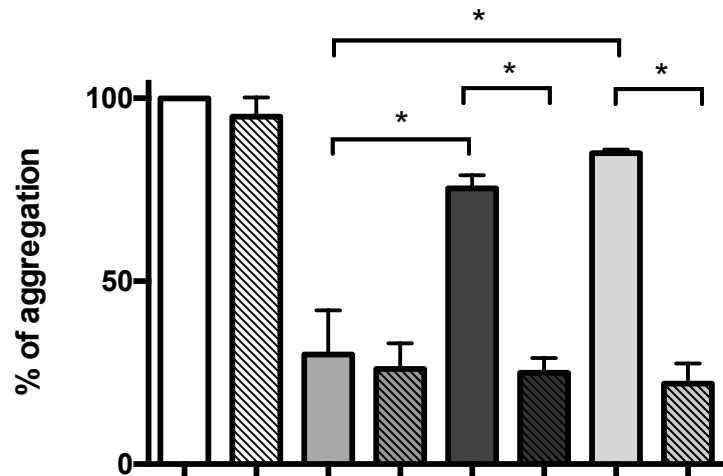
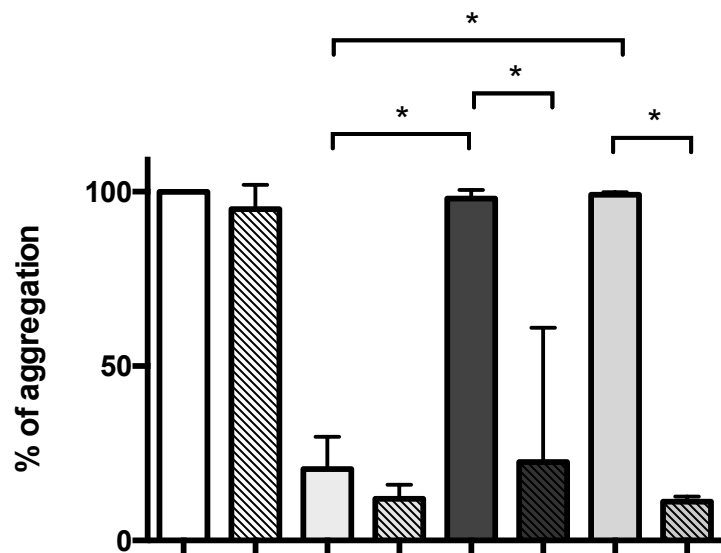
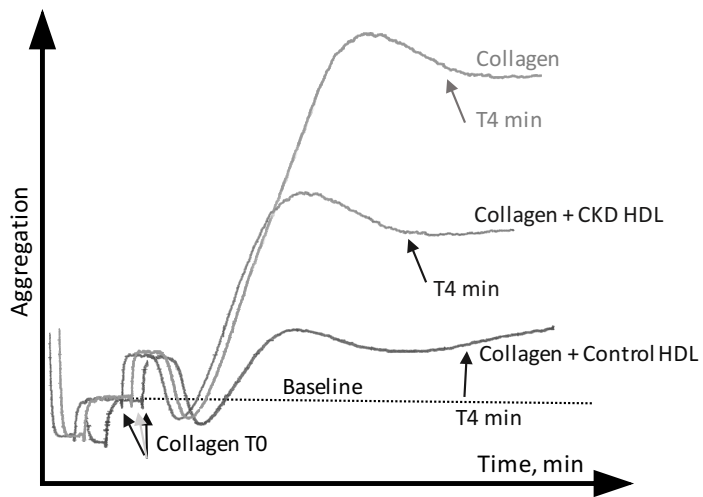
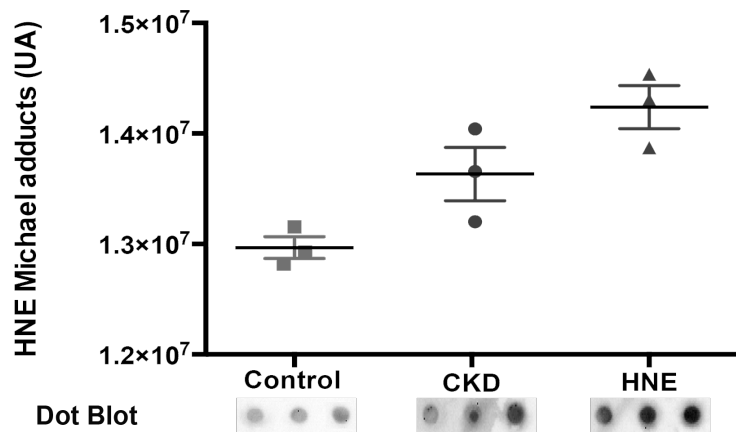


