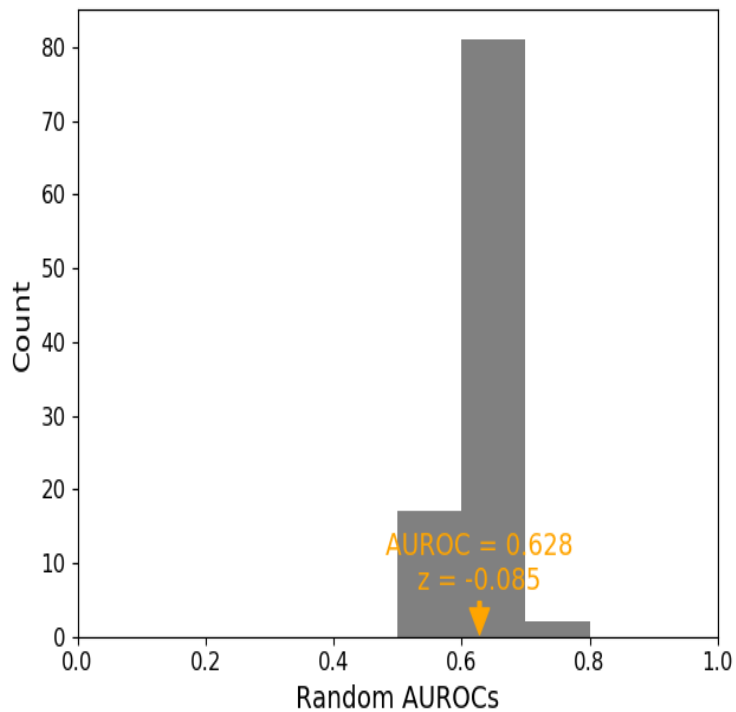
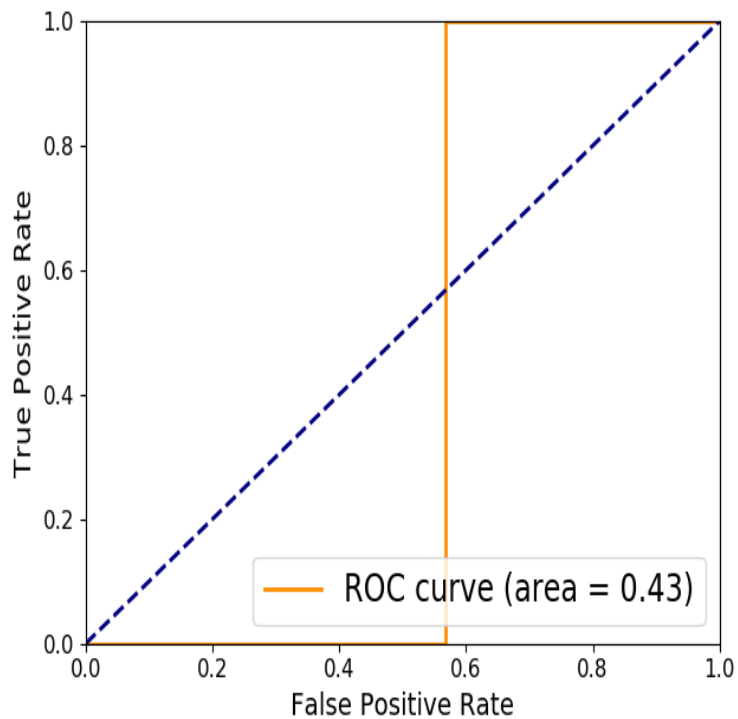


