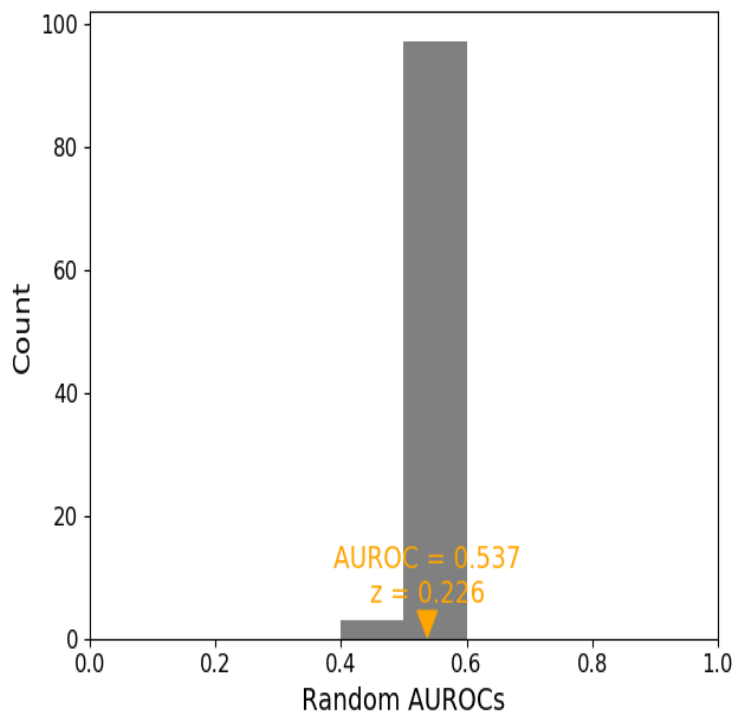
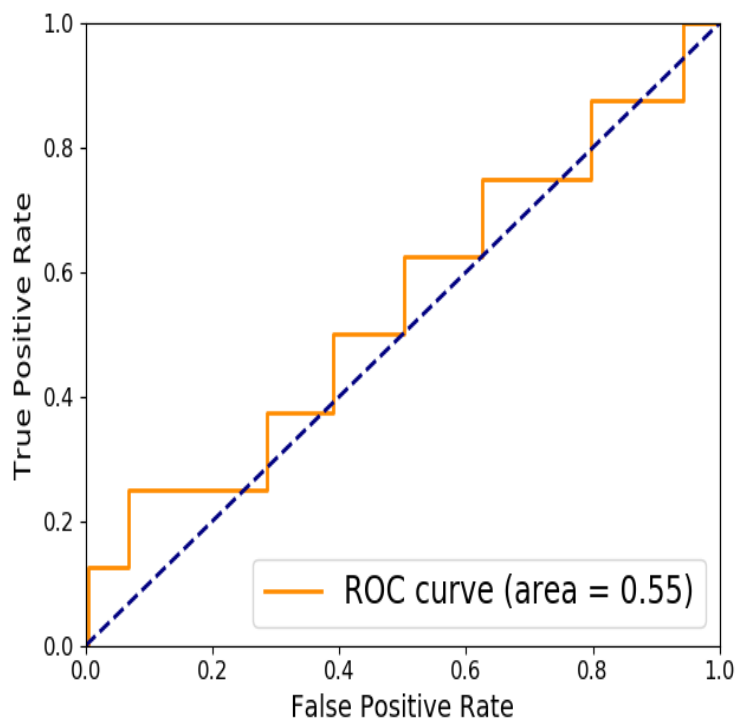