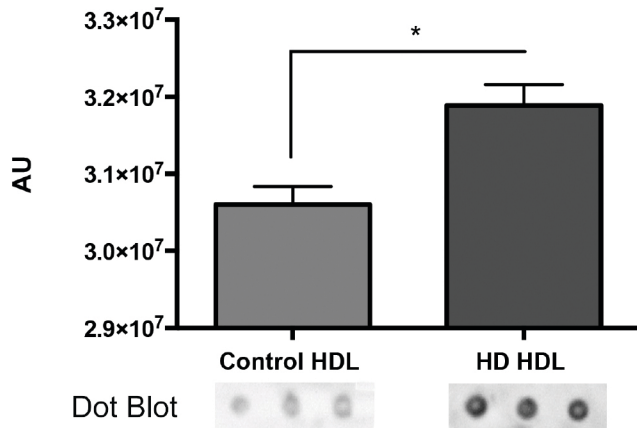
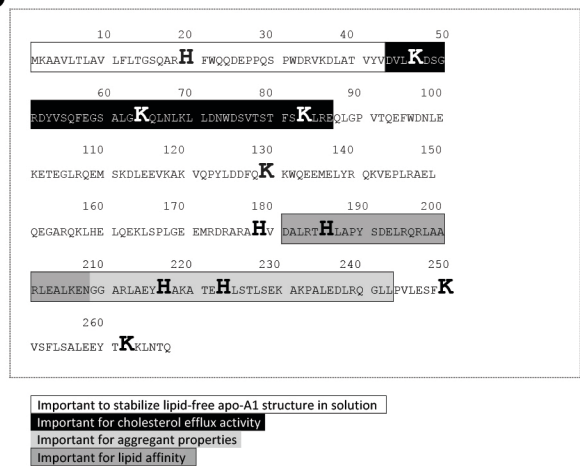
A

Collagen	+	+	+	+	+	+	+
CD36 Ab	-	+	-	+	-	+	-
Control HDL	-	-	+	+	-	-	-
CKD HDL	-	-	-	-	+	+	-
HNE HDL	-	-	-	-	-	-	+

B

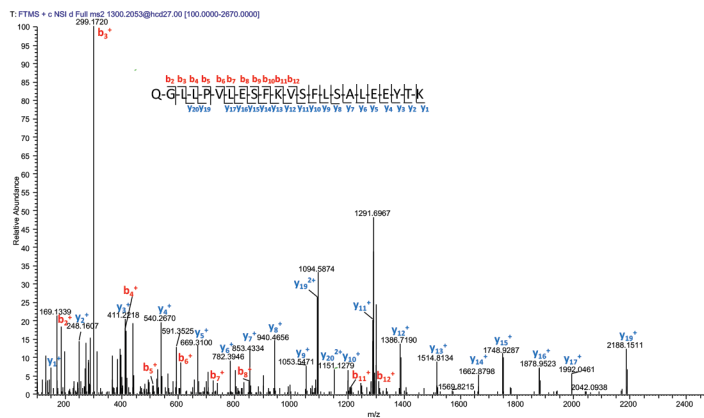
Collagen	+	+	+	+	+	+	+
CD36 Ab	-	+	-	+	-	+	-
Control LPP	-	-	+	+	-	-	-
CKD LPP	-	-	-	-	+	+	-
HNE LPP	-	-	-	-	-	-	+

