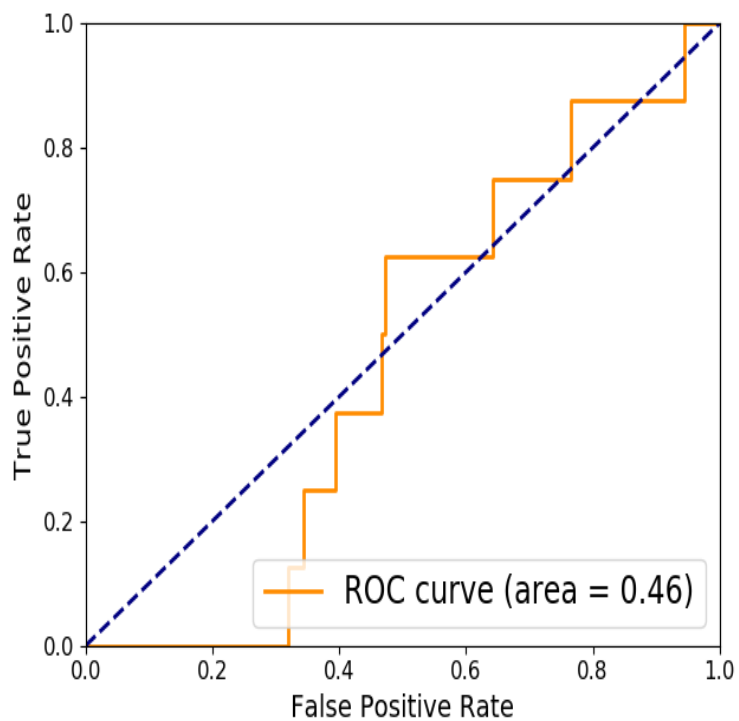
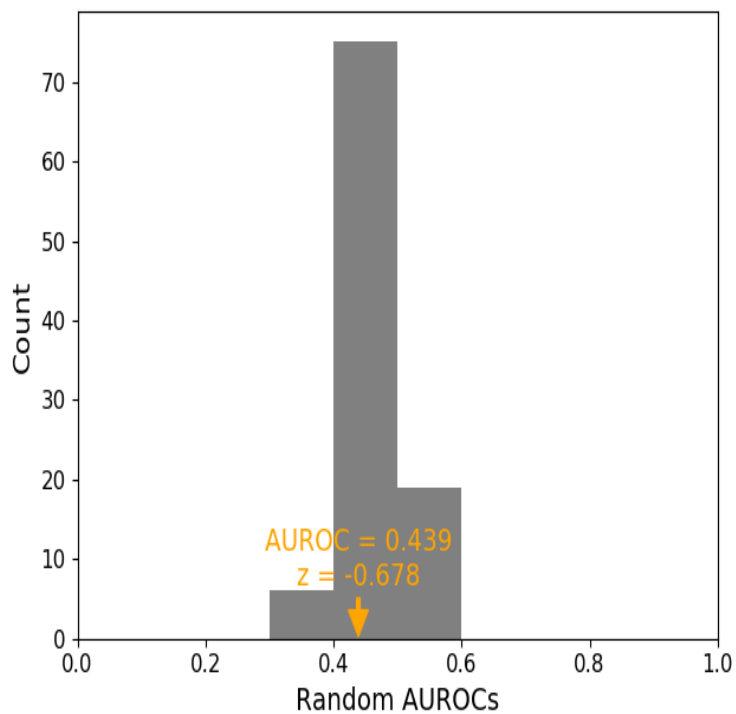


