

## Profile of Prof. Faizan Ahmad

S.No.	Name	Prof. Faizan Ahmad	
1	Designation	INSA Honorary Scientist	
2	Qualifications	M.Sc., M.Phil., Ph.D.	
3	State of Domicile	New Delhi	
4	Department & School	Department of Biochemistry, School of Chemical and Life Sciences Jamia Hamdard Jamia Nagar, New Delhi, India-110062	
5	Specialization	Molecular Biophysics, Mechanism of Protein Folding, Protein Folding Intermediates, Protein Stability Determination, Mechanism of Protein Stabilization by Osmolytes, Urea Stress, and Alcohol Stress.	
6	Home Tel	011-2692-0954	
7	Cell	+91-9810413115	
8	E-mail	<a href="mailto:faizanahmad@jamiahamdard.ac.in">faizanahmad@jamiahamdard.ac.in</a> ; <a href="mailto:Faizan.ahmad.jmi@gmail.com">Faizan.ahmad.jmi@gmail.com</a> ;	
	URL	<a href="https://jmi.ac.in/Centre-For-Interdisciplinary-Research-In-Basic-Sciences/Faculty-Members/1631/Faizan_Ahmad">https://jmi.ac.in/Centre-For-Interdisciplinary-Research-In-Basic-Sciences/Faculty-Members/1631/Faizan_Ahmad</a>	
9	Date of Birth	July 10, 1949	
<b>PUBLICATIONS</b>			
10	Publications in peer reviewed National & International journals, Citations, H-index	328 Citations: 12400; h-index: 55; i10 : 297 Link : <a href="#">Professor Faizan Ahmad - Google Scholar</a>	
<b>BOOKS &amp; BOOK CHAPTERS PUBLISHED</b>			
	Number of Book Chapters in Edited books of International Publishers	04	
<b>RESEARCH GRANTS AND CONSULTANCY PROJECTS</b>			

	Number of research grants from govt. funding agencies as PI	16 Research Projects as PI 04 Research Projects as Co-PI
	<b>RESEARCH SUPERVISION</b>	
	<b>As Supervisor</b>	
	Number of Ph.D. Guided	53
	<b>PRESENTATIONS IN CONFERENCES AS SPEAKER/ RESOURCE PERSON</b>	
	Number of presentations in National or International Conferences in India	53
	Number of presentations in International Conferences abroad	20
	<b>NATIONAL AWARDS, HONOURS AND FELLOWSHIPS RECEIVED</b>	
	2021 2020 2008: till date 1997 1997 1996: till date 1996: till date	<b>Selected among top 2% scientists of the world</b> <b>Fourth top scientist in India in the field of Biophysics</b> <b><i>Fellow of Indian National Science Academy</i></b> <b><i>Pride of Delhi</i></b> , honored by the Government of Delhi. <b><i>Norman H. Dill Memorial Gold Medal</i></b> <b><i>Member of Guha Research Conference</i></b> <b><i>Fellow of National Academy of Sciences</i></b>
	<b>INTERNATIONAL AWARDS, HONOURS AND FELLOWSHIPS RECEIVED</b>	
	1) 2013 - 2022 2) 1996 3) 1978-1979	Adjunct Professor, IBB, University of Tehran, Iran DAAD Fellow, Centre for Molecular Medicine, Buch, Berlin, Germany National level award by National Research Council, Washington
	<b>MEMBER(S) OF NATIONAL COMMITTEES</b>	
	1) American Society of Biochemistry & Molecular Biology (1983-1989: 2005-to-date) 2) Canadian Biochemical Society (1981-1987) 3) Society of Biological Chemists (India)(1995-todate) 4) Indian Biophysical Society(1996-todate) 5) ITCB, Trieste (1996-todate) 6) Guha Research Conference(1996-todate) 7) Society of Biosciences(1996-todate)	

	8) Advisory Member of the Editorial Board, Journal of Iranian Chemical Society 9) On the Editorial Board of the Journal of Iranian Chemical Society
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## Curriculum Vitae

Name : Prof. **Faizan Ahmad**  
Designation : INSA Honorary Scientist  
Office Address : Department of Biochemistry, School of Chemical and Life Sciences,  
Jamia Hamdard Jamia Nagar, New Delhi, India-110062  
Home Tel : 011-2692-0954  
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Date of Birth : July 10, 1949

### Education:

1975 Ph.D.(Chemistry), Aligarh Muslim University.  
1972 M. Phil (Physical Chemistry), Aligarh Muslim University  
1970 M.Sc. (Chemistry), Aligarh Muslim University.  
1968 B.Sc. (Chemistry), Aligarh Muslim University.

### Work Experience:

2023- Till date **INSA Honorary Scientist**, Department of Biochemistry, SCLS, Jamia Hamdard, New Delhi  
2021- 2023 **INSA Senior Scientist**, Dept. Biochemistry, SCLS, JH, New Delhi  
2017- 2021 **INSA Senior Scientist**, CIRBSc, Jamia Millia Islamia (JMI), New Delhi  
2006- 2017 **Professor**, CIRBSc, JMI, New Delhi.  
2006-2012 **Professor & Director**, CIRBSc, JMI, New Delhi.  
2000-2006 **Dean of Students' Welfare**, JMI, New Delhi.  
1998-2006 **Professor**, Department of Biosciences, JMI, New Delhi.  
1994-1998 **Professor & Head**, Department of Biosciences, JMI, New Delhi.  
1993-1994 **Professor**, Department of Chemistry, JMI, New Delhi.  
1990-1993 **Professor & Head**, Department of Chemistry, JMI, New Delhi.  
1987-1990 **Reader**, Department of Biosciences, JMI, New Delhi.  
1981-1987 **Research Associate** for Dr. C.C. Bigelow, Chemistry Department, University of Manitoba, Winnipeg, Manitoba, Canada.  
1979-1980 **Research Associate** for Dr. R. S. Roche, Chemistry Department, University of Calgary, Calgary, Alberta, Canada.  
1979 (Jan-Aug) **NRC Research Associate**, Naval Medical Research Institute, Department of Navy, Bethesda, Maryland 20014, U.S.A.  
1976-1978 **Visiting Fellow** with Dr. P. McPhie, National Institutes of Health, Bethesda, Maryland, U.S.A.  
1975-1976 **Postdoctoral Fellow** for Dr. C.C. Bigelow, Biochemistry Department, M.U.N., St. John's, Newfoundland, Canada.  
1974-1975 **Research Assistant** for Dr. V.S. Anathanarayanan, Molecular Biophysics Unit, Indian Institute of Science, Bangalore 560012, India

### Research Interest

### **Specialization:**

Molecular Biophysics, Protein Folding: Protein Stability determination/Mechanism of Protein stabilization by osmolytes/Urea and Alcohol stresses/Protein folding intermediates

*We have contributed to the problems relating to the "protein folding and stability", namely:*

1. Development of methods of the structural characterization of the random coil state native state, of the folding equation, native state  $\leftrightarrow$  denatured state (Biochemistry (USA), 1974; Intl. J. Pept. Protein Res., 1975, 1980; Intl. J. Biochem., 1978; etc.);
2. conformational and thermodynamic characterization of partial denatured states (BBA, 1977, 1994, 1996; J. Mol. Biol., 1979; J. Biol. Chem. 1983a, 1983b, 1984, 1985; Biochemistry (USA), 2003, 2006; PlosOne, 2015; etc.);
3. Development of methods of estimation of  ${}_D\Delta G^0_D$ , the protein stability under physiological conditions (J. Biol. Chem., 1982; Biopolymers, 1986, 1990; Biochem. J., 1992, 2000; J. Biochem., 1994; Biochemistry (USA), 1996, 1999, 2003, 2006; Anal. Biochem., 2000; PINSA, 2002; BBA, 2003; JICS, 2004; Biophys. Chem., 2008; J. Chem. Therm., 2013; Front. Mol. Biosc., etc.);
4. Estimation of  $\Delta G^0_D$  (Biochemistry (USA), 1976, 1978, 1996, 1999, 2003, 2006, BBA, 2008; J. Biol. Inorg. Chem., 2009, 2010; JBDS, 2013, 2015; ABB, 2014; etc.)
5. Characterization of the phenomenon of stabilization of proteins by osmolytes against environmental stresses (Biochem. J., 1994, 1998; BBA, 2000, 2007, 2017; J. Biol. Chem., 2005; FEBS Lett., 2005, Biophys. Chem., 2005, 2006; FEBS J., 2009; JBSD, 2009; JICS, 2011; PINSA, 2013; PLOS ONE, 2013, 20015; Biochemistry, 2015; Int JBiol Macromol, 2017, etc.)
6. Characterization of folding intermediates (pre-molten and molten globules) of proteins (BBA, 2003, 2007; Biochemistry (USA), 2003, 2006; BBA, 2007; Boiphys. Chem., 2007; J. Biol. Inorg. Chem, 2009, 2010; Metallomics, 2011; I. J. Biol. Macromol., 2015, 2016, 2017; PLOS ONE, 2015; JBSD, 2015; etc.).

### **Grants/Projects (16)**

#### **Principal Investigator**

1. In Silico (Molecular Dynamics Simulation) and In Vitro Approaches to Understand Counteraction of Urea's Effects on Proteins by Kidney Osmolytes [April 2016 - till date] **Principal Investigator** (ICMR submission id 2014-3313)
2. Mechanism of Survival of Ethanol Producing Organisms: Role of Cellular Compatible Osmolytes in Counteracting the Deleterious Effects of Ethanol on Structure, Stability and Functions of Proteins [December, 2013- 2016] **Principal Investigator** (CSIR Ref. No. 3809/NS-EMR-II).
3. Protein Structural Biology (Protein Folding) [September 2013 till Five Years]. **Principal Coordinator:** FIST (Ref. No. PDB/PC-402/2013/FISR#: 185826)
4. The Critical Role of Five N-terminal Residues in the Folding and Stability of Yeast

- Iso-1 Cytochrome-C [ April 1, 2011 – Nov 30, 14] **Principal Investigator** - DST ( Ref. No. SR/SO/BB-7/2010)
5. Investigating the Involvement of Other Osmolytically Active solutes (Non-Methylamines) of urea Rich Cells in Counteracting the Urea's Effect on Protein Stability and Function [Dec 1, 2009 – Nov 2012] **Principal Investigator** - CSIR (Ref. No. 37(1377)/09/EMR-II)
  6. PROTEIN FOLDING: Would the Heat/Acid Denatured State Serve As Reference State for Protein Folding [May 1, 2009- Apr 30, 2012] **Principal Investigator** - UGC (Ref No. F. No. 36-112/2008(SR))
  7. In Vivo and In Vitro Paradox of the Compensatory Effect of Methylamines: Urea (1:2) on Stability and Function of Proteins [ April 1, 2006 – 2010], **Principal Investigator** - DST (Ref. No. SR/SO/BB-80/2004)
  8. PROTEIN FOLDING: Conformational and Thermodynamic Studies of the Effect of Amino Acid Substitution on Equilibrium, Native State  $\leftrightarrow$  Molten Globule State  $\leftrightarrow$  Denatured State of Cytochrome C [ Oct 1, 2005 – Sept 2009] **Principal Investigator** - CSIR ( Ref. No. 37(1232)/05/EMR-II)
  9. Stabilization of Industrial and Therapeutic Enzymes by Osmolytes [March 1, 2005 – 2008] **Principal Investigators-II** - Iran National Science Foundation
  10. Mechanism of Stabilization of Proteins by Naturally Occurring Polyol Osmolytes Accumulated in Response to Environmental Stresses [April 1, 2001 – Sept 30, 2005] **Principal Investigator** - CSIR (ref no. 37(1078)/01/EMR-II)
  11. Protein Folding: Determination of Stability of Molten Globules. [From April 1, 1998- March 31, 2002] **Principal Investigator** - CSIR (Ref no. 37(976)/98 EMR-II)
  12. Protein Folding: Estimation of Protein Stability from Conformational Transition Curves [October 1997-November 30, 2001], **Principal Investigator** - DST (Ref. no. SP/SO/D26/96)
  13. Mechanism of Stabilization of Proteins by amino acids and other Additives Accumulated in Response to Heat Stress [1-11-94 – 31-10-98] **Principal Investigator** - U.G.C. (Ref. no. 12-4/94(SR-I))
  14. Roll of Building Block Molecules Induced by Heat in the Stabilization of Proteins Under Hyperthermia. [1-4-94 – 31-3-97] **Principal Investigator** - CSIR (Ref. no. 37\0841\94\EMR-II)
  15. Fast reaction Kinetics of ATP Hydrolysis by Myosin ATPase. [17-11-79 - 17-5-92] **Principal Investigator** - DST (Ref. no. SP\SO\D51\87)
  16. Mechanism of Protein Denaturation: Effect of Physical and Chemical Denaturants on Cytochrome-c and Myoglobin. [1-4-89 - 30-6-92] **Principal Investigator** - CSIR (Ref. No. 9\295)