C**D**

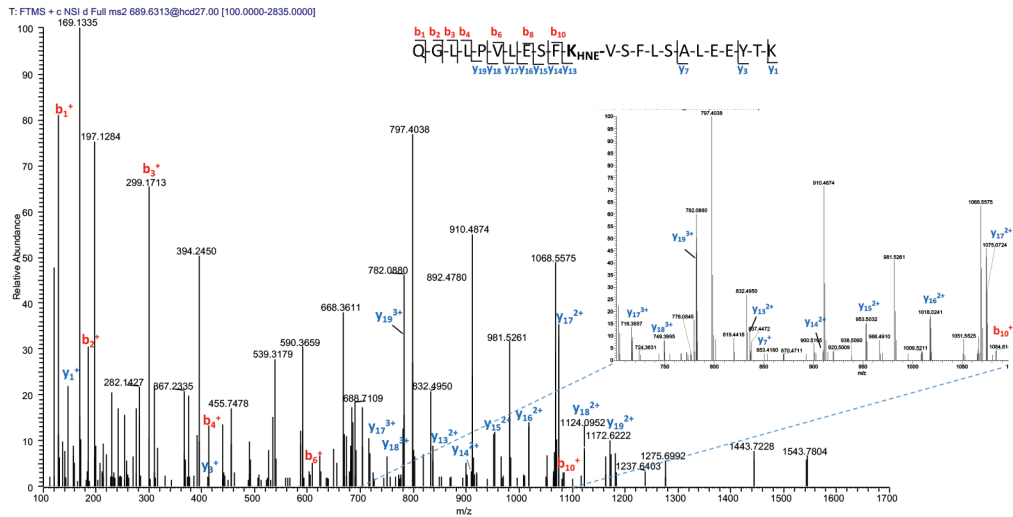
A**B****C**

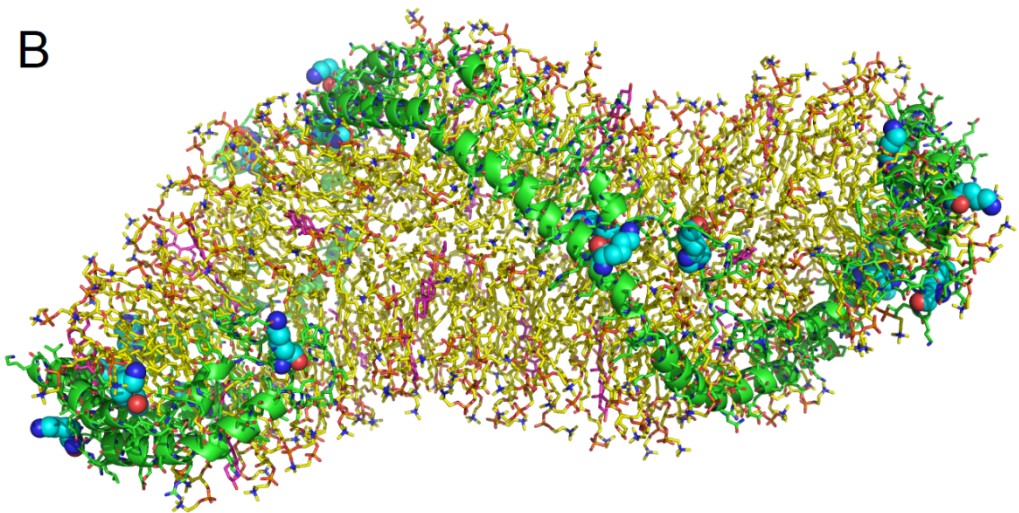
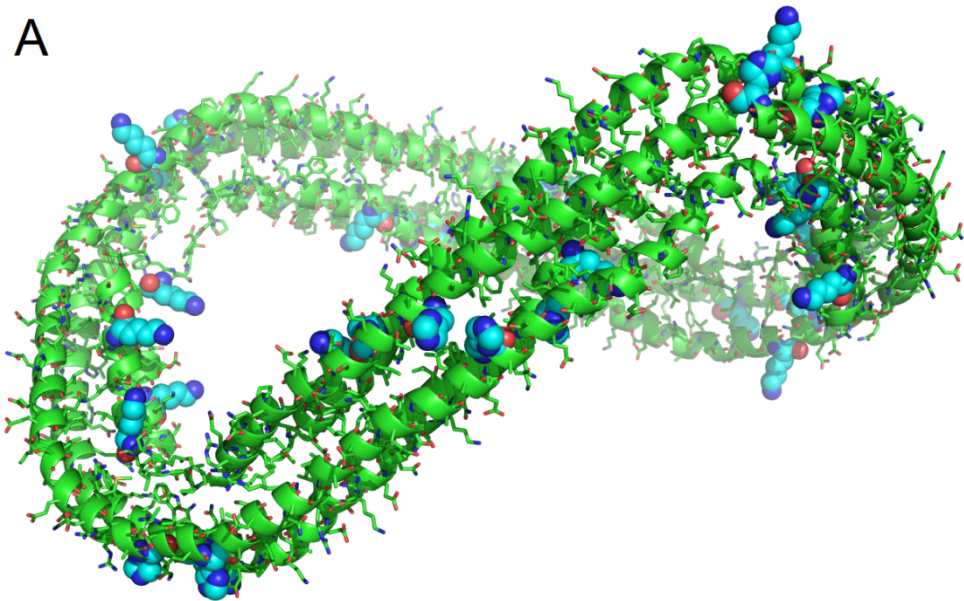
Sequence	Modifications	Theo. MH+ [Da]	Abundances (Control)	Abundances (HD)	Abundance Ratio	P-Value
[R]LQLLPVLESFKVLSLSALEEYTK[K]	1xHNE [K11]	2754.52156	28.2	171.8	5.3	0.009