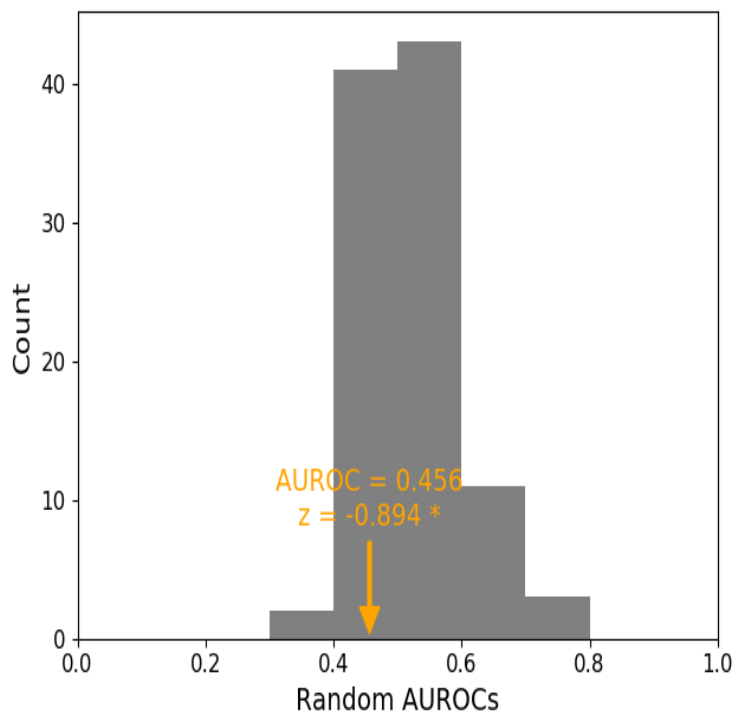
show_1_Diffusion ROC from ADNI to Marttinen_protein_decisionTree STR



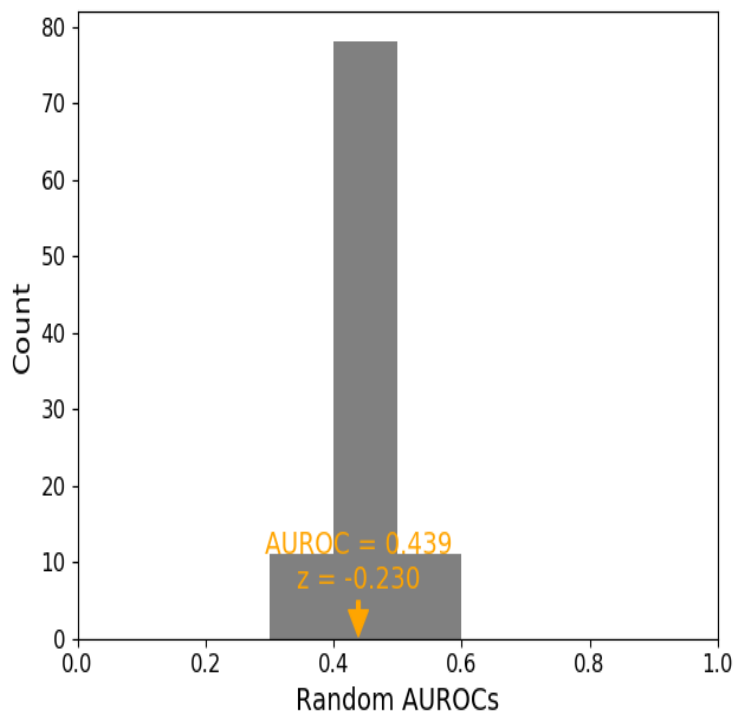
show_22_Zf_degree-matched_from Marttinen_protein_decisionTree to AD



