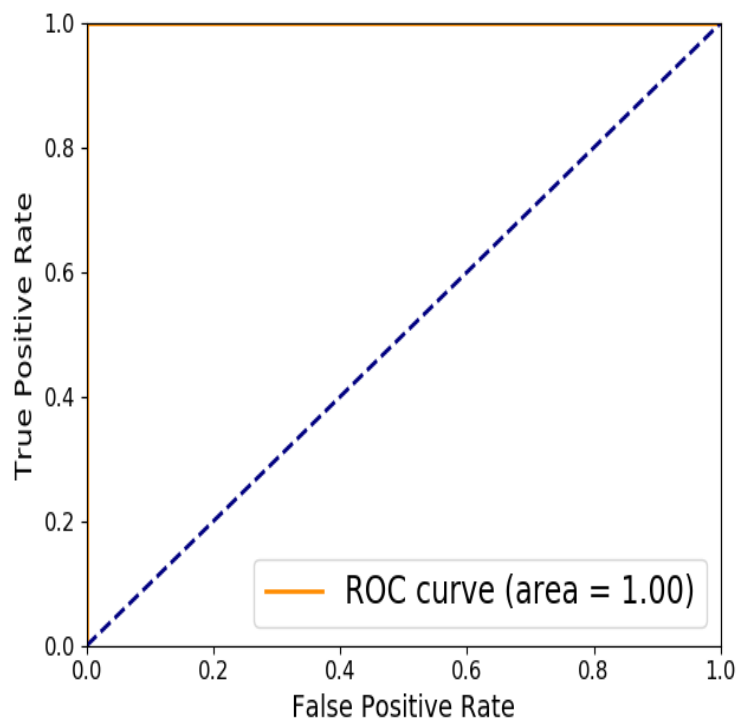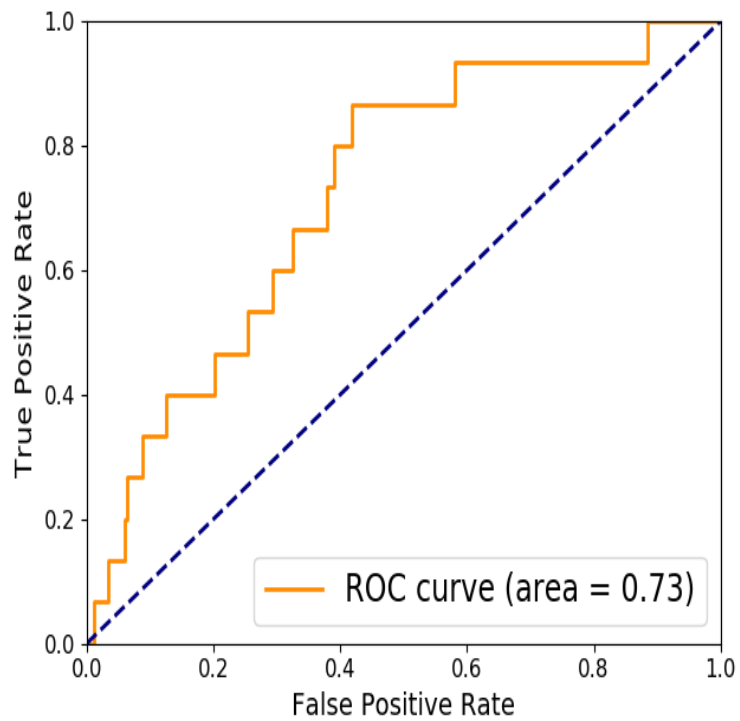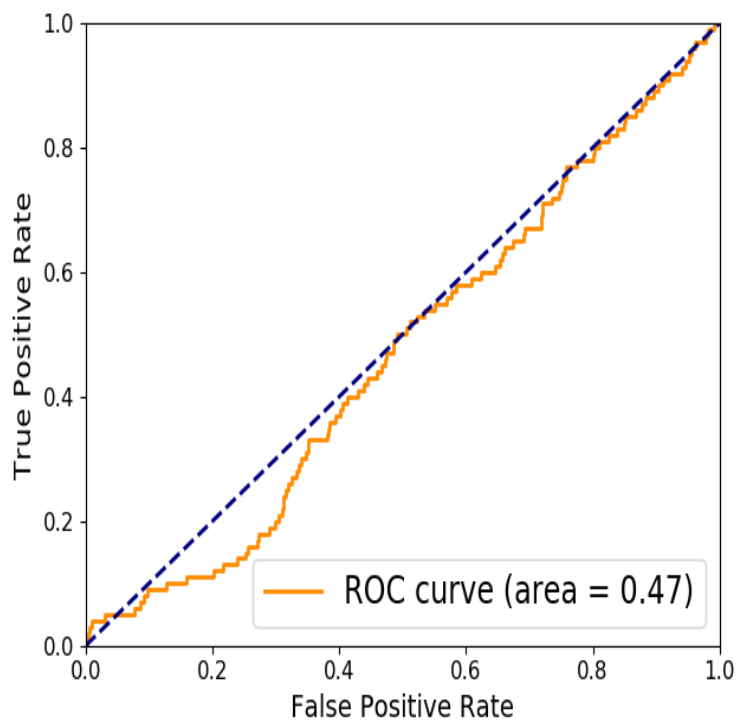# Diffusion ROC from fGSonly_tOverlap STRING10_combined.png
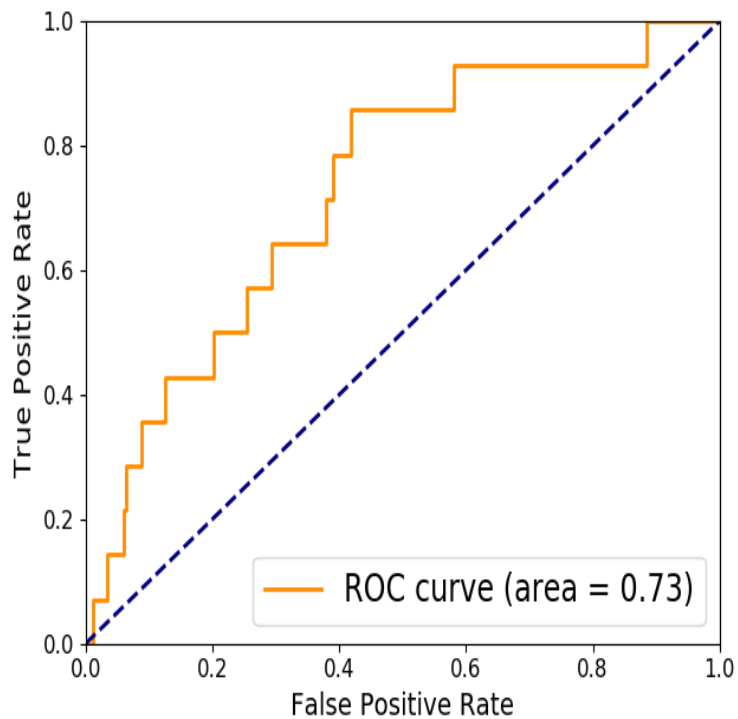
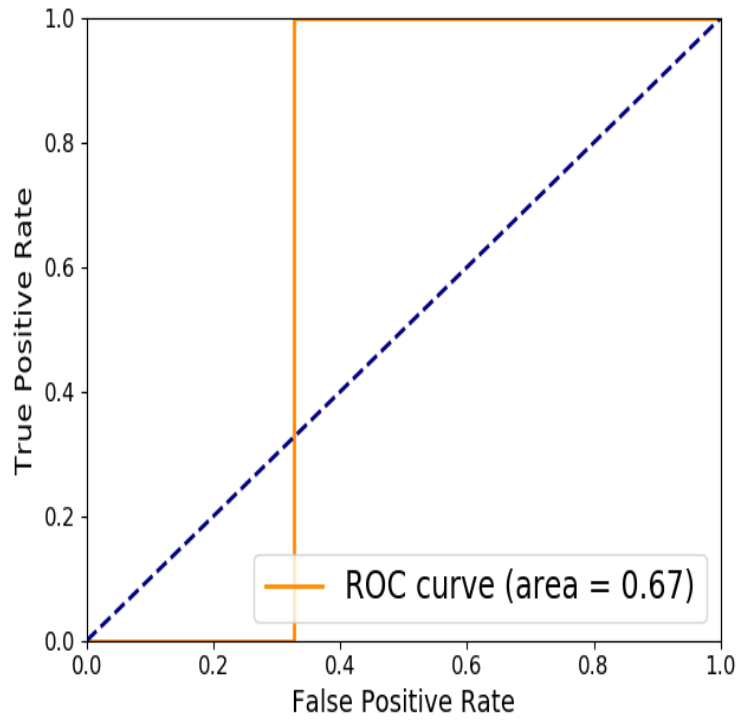# Diffusion ROC from fTestOnly_tGS STRING10_combined.png

