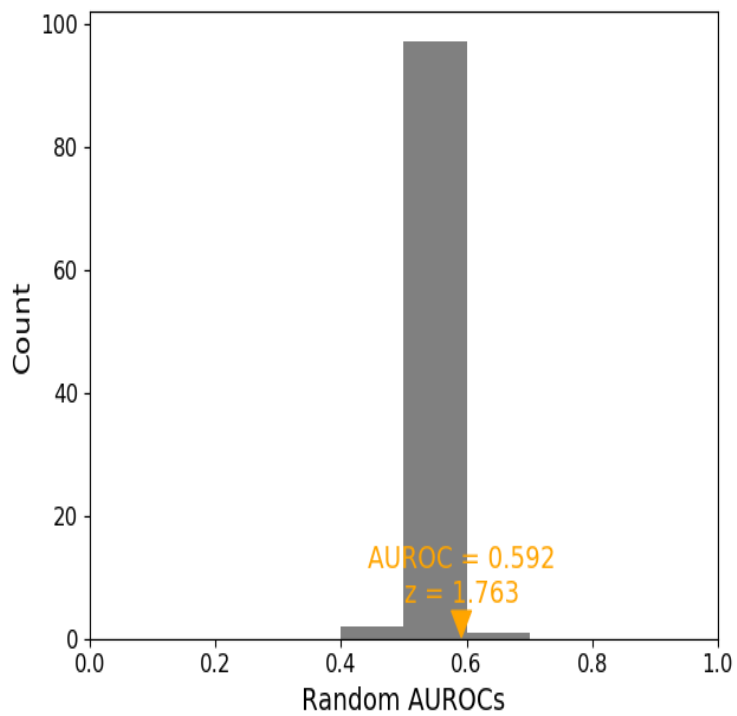
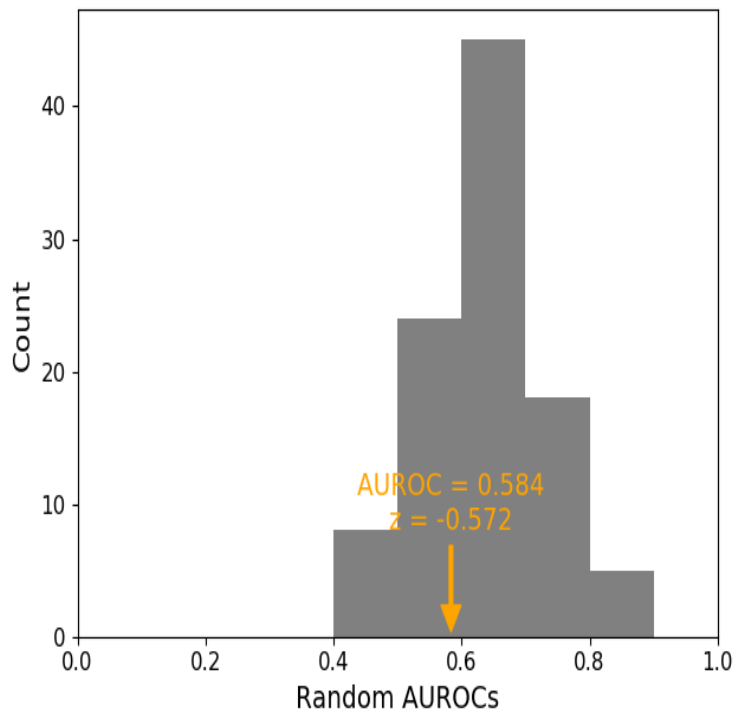