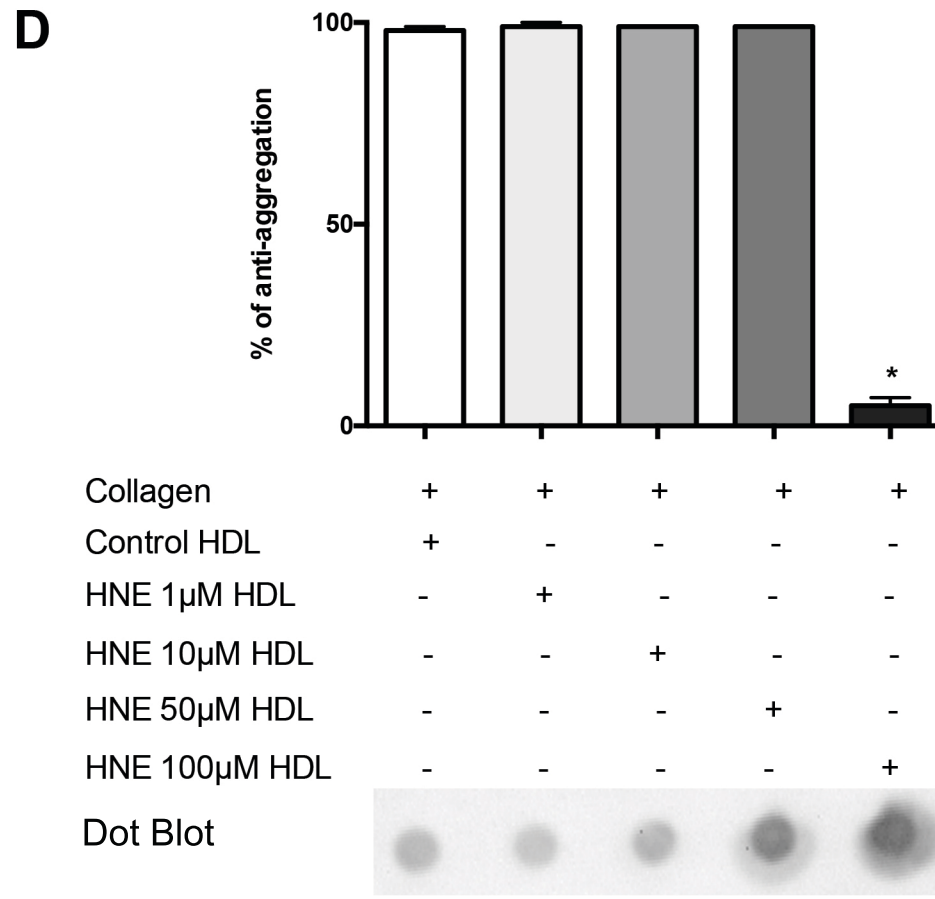
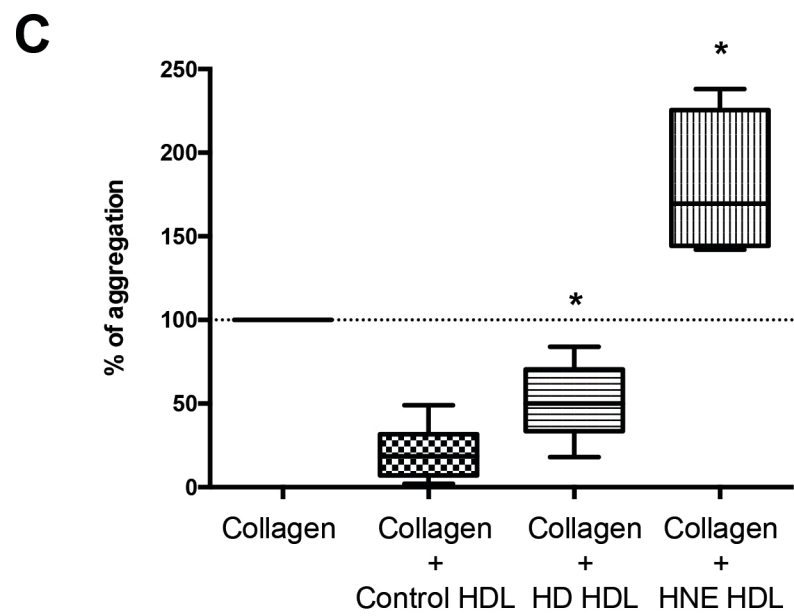
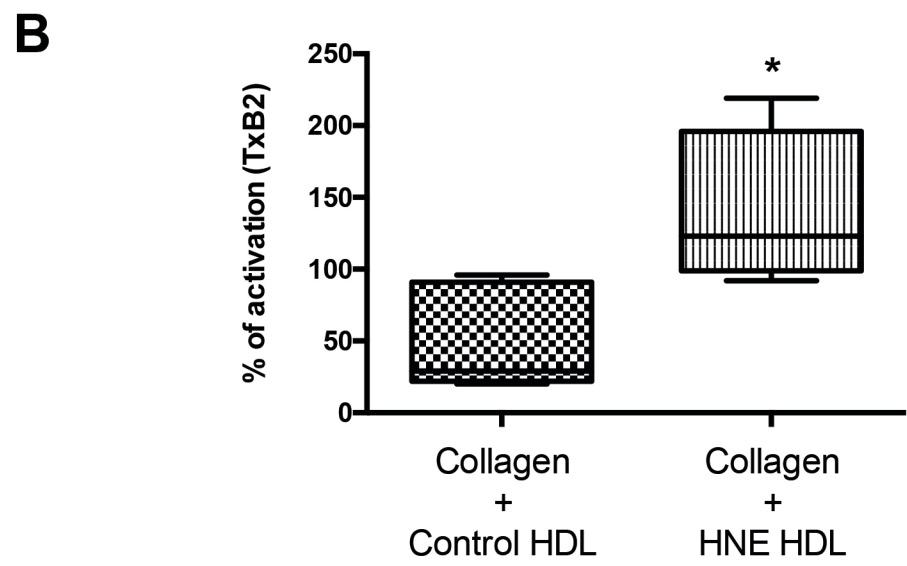