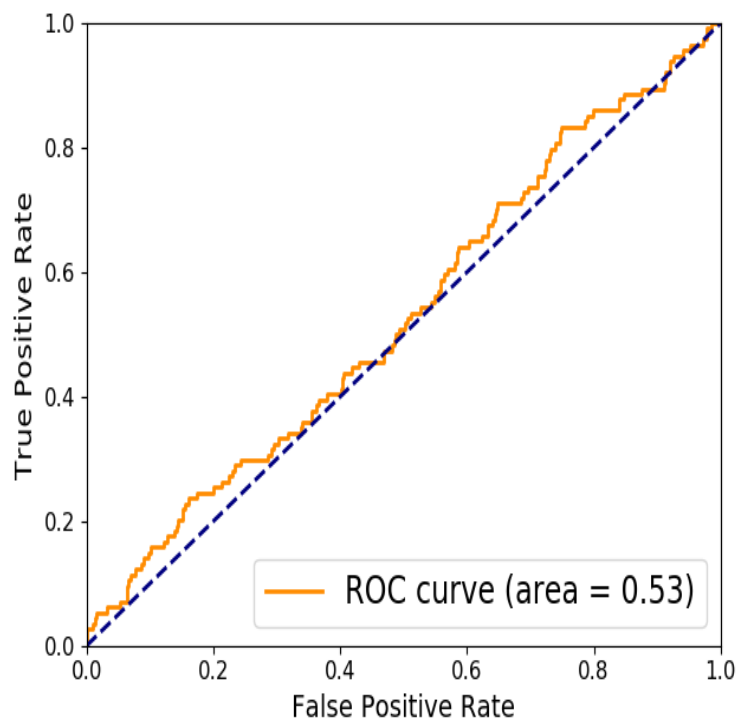# Diffusion ROC from fGSonly_tTestOnly STRING10_combined.png