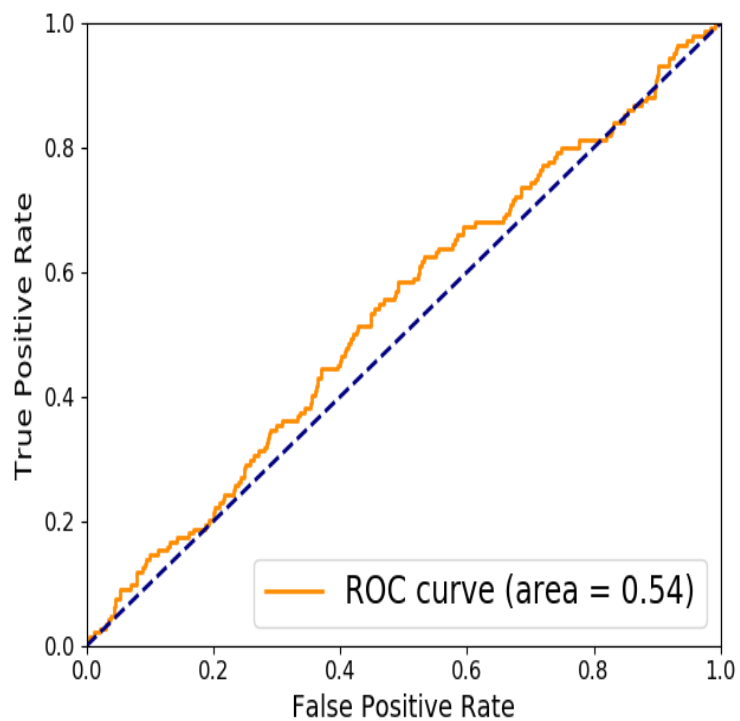
show_21_Zt_degree-matched_from Marttinen_protein_decisionTree to AD



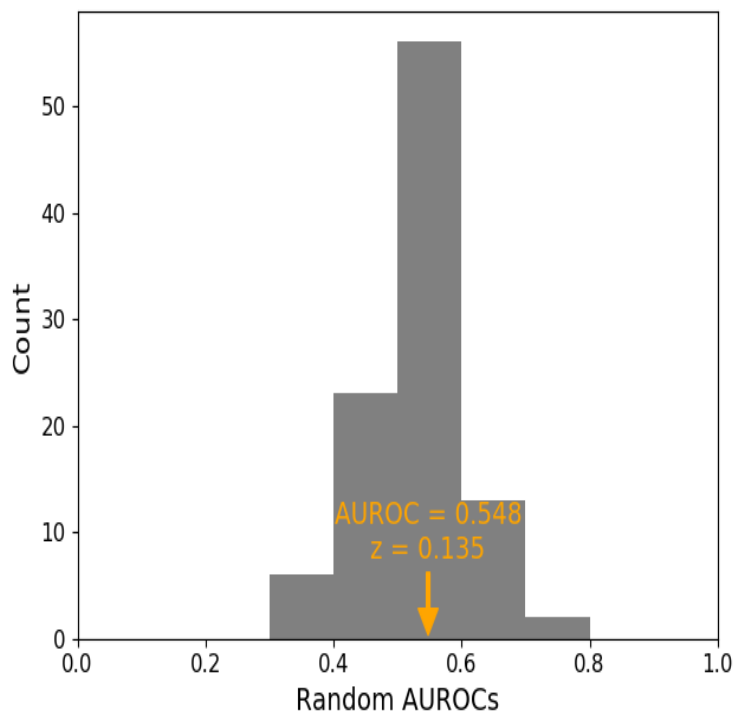
show_1_Diffusion ROC from ADSP_iDEAL to Marttinen_protein_decisionT