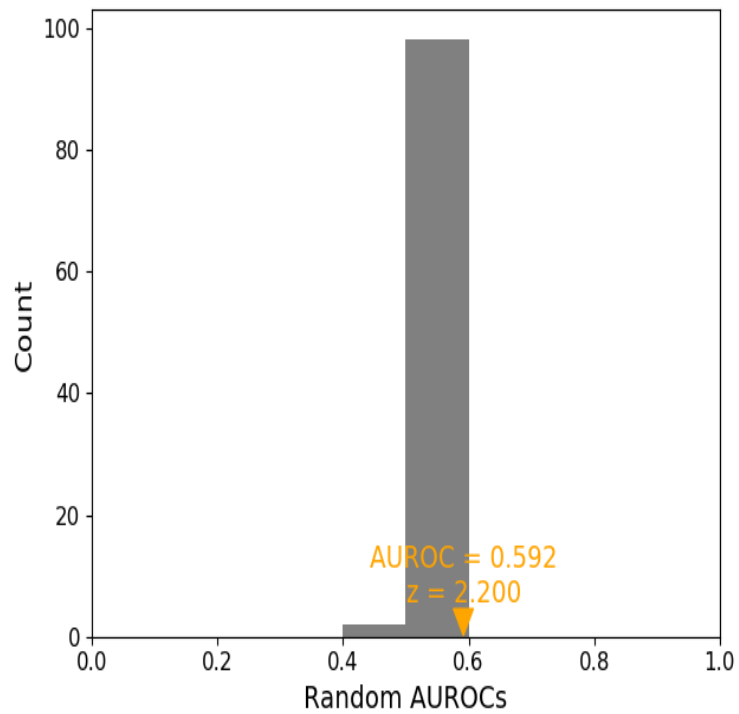
# show\_21\_Zt\_degree-matched\_from Ismael to iDEAL.png



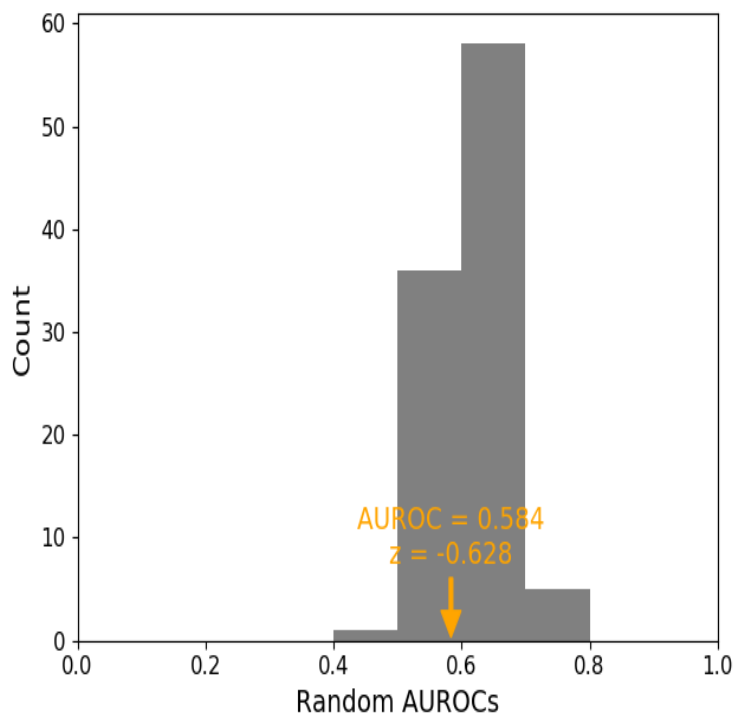
# show\_12\_Zf\_degree-matched\_from iDEAL to Ismael.png