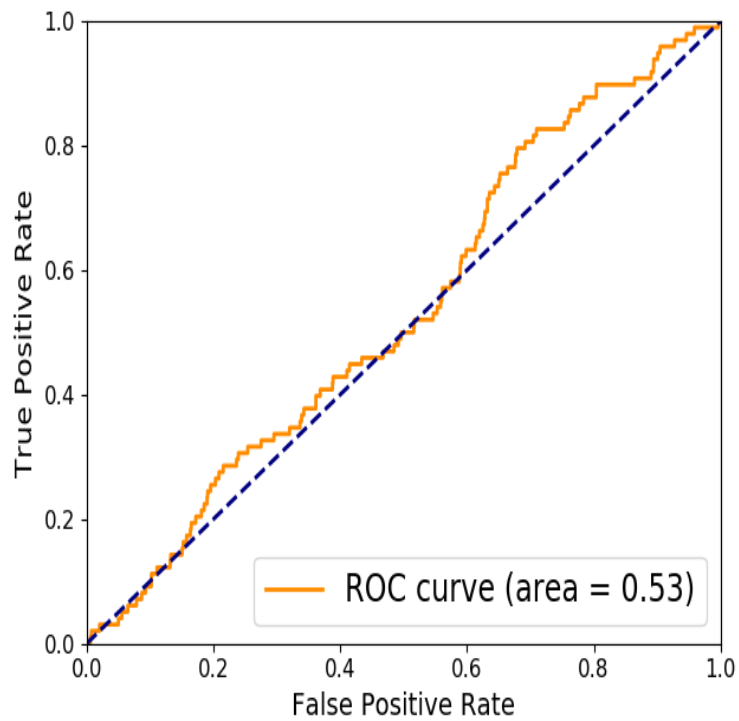
show_22_Zf_degree-matched_from ADSP Only to Martinet.png



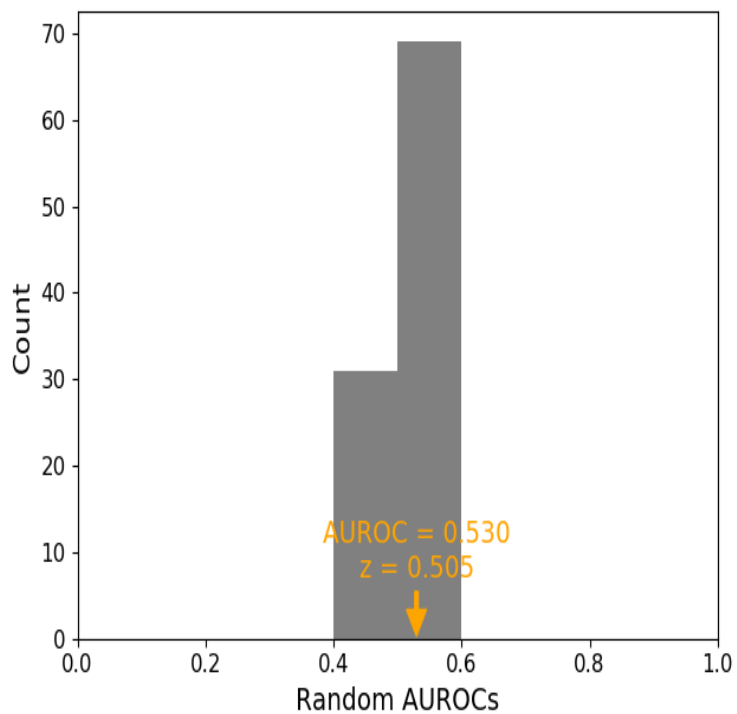
Diffusion ROC from ADSP Only to Overlapped STRING10_combined.png



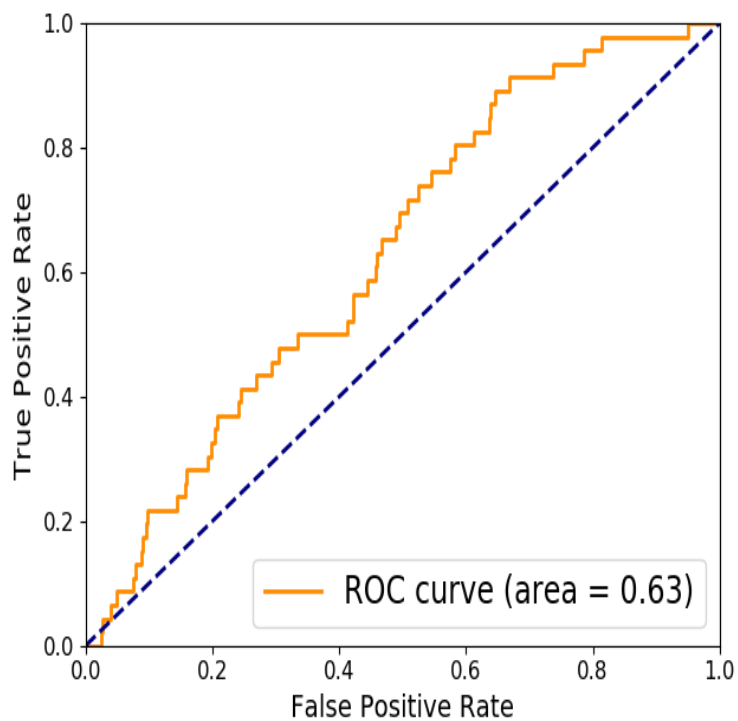
Diffusion ROC from Overlapped to Non-overlapped in both groups STRING



