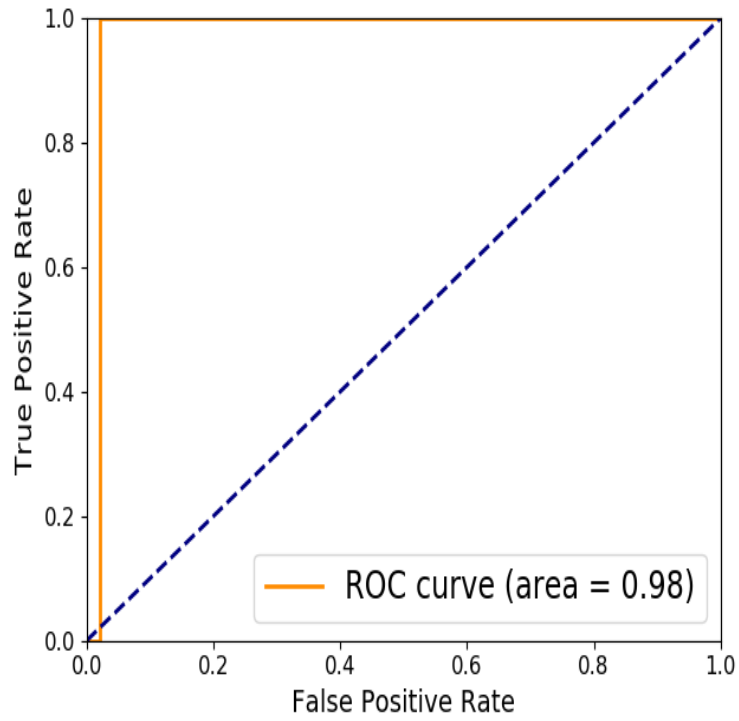
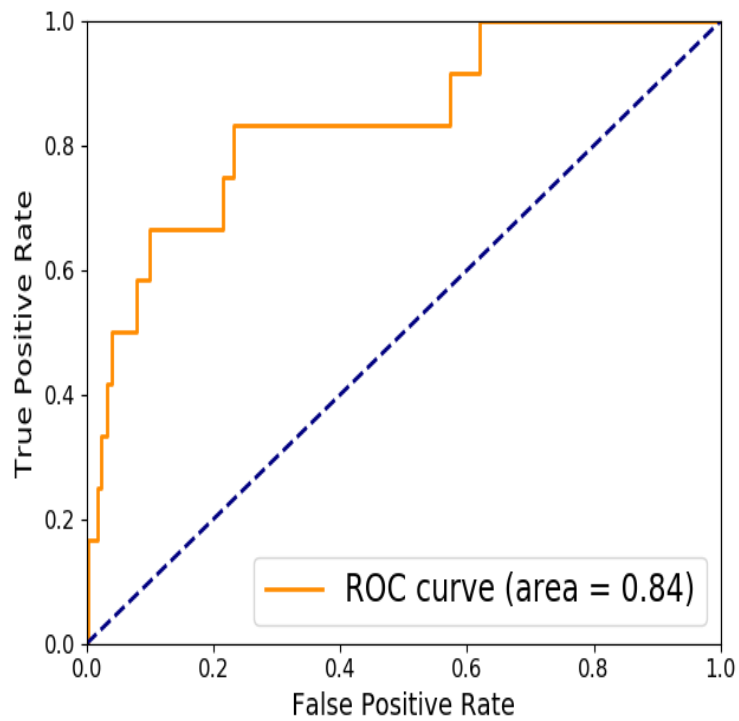


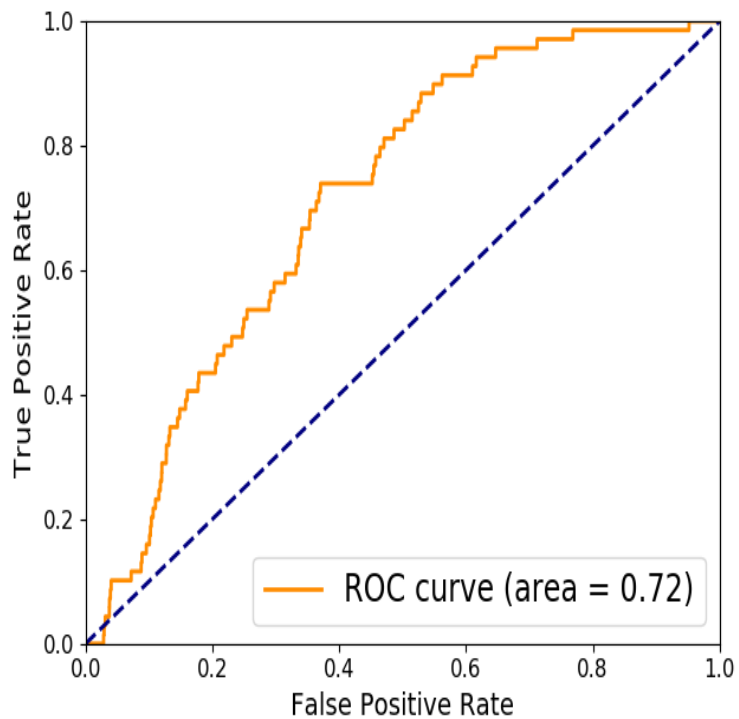
Diffusion ROC from fGOnly_tOverlap STRING10_combined.png