#### **Co- Principal Investigator**

17. Relation between Stability and Functional Activity of Proteins in the Presence of Different Sizes of Sugar Osmolytes. [2013-2017] – CSIR
18. *In Silico* (Molecular Dynamic Simulation) and *In Vitro* Approaches to Understand Counteraction of Urea's Effects on Proteins by Kidney Osmolytes. [2016-2019] -ICMR BIC/12(16)/2014).
19. Compilation of Useful Information for Indian Diseases through an Online Database Management: a Useful Resource for Researcher and Public Awareness. [01-11-2011 to 30-10-2014] – UGC

20. Structure and Functional Analysis of Putative Conserved Proteins from Common Indian Pathogens. [1-02-2012 to 31-01-2015] –ICMR
21. Folding and stability of naturally truncated photosynthetic pigment, C-phycoerythrin from cyanobacteria *Phormidium tenue*. [01-09-2012 to 30-10-2015]- DST

### Award and Honors

- 2023 **World's Best Chemistry Scientist (Top 1% as on 21 Nov 2023)**  
[<https://research.com/scientists-rankings/chemistry/in>]
- 2021 **Selected among top 2% scientists of the world**
- 2020 **Fourth top scientist in India in the field of Biophysics**
- 2013 - 2025 **Adjunct Professor**, IBB, University of Tehran, Iran
- 2008: till date **Fellow of Indian National Science Academy**
- 1997 **Pride of Delhi**, honored by the Government of Delhi.
- 1997 **Norman H. Dill Memorial Gold Medal**
- 1996 **DAAD Fellow**, Center for Molecular Medicine, Buch, Berlin, Germany
- 1996: till date **Member of Guha Research Conference**
- 1996: till date **Fellow of National Academy of Sciences**
- 1978-1979 National level award by **National Research Council, Washington**

### Member of Societies

- American Society of Biochemistry & Molecular Biology (1983-1989: 2005- to-date)
- Canadian Biochemical Society(1981-1987)
- Society of Biological Chemists (India)(1995-todate)
- Indian Biophysical Society(1996-todate)
- ITCP, Trieste (1996-todate)
- Guha Research Conference(1996-todate)
- Society of Biosciences(1996-todate)
- Advisory Member of the Editorial Board, Journal of Iranian Chemical Society
- On the Editorial Board of the Journal of Iranian Chemical Society

### PhD Degree Awarded: 53

\*Main Supervisor

S. No.	Name	Year	S. No.	Name	Year
1.	Luqman A Khan	1991	35	Sobia Zaidi	2015
2.	Zulfiqar Ahmad*	1992	36	Charu Thaplial	2015
3.	Sushma Yadav*	1992	37	Sheeza Khan*	2015
4.	Sperna Taneja*	1992	38	Md. Anzarul Haque*	2016
5.	Amita Sinha*	1995	39	Farha Naz	2016
6.	Syed Ehtaishamul Haque*	1997	40	Parvez Khan	2016
7.	Reena Gupta*	1998	41	Huma Naz	2016
8.	Tapas Saha*	1998	42	Danish Idreess	2016

9.	Viks Rishi*	1999	43	Syed Ausaf Ali*	2017
10.	Sunita Yadav*	2000	44	Mohd. Aasif Dar*	2017
11.	Farah Anjum*	2000	45	Shabab H. Khan *	2017
12.	Hanief M.Shahjee*	2001	46	Ilyas Beg	2017
13.	Mohammad Zaffrullah*	2002	47	Wahiduzzaman*	2017
14.	Beenu Moza*	2002	48	Moin Ishrat	2017
15.	Shabir H. Qureshi*	2003	49	Abdullah Naiyer*	2018
16.	Akalank Jain*	2005	50	Sumra Shahid	2018
17.	Ritu Singh*	2005	51	Khalida Nasreen	2019
18.	Inamul Haque*	2005	52	Shagufta Khan	2019
19.	L.R. Singh*	2006	53	Mohd. Amir	2020
20.	Madhvi Gupta*	2007			
21.	Humaira Farooqi	2008			
22.	Gul M. Mustafa	2008			
23.	Tanveer A. Dar*	2009			
24.	Md. H. Rahaman*	2009			
25.	Nitesh Poddar*	2009			
26.	Md. K.A.Khan*	2010			
27.	Asimul Islam*	2010			
28.	Rinky Minakshy*	2010			
29.	Shazia Jamal*	2011			
30.	Mohd. Wahid*	2011			
31.	Md. T. Rehman*	2012			
32.	Syed I. Hassan*	2012			
33.	Shafikur Rahman*	2012			
34.	Shah Ubaidullah*	2014			

#### **Invited talk delivered abroad**

2023	IBB, University of Tehran, Tehran, Iran
2022	IBB, University of Tehran, Tehran, Iran
2021	IBB, University of Tehran, Tehran, Iran
2021	ISOBC & IASBS, Iran
2020	Institute of Biochemistry & Biophysics, Univ. Tehran, Iran
2016	Institute of Biochemistry & Biophysics, Univ. Tehran, Iran
2015	Biophysical Society of China, Hangzhou, China
2012	International Conference of Biophysical Chemistry, Ardabil, Iran.
2010	Gordon research Conferenc, N.C., USA.
2009	IBB, University of Tehran, Iran
2007	IBB, University of Tehran, Iran.
2006	Society of the Biophysical Chemistry (Iran), University of Tabriz, Tabriz, Iran
2006	Workshop on Protein Characterization, Institute of Biochemistry & Biophysics, University of Tehran, Iran
2005	University of Terbiat Mudarris, Iran
2005	University of Tehran, Iran
2004	University of Mashhad, Iran

2004 University of Tehran, Iran  
2002 "The 2002 Colorado Protein Stability Conference" in Colorado, USA.  
2000 "NIDKD, National Institutes of Health, Bethesda, MD, USA.  
1999 "University of Texas Medical Centre, Galveston, TX, USA.  
1999 "Dept. Biophysics and Biochemistry, A&M Univ. College Station, USA.

BIBLIOGRAPHY-2023

**Total no. of publications: 335**

S.No.	<u>Publications</u>
1.	<b>Ahmad, F.</b> and Salahuddin, A. (1974), Influence of Temperature on the Intrinsic Viscosities of Proteins in Random Coil Conformation. <i>Biochemistry</i> <b>13</b> , 245-249.
2.	<b>Ahmad, F.</b> and Salahuddin, A. (1975), Intrinsic Viscosity of Ovomuroid in Random Coil Conformation. <i>Intl. J. Pept. Protein Res.</i> <b>7</b> , 417-421.
3.	<b>Ahmad, F.</b> and Salahuddin, A. (1976), Reversible Unfolding of the Major Fraction of Ovalbumin by Guanidine Hydrochloride. <i>Biochemistry</i> <b>15</b> , 5168-5175.
4.	Ananthanarayanan, V. S. and <b>Ahmad, F.</b> (1977), Evidence from Rotatory Measurements for an Intermediate State in the Guanidine Hydrochloride Denaturation of $\beta$ -Lactoglobulin. <i>Can. J. Biochem.</i> <b>55</b> , 239-243.
5.	Ananthanarayanan, V.S., <b>Ahmad, F.</b> , and Bigelow, C. C. (1977), The Denaturation of $\beta$ -Lactoglobulin-A at pH 2. <i>Biochim. Biophys. Acta</i> <b>492</b> , 194-203.
6.	<b>Ahmad, F.</b> and McPhie, P. (1978), Thermodynamics of the Denaturation of Pepsinogen by Urea. <i>Biochemistry</i> <b>17</b> , 241-246.
7.	<b>Ahmad, F.</b> and McPhie, P. (1978), Intrinsic Viscosities of Randomly Coiled Glycoproteins. <i>Intl. J. Biochem.</i> <b>9</b> , 647-651.
8.	<b>Ahmad, F.</b> and McPhie, P. (1978), The Denaturation of Covalently Inhibited Swine Pepsin. <i>Intl. J. Pept. Protein Res.</i> <b>12</b> , 155-163.
9.	<b>Ahmad, F.</b> and McPhie, P. (1978), Spectrophotometric Titration of Phenolic Groups of Pepsin. <i>Biochim. Biophys. Acta</i> <b>537</b> , 247-254.
10.	<b>Ahmad, F.</b> and Bigelow, C. C., (1978), Inorganic Salt Denaturants Stabilize Ribonuclease Against Denaturation by Urea. <i>Can. J. Biochem.</i> <b>56</b> , 1003-1005.
11.	<b>Ahmad, F.</b> and Salahuddin, A. (1979), The pH Dependence of Reversible Unfolding of Ovalbumin by Guanidine Hydrochloride. <i>Biochim. Biophys. Acta</i> <b>576</b> , 333-338.
12.	<b>Ahmad, F.</b> and Bigelow, C. C. (1979), The Denaturation of Ribonuclease-A by Combinations of Urea and Salt Denaturants. <i>J. Mol. Biol.</i> <b>131</b> , 607-617.
13.	<b>Ahmad, F.</b> and McPhie, P. (1979), Characterization of a Stable Intermediate in the Unfolding of DAG-Pepsin by Urea. <i>Can. J. Biochem.</i> <b>57</b> , 1090-1092.
14.	<b>Ahmad, F.</b> and McPhie, P. (1980), The Intrinsic Viscosity of Glycoproteins. <i>Intl. J. Biochem.</i> <b>11</b> , 91-96.
15.	<b>Ahmad, F.</b> (1981), Stability of Acetylcholinesterase in Guanidine Hydrochloride Solution. <i>Can. J. Biochem.</i> <b>59</b> , 551-555.
16.	<b>Ahmad, F.</b> and Bigelow, C. C. (1982), Estimation of Free Energy of Stabilization of Ribonuclease-A, Lysozyme, $\alpha$ -Lactalbumin and Myoglobin. <i>J. Biol. Chem.</i> <b>257</b> , 12935-12938.
17.	<b>Ahmad, F.</b> , Contaxis, C. C., and Bigelow, C. C. (1983), Free Energy Changes in Lysozyme denaturation. <i>J. Biol. Chem.</i> <b>258</b> , 7960-7963.

18.	<b>Ahmad, F.</b> (1983), Free Energy Changes in Ribonuclease-A denaturation: Effect of urea, guanidine hydrochloride and lithium salts. <i>J. Biol. Chem.</i> <b>258</b> , 11143-11146.
19.	<b>Ahmad, F.</b> (1984), Free Energy Changes on Denaturation of Ribonuclease-A by Mixed Denaturants: Effects of combination of guanidine hydrochloride and one of the denaturants, LiBr, LiCl and NaBr. <i>J. Biol. Chem.</i> <b>259</b> , 4183-4186.
20.	<b>Ahmad, F.</b> (1985), Complexities in the Denaturation of Horse Metmyoglobin by Guanidine Hydrochloride. <i>J. Biol. Chem.</i> <b>260</b> , 10458-10461.
21.	<b>Ahmad, F.</b> (1985), Thermodynamic Characterization of the Partially Denatured States of Ribonuclease-A in Calcium Chloride and Lithium Chloride. <i>Can. Biochem. Cell Biol.</i> <b>63</b> , 1058-1060.
22.	<b>Ahmad, F.</b> and Bigelow, C. C. (1986), Estimation of the Stability of Globular Proteins. <i>Biopolymers</i> <b>25</b> , 1623-1633.
23.	<b>Ahmad, F.</b> and Bigelow, C. C. (1986), Thermodynamic Stability of Proteins in Salt Solutions: a comparison of the effectiveness of protein stabilizers. <i>J. Protein Chem.</i> <b>5</b> , 355-367.
24.	Corbett, R. J. T., <b>Ahmad, F.</b> , and Roche, R.S. (1986), Domain Unfolding and Stability of Thermolysin in Guanidine Hydrochloride. <i>Can Biochem. and Cell Biol.</i> <b>64</b> , 953-961.
25.	<b>Ahmad, F.</b> and Khan, L.A. (1989), The Denatured States of Ribonuclease-A: Mechanism of Denaturation by Lithium Chloride. <i>Ind. J. Biochem. Biophys.</i> <b>26</b> , 301-304.
26.	<b>Ahmad, F.</b> and Bigelow, C. C. (1990), Thermodynamics of Solvation of Proteins in Guanidine Hydrochloride. <i>Biopolymers</i> <b>29</b> , 1593-1598.
27.	<b>Ahmad, F.</b> (1991), Protein Stability from Denaturation Transition Curves. <i>Ind. J. Biochem. Biophys.</i> <b>28</b> , 168-173.
28.	<b>Ahmad, F.</b> , Yadav, S. and Taneja, S. (1992), Determining Stability of Proteins from Guanidinium Chloride Transition Curves. <i>Biochem. J.</i> <b>287</b> , 481-485.
29.	Ahmad, Z. and <b>Ahmad, F.</b> (1992), Mechanism of Denaturation of Cytochrome-c by Lithium Salts, <i>Ind. J. Chem.</i> <b>31B</b> , 874-879.
30.	Yadav, S., Taneja, S. and <b>Ahmad, F.</b> (1992), Measuring the Conformational stability of Proteins. <i>Ind. J. Chem.</i> <b>31B</b> , 859-864.
31.	<b>Ahmad, F.</b> (1993), Measuring the Conformational Stability of Enzymes <i>in the Thermostability of Enzymes</i> (M.N. Gupta, ed.), pp 95-112, Narosa Publishing House, India.
32.	<b>Ahmad, F.</b> , Taneja, S., Yadav, S. and Haque, S. E. (1994), A New Method for Testing the Functional Dependence of Unfolding Free Energy Changes on Denaturant Concentration, <i>J. Biochem.</i> <b>115</b> , 322-327.
33.	Ahmad, Z. and <b>Ahmad, F.</b> (1994), Physico-Chemical Characterization of Products of Unfolding of Cytochrome-C by Calcium Chloride. <i>Biochim. Biophys. Acta</i> <b>1207</b> , 223-230.
34.	Taneja, S. and <b>Ahmad, F.</b> (1994), Increased Thermal Stability of Proteins in Presence of Amino Acids. <i>Biochem. J.</i> <b>303</b> , 147-153.
35.	Ahmad, Z., Yadav, S., <b>Ahmad, F.</b> and Khan, N. Z. (1996), Effects of Salts of Alkali Earth Metals and Calcium chloride on the Stability of Cytochrome-C and Myoglobin. <i>Biochim. Biophys. Acta</i> <b>1294</b> , 63-71.