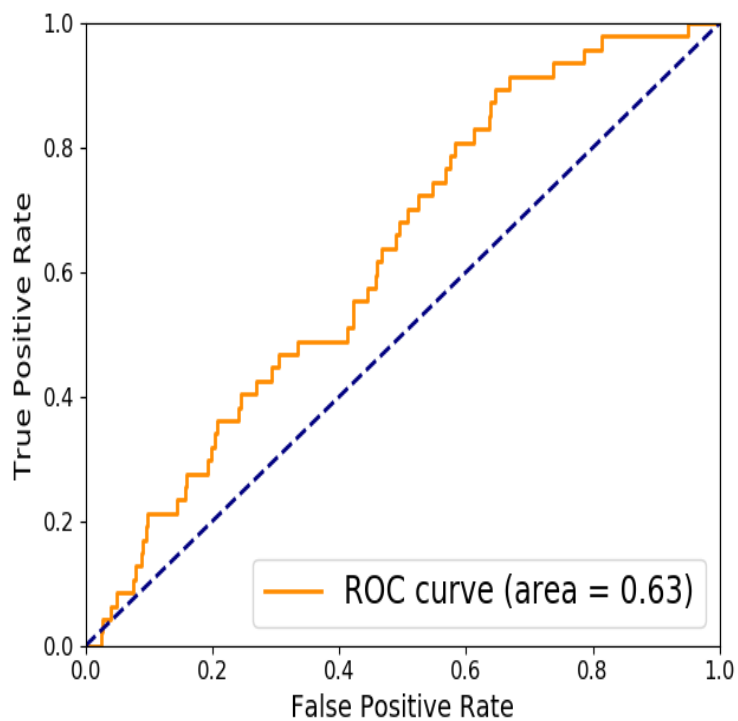
show_11_Zt_degree-matched_from Martinet Only to ADSP.png



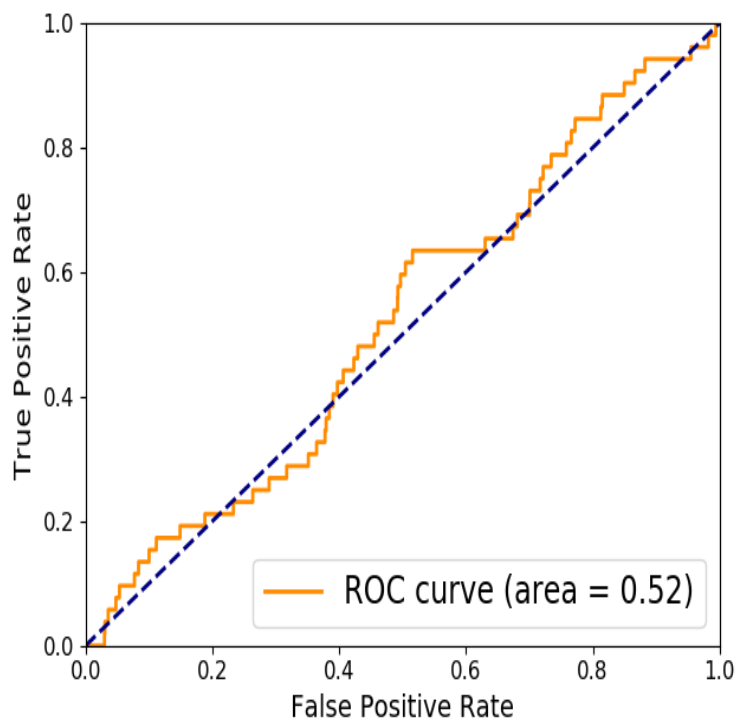
Diffusion ROC from ADSP Only to Martinet Only STRING10_combined.png