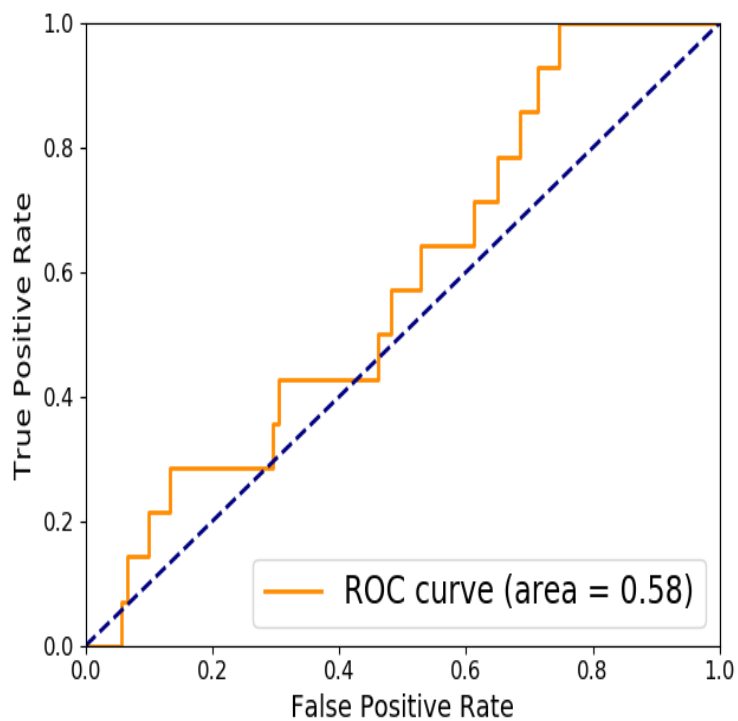
# show\_22\_Zf\_degree-matched\_from Ismael to iDEAL.png



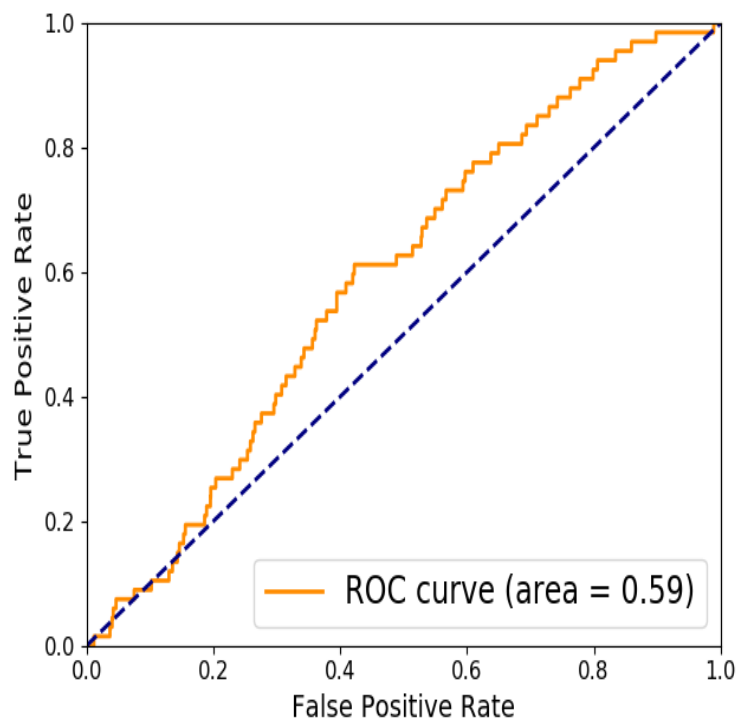
# show\_11\_Zt\_degree-matched\_from iDEAL to Ismael.png



# show\_1\_Diffusion ROC from iDEAL to Ismael STRING10\_combined.png



## show\_2\_Diffusion ROC from Ismael to iDEAL STRING10\_combined.png



Diffusion	Diffusion	AUROC	Zt	Zt	Zf	Zf (unif
FROM	TO		(degree-	(uniform)	(degree-	orm)
		matched)		matched)		
iDEAL	Ismael	0.584	-0.628	1.001	-0.572	-0.452
Ismael	iDEAL	0.592	1.763	2.585	2.200	2.181 *

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
iDEAL	67	68	SLC35G6   N/A
Ismael	14	14	N/A

(end of file)