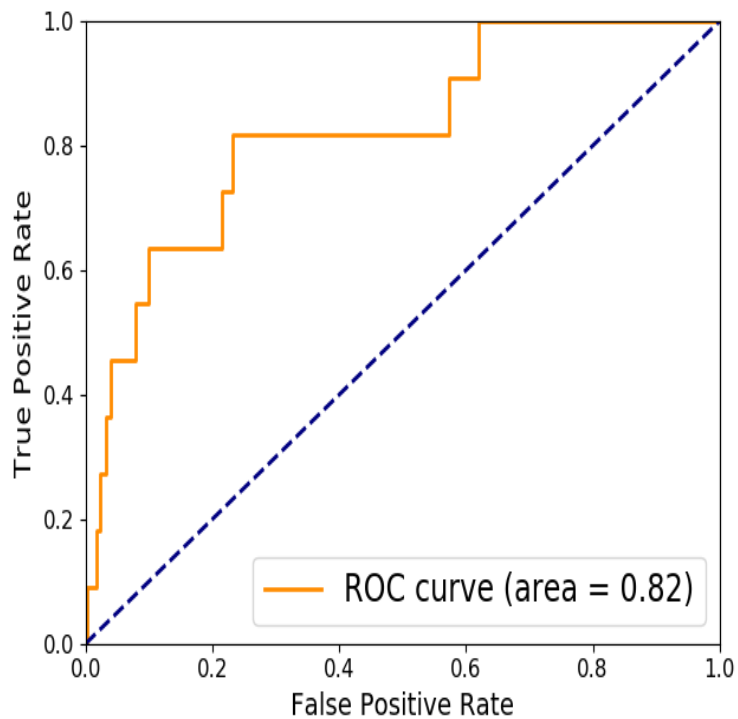
Diffusion ROC from fTestOnly_tGS STRING10_combined.png



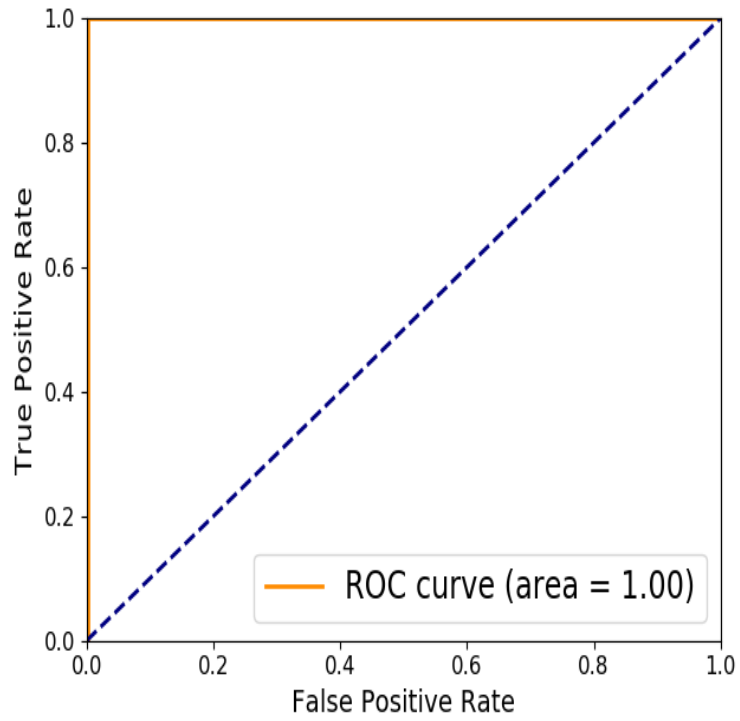
Diffusion ROC from fGOnly_tTestOnly STRING10_combined.png



