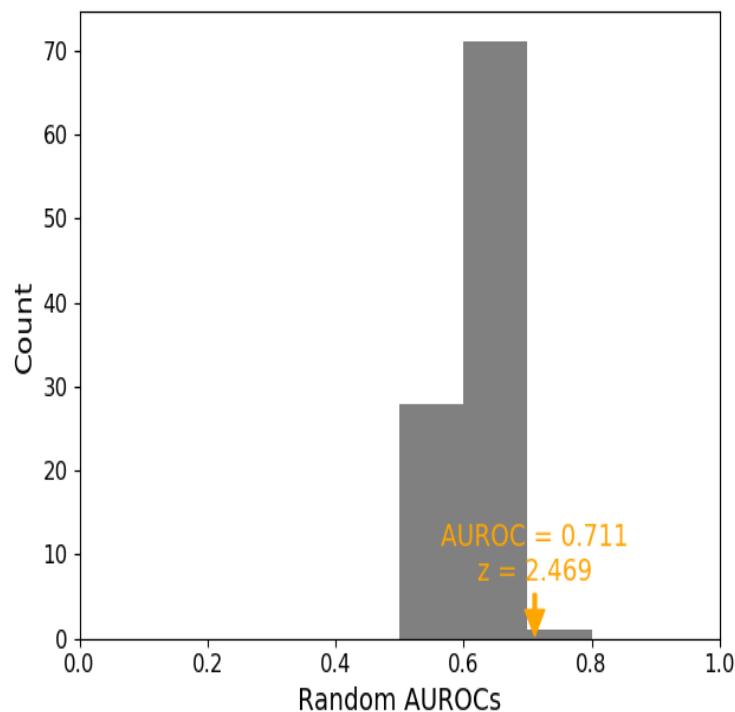
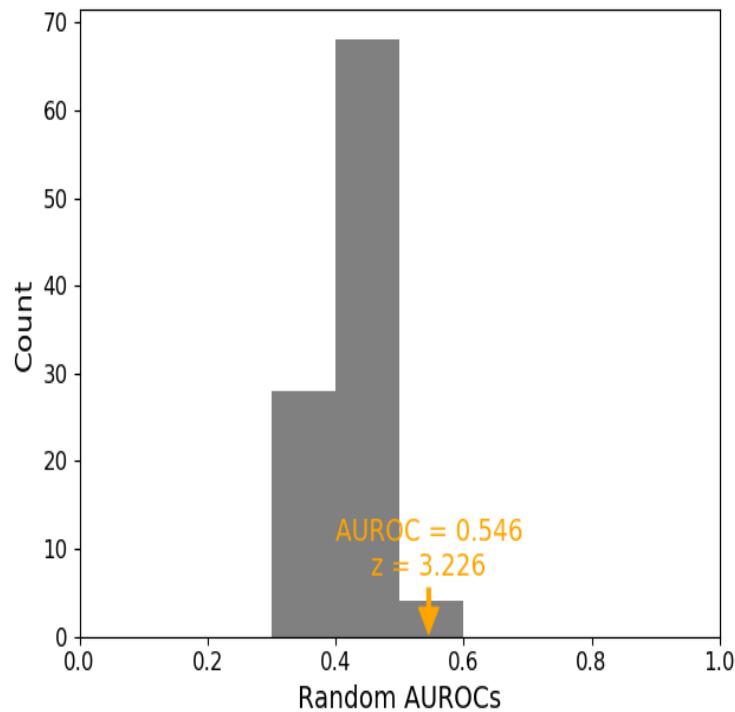