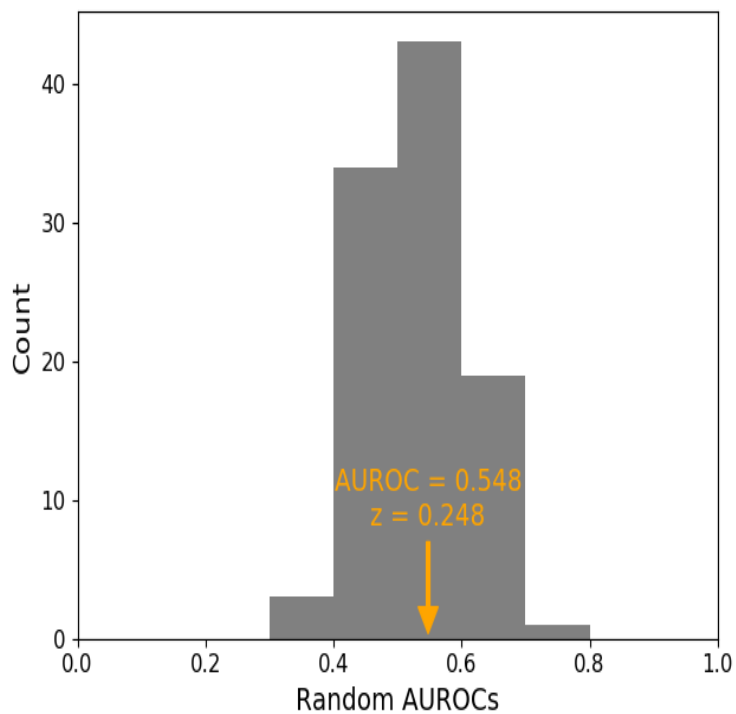
show_2_Diffusion ROC from Marttinen_protein_decisionTree to ADSP_iDE



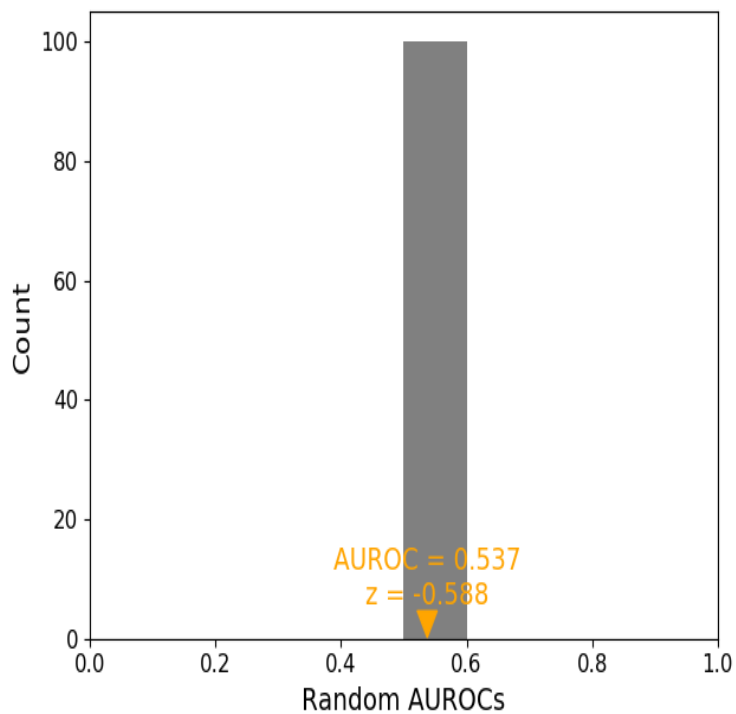
show_12_Zf_degree-matched_from ADSP_iDEAL to Marttinen_protein_dec



show_11_Zt_degree-matched_from ADSP_iDEAL to Marttinen_protein_dec



show_22_Zf_degree-matched_from Marttinen_protein_decisionTree to AD



Diffusion FROM	Diffusion TO	AUROC	Zt (degree-matched)	Zt (uniform)	Zf (degree-matched)	Zf (uniform)
ADSP_iDEA	Marttinen	0.548	0.248	0.226	0.135	0.234
L	_protein_					
	decisionT					
	ree					
Marttinen	ADSP_iDEA	0.537	0.226	1.438	-0.588	-0.048
protein	L					
decisionT						
ree						

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
ADSP_iDEAL	144	148	MAL2;VSTM5;DCDC1;MGME1 N/A
Marttinen_protein_decision	8	8	N/A
Tree			

(end of file)