Table S1. $A\beta$ -binding A and B chains of FDH by H-bond. Donor = H donor; Receptor = H receptor.

Protein (P) Ligand (L) Donor		Donor	Receptor	Bond Length		
		P_A: ASN 225:HND21	L: GLU11:OE2	2.375		
		P_A: LYS 226: HNZ1	L: VAL12: O	1.770		
1TEH_AB		P_A: HIS 363:HD1	L: GLU3:OE2	2.437		
		P_B: LYS 330:HZ1	L: ARG5: O	1.996		
		P_B: LYS 330:HZ2	L: ARG5: O	1.659		
		L: ARG5: HE	P_B: GLU326:OE2	2.292		
		L: HIS6:HD1	P_B: GLU326:OE1	2.292		
	1700	L: HIS13:HE2	P_B: ASP 161:OD1	1.540		
	1202	L: HIS14:HD1	P_B: LYS 82: O	2.466		
	(Under	L:GLN15:HE21	P_A: ASP 227:OD2	1.671		
	vacuum	P_A: ASN 225:ND2	L: GLU11:OE2	2.973		
	conditions)	P_A: LYS 226:NZ	L: VAL12: O	2.572		
	conditions)	P_B: GLU 326: N	L: TYR10: OH	3.054		
		P_B: LYS 330:NZ	L: ARG5: O	2.216		
1TEH AB-Zn		P_B: LYS 330:NZ	L: SER8: OG	3.198		
		L: ARG5: HE	P_B: GLU326:OE2	2.289		
		L: HIS6:HD1	P_B: GLU326:OE1	2.292		
		L: HIS13:HE2	P_B: ASP 161:OD1	1.537		
		L: HIS14:HD1	P_B: LYS 82: O	2.462		
		L:GLN15:HE21	P_A: ASP 227:OD2	1.670		
		P_A: HIS 47:HE2	L: GLU3: O	2.406		
		P_A: SER 54: HG	L: ASP7:OD1	2.127		
1TEH_AB-eq		P_B: LYS 82:HZ2	L: HIS13:NE22	2.335		
		L: SER8: HG	P_A: HIS 363: O	1.757		
		L: HIS14:HD1	P_A:GLY18:O	2.357		
		P_A: LYS 226:HZ1	L: VAL12: O	1.776		
		P_A: LYS 366:HZ1	L: GLU11:OE1	2.169		
	1Z0Q-eq	P_A: LYS 366:HZ1	L: GLU11:OE2	1.851		
		P_A: LYS 366:HZ2	L: GLU11:OE1	1.556		
	(Under the	P_A: LYS 366:HZ2	L: GLU11:OE2	1.333		
	condition of	P_A: LYS 366:HZ3	L: GLU11:OE2	1.579		
1TEH_AB-Zn-eq	ion	P_B: LYS 82:HZ1	L: ALA2: O	1.995		
	equilibrium)	P_B: LYS 82:HZ2	L: ASP1:OD2	2.487		
		P_B: GLY 98:HN	L:ASP1: O	2.061		
		P_B: LYS159:HZ3	L: HIS14:NE2	1.752		
		P_B: GLU 326:HN	L: HIS63:NE2	1.619		
		L: ASP1:HT1	P_B: GLN96: O	0.819		
		L: ASP1:HT2	P_B: GLN96: O	1.294		
		L: ASP1:HT3	P_B: GLN96: O	1.730		
		L:GLN15:HE21	P_A: ASP 227:OD2	2.393		

Table S2: Changes in the cysteine residues of CYS45, CYS90, CYS100, and CYS111 residues of the samples of hFDH, A β +FDH, and RL+A β +FDH.

Cysteine status	Sample 1: FDH	Sample 2: Aβ+FDH	Sample 3: RL+Aβ+FDH
Cys 43	%	%	%
SH	96.8	95.3	94.6
MTHIO	3.2	4.2	4.3
SOxH		0.4	1.1
Cys 97,100			
SH	100.0	100.0	100.0
MTHIO			
SOxH			0.3
S-S		1.4	
Cys 111	%	%	%
SH	76.7	93.2	92.1
MTHIO	53.2	6.8	5.3
S-S	5.9		2.6

Table S3: Changes in the cysteine residues of CYS45, CYS90, CYS100, and CYS111 residues of sample 1-hFDH.