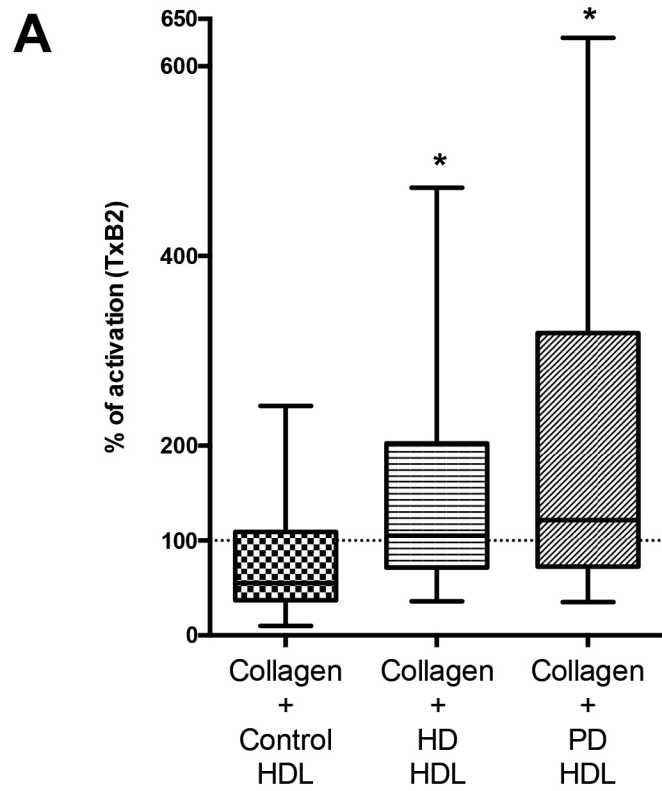
Peptide [241-262] : MH⁺ : 2598.4056 Th - MSMS on m/z 1299.7064 (2⁺; -0.37 ppm)