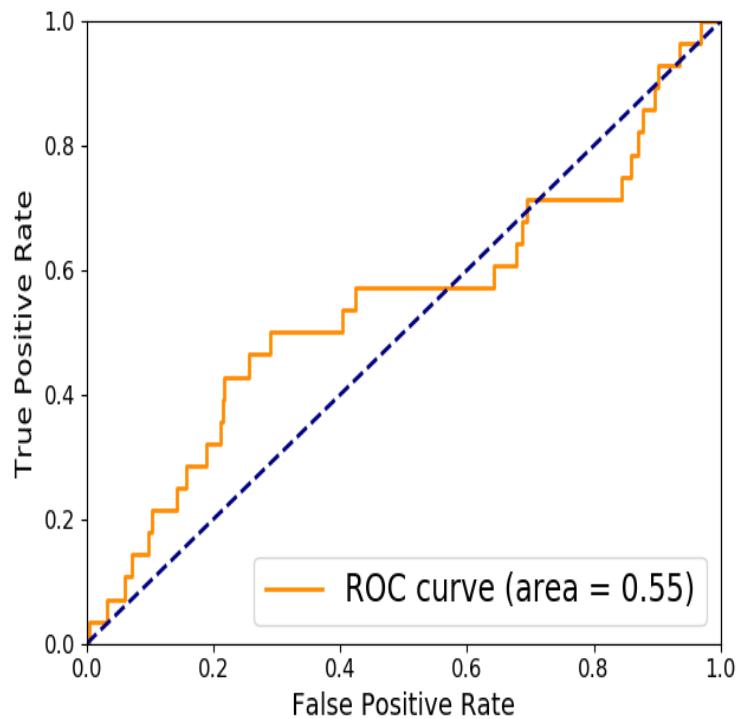
**show\_22\_Zf\_degree-matched\_from ADNI to Martinet.png**



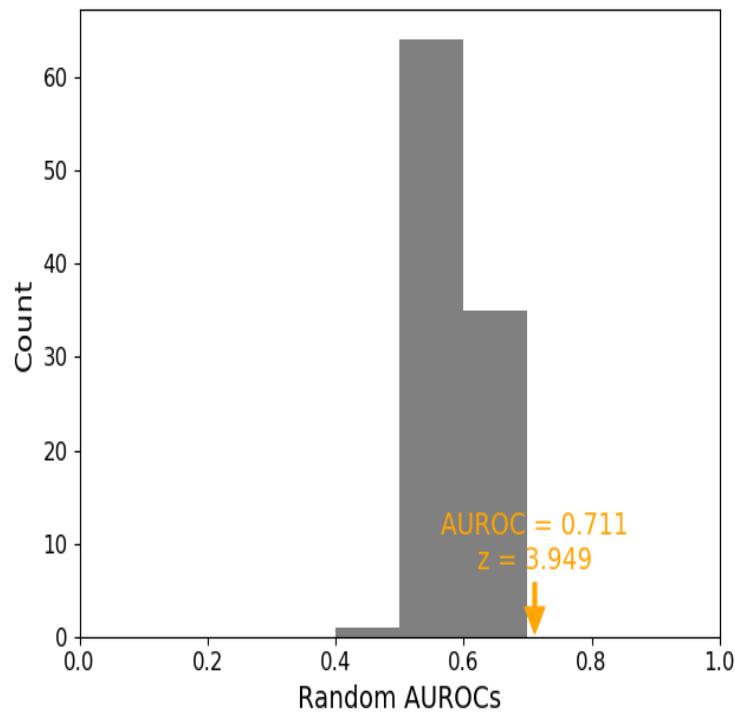
## show\_12\_Zf\_degree-matched\_from Martinet to ADNI.png