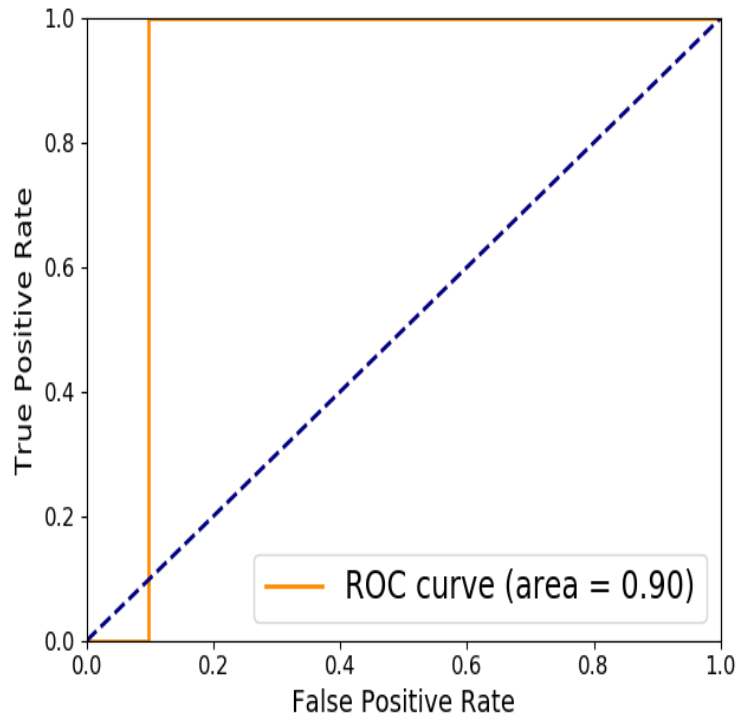
show_2_Diffusion ROC from ADSP Only to Martinet STRING10_combined.



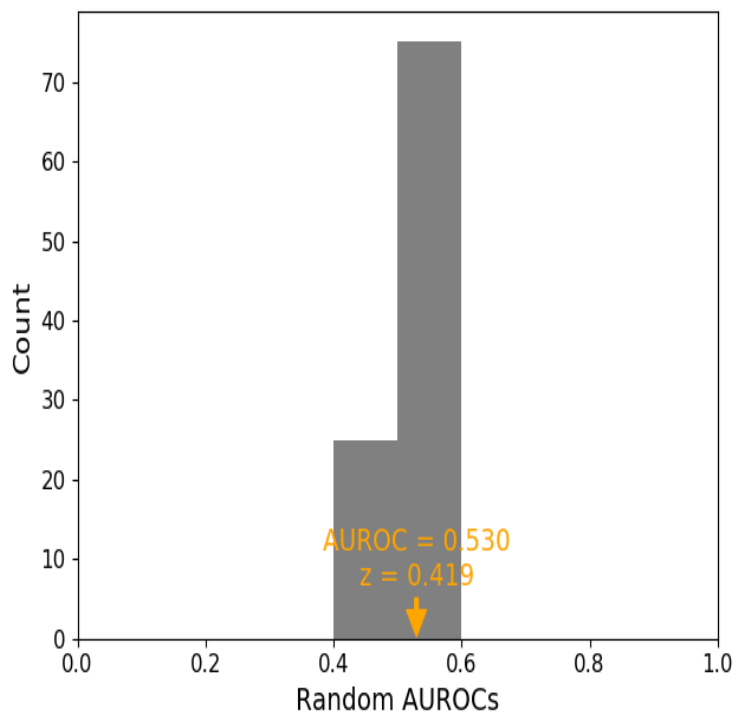
Diffusion ROC from Martinet Only to ADSP Only STRING10_combined.png