36.	Shereghi, B., <b>Ahmad, F.</b> and Moosavi-Movahedi, A. A. (1996), Stability of D-Amino Acid Oxidase: Denaturation by Guanidine Hydrochloride and Urea. <i>Ind. J. Biochem. Biophys.</i> <b>33</b> , 357-362.
37.	Gupta, R., Yadav, S. and <b>Ahmad, F.</b> (1996), Protein Stability: Urea- induced Versus Guanidine-induced Unfolding of Metmyoglobin. <i>Biochemistry</i> <b>35</b> , 11925-11930.
38.	Rishi, V., Anjum, F., <b>Ahmad, F.</b> and Pfeil, W. (1998), Role of Non-compatible Osmolytes in the Stabilization of Proteins during Heat Stress, <i>Biochem. J.</i> <b>329</b> , 137-143.
39.	Gupta, R. and <b>Ahmad, F.</b> (1999), Protein Stability: Functional Dependence of Denaturational Gibbs Energy on Urea Concentration. <i>Biochemistry</i> <b>38</b> , 2471-2479.
40.	Anjum, F., Rishi, V. and <b>Ahmad, F.</b> (2000), Compatibility of Osmolytes with Gibbs Energy of Stabilization of Proteins. <i>Biochim. Biophys. Acta</i> <b>1476</b> , 75-84
41.	Sinha, A., Yadav, Y., Ahmad, R. and <b>Ahmad, F.</b> (2000), A Possible Origin of Differences Between Calorimetric and Equilibrium Estimates of Stability Parameters of Proteins. <i>Biochem. J.</i> <b>345</b> , 711-717.
42.	Yadav, S. and <b>Ahmad, F.</b> (2000), A New Method for the Determination of Stability Parameters of Proteins from Their Heat-induced Denaturation Curves. <i>Anal. Biochem.</i> <b>283</b> , 207-213.
43.	Xiaofang, L., Zafrullah, M., <b>Ahmad, F.</b> and Jameel, S. (2001), A C-Terminal Hydrophobic Region is Required for Homo-Oligomerization of the Hepatitis E Virus Capsid (ORF2) Protein. <i>J. Biomed. Biotech.</i> <b>1:3</b> , 122-128.
44.	Jain, A., Rajeswari, M. R. and <b>Ahmad, F.</b> (2002), Formation and Thermodynamic Stability of Intermolecular (R*R.Y) DNA Triplex in GAA/TTC Repeats Associated with Freidreich's Ataxia. <i>J. Biomol. Stru. Dyna.</i> <b>19</b> , 691-699.
45.	<b>Ahmad, F.</b> (2002), Protein Folding: Estimates of Stability Parameters from Heat-Induced Conformational Transition curves of Proteins. <i>Proc. Ind. Natl. Sci. Acad.</i> <b>68, A</b> , 385-390.
46.	Shahjee, H. M., Banerjee, K. and <b>Ahmad, F.</b> (2002), Comparative Analysis of Naturally Occurring L-Amino Acid Osmolytes and their D-Isomers on Protection of <i>Escherichia Coli</i> Against Environmental Stresses. <i>J. Biosc.</i> <b>27</b> , 515-520.
47.	Shahjee, H. M., Rishi, V. and <b>Ahmad, F.</b> (2002), Effect of D- Amino Acids on the Functional Activity and Conformational Stability of Ribonuclease-A. <i>Ind. J. Biochem. Biophys.</i> <b>39</b> , 368-376.
48.	Jain, A., <b>Ahmad, F.</b> and Rajeswari, M. R. (2003), Structural Studies on DNA Triplet Helix Formed by Intronic GAA Triplet Repeat Expansion in Freidreich's A.toxia. <i>Nucleoside, Nucleotides and Nucleic Acids</i> <b>22</b> , 1517-1519.
49.	Moza, B., Qureshi, S. H. and <b>Ahmad, F.</b> (2003), Equilibrium Studies of the Effect of Difference in Sequence Homology on the Mechanism of Denaturation of Bovine and Horse Cytochromes-c. <i>Biochim. Biophys. Acta</i> <b>1646</b> , 49-56.
50.	Qureshi, S. H., Moza, B., Yadav, S. and <b>Ahmad, F.</b> (2003), Conformational and Thermodynamic Characterization of the Molten Globule State Occurring During

	Unfolding of Cytochromes-C by Weak Salt Denaturants. <i>Biochemistry</i> <b>42</b> , 1684-1695.
51.	Zafrullah, M., Khursheed, Z., Yadav, S., Sahgal, D., Jameel, S. and <b>Ahmad, F.</b> (2004), Acidic pH Enhances Structure and Structural Stability of the Capsid Protein of Hepatitis E virus. <i>Biochem. Biophys. Res. Commun.</i> <b>313</b> , 67-73.
52.	<b>Ahmad, F.</b> (2004), On the Estimation of Stability Parameters from Heat-induced Conformational Transition Curves of Proteins. <i>J. Iran. Chem. Soc.</i> <b>1</b> , 99-105.
53.	Singh, R., Haque, I. and <b>Ahmad, F.</b> (2005), COUNTERACTING OSMOLYTE TRIMETHYLAMINE N-OXIDE DESTABILIZES PROTEINS AT pH BELOW ITS pKa: Measurements of Thermodynamic Parameters of Proteins in the Presence and Absence of Trimethylamine N-Oxide. <i>J. Biol. Chem.</i> <b>280</b> , 11035-11042.
54.	Amani, M., Moosavi-Movahedi, A.A., Floris, G., Longu, S. Mura, A., Moosavi-Nijad, S.Z., Saboury, A. A. and <b>Ahmad, F.</b> (2005), Comparative Study of the Conformational Lock, Dissociative Thermal Inactivation and Stability of <i>Euphorbia</i> Latex and Lentil Seedling Amine Oxidases. <i>The Protein J.</i> <b>24</b> , 183-191.
55.	Haque. I., Singh, R., Moosavi-Movahedi, A.A. and <b>Ahmad, F.</b> (2005), Effect of Polyol Osmolytes on $\Delta G$ , the Gibbs Energy of Stabilization of Proteins at Different pH Values. <i>Biophys. Chem.</i> <b>117</b> , 1-12.
56.	Haque. I., Singh, R., <b>Ahmad, F.</b> and Moosavi-Movahedi, A.A. (2005), Testing Polyols' Compatibility Gibbs Energy of Stabilization of Proteins Under Conditions in Which They Behave as Compatible Osmolytes. <i>FEBS Letter</i> <b>579</b> , 3891-3898.
57.	Haque. I., Islam, A., Singh, R., Moosavi-Movahedi, A.A. and <b>Ahmad, F.</b> (2006), Stability of Proteins in the Presence of Polyols Estimated from Their Guanidinium Chloride-induced Transition curves at Different pH Values and 25 °C. <i>Biophys. Chem.</i> <b>119</b> , 224 – 233.
58.	Moza, B., Qureshi, S. H., Islam, A., Singh, R., Anjum, F., Moosavi-Movahedi, A. A., and <b>Ahmad, F.</b> (2006), A Unique Molten Globule State Occurs during Unfolding of Cytochrome C by LiClO <sub>4</sub> Near Physiological pH and Temperature: Structural and Thermodynamic Characterization. <i>Biochemistry</i> <b>45</b> , 4695 – 4702
59.	Moosavi-Movahedi, A. A., Gharanfoli, M., Jalil, S., <b>Ahmad, F.</b> , Chamani, J., Hakimelahi, G. H., Sadeghi, M., Amani, M. and Saboury, A. A. (2006), The Correlation of RNase- A Enzymatic Activity with the Changes in Distance Between N <sub>82</sub> -His <sub>12</sub> and N <sub>81</sub> -His <sub>119</sub> Upon Addition of Stabilizing and Destabilizing Salts, <i>The Protein J.</i> <b>25</b> , 117 - 125.
60.	Pirzadeh, P., Moosavi-Movahedi, A. A., Hemmateenejad. B., <b>Ahmad, F.</b> , Shamsipur, M., and Saboury, A. A. (2006), Chemometric Studies of Lysozyme upon Interaction with Sodium Dodecyl Sulfate and $\beta$ -Cyclodextrin, <i>Colloid and Surfaces B: Biointerfaces</i> <b>52</b> , 31 – 38.
61.	Hashemia, S., Moosavi-Movahedi, A. A., Ghourchian, H., <b>Ahmad, F.</b> , Hakimelahi, G. H., and Saboury, A. A. (2006), Diminishing Aggregation for Bovine Liver Catalase Through Acidic Residue Modification, <i>Int. J. Biol. Macromol.</i> <b>40</b> , 47-53.

62.	Hadi-Alijanvand, H., <b>Ahmad, F.</b> , and Moosavi-Movahedi, A. A. (2007), The Correlation of Cold Denaturation Temperature with Surface Stability Factor of Proteins, <i>Protein J.</i> <b>26</b> , 395-402.
63.	Amani, M., Moosavi-Movahedi, A. A., Floris, G., Mura, A., Kurgananov, B. I., <b>Ahmad, F.</b> , and Saboury, A. A. (2007), Two-state Irreversible Thermal Denaturation of <i>Euphorbia characias</i> Letex Amine Oxidase, <i>Biophys. Chem.</i> , <b>125</b> , 254-259.
64.	Barzegar, A., Moosavi-Movahedi, A. A., Sattarahmady, N., Hosseinpour_Faizi, M. A., Aminbakhsh, M., <b>Ahmad, F.</b> , Saboury, A. A., Ganjali, M. R., and Norouzi, P. (2007), Spectroscopic Studies of the Effects of Glycation of Human Serum Albumin on L-Trp Binding, <i>Protein Pept. Lett.</i> , <b>14</b> , 13-18.
65.	Sattarahmady, N., S., Moosavi-Movahedi, A.A., <b>Ahmad, F.</b> , Hakimelahi, G. H., Habibi_Razaei, Saboury, A. A. and Sheibani, N. (2007), Formation of Molten Globule-Like State During Prolonged Glycation of Human Serum Albumin, <i>Biochim. Biophys. Acta</i> <b>1770</b> , 933-942.
66.	Dar, T. A., Singh, L.R., Islam, A., Anjum, F., Moosavi-Movahedi, A. A. and <b>Ahmad, F.</b> (2007), Guanidinium Chloride and Urea Denaturations of $\beta$ -Lactoglobulin A at pH 2.0 and 25 °C: The equilibrium intermediate contains non-native structures (helix, tryptophan and hydrophobic patches), <i>Biophys. Chem.</i> <b>127</b> , 140-148.
67.	Moosavi-Movahedi, A. A., Peerzada, P., Hasemnia, S., Ahmadian, S., Hemmatinejad, B., Amani, M., Sabouri, A. A., <b>Ahmad, F.</b> , Shamshipur, M., Hakimelahi, B., Tsai, F., Alijavand, H. H., and Yusefi, R. (2007), Fibril Formation of Lysozyme upon Interaction with Sodium Dodecyl Sulphate at pH 9.2. <i>Colloides Surf.: Biointerfaces</i> <b>60</b> , 55-61.
68.	Moosavi-Nizad, S. Z., Moosavi-Movahedi A. A., Floris, G., Padiglia, A., Rezaei-Tavirani, M., <b>Ahmad, F.</b> , and Amani, M. (2007), Thermal Dissection of Lentil seedling Amine Oxidase Domains by Differential Scanning Calorimetry, <i>Biosc. Biotech. Biochem.</i> <b>71</b> , 1644-1649.
69.	Dey, P., Islam, A., <b>Ahmad, F.</b> , and Batra, K. K., (2007), Role of unique basic residues of human pancreatic ribonuclease in its catalysis and structural stability. <i>Biochem. Biophys. Res. Commu.</i> <b>360</b> , 809-814.
70.	Shee, C., Islam, A., <b>Ahmad, F.</b> , and Sharma, A. K., (2007) Structure-function studies of <i>Murraya koenigii</i> trypsin inhibitor revealed a stable core beta sheet structure surrounded by $\alpha$ -helices with a possible role for $\alpha$ -helix in inhibitory function. <i>Intl. J. Biol. Macromol.</i> <b>41</b> , 410-414.
71.	Heli, A., Moosavi_Movahedi, A. A., <b>Ahmad, F.</b> (2007) An Electrochemical Study of Safranin O Binding to DNA at the Surface. <i>J. Solid State Electrochem.</i> <b>11</b> , 593-599.
72.	Singh, L. R., Dar, T. A., Haque, I., Anjum, F., Moosavi-Movahedi, A. A., <b>Ahmad, F.</b> (2007) Testing the Paradigm that the Denaturing Effect of Urea on Protein stability is Offset by Methylamines at Physiological Ratio of 2:1 (Urea:Metylyamines). <i>Biochim. Biophys. Acta</i> <b>1774</b> , 1555-1562.