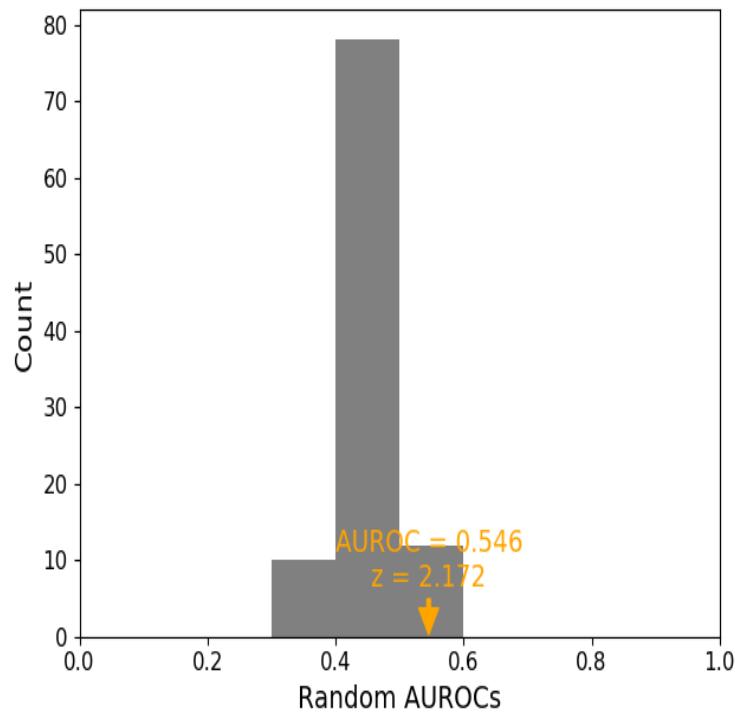
**show\_1\_Diffusion ROC from Martinet to ADNI STRING10\_combined.png**



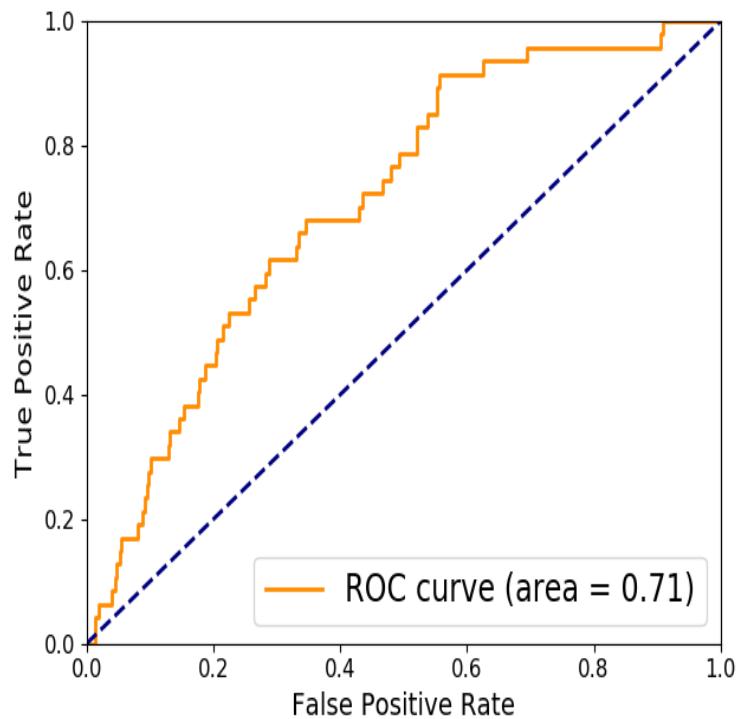
**show\_21\_Zt\_degree-matched\_from ADNI to Martinet.png**



## show\_11\_Zt\_degree-matched\_from Martinet to ADNI.png



## show\_2\_Diffusion ROC from ADNI to Martinet STRING10\_combined.png



<i>Diffusion</i>	<i>Diffusion</i>	<i>AUROC</i>	<i>Zt</i>	<i>Zt</i>	<i>Zf</i>	<i>Zf (unif)</i>
<i>FROM</i>	<i>TO</i>		(degree- matched)	(uniform)   matched)	(degree- matched)	orm)
<i>Martinet</i>	<i>ADNI</i>	0.546	2.172	0.622	3.226	2.443
<i>ADNI</i>	<i>Martinet</i>	0.711	3.949	5.626	2.469	2.405 *

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

<i>**</i>	<i>#Mapped</i>	<i>#Total</i>	<i>Not mapped genes</i>	
<i>Martinet</i>	47	48	cDNA FLJ60385	N/A
<i>ADNI</i>	28	28		N/A

(end of file)