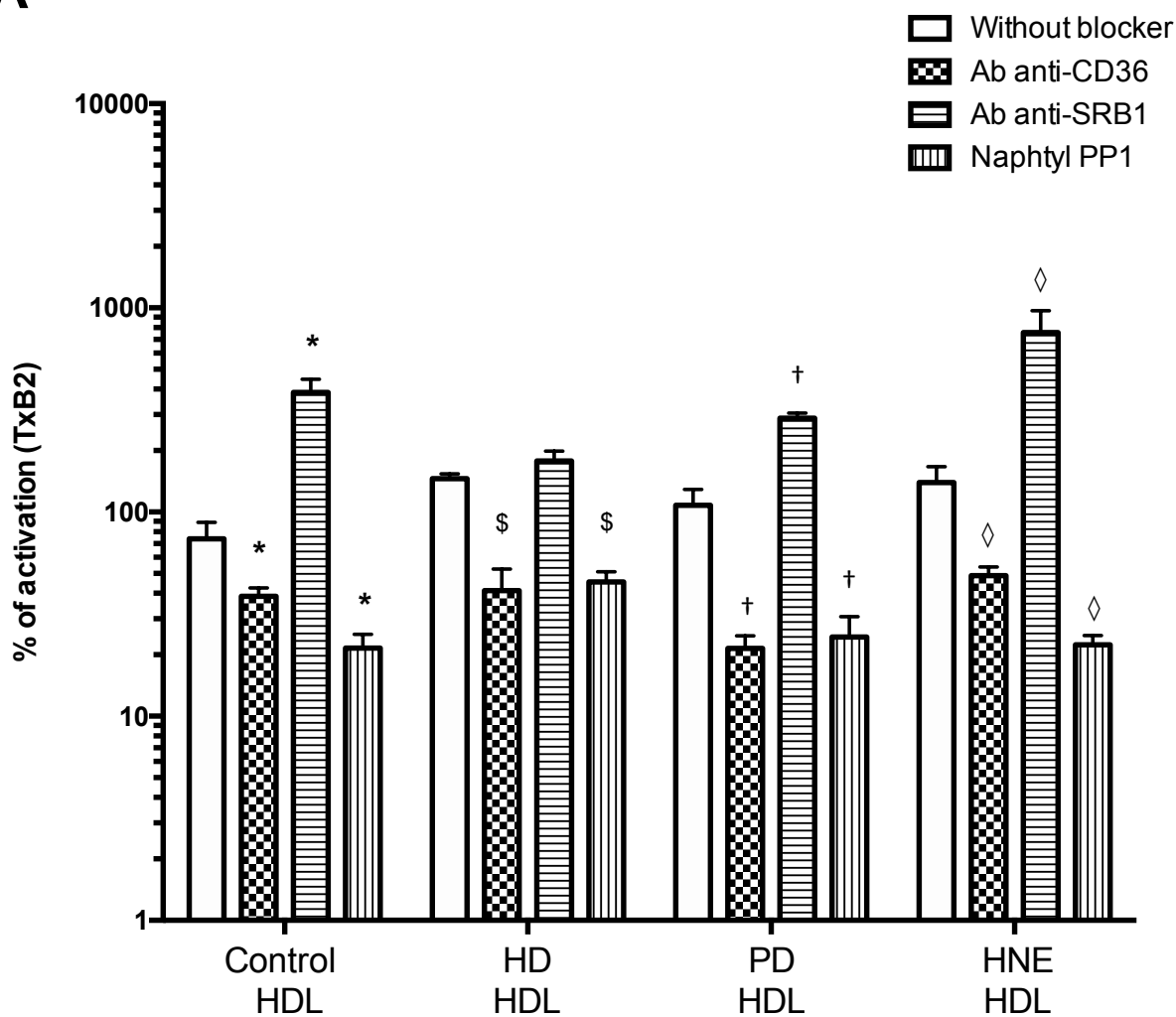


Peptide [241-262] HNE Modification : MH⁺ : 2754.5001 Th - MSMS on m/z 689.3805 (4⁺; -7.78 ppm)







A**B**