73.	Singh, R., Dar, T. A., Ahmad, S., Moosavi-Movahedi, A. A., <b>Ahmad, F.</b> (2008) A New Method for the Determining the Constant-Pressure Heat Capacity Change Associated with the Protein Denaturation Induced by Guanidinium Chloride (or Urea). <i>Biophys. Chem.</i> <b>133</b> , 81-89.
74.	Hassan, M. I., Waheed, A., Yadav, S., Singh, T. P., <b>Ahmad, F.</b> (2008) Zink $\alpha$ 2-Glycoprotein: A Multidisciplinary Protein. <i>Mole. Cancer Res.</i> <b>6</b> , 892-906.
75.	Mahnam, K., Moosavi-Movahedi, A. A., Behrami, H., Hakimelahi, H. G., Atai, G., Jalali, S., Saboury, A. A., <b>Ahmad, F.</b> , Safarian, S. and Amanlou, M. (2008) Efficient Factor in Protein Modification: Adenosine Deaminase Esterfication by Woodward Reagent K. <i>J. Iran. Chem. Soc.</i> <b>5</b> , 464-475.
76.	<u>Waheed A, Hassan MI, Etten RL, Ahmad, F.</u> (2008) Human seminal proteinase and prostate-specific antigen are the same protein. <i>J. Biosci.</i> <b>33</b> , 195-207.
77.	<u>Chaudhary NS, Shee C, Islam A, Ahmad, F, Yernool D, Kumar P, Sharma AK.</u> (2008). Purification and characterization of a trypsin inhibitor from Putranjiva roxburghii seeds. <i>Phytochemistry</i> <b>69</b> , 2120-2126.
78.	<u>Gupta M, Acharya R, Mishra A, Ramakumar S, Ahmad F, Chauhan VS</u> (2008). Dehydrophenylalanine (DeltaPhe) as a beta breaker: extended structure terminated by a DeltaPhe-induced turn in the pentapeptide Boc-Phe1-Ala2-Ile3-DeltaPhe4-Ala5-OMe. <i>Chembiochem.</i> <b>9</b> ,1375-8.
79.	Hekmat Azadeh, Saboury AA, Moosavi-Movahedi AA, Ghourchian H, <b>Ahmad F</b> (2008). Effects of pH on the activity and structure of choline oxidase from <i>Alcaligenes</i> species. <i>Acta Biochemica Polonica.</i> <b>55</b> , 549-557.
80.	Rahamam, H. M. Khan, M. K. A., Hassan, M. I., Wahid, M., Singh, S. B., Singh, T. P., Moosavi-Movahedi, A. A., <b>Ahmad, F.</b> (2008) Sequence and Stability of Goat Cytochrome C. <i>Biophys. Chem.</i> <b>138</b> , 23-28.
81.	Poddar, N. K., ansari, z. A., Singh, R. K. B, Moosavi-Movahedi, A. A., <b>Ahmad, F.</b> (2008) Effect of Monomeric and Oligomeric sugar osmolytes on $\Delta G_D^0$ , the Gibbs energy of Stabilization of the Protein at Different pH Values: Is the sum effect of monosaccharide individually additive in a mixture. <i>Biophys. Chem.</i> <b>138</b> ,120-129
82.	Aruna, B., Islam, A., Ghosh, S., Singh, A. K., Vijayalakshmi, M., <b>Ahmad, F.</b> , Ehtesham, N. Z. (2008) Biophysical Analyses of Human resistin: Oligomer Formation Suggests Noval biological Function. <i>Biochemistry</i> <b>47</b> , 12457-12466.
83.	Hassan, M. I., Waheed, A., Yadav, S., Singh, T. P., <b>Ahmad, F.</b> (2009) Prolactin Inducible Protein in Cancer, Fertility and Immunoregulation: Structure, function and its clinical applications. <i>Cell Mol. Life Sci.</i> <b>66</b> . 447-459.
84.	S. J. Mousavy, G.H. Razi, M. Kamarie, H. Aliakbarian, N. Sattarahmady, A. Sharifizadeh, S. Safarian, F. <b>Ahmad, F.</b> , A. A. Moosavi-Movahedi (2009) Effect of Mobile Phone Radiofrequency on the Structure and Function of the Normal Human Hemoglobin. <i>Intl. J. Biol. Macromol</i> <b>44</b> , 278-285.
85.	Singh, L.R., Dar, T.A., <b>Ahmad, F.</b> (2009) Living with Urea Stress. <i>J. Biosc.</i> <b>34</b> , 321-331.
86.	Singh, L.R., Dar, T.A., Ahmad, S., Jamal, S., <b>Ahmad, F.</b> (2009) Methylated Glycine Have Opposite Effects on Proteins at low pH Values. <i>Biochim. Biophys. Acta</i> <b>1794</b> , 929-935.

87.	Khan, MKA, Das, U., Rahaman, MH, Hassan, MI, Srinivasan, A., Singh, TP, and <b>Ahmad, F.</b> (2009) A single mutation induces molten globule formation and a drastic destabilization of the wild type cytochrome <i>c</i> at pH 6.0. <i>J. Biol. Inorg. Chem.</i> <b>14</b> , 751-760.
88.	Ojha, H., Murari, B. M., Anand, S., Hassan, M.I., <b>Ahmad, F.</b> , and Chaudhury, N. K. (2009) Interaction of DNA minor groove binder Hoechst 33258 with BSA. <i>Chem. Pharm. Bull.</i> <b>57</b> , 481-486.
89.	Salami, M., Yousefi, REhsani, M. R., Razavi, S. H., Chobert, J-M, Haertlé, T., Saboury; A.A., Atri, M. S A., Niasari-Naslaji, A., <b>Ahmad, F.</b> and Moosavi-Movahedi, A.A. (2009) Enzymatic Digestion and Antioxidant Activity of Native and MG State of Camel $\alpha$ -Lactalbumin: Possible use in infant formula. <i>Intl. Dairy J.</i> <b>19</b> , 518-523.
90.	Divsalar, A., Sabouryl, A.A., Mansooti-Torshizi, H., Moghaddam, M.I., <b>Ahmad, F.</b> , Hakimelahi, G. H. (2009) Comparative Studies on the Interaction Between Bovine $\beta$ -Lactoglobulin A and B and a New Designed Pd(II) Complex with Anti-tumor Activity at Different Temperatures, <i>J. Biomol., Stru. Dyna.</i> <b>26</b> , 587-597.
91.	Badraghi, B., Yousefie, R., Saboury, A. A., Sharifzadeh, A., Heartle T., <b>Ahmad, F.</b> And Moosavi-Movahedi, A.A. (2009) Effects of Salt and Sodium dodecyl Sulphate on Cheperone Activity of $\alpha$ S <sub>1</sub> -CN: Insulin as the Target Protein, <i>Colloid Surfaces B: Biointerfaces</i> <b>71</b> , 300-305.
92.	Sharifi, E., Sattarahmady, N., Habibi-Rezaei, M., Farhadi, M., Sheibani, N., <b>Ahmad, F.</b> and Moosavi-Movahedi, A. A. (2009) Inhibitory effects of beta-cyclodextrin and trehalose on nanofibril and AGE formation during glycation of human serum albumin, <i>Protein Pept. Lett.</i> <b>16</b> , 653-659.
93.	Jamal, S., Poddar, N. K., Singh, L. R., Dar, T. A., Rishi, V. and <b>Ahmad F.</b> (2009) Relationship between functional activity and protein stability in the presence of all classes of stabilizing osmolytes, <i>FEBS J.</i> <b>276</b> , 6024-6032.
94.	Moosavi-Movahedi, A. A., Mousavy, S. J., Divsalar, A., Babaahmadi, A., Karimian, K., Shafiee, A., Kamarie, M., Poursasan, N., Farzami, B., Riazzi, G. H., Hakimelahi, G. H., Tsai, F.-Y., <b>Ahmad, F.</b> , Amani, M. and Saboury, A. A. (2009) The effects of Deferipone and Deferasirox on the structure and function of $\square$ -Thalassemia Hemoglobin, <i>J. Biomol. Strc. Dyna.</i> <b>27</b> , 319-329.
95.	Minakshi, R., Padhan, K., <b>Ahmad, F.</b> and Jameel, S. (2009) The SARS coronavirus 3a protein causes endoplasmic reticulum stress and induces ligand-independent downregulation of the type 1 interferon receptor, <i>PLose ONE</i> <b>4</b> , 1-10
96.	Divsalar, A., Saboury, A. A, <b>Ahmad, F.</b> and Moosavi-Movahedi, A. A. (2009) Effects of temperature and chromium (III) ion on the structure of bovine $\square$ -lactoglobulin-A, <i>J. Braz. Chem. Soc.</i> <b>20</b> , 1782-1789.
97.	Kumar, P., Islam, I, <b>Ahmad, F.</b> and Satyanarayana, T. (2009) Characterization of a Neutral and Thermostable Glucoamylase from the Thermophilic Mould <i>Thermomucor indicae-seudaticae</i> : Activity, Stability and Structural Correlation, <i>Applied Biochem. Biotechnol.</i> <b>160</b> , 879-890.