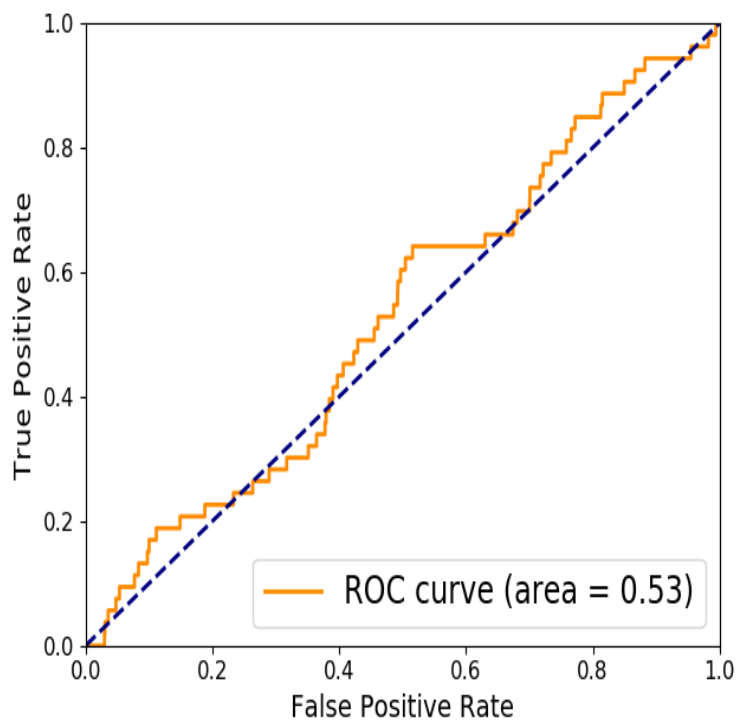
Diffusion ROC from Martinet Only to Overlapped STRING10_combined.png



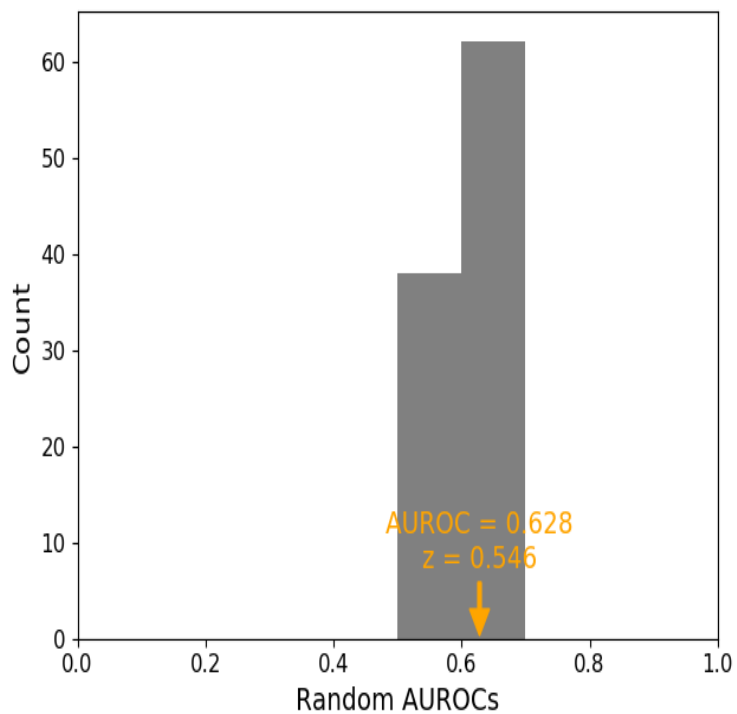
show_12_Zf_degree-matched_from Martinet Only to ADSP.png



show_1_Diffusion ROC from Martinet Only to ADSP STRING10_combined.



show_21_Zt_degree-matched_from ADSP Only to Martinet.png



Diffusion FROM	Diffusion TO	AUROC	Zt (degree-matched)	Zt (uniform)	Zf (degree-matched)	Zf (uniform)
Martinet only	ADSP	0.530	0.505	0.584	0.419	0.857
ADSP only	Martinet	0.628	0.546	3.187	-0.085	0.207
Martinet only	ADSP only	0.522	0.500	0.829 *	0.172	0.928
ADSP only	Martinet only	0.632	0.901 *	3.073	0.028	0.403
Martinet only	Overlappe d	0.902	1.395 *	1.300 *	0.686 *	1.177 *
ADSP only	Overlappe d	0.432	-1.561 *	-0.381 *	-2.000 *	-1.862 *
Overlappe d	Non-overlapped genes in both groups	0.534	-0.335	1.248	-0.768 *	-0.908

Z-scores are computed for the experimental AUROC based on distributions of the random AUROCs
Zf : when randomizing source genes (Diffusion FROM)
Zf : when randomizing recipient genes (Diffusion TO)
Random genes are selected either uniformly or degree matched
* : Distribution of random AUROC is not Gaussian

**	#Mapped	#Total	Not mapped genes
Martinet Only	46	47	cDNA FLJ60385 N/A
ADSP Only	52	53	IQCF6 N/A
Overlapped	1	1	NOP56

(end of file)