Modifications	Area					
C4(Carbamidomethyl)	1525305SH		Cys 43	Area	%	
C4(Carbamidomethyl)	4162714SH		SH	471776	6 0	96.80317
C4(Methylthio)	1557996MTHIO		MTHIO	155799	96	3.196829
C4(Carbamidomethyl)	41489641SH					
			Cys 97,100			
C5(Carbamidomethyl); M6(Oxidation); C8(Carbamidomethyl)	3868177SH		SH	437313	6	100
C6(Methylthio)	218750.3MTHIO		Cys 111	Area	%	
C1(Dehydro); C9(Dehydro)	286208.3S-S		SH	122435	3 9	76.67404
C1(Dehydro)	244006S-S		MTHIO	849088	34	53.17338
C1(Methylthio); C9(Methylthio)	2293431MTHIO		S-S	949382	.4	5.94542
C1(Methylthio); C9(Methylthio)	364871.5MTHIO					
C1(Methylthio); C9(Carbamidomethyl)	4346956SH	MTHIO				
C1(Carbamidomethyl); C9(Methylthio)	803793.8SH	MTHIO				
C1(Carbamidomethyl); C9(Methylthio)	145588.7SH	MTHIO				
	1422834SH					
C6(Methylthio)	317492.7MTHIO					
C6(Carbamidomethyl)	5524366SH					

Table S4: Changes in the cysteine residues of CYS45, CYS90, CYS100, and CYS111 residues of sample 2- $A\beta$ +FDH.

Modifications	Area					
C4(Methylthio)	3593866	MTHIC)	Cys 43	Area	%
C4(Carbamidomethyl)	92585621	SH		SH	2.28E+	0895.31692
C4(Methylthio)	4375416	MTHIC)	MTHIO	10162	90 4 4.24385
C4(Methylthio)	2193622	MTHIC)	SOxH	10518	460.439232
C4(Trioxidation)	1051846	SOxH		S-S	ND	
C4(Carbamidomethyl)	1.02E+08	SH				
C4(Carbamidomethyl)	33906144	SH				
C5(Carbamidomethyl); C8(Dehydro)	157720.2	SH	S-S	Cys 97,100		
C5(Carbamidomethyl); C8(Carbamidomethyl)	586814.6	SH	SH	SH	11076	575 4 100
C5(Carbamidomethyl); M6(Oxidation); C8(Carbamidomethyl)	9845248	SH	SH	S-S	157720	0.21.423884
C4(Carbamidomethyl); M5(Oxidation); C7(Carbamidomethyl)	486971.5	SH	SH			
C6(Methylthio)	507735.2	MTHIC)	Cys 111	Area	%
C6(Carbamidomethyl)	3917935	SH		SH	16999	28 193.20222
	1097808	SH		MTHIO	12398	576.797781
C6(Methylthio)	732121.5	MTHIC)	SOxH	ND	
C6(Carbamidomethyl)	11983539	SH		S-S	ND	

Table S5: Changes in the cysteine residues of CYS45, CYS90, CYS100, and CYS111 residues of sample 3-RL+A β +FDH.

Modifications	Area				
C4(Dioxidation)	6033516SOxH		Cys 43	Area 9	%
C4(Methylthio)	21570859MTHIO		SH	7.33E+08	94.61604
C4(Methylthio)	8948802MTHIO		MTHIO	33327636	4.30004
C4(Trioxidation)	2367469SOxH		SOxH	8400985	1.083922
C4(Carbamidomethyl)	4.63E+08SH		S-S	ND	
C4(Carbamidomethyl)	1.67E+08SH				
C4(Methylthio)	2807975MTHIO				
C4(Carbamidomethyl)	1.03E+08SH				
C5(Dioxidation); M6(Oxidation); C8(Carbamidomethyl)	69652.33SH	SOxH	Cys 97,100		
C5(Carbamidomethyl); C8(Carbamidomethyl)	9525196SH	SH	SH	2061331 8	100
C5(Carbamidomethyl); M6(Oxidation); C8(Carbamidomethyl)	9729114SH	SH	SOxH	69652.33	0.3379
C4(Carbamidomethyl); C7(Carbamidomethyl)	797207.1SH	SH			
C4(Carbamidomethyl); M5(Oxidation); C7(Carbamidomethyl)	492148.6SH	SH			
C6(Methylthio)	323035.8MTHIO		Cys 111	Area 9	%
C6(Carbamidomethyl)	3874670SH		SH	1504216 5	92.05637
C1(Dehydro); C9(Dehydro)	432942.9S-S		MTHIO	865060	5.294071
C6(Methylthio)	542024.2MTHIO		SOxH	ND	
C6(Carbamidomethyl)	11167495SH		S-S	432942.9	2.649562