98.	Hassan, M. I., Naiyer, A. and <b>Ahmad, F.</b> (2010) Fragile histidine triad protein: structure, function, and its association with tumorigenesis, <i>J. Cancer Res. Clin. Oncol.</i> <b>136</b> , 333-350.
99.	Hassan, M. I., Toor, A. and <b>Ahmad, F.</b> (2010) Progastriscin: structure, function and its role in tumour progression, <i>J. Mol. Cell Biol.</i> <b>2</b> , 118-127.
100.	Hassan, M. I., Aijaz, A. and <b>Ahmad, F.</b> (2010) Structural and functional analysis of human prostatic acid phosphatase, a prognostic biomarker for prostate carcinoma, <i>Expert Rev. Anticancer Therapy</i> <b>10</b> , 1055-1068.
101.	Moosavi-Movahedi, A.A., Terani, H.S., Amanlou, M., Rad, M.N.S., Hakimelahi, G.H., Tsai, F.-Y., Ataie, G., Saboury, A. A., and <b>Ahmad, F.</b> , Khalafi-Nezhad, A., Poursasan, N., and Sharifzadeh, A. (2010) Kinetics and Conformational Studies of Adenosine Daminase upon Interaction with Oxazepam and Lorazepam, <i>Protein Pept. Lett.</i> <b>17</b> , 197-205.
102.	Divasalar, A, Saboury, A.A., Mansoor-Torshizi, H., and <b>Ahmad, F.</b> (2010) Design, Synthesis and Biological Evaluation of a New Palladium(II) Complex: Lactoblobulin and K562 as Targets, <i>J. Phys. Chem.. B</i> <b>114</b> , 3639-3647.
103.	Ali, S., Farooqi, H., Prasad, R., Naim, M., Routray, I., Yadav, S., and <b>Ahmad, F.</b> (2010) Boron stabilizes Peroxide Mediated Changes in the Structure of Heme Proteins, <i>Intl J. Biol. Macromol</i> <b>47</b> , 109-115.
104.	Khan, M.K.A., Rahaman, M. H., Hassan, M.I., Singh, T.P., Moosavi-Movahedi, A.A., and <b>Ahmad, F.</b> (2010), Conformational and Thermodynamic Characterization of the Premolten Globule State Occurring During Unfolding of Molten Globule State of Cytochrome- C, <i>J. Biol. Inorg. Chem.</i> <b>15</b> , 1319-1329.
105.	Poddar, N.K., Ansari, Z.A., Singh, B.K., Moosavi-Movahedi, A.A., <b>Ahmad, F.</b> (2010), effect of Oligosaccharide and Their Monosaccharide Mixtures on the Stability of Proteins: A Scaled Particle Theory, <i>J. Biomol. Struc. Dyna.</i> <b>28</b> , 331-334.
106.	Khan, S. H., Ahmad, N., <b>Ahmad, F.</b> and Kumar, R. (2010), naturally Occurring Osmolytes: From Cell Physiology to Disease Prevention, <i>IUBMB Life</i> <b>62</b> , 891-895.
107.	Singh, R.L., Poddar, N.K., Dar, T.A., Kumar, R. and <b>Ahmad, F.</b> (2011), Protein and DNA Destabilization by Osmolytes: the other side of the coin, <i>Life Sciences</i> <b>88</b> , 117-125.
108.	Farooqui, H., <b>Ahmad, F.</b> and Ali, S. (2011), Bron Increases the Transition temperature and Enhances Thermal Stability of heme Proteins, <i>J. thermal Analysis and Calorimetry</i> <b>104</b> , 339-342.
109.	Khan, S. H., <b>Ahmad, F.</b> , Ahmad N., Flynn, D. C. and Kumar, R. (2011), Protein-protein interactions: Principles, techniques, and their potential role in new drug development, <i>J. Biomol. Struc. Dyna.</i> <b>28</b> , 929-938.
110.	Singh, L. R., Poddar, N. K., Dar, T. A., Rahman, S., Kumar, R. and <b>Ahmad, F.</b> (2011), Forty Years of Research on Osmolyte-induced Protein Folding and Stability, <i>Iran. J. Chem. Soc.</i> <b>8</b> , 1-23.
111.	<b>Hassan, M. I. and Ahmad, F.</b> (2011), Structural Specificities of Classical and Non-classical MHCs Towards Ligand and Beta2-Microglobulin Binding, <i>Adv. Protein Chem. Stru. Biol.</i> <b>83</b> , 223-270.

112.	Khan, M. K. A., Rahaman, M. H., and <b>Ahmad, F.</b> (2011), Conformation and thermodynamic stability of pre-molten and molten globule states of mammalian cytochromes-c, <i>Metalomics</i> <b>3</b> , 327-2338.
113.	Rehman, M.T., Dey, P., Hassan, M.I., <b>Ahmad, F.</b> , and Batra, J.K. (2011). Functional role of glutamine 28 and arginine 39 in double stranded RNA cleavage by human pancreatic ribonuclease. <i>PLoS One</i> <b>6</b> , e17159.
114.	Moosavi_Movahedi, A. A., Rajabzaheh, H., Amani, M., Nourrouzian, D., Zare, K., Hadi, H., Sharifzadeh, A., Poursasan, N., <b>Ahmad, F.</b> , and Sheibani, N. (2011), Acidic Residue Modifications Restores Chaperone Activity of B-casein Interacting with Lysozyme, <i>Intl. J. Biol. Macromol.</i> <b>49</b> , 616-621.
115.	Ariaeenejad A, Habibi-Rezaei M, Jamili S, Fatemi MR, Poursasan, N, <b>Ahmad F</b> , Sheibani N, Kavousi K, and Moosavi-Movahedi AA (2012) Biochemical Characterization of Hemoglobins from Caspian Sea Sturgeons (Acipenser percicus and Acipenser stellatus), <i>Cell Biochem. Biophys.</i> <b>62</b> , 73-81
116.	Hassan MI, Saxena A and <b>Ahmad F</b> (2012) Structure and function of von Willebrand factor, the protein that is deficient and/or abnormal in inherited von Willebrand disease, <i>Blood Coagul Fibrinolysis</i> , <b>23</b> , 11-22.
117.	Alaei, L, Moosavi-Movahedi, AA, Hadi, H, Saboury, AA, <b>Ahmad, F</b> and Amani, M (2012), Thermal inactivation and conformational lock of bovine carbonic anhydrase, <i>Protein Pept. Lett</i> <b>19</b> , 852-858.
118.	Singh, L.R. and <b>Ahmad, F.</b> (2013) Compatible Osmolytes Are Like of S. O. S. Thing in the Living Cells, <i>Proc. INSA</i> <b>78</b> , 701-711.
119.	Hassan, M. I., Waheed, A., Chang, Y-H., Sly, W. S., <b>Ahmad, F.</b> , and Fleming, R. E. (2013). Characterization and Quantization of Hepcidin from Human and Mouse Serum and Secretion Medium from Human Cell Line, Hu.7 by Malditof Mass Spectrometry, <i>J. Protein Proteomics</i> <b>3</b> , 177-185.
120.	Hassan, M. I., Shaji, B., Waheed, A., <b>Ahmad, F.</b> And Sly, W. S. (2013), Structure, Function and Applications of Carbonic Anhydrase Isozymes, <i>Bioinorg. Med. Chem.</i> <b>21</b> , 1570 - 1582.
121.	Khan, S., Bano, Z., Singh, L. R., Hassan, M. I., Islam, A. and <b>Ahmad, F.</b> (2013), Why Is Glycine not a Part of Osmoticum in the Urea Rich Cell, <i>Protein Pept. Lett.</i> <b>20</b> , 61-70.
122.	Rahaman, H., Khan, M. K. A, Hassan, Md. I., Islam, A., Moosavi-Movahedi, A. A., <b>Ahmad, F.</b> (2013). Evidence of non-coincidence of normalized sigmoidal curves of two different structural properties for two-state protein folding/unfolding. <i>J. Chm. Therm.</i> <b>58</b> , 351-358.
123.	Bohlooli, M., Moosavi-Movahedi, A. A., Taghavi, F., Habibi-Rezaei, M., Seyedarabi, A., Saboury, A. A., <b>Ahmad, F.</b> (2013). Thermodynamics of a molten globule state of human serum albumin by 3- $\beta$ -hydroxybutyrate as a ketone body, <i>Intl. J. Biol. Macromol.</i> <b>54</b> , 258-263.
124.	Naz, F., Anjum, F., Islam, A., <b>Ahmad, F.</b> , and Hassan, M. I. (2013). Microtubule Affinity-Regulating Kinase 4: Structure, Function and Regulation, <i>Cell. Biochem. Biophys.</i> <b>67</b> , 485 – 499.
125.	Hassan, Md. H., Waheed, A., <b>Ahmad, F.</b> , and Van Etten, R. L. (2013). Fluorescent Dye Conjugates of Rabbit Arylsulfatase A as a Biological Tracer for Protein Endocytosis, <i>Appl. Biochem. Biotechnol.</i> <b>170</b> , 972-979.

126.	Das, A., Saha, T., <b>Ahmad, F.</b> , Roy, K. B., and Rishi, V. (2013), Dodecamer d-AGATCTAGATCT and a Homologous Hairpin form Triplex in the Presence of peptide REWER. <i>PLOS ONE</i> 8, e65010. doi: 10.1371/journal.pone.0065010
127.	Naz, H., Islam, A., Waheed, A., <b>Ahmad, F.</b> , Sly, W.S., and Hassan, M.I. (2013), Human Glucuronidase: Structure, Function and Application in Enzyme Replacement Therapy. <i>Rejuvenation Res.</i> 16,352-63.
128.	Khan, S., Bano, Z., Singh L. R., Hassan, M. I., Islam, A., <b>Ahmad, F.</b> (2013), Testing the Ability of Non-Methylamine Osmolytes Present in Kidney Cells to Counteract the Deleterious Effects of Urea on Structure, Stability and Function of Proteins. <i>PLOS ONE</i> 8, e72533.
129.	Ubaid-ullah, S., Haque, M. A., Zaidi, S., Hassan, M. I., Islam, A., Batra, J. K., Singh, T. P., and Ahmad, F. (2014) Effect of sequential deletion of extra N-terminal residues on the structure and stability of yeast iso-1-cytochrome-c, <i>J. Biomol. Stru. Dyna.</i> 32, 2005-2016.
130.	Shahbaz M, Ahmad F and Hassan MI (2013) Functional Annotation of Conserved Hypothetical Proteins from Haemophilus influenzae Rd KW20. <i>PloS ONE</i> 8: e84263.
131.	Thakur P, Prakash A, Fleming RE, Waheed A, <b>Ahmad F</b> and Hassan MI, (2014) Identification of interfacial residues involved in Heparin-Ferroproteom interaction and their role in Iron Homeostasis. <i>Lett. Drug Design Discovery</i> 11, 363-374.
132.	Tehrani, H. S., Moosavi-Movahedia, A. A., Ghourchiana, H., Ahmad, F., Kianya, A., Atria, M. S., Ariaeenejada, Sh., Kavousid, K., and Saboury, a. A. (2013), Effect of compatible and noncompatible osmolytes on the enzymatic activity and thermal stability of bovine liver catalase, <i>J. Biomol. Struc. Dyna.</i> 31, 1440 – 1454
133.	Naz, F., Asad, M., Malhotra, P., Islam, A., Ahmad, F., and Hassan, M. I. (2014) Cloning, Expression, Purification and Refolding of Microtubule Affinity-Regulating Kinase 4 Expressed in Escherichia coli, <i>Appl. Biochem. Biotech.</i> 172, 2838-2848.
134.	Taghavi, F., Moosavi-Movahedi, A. A., Bohlli, M, Habibi-Razaei, M., Alijavand, H. H., Amanlou, M, Sheibani, N., Sabouri, A. A., and Ahmad, F. (2014). Energetic Domains and Conformational Analysis of Human Serum Albumin upon Co-incubation with Sodium Benzoate and Glucose, <i>J. Biomol. Struct. Dyn.</i> 32, 438-447.
135.	Zaidi, S., Hassan, M. H., Islam, A., and Ahmad, F. (2014). The role of key residues in structure and stability of cytochrome c, <i>Cell. Mol. Life Sci</i> 71, 229-255.
136.	Prakash A, Islam A, <b>Ahmad F</b> and Hassan MI (2014) Development of Novel and Potent Carbonic Anhydrase-IX Inhibitor using Pharmacophore Modelling. <i>J. Carcinogenesis &amp; Mutagenesis</i> S8:003.
137.	Anwar, K., Parmar, A., Rahman, S., Kaushal, A., Madamwar, D., Islam, A, Hassan. M. I., and <b>Ahmad, F.</b> (2014), Folding and stability studies on C-PE and its natural N-terminal truncant, <i>Arch. Biochem. Biophys.</i> 545, 9-21.
138.	Bohlooli M, Moosavi-Movahedi AA, Taghavi F, Saboury AA, Maghami P, Seyedarabi A, Moosavi-Movahedi F, <b>Ahmad F</b> , Shockravi A, Habibi-Rezaei M.(2014), <u>Inhibition of fluorescent advanced glycation end products (AGEs) of</u>