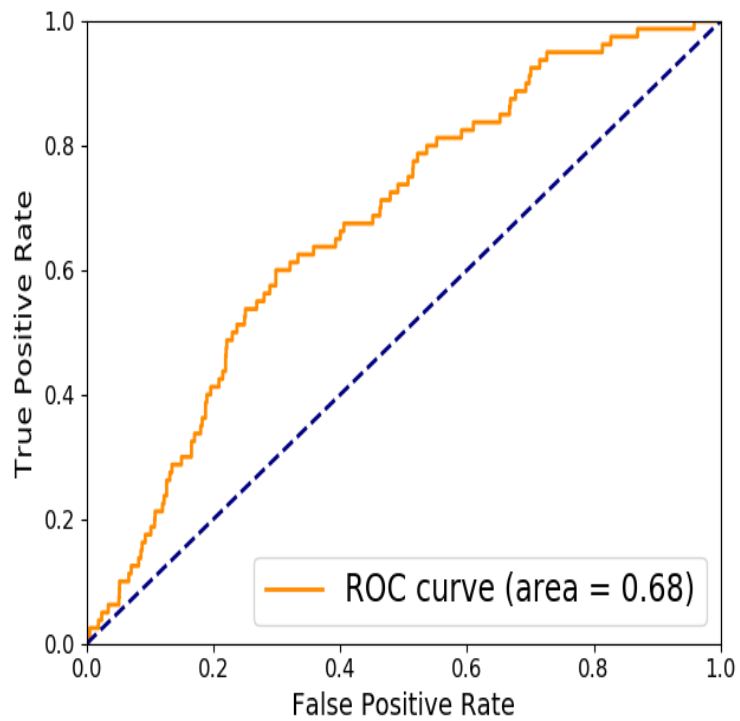
Diffusion ROC from fTestOnly_tGOnly STRING10_combined.png



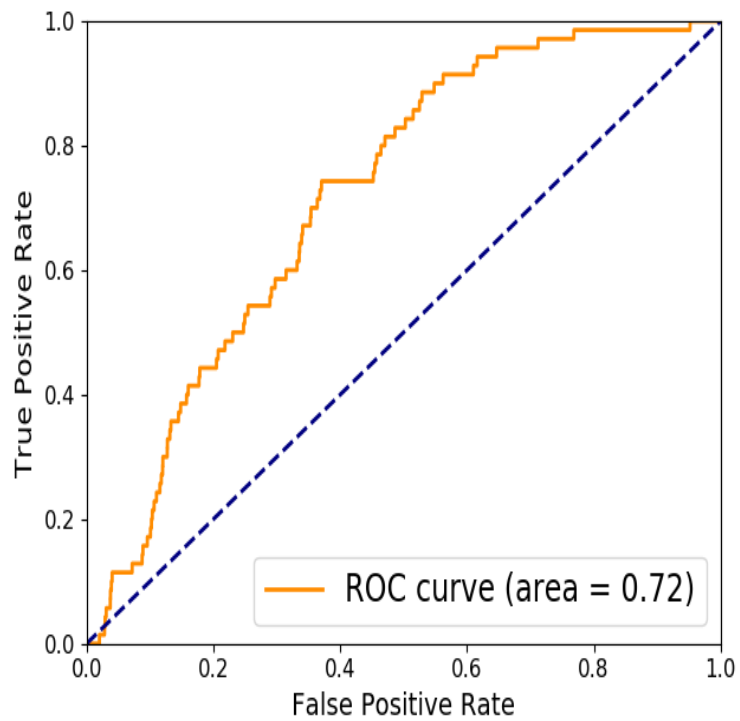
Diffusion ROC from fTestOnly_tOverlap STRING10_combined.png



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 | 0.715 | 4.190 | 6.100 | 4.079 | 3.301 |
| Test_only | GS_only | 0.824 | 1.767 | 3.8711956 | 1.410 | 1.638 |
| | | | 74653578 | | | |
| | | | (invalid) | | | |
| GS_only | Overlap | 0.980 | -68875275 | 1.6041022 | 0.5950386 | 1.298550 |
| | | | 200110.0 | 320118739 | 66321028 | 64943441 |
| | | | (invalid) | (invalid) | (invalid) | 68 (inva |
| | | | | | | lid) |
| GS_only | Test | 0.719 | 4.074 | 6.541 | 3.783 | 2.939 |
| Test_only | Overlap | 0.997 | -11748929 | 1.6860494 | 0.6642160 | 1.047940 |
| | | | 425469.0 | 66290462 | 600829407 | 42955062 |
| | | | (invalid) | (invalid) | (invalid) | 5 (inval |
| | | | | | | id) |
| Test_only | GS | 0.839 | 1.995 | 3.932 | 1.475 | 1.861 |
| Overlap | Onlies | 0.677 | 1.918 | 5.131 | -20167156 | 1.144 |
| | | | | 7495296.5 | | |
| | | | | (invalid) | | |

| ** | #Mapped | #Total | Not mapped genes |
|-----------|---------|--------|----------------------------|
| GS | 11 | 12 | MYCL |
| | | | N/A |
| Test_only | 69 | 73 | PCDHGA6CFAP44;MK167;GPRIN2 |
| | | | N/A |
| Overlap | 1 | 1 | TP53 |

(end of file)