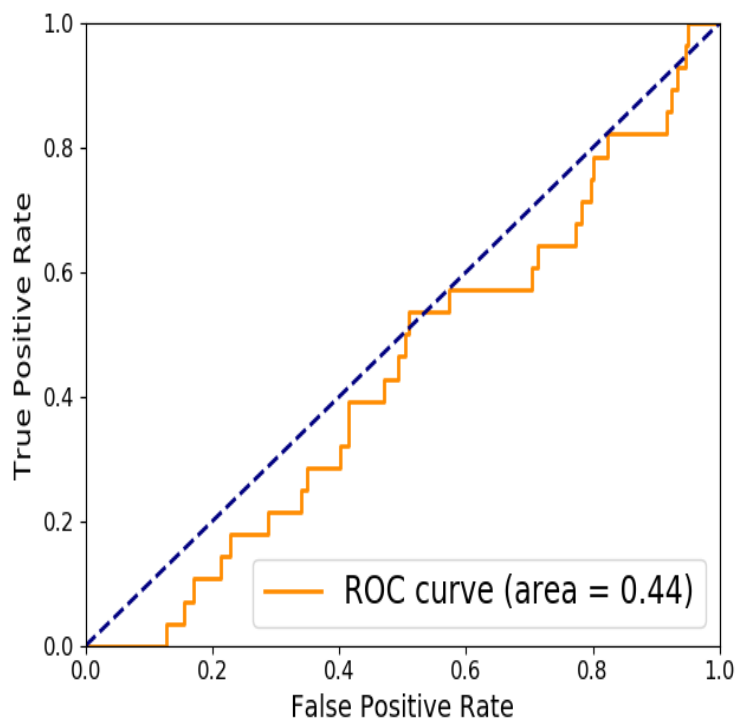
show_12_Zf_degree-matched_from ADNI to Marttinen_protein_decisionTree



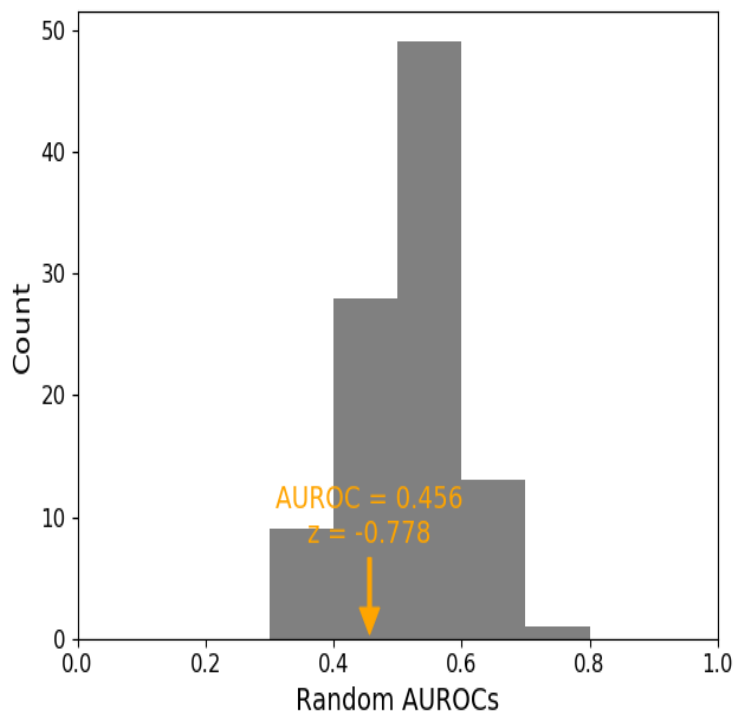
show_21_Zt_degree-matched_from Marttinen_protein_decisionTree to AD



show_2_Diffusion ROC from Marttinen_protein_decisionTree to ADNI STR



show_11_Zt_degree-matched_from ADNI to Marttinen_protein_decisionTree



Diffusion FROM	Diffusion TO	AUROC	Zt (degree-matched)	Zt (uniform)	Zf (degree-matched)	Zf (uniform)
ADNI	Marttinen	0.456	-0.778	-0.602	-0.894 *	-0.975 *
protein	decisionT					
ree						
Marttinen	ADNI	0.439	-0.230	-1.186	-0.678	-0.392
protein						
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
ADNI	28	28	N/A
Marttinen_protein_decisionTree	8	8	N/A

(end of file)