	human serum albumin upon incubation with 3- $\beta$ -hydroxybutyrate, <i>Mol. Biol. Rep.</i> <b>41</b> , 3705-13. doi: 10.1007/s11033-014-3235-1
139.	Khan P, Idress D, Moxley MA, Corbett, JA, Ahmad, F., von Figura G, Sly WS, Waheed A and Hassan, M. I. (2014), Luminol-based Chemiluminescent Signals: Clinical and Non-clinical Application and Future Uses", <i>Applied Biochemistry and Biotechnology</i> . <i>Appl. Biochem. Biotechnol.</i> <b>173</b> , 333-55.
140.	Sinha A, Ahmad F, Hassan M I (2015), Structure based functional annotation of putative conserved proteins from <i>Treponema pallidum</i> : search for a potential drug target. <i>Lett Drug Des Disco</i> <b>12</b> , 46-59.
141.	Ali, S. A., Hassan, M I., Islam, A., and <b>Ahmad, F.</b> (2014) A Review of Methods Available to Estimate Solvent-Accessible Surface Areas of Soluble Proteins in the Folded and Unfolded States, <i>Curr. Protein Pept. Sc.</i> <b>15</b> , 456-476.
142.	Kumar, K., Prakash, A., Tasleem, M., Islam, A., <b>Ahmad, F.</b> , and Hassan, M. I. (2014) Functional annotation of putative hypothetical proteins from <i>Candida dubliniensis</i> , <i>Gene</i> <b>543</b> , 93-100.
143.	Minakshi, R., Padhan, K., Rehman, S., Hassan, M. I., and Ahmad, F. (2014) The SARS Coronavirus 3a protein binds calcium in its cytoplasmic domain, <i>Virus Res.</i> <b>191</b> , 180-183.
144.	Sinha, A., Ahmad, F., and Hassan, I. (2015) Structure based functional annotation of putative conserved proteins from search for a potential drug target, <i>Lett. Drug Design Discovery</i> <b>12</b> , 46-59.
145.	Tasleem, M., Ishrat, R., Islam, A., Ahmad, F., and Hassan, I. (2014) Structural characterization, homology modeling and docking studies of ARG674 Mutation in MyH8 Gene associated with trismus-pseudocamptodacty syndrome, <i>Lett. Drug Design Discovery</i> <b>11</b> , 1177-1187.
146.	Dar, M. A., Islam, A., Hassan, M. I., and Ahmad, F. (2014) Purification and Characterization of Calreticulin: a Ca <sup>2+</sup> -Binding Chaperone from Sheep Kidney, <i>Appl. Biochem. Biotech.</i> <b>174</b> , 1771-1783.
147.	Divsalar A, Razmi, M, Saboury A A, Mansouri-Torshizi, H, and Ahmad, F (2014) Biological Evaluation of a New Synthesized Pt(II) Complex by Cytotoxic and Spectroscopic Studies, <i>Cell Biochem. Biophys.</i> , DOI 10.1007/s12013-014-0364-z
148.	Goodarzi, M., Moosavi-Movahedi, A.A., Habibi-Rezaei, M., Shouriana, M., Ghourchian, H., <b>Ahmad, F.</b> , Farhadi, M., Saboury, A.A., and Sheibani, N (2014), Hemoglobin fructation promotes heme degradation through the generation of endogenous reactive oxygen species, <i>Spectrochim. Acta Part A: Mol. Biomol. Spectrosc.</i> <b>130</b> , 561–567.
149.	Haque A, Ubaidullah S, Zaidi S, Hassan MI, Islam A, Batra JK and Ahmad F (2015) In vitro and in silico studies of urea-induced denaturation of yeast iso-1-cytochrome c and its deletants at pH 6.0 and 25 °C. <i>J. Biomol. Struc. Dyna.</i> <b>33</b> ,1493-1502.
150.	Shahbaz M, <b>Ahmad F</b> and Hassan MI (2015) Sequence and Structure Analysis Putative Conserved Proteins having Lyase Activity from <i>Haemophilus influenzae</i> Rd KW20. <i>3Biotech</i> <b>5</b> , 317-336.

151.	Haque, M. A., Ubaid-ullah, S., Zaidi, S., Hassan, M. I., Islam, A., Batra, J. K., and Ahmad, F. (2015) Characterization of pre-molten globule state of yeast iso-1-cytochrome c and its deletants at pH 6.0 and 25° C, <i>Intl. J. Biol. Macromol.</i> 72, 1406-1418.
152.	Kumar K, Prakash A, Islam A, <b>Ahmad F</b> and Hassan MI (2015) Structure based Functional Annotation of Hypothetical Proteins from <i>Candida dubliniensis</i> : A Quest for Novel Drug Target, <b>3Biotech</b> 5, 561–576
153.	Naiyer A., Hassan M.I., Islam A.; Sundd M., Ahmad F. (2015) Structural characterization of MG and pre-MG states of proteins by MD simulations, NMR, and other techniques; <i>J. Biomol. Struc. Dyna.</i> 33(10):2267-2284
154.	Anwer, K., Sonani, R., Madamwar, D., Singh, P., Khan, F., Bisetty, K., Ahmad, F., and Hassan, M. I. (2015) Role of N-terminal residues on folding and stability of C-phycoerythrin: simulation and urea-induced denaturation studies, <i>J. Biomol. Struc. Dyna.</i> 33, 121-133.
155.	Khatibi A, Ma'mani L, Khodarahmi R, Shafiee A, Maghami P, Ahmad F, Sheibanif N, and Moosavi-Movahedi, AA (2015) Enhancement of thermal reversibility and stability of human carbonic anhydrase II by mesoporous nanoparticles, <i>Intl. J. Bio. Macromol.</i> 75, 67–72.
156.	Rahman S, Rehman M T, Singh, L R, Warepam, M, <b>Ahmad F</b> and Dar T A (2015) Salt Potentiates Methylamine Counteraction System to Offset the Deleterious Effects of Urea on Protein Stability and Function, <b>PLOS ONE</b> 10(3):e0119597
157.	Atri, M S, Saboury, A A, and Ahmad, F (2015) Biological Applications of Isothermal Titration Calorimetry, <i>Phys. Chem. Res.</i> DOI: 10.22036/PCR.2015.11066
158.	Shahbaaz M., Ahmad F., Hassan M.I. (2015) Structure-based function analysis of putative conserved proteins with isomerase activity from <i>Haemophilus influenzae</i> ; <b>3 Biotech</b> 5, 741–763
159.	Naqvi, AAT., Shahbaaz, M., <b>Ahmad, F.</b> , and Hassan, M.I. (2015) Identification of functional candidates amongst hypothetical proteins of <i>Treponema pallidum</i> ssp. pallidum. <b>PLoS One.</b> 10(4): e0124177.
160.	Rahaman, H., Alam, K. M., Hassan, M., Islam, A., Moosavi-Movahedi, A., and <b>Ahmad, F.</b> (2015) Heterogeneity of Equilibrium Molten Globule State of Cytochrome c Induced by Weak Salt Denaturants under Physiological Condition, <b>PloS One</b> 10, e0120465.
161.	Shahbaaz, M., Bisetty K., <b>Ahmad, F.</b> , and Hassan, M.I. (2015) Functional prediction of putative uncharacterized proteins of <i>Neisseria meningitidis</i> MC58 and its virulence characterization. <b>OMICS: J. Integrative Biol.</b> 19, 416-34.
162.	Zaidi S., Hassan M.I., Islam A., <b>Ahmad F.</b> (2015) Structural Characteristics of Stable Folding Intermediates of Yeast Iso-1-Cytochrome-c, <b>Biomacromole. J.</b> 1, 19-45.

163.	Naqvi A.A.T., <b>Ahmad F.</b> , Hassan M.I. (2015) Identification of functional candidates amongst hypothetical proteins of Mycobacterium leprae Br4923, a causative agent of leprosy; <i>Genome</i> <b>58</b> , 25-42.
164.	Beg I, Minton AP, Hassan M.I., Islam A, <b>Ahmad F.</b> (2015) Thermal Stabilization of Proteins by Mono- and Oligosaccharides: Measurement and Analysis in the Context of an Excluded Volume Model. <i>Biochemistry</i> <b>54</b> , 3594-603.
165.	Singh R, Hassan MI, Islam A and <b>Ahmad F.</b> (2015) Cooperative Unfolding of Residual Structure in Heat Denatured Proteins by Urea and Guanidinium Chloride, <i>Plos One</i> <b>10</b> (6): e0128740
166.	Shahbaaz, M., Bisetty K., <b>Ahmad, F.</b> , and Hassan, M.I. (2015) Functional Insight into Putative Uncharacterized Proteins of Rickettsia rickettsii and its Virulence Characterization. <i>Curr. Proteomics</i> <b>12</b> (2): 101 - 116.
167.	Naz F, Singh P, Islam A, <b>Ahmad F</b> , Hassan MI. (2015) Human microtubule affinity-regulating kinase 4 is stable at extremes of pH. <i>J. Biomol. Struc. Dyna.</i> <b>21</b> , 1-11. PMID: 26208600.
168.	Naz F, Islam A, <b>Ahmad F</b> , Hassan MI. (2015), A typical PKC phosphorylates microtubule affinity-regulating kinase 4 in vitro. <i>Mol. Cellular Biochem.</i> <b>410</b> (1-2):223-8.
169.	Shahid S, Hassan MI, <b>Ahmad F</b> , Islam A (2015) Relationship between protein stability and functional activity in the presence of macromolecular crowding agents alone and in mixture: an insight into stability-activity trade-off. <i>Arch. Biochem. Biophys.</i> <b>584</b> , 42-50.
170.	Shahbaaz, M., Bisetty, K., <b>Ahmad, F.</b> , and Hassan, M.I. (2015) In silico approaches for the identification of virulence candidates amongst hypothetical proteins of Mycoplasma pneumoniae 309. <i>Comput. Biol. Chem.</i> <b>59</b> (Pt A):67-80.
171.	Naz F, Shahbaaz, M., Khan S, Bisetty K., Islam A, <b>Ahmad F</b> , Hassan MI. (2015) PKR-inhibitor binds efficiently with human microtubule affinity-regulating kinase 4. <i>J. Mol. Grap. Model.</i> <b>62</b> , 245-52.
172.	Naz F, Shahbaaz, M., Bisetty K., Islam A, <b>Ahmad F</b> , Hassan MI. (2015) Designing new kinase inhibitor derivatives as therapeutics against common complex diseases: Structural basis of microtubule affinity-regulating kinase 4 (MARK4) inhibition. <i>OMICS: J. Integ. Biol.</i> <b>19</b> , 700-11
173.	Tasleem M., Ishrat R, Islam A, <b>Ahmad F</b> and Hassan MI (2015) Human Disease Insight: An integrated knowledge-based platform for disease-gene-drug information. <i>J. Infec. Public Health</i> <b>S1876-0341</b> , 00202-6.

174.	Khan S, Shhahbaaz M, Bisetty K, Islam A, <b>Ahmad F</b> and Hassan MI (2015) Classification and functional analyses of putative conserved proteins from Chlamydophila pneumoniae CWL029. <i>Interdisciplinary S. Comput. Life Sci.</i> PMID: 26649559.
175.	Khan W H, Raghuram VLN, Srungaram, VLNR, Islam A, Beg I, Haider MSH, <b>Ahmad F</b> , Broor S, and Parveen S (2015), Biophysical Characterization of G Protein Ectodomain of Group B Human Respiratory Syncytial Virus from E. Coli, <i>Prep. Biochem. Biotech.</i> <b>46</b> , 483–488.
176.	Hoda N, Naz H, Jameel E, Shandilya A, Dey S, Hassan M I, <b>Ahmad F</b> , Jayaram B (2016) Curcumin Specifically binds to the Human Calcium-calmodulin Dependent Protein Kinase IV: Fluorescence and Molecular Dynamics Simulation Studies, <i>J. Biomol. Struc. Dyna.</i> 1-38 (PMID: 25929263).
177.	Idrees D, Parkash A, Islam A, <b>Ahmad F</b> and Hassan MI (2016) Spectroscopic and MD simulation studies on unfolding processes of mitochondrial carbonic anhydrase VA induced by urea. <i>J. Biomol. Struc. Dyna.</i> 34(9),1987-1997.
178.	Shahbaaz, M., Bisetty K., <b>Ahmad, F.</b> , and Hassan, M.I. (2016) Current advances in the identification and characterization of putative drug and vaccine targets in the bacterial genomes. <i>Curr. Topics Medi. Chem.</i> <b>16</b> (9), 1040-1069.
179.	Naz H, Islam A, <b>Ahmad F</b> , Hassan MI (2016) Calcium/Calmodulin-dependent protein kinase IV: A multifunctional enzyme and potential therapeutic target. <i>Prog. Biophys. Mol. Biol.</i> <b>121</b> , 54-65.
180.	Khan FI, Aamir M, Wei DQ, <b>Ahmad F</b> , Hassan MI. (2017) Molecular mechanism of Ras-related protein Rab-5A and effect of mutations in the catalytically active phosphate-binding loop. <i>J. Biomol. Struc. Dyna.</i> 1-36. PMID: 26727234
181.	Naz H, Shahbaaz M, Bisetty K, Islam A, <b>Ahmad F</b> , Hassan MI (2016) Effect of pH on the structure, function and stability of human calcium/calmodulin-dependent protein kinase IV: A combined spectroscopic and MD simulation studies. <i>Biochem. Cell Biol.</i> <b>94</b> , 221-228.
182.	Anwer K, Rahman S, Sonani R, Khan FI, Islam A, Madamwar D, <b>Ahmad F</b> and Hassan MI (2016) Probing pH sensitivity of $\alpha$ C-phycoerythrin and its natural truncant: A comparative study. <i>Intl. J. Biol. Macromol.</i> <b>86</b> , 18-27.
183.	Khan FI, Shahbaaz M, Bisetty K, Waheed A, Sly WS, <b>Ahmad F</b> and Hassan MI (2016) Large scale analysis of the mutational landscape in $\beta$ -glucuronidase: A major player of mucopolysaccharidosis type VII, <i>Gene</i> <b>576</b> (1 Pt 1), 36-44.

184.	Naz H, Jameel E, Hoda N, Shandilya A, Khan P, Islam A, <b>Ahmad F</b> , Jayaram B, Hassan MI (2016) Structure guided design of potential inhibitors of human calcium-calmodulin dependent protein kinase IV containing pyrimidine scaffold, <i>Bioorg. Medi. Chem. Lett.</i> <b>26</b> , 782-788.
185.	Idrees D, Kumar D, Rehman SA, Gourinath S, Islam A, <b>Ahmad F</b> and Hassan MI (2016) Cloning, Expression, Purification of Characterization of Human Mitochondrial Carbonic Anhydrase VA. <i>3Biotech</i> <b>6</b> , 1-8.
186.	Amir M, Wahiduzzaman, Dar MA, Haque MA, Islam A, <b>Ahmad F</b> and Hassan MI (2016) Purification and characterization of oligonucleotide binding (OB)-fold protein from medicinal plant <i>Tinospora cordifolia</i> . <i>J Chromatogr B, Analyt. Technol. Biomed. Life Sci.</i> <b>1008</b> , 38-44.
187.	Khan P, Parkash A, Islam A, <b>Ahmad F</b> , Hassan MI. (2016) Molecular basis of the structural stability of hemochromatosis factor E: A combined molecular dynamic simulation and GdmCl-induced denaturation study. <i>Biopolymers</i> <b>105</b> , 133-142.
188.	Rahman S, Ali SA, Hassan MI, Islam A and <b>Ahmad F</b> . (2016) Testing the dependence of stabilizing effect of osmolytes on the fractional increase in the accessible surface area on thermal and chemical denaturations of proteins. <i>Arch. Biochem. Biophys.</i> <b>12</b> , 591:7-17.
189.	Amir M, Wahiduzzaman, Dar MA, Haque MA, Islam A, <b>Ahmad F</b> and Hassan MI (2016) Purification and Characterization of Ras related protein, Rab5a from <i>Tinospora Cordifolia</i> . <i>Intl. J. Biol. Macromol.</i> <b>82</b> , 471-479.
190.	Ishrat M, Hassan MI, <b>Ahmad F</b> , Moosavi-Movahedi AA, Islam A (2016) Effect of dextran macromolecular crowding on the thermodynamic stability and structure of ribonuclease A. <i>J. Iran. Chem. Soci.</i> <b>13</b> (1), 181-189.
191.	Amir M, Dar MA, Wahiduzzaman, Islam A, <b>Ahmad F</b> , Hassan MI (2016) Purification and characterization of RGA2, a Rho2 GTPase-activating protein from <i>Tinospora cordifolia</i> . <i>3 Biotech</i> <b>6</b> , 85-.
192.	Kumari S, Idrees D, Mishra CM, Prakash P, Wahiduzzaman, <b>Ahmad F</b> , Hassan MI, Tiwari M (2016) Design and synthesis of a novel class of carbonic anhydrase-IX inhibitor 1-(3-(phenyl/4-fluorophenyl) -7-imino-3H-[1,2,3] triazolo[4,5d]pyrimidin 6 (7H)yl)urea, <i>J. Mol. Graph. Model.</i> <b>64</b> , 101-109.
193.	Idrees D, Shahbaaz M, Bisetty K, Islam A, <b>Ahmad F</b> , Hassan MD (2016), Effect of pH on structure, function and stability of mitochondrial carbonic anhydrase VA. <i>J. Biomol. Struc. Dyna.</i> <b>4</b> , 1-13.

194.	Naz, H, Shahbaaz M, Haque MA, Bisetty K, Islam A, <b>Ahmad F</b> , and Hassan MH, Urea-induced denaturation of human calcium-calmodulin dependent protein kinase IV: A combined spectroscopic and MD simulation studies. <i>J. Biomol. Struc. Dyna.</i> <b>8</b> , 1-13.
195.	Khan, S, Jamal, M S, Anjum, F, Rasool, M, Ansari, A, Islam, A, <b>Ahmad, F</b> , and Hassan, M I (2016), Functional annotation of putative conserved proteins from <i>Borrelia burgdorferi</i> to find potential drug targets. <i>Int. J. Computational Biol. Drug Design</i> <b>9</b> , 295-317.
196.	Prakash, A, drees, D, Haque M A, Islam, A, <b>Ahmad, F</b> , and Md. Imtaiyaz Hassan (2017), GdmCl-induced unfolding of studies of human carbonic Anhydrase IX: A combined spectroscopic and MD simulation approach. <i>J. Biomol. Struc. Dyna.</i> <b>35</b> , 1295-1306.
197.	Zaidi, S, Haque, M A, Ubaid-ullah, S, Prakash, A, Hassan, M I, Islam, A, Batra, J K, and <b>Ahmad, F</b> (2017), Denatured states of yeast cytochrome <i>c</i> induced by heat and guanidinium chloride are structurally and thermodynamically different. <i>J. Biomol. Struc. Dyna</i> <b>35</b> , 1420-1435.
198.	Dar, M A, Wahiduzzaman, Haque, M A, Islam, A, Hassan, M I, and <b>Ahmad, F</b> (2016), Characterisation of molten globule-like state of sheep serum albumin at physiological pH. <i>Int. J. Biol. Macromol.</i> <b>89</b> , 605-613.
199.	Khan, P, Shandilya, A, Jayaram, B, Islam, A, <b>Ahmad, F</b> , and Hassan, M. D (2017), Effect of pH on the stability of hemochromatosis factor E: A spectroscopic and molecular dynamics simulation-based approach <i>J. Biol. Struc. Dyna.</i> <b>35</b> , 1582-1598.
200.	Kumar, V, Islam, A, Hassan, MI, and <b>Ahmad, F</b> (2016), Therapeutic progress in amyotrophic lateral sclerosis- beginning to learning. <i>Eur. J. Med. Chem.</i> <b>121</b> , 903-917.
201.	Kumar, V, Islam, A, Hassan, M I*, and <b>Faizan Ahmad</b> (2016), Delineating the relationship between amyotrophic lateral sclerosis and frontotemporal dementia: Sequence and Structure based predictions. <i>Biochim. Biophys. Acta. - Mole. Basis Diseases.</i> <b>1862</b> , 1742–1754.
202.	Khan, S, Islam, A, Hassan, M I, <b>Ahmad, F</b> (2016), Purification and Structural Characterization of Mce4A from <i>Mycobacterium tuberculosis</i> . <i>Int. J. Biol. Macromol.</i> <b>93</b> , 235–241.
203.	Khan, P, Prakash, A, Haque, M A, Islam, A, Hassan, M I, and Ahmad, F (2016), Structural Basis of Urea-induced Unfolding: Unraveling the Folding Pathway of Hemochromatosis Factor E. <i>Int. J. Biol. Macromol.</i> <b>91</b> , 1051–1061
204.	<i>Chowhan, R K, Ali, F, Bhat, M Y, Rahman, S, Singh, L R, Ahmad, F and Dar, T A</i> (2016), Alanine Counteracts the Destabilizing Effect that Urea has on RNase-A. <i>Protein Pept. Lett.</i> <b>23</b> , 795-9.

205.	Idrees, D, Prakash, A, Haque, M A, Islam, A, Hassan, M I, and <b>Ahmad, F</b> (2016), GdnHCl-induced unfolding intermediate in mitochondrial carbonic anhydrase VA. <i>Int. J. Biol. Macromol.</i> <b>91</b> , 1151–1160.
206.	Khan S H, Kumar A, Prakash A, Taneja B, Islam A, Hassan I H, and <b>Ahmad F</b> (2016), Structural and thermodynamic characterisation of L94F mutant of horse cytochrome <i>c</i> . <i>Int. J. Biol. Macromol.</i> <b>92</b> , 202–212.
207.	Kumar V, Kashav T, Islam A, Ahmad F, Hassan MI. (2016) Structural insight into C9orf72 hexanucleotide repeat expansions: Towards new therapeutic targets in FTD-ALS. <i>Neurochem. Int.</i> <b>100</b> , 11-20.
208.	<i>Kumar, V, Sami, N, Kashav, T, Islam, A, Ahmad, F, Hassan, M I (2016), Protein Aggregation and Neurodegenerative Diseases: From Theory to Therapy. Eur. J. Med. Chem. 124, 1105-1120.</i>
209.	Naqvi, A A T, Khan, F I, Anjum, F, Islam, A, Ahmad, <b>F</b> , and Hassan, M I (2016), Sequence analysis of hypothetical proteins from <i>Halobacterium salinarum</i> R1 to identify potential virulence factors. <i>Genomics Inform.</i> <b>14</b> , 125-135.
210.	Naz, F, Sami, N, Islam, A, Ahmad, F, and Hassan, M I (2016), Ubiquitin associated domain of MARK4 provide stability at physiological pH. <i>Int. J. Biol. Moacromol.</i> <b>93</b> , 1147-1154.
211.	Rahman, S., Ali, S A, Islam, A, Hassan, M I, <b>Ahmad, F</b> (2017) Data on the role of accessible surface area on osmolyte-induced protein stabilization. <i>Data in Brief</i> <b>10</b> , 47-56.
212.	Valipour M, Maghami P, Habibi-Rezaei M, Sadeghpour M, Khademian M A, Mosavi K, <b>Ahmad F</b> , Moosavi-Movahedi A A (2017) Counteraction of the deleterious effects of reactive oxygen species on hemoglobin structure and function by ellagic acid, <i>J. Luminescence.</i> <b>182</b> , 1–7.
213.	Wahiduzzaman , Dar, MA, Amir, Md, Islam, A, Hassan, M I, <b>Ahmad, F</b> (2017) Purification, preliminary X-ray crystallography and biophysical studies of triose phosphate isomerase- $\beta$ -globin subunit complex. <i>I. J. Biol. Moacromol.</i> <b>94</b> , 746-753.
214.	Shahid, S, Hassan, MI, Islam A, <b>Ahmad, F</b> (2017) Size-dependent studies of macromolecular crowding on the thermodynamic stability, structure and functional activity of the protein: <i>In vitro</i> and <i>in silico</i> approaches. <i>Biochim. Biophys. Acta (General)</i> <b>1861</b> , 178-197.

215.	Beg, I, Minton, A P, Islam, A, Hassan M I, and <b>Ahmad, F</b> (2017), pH-dependence of thermal stabilization of proteins by saccharides: analysis in the text of an excluded volume model. <i>J. Biol. Chem.</i> <b>292</b> , 505-511.
216.	Jameel E, Naz H, Khan P, Tarique M, Kumar J, Mumtazuddin S, Ahamad S, Islam A, <b>Ahmad F</b> , Hoda N, Hassan MI (2017) <u>Design, synthesis and biological evaluation of pyrimidine derivatives as potential inhibitors of human CAMKIV.</u> <i>Chem. Biol. Drug Des.</i> <b>89</b> , 741-754.
217.	Wahiduzzaman, Dar, M A, Haque, M A, Idrees, D, Hassan, M I, Islam A*, and <b>Ahmad, F</b> (2017), Characterization of folding intermediates during urea-induced denaturation of human carbonic anhydrase II. <i>Int. J. Biol. Moacromol.</i> <b>95</b> , 881-887.
218.	Naz H, Khan P, Tarique M, Rahman S, Meena A, Ahamad S, Luqman S, Islam A, <b>Ahmad F</b> , and Hassan MI (2017) Binding studies and biological evaluation of $\beta$ -carotene as a potential inhibitor of human calcium/calmodulin-dependent protein kinase IV. <i>Int. J. Biol. Macromol.</i> <b>96</b> , 161-170.
219.	Sami, N, Rehman S, Kumar, V, Zaidi, S, Islam, A, Ali, S, <b>Ahmad, F</b> , and Hassan, M I (2017), Protein, Aggregation and consequential Human Neurodegenerative Diseases. <i>Int. J. Neurosc.</i> <b>127</b> , 1047-1057.
220.	<u>Dar</u> , MA, Islam, A, Hassan, M I, <b>Ahmad, F</b> (2017), Effect of mammalian kidney osmolytes on the folding pathway of sheep serum albumin. <i>Int. J. Biol. Macromol.</i> <b>97</b> , 625-634.
221.	Parray Z A, <b>Ahmad F</b> , Hassan M I and, Islam, A (2017), Characterization of Intermediate State of Myoglobin in the Presence of PEG 10 under Physiological Conditions. <i>Int. J. Biol. Macromol.</i> <b>99</b> , 241-248.
222.	Idrees D, Rahman S, Shahbaaz M, Haque A, Islam A <sup>1</sup> , <b>Ahmad F</b> , Kim J, and Hassan, M I (2017), Estimation of thermodynamic stability of human carbonic anhydrase IX from urea-induced denaturation and MD simulation studies. <i>Int. J. Biol. Macromol.</i> <b>105</b> , 183-189.
223.	Srivastava S, Syed S B, Kumar V, Islam A, <b>Ahmad F</b> , Hassan, M I (2017), Fas-activated serine/threonine kinase: Structure and function. <i>Gene Reports</i> <b>8</b> , 117-27.
224.	Naz H, Tarique M, Khan P, Luqman S, Ahamad S, Islam A, <b>Ahmad F</b> and Hassan MI (2017) Evidence of vanillin binding to CAMKIV explains the anticancer mechanism in human hepatic carcinoma and neuroblastoma cells. <i>Mol. Cellular Biochem.</i> <b>438</b> , 35-45.