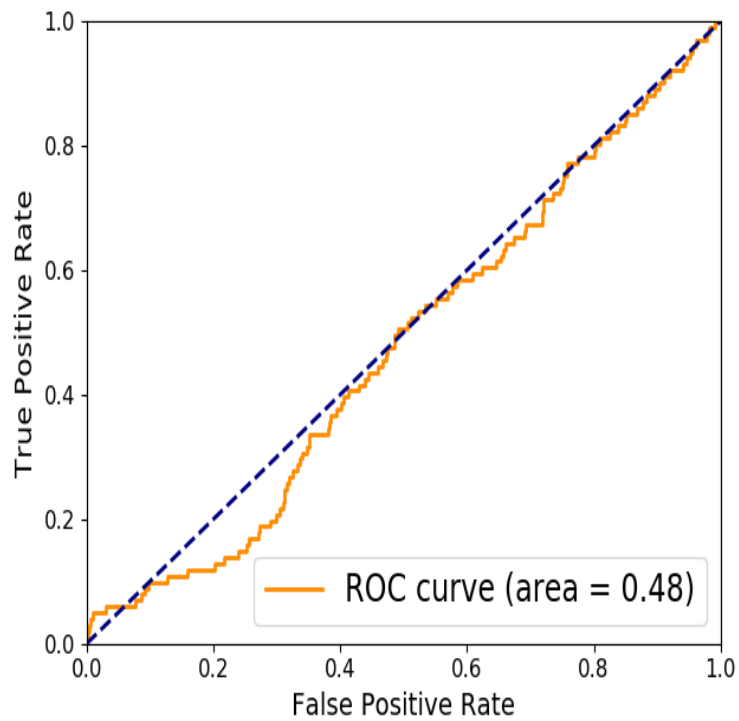# Diffusion ROC from fTestOnly_tGSonly STRING10_combined.png

Diffusion ROC from fTestOnly_tOverlap STRING10_combined.png

Diffusion ROC from fOverlap_tOnlies STRING10_combined.png

Diffusion ROC from fGSonly_tTest STRING10_combined.png

Diff_FROM	Diff_TO	AUC	Zt_wdegree	Zt	Zf_wdegree	Zf
GS_only	Test_only	0.472	-1.150	-1.098	-1.846	-0.785
Test_only	GS_only	0.730	1.063	3.192	0.782	0.592
GS_only	Overlap	0.999	1.3844373270975416 (invalid)	1.6083459629347872 (invalid)	2.1942578362474046 (invalid)	2.1053452815615 9 (invalid)
GS_only	Test	0.477	-0.937	-1.041	-1.485	-0.730
Test_only	Overlap	0.674	-5.024417218272335 (invalid)	0.5836428763207943 (invalid)	-1.952904709790336 3 (invalid)	-2.7737436775097 972 (invalid)
Test_only	GS	0.726	0.9808369077157943 (invalid)	3.030	0.5383208163200197 (invalid)	0.327
Overlap	Onlies	0.530	0.395	1.069	2.7233947298945407 (invalid)	0.479

**	#Mapped	#Total	Not mapped genes	
GS	14	15	CAMK1V	N/A
Test_only	100	102	AMER2;ERVFRD.1	N/A
Overlap	1	1		APOE

(end of file)