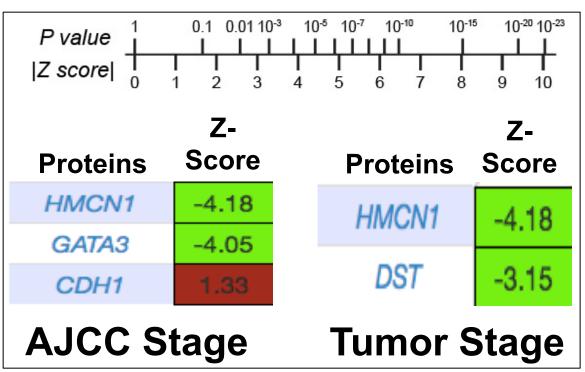
P value 1	0.1 0.01	 10 ⁻⁷ 10 ⁻¹⁰	10 ⁻¹⁵ 10 ⁻²⁰ 10 ⁻²³ 8 9 10
Genes	Z- Score	Genes	Z- Score
GABARAP	-4.45	KIAA0040	-6.05
CACNA1D	-1.96	RTN4	-1.11
LASS2	-1.6	ACADVL	-0.81
KIAA0556	-1.48	NME4	1.82
FBLN2	-1.1	B3GNT3	1.98
ARSA	-1.08	SFPQ	2.04
EFEMP2	-1.04	ARPC1A	3.41
PPIE	0.01		
ZNF37A	0.22	POR	4.44
KIAA1609	4.18	CKS2	5.11
POR	4.44	KARS	7.12
AJCC St		umor	Stage

Key Genes identified by RFE analysis critical to transcriptomics models predicting AJCC Stage and Tumor Stage. Above are the correlations with clinical outcome as defined by the PRECOG database



Key Proteins identified by RFE analysis critical to proteomics models predicting AJCC Stage and Tumor Stage. Above are the correlations with clinical outcome as defined by the PRECOG database