225.	Shahbaaz M, Amir M, Rahman S, Hasan G M, Dohare R, Bisetty K, Kim J, <b>Ahmad F</b> , Hassan M I (2017) Structural insights into Rab21 GTPase activation mechanism by molecular dynamics simulations, <i>Mol. Simulation</i> <a href="http://dx.doi.org/10.1080/08927022.2017.1357813">http://dx.doi.org/10.1080/08927022.2017.1357813</a> .
226.	Nasreen K, Ahamad S, <b>Ahmad F</b> , Hassan M I, and Islam A (2017) Macromolecular crowding induces molten globule state in the native myoglobin at physiological pH, <i>Int. J. Biol. Macromol.</i> <b>106</b> ,130-139.
227.	Shafaei Z, Abazari O, Divsalar A, Ghalandari B, Poursoleiman A, Saboury A A, and <b>Ahmad F</b> (2017) Effect of a Synthesized Amyl-Glycine1, 10-Phenanthroline Platinum Nitrate on Structure and Stability of Human Blood Carrier Protein, Albumin: Spectroscopic and Modeling Approaches, <i>J. Fluoresc.</i> <b>27</b> , 1829–1838
228.	Khan S H, Islam A, Hassan M I, Sharma S, Singh T P, and Ahmad F (2017) Effect of conservative mutations (L94V and L94I) on the structure and stability of horse cytochrome <i>c</i> , <i>Ach. Biochem. Biophys.</i> <b>633</b> , 40-49.
229.	Khan P, Manzoor S, Rahman S, Queen A, Naz F, Hasan GM, Luqman S, Kim J, Islam A, <b>Ahmad F</b> , Hassan MI (2017) <u>Elucidation of Dietary Polyphenolics as Potential Inhibitor of Microtubule Affinity Regulating Kinase 4: In silico and In vitro Studies.</u> <i>Sci. Report</i> <b>7</b> , 9470.
230.	Kumar V, Rahman S, Choudhry H, Zamzami MA, Sarwar Jamal M, Islam A, <b>Ahmad F</b> , Hassan MI (2017) Computing disease-linked SOD1 mutations: deciphering protein stability and patient-phenotype relations, <i>Sci. Rep.</i> <b>7(1)</b> :4678.
231.	Naz F, Sami N, Naqvi AT, Islam A, <b>Ahmad F</b> , Imtaiyaz Hassan M (2017) Evaluation of human microtubule affinity-regulating kinase 4 inhibitors: fluorescence binding studies, enzyme, and cell assays, <i>J. Biomol. Struc. Dyna.</i> <b>35</b> , 3194-3203.
232.	Wahiduzzaman, Dar MA, Haque MA, Idrees D, Hassan MI, Islam A, <b>Ahmad F</b> (2017) Characterization of folding intermediates during urea-induced denaturation of human carbonic anhydrase II. <i>Int. J. Biol. Macromol.</i> <b>95</b> , 881-887.
233.	Sami N, Kumar V, Islam A, Ali S, <b>Ahmad F</b> , Hassan I (2017) Exploring Missense Mutations in Tyrosine Kinases Implicated with Neurodegeneration, <i>Mol. Neurobiol.</i> <b>54</b> , 5085-5106.

234.	Khan P, Shandilya A, Jayaram B, Islam A, <b>Ahmad F</b> , Hassan MI (2017) Effect of pH on the stability of hemochromatosis factor E: a combined spectroscopic and molecular dynamics simulation-based study, <i>J. Biomol. Struc. Dyna.</i> <b>35</b> ,1582-1598.
235.	Prakash A, Idrees D, Haque MA, Islam A, <b>Ahmad F</b> , Hassan MI (2017) GdmCl-induced unfolding studies of human carbonic anhydrase IX: a combined spectroscopic and MD simulation approach, <i>J. Biomol. Struc. Dyna</i> <b>35</b> ,1295-1306.
236.	Naz H, Shahbaaz M, Haque MA, Bisetty K, Islam A, <b>Ahmad F</b> , Hassan M I (2017) Urea-induced denaturation of human calcium/calmodulin-dependent protein kinase IV: a combined spectroscopic and MD simulation studies, <i>J. Biomol. Struc. Dyna</i> <b>35</b> , 463-475.
237.	Idrees D, Shahbaaz M, Bisetty K, Islam A, <b>Ahmad F</b> , Hassan MI (2017), <u>Effect of pH on structure, function, and stability of mitochondrial carbonic anhydrase VA.</u> <i>J. Biomol. Struc. Dyna.</i> <b>2017</b> , 449-461.
238.	Khan S, Shhahbaaz M, Bisetty K, Islam A, <b>Ahmad F</b> and Hassan MI (2017) Classification and functional analyses of putative conserved proteins from Chlamydophila pneumoniae CWL029. <i>Interdisciplinary Sc. Computational Life Sc.</i> <b>9</b> , 96–106.
239.	Khan FI, Aamir M, Wei DQ, <b>Ahmad F</b> , Hassan MI. (2017) Molecular mechanism of Ras-related protein Rab-5A and effect of mutations in the catalytically active phosphate-binding loop. <i>J. Biomol. Struc. Dyna.</i> <b>35</b> , 105-118.
240.	Zaidi S, Haque MA, Ubaid-Ullah S, Prakash A, Hassan MI, Islam A, Batra JK, <b>Ahmad F</b> . (2017) Denatured states of yeast cytochrome c induced by heat and guanidinium chloride are structurally and thermodynamically different. <i>J. Biomol. Struc. Dyna.</i> <b>35</b> , 1420-1435.
241.	Dar M A, Wahiduzzaman, Islam A, Hassan M I, and Ahmad F (2018) Counteraction of the deleterious effects of urea on structure and stability of mammalian kidney proteins by osmolytes, <i>Int. J. Biol. Macromol.</i> <b>107 (Pt B)</b> , 1659-1667.
242.	Naz F, Khan F I, Mohammad M, Khan P, Hasan, G M, Lobb A, Luqman S, Islam A, <b>Ahmad F</b> and Hassan M I (2018) Investigation of Molecular Mechanism of Recognition between Citral and MARK4: A Newer Therapeutic Approach to Attenuate Cancer Cell Progression, <i>Int. J. Biol. Macromol.</i> <b>107 (Pt B)</b> , 2580-2589.

243.	Amir M, Kumar V, Dohare R, Hussain A, Rehman MT, Alajami MF, Islam A, <b>Ahmad F</b> , Hassan MI. (2018) Investigation of deleterious effects of nsSNPs in POT1 gene: A structural genomics-based approach to understand mechanism of cancer development. <i>J. Cell. Biochem.</i> DOI: 10.1002/jcb.28312
244.	Syed SB, Khan FI, Khan SH, Srivastava S, Hasan GM, Lobb KA, Islam A, Ahmad F and Hassan MI (2018) Mechanistic insights into the urea-induced denaturation of kinase domain of human Integrin linked kinase, <i>Int. J. Biol. Macromol.</i> <b>111</b> , 208-211.
245.	Khan S, Khan FI, Lobb KA, Hasan MG, Islam A, <b>Ahmad F</b> and Hassan MI (2018) Exploring molecular insights into the interaction mechanism of cholesterol derivatives with the Mce4A: A combined spectroscopic and molecular dynamic simulation studies. <i>Int. J. Biol. Macromol.</i> <b>111</b> , 548-560.
246.	Beg I, Islam A, Minton AP, Hassan MI, <b>Ahmad F</b> (2018) Comparison of the thermal stabilization of proteins by oligosaccharides and monosaccharide mixtures: measurement and analysis in the context of excluded volume theory. <i>Biophys. Chem.</i> <b>237</b> , 31-37.
247.	Syed SB, Khan FI, Srivastava S, Khan SH, Hasan MG, Lobb KA, Islam A, <b>Ahmad F</b> and Hassan MI (2018) Unravelling the unfolding mechanism of human integrin linked kinase by GdmCl-induced denaturation. <i>Int. J. Biol. Macromol.</i> <b>117</b> , 1252-1263.
248.	Ghaeidamini M, Kharat AN, Haertlé T, <b>Ahmad F</b> , Saboury AA (2018) <u><math>\beta</math>-Cyclodextrin-Modified Magnetic Nanoparticles Immobilized on Sepharose Surface Provide an Effective Matrix for Protein Refolding</u> , <i>J. Phys. Chem. B.</i> <b>122</b> , 9907-9919.
249.	Bayat M, Gourabi H, Khammari A, <b>Ahmad F</b> , Saboury AA (2018) <u>A comparative study of structure, stability and function of sc-tenectopase in the presence of stabilizing osmolytes</u> . <i>J. Biotechnol.</i> <b>280</b> , 1-10.
250.	Ishrat M, Hassan MI, <b>Ahmad F</b> , Islam A. (2018) Sugar osmolytes-induced stabilization of RNase A in macromolecular crowded cellular environment. <i>Int. J. Biol. Macromol.</i> <b>115</b> , 349-357.
251.	Amir M, Kumar V, Dohare R, Islam A, <b>Ahmad F</b> , Hassan MI (2018) Sequence, structure and evolutionary analysis of cold shock domain proteins, a member of OB fold family. <i>J Evol Biol</i> Sep 29. doi: 10.1111/jeb.13382
252.	<u>Syed B S, Shahbaaz M, Khan SH, Srivastava S, Islam A, Ahmad F and Hassan MI</u> (2017) Estimation of pH effect on the structure and stability of kinase domain of human Integrin linked kinase <i>J. Biomol. Struc. Dyna</i> doi: 10.1080/07391102.2017.1420492

253.	Ali N, Amir M, Hassan MI, <b>Ahmad F</b> , Islam A. (2019) Purification, modeling and structural insights of calmodulin-binding receptor like cytoplasmic kinase 2 from <i>Oroxylum Indicum</i> . <i>Int. J. Biol. Macromol.</i> <b>123</b> , 704-712.
254.	Khan SH, Prakash A, Pandey P, Lynn AM, Islam A, Hassan MI, <b>Ahmad F</b> . (2019) Protein folding: Molecular dynamics simulations and in vitro studies for probing mechanism of urea- and guanidinium chloride-induced unfolding of horse cytochrome- <i>c</i> . <i>Int. J. Biol. Macromol.</i> <b>122</b> , 695-704
255.	Parray ZA, Ahamad S, Hassan MI, <b>Ahmad F</b> and Islam A (2019) First Evidence of Formation of Pre-Molten Globule State in Myoglobin: A macromolecular crowding approach towards protein folding in vivo. <i>Int. J. Biol. Macromol.</i> <b>126</b> , 1288-1299.
256.	Khan S, Khan P, Hassan MI, <b>Ahmad F</b> and Islam A (2019) Protein Stability: Determination of structure and stability of the transmembrane protein Mce4A from <i>M. tuberculosis</i> in membrane-like environment. <i>Int. J. Biol. Macromol.</i> <b>126</b> , 488-495.
257.	Ahamad S, Islam A, <b>Ahmad F</b> , Dwivedi N, Hassan MI (2019) 2/3D-QSAR, molecular docking and MD simulation studies of FtsZ protein targeting benzimidazoles derivatives. <i>Computational Biol. Chem.</i> <b>78</b> , 394-413.
258.	Bayat M, Karami L, Hamid Gourabi H, <b>Ahmad F</b> , Dormiani K, Esfahani MHN, Saboury AA (2019) Stabilizing osmolytes' effects on the structure, stability and function of tc-tenecteplase: A one peptide bond digested form of tenecteplase. <i>J Biol Macromol xx</i> ,
259.	Mohammad T, Khan FI, Lobb KA, Islam A, <b>Ahmad F</b> , Hassan MI. (2017) Identification and evaluation of bioactive natural products as potential inhibitors of human microtubule affinity-regulating kinase 4 (MARK4). <i>J. Biomol. Struct. Dyna.</i> <b>37</b> , 1813-1819.
260.	Beg A, Khan FI, Lobb KA, Islam A, <b>Ahmad F</b> , Hassan MI. (2017) High throughput screening, docking and molecular dynamics studies to identify potential inhibitors of human calcium/calmodulin-dependent protein kinase IV. <i>J. Biomol. Struct. Dyna.</i> <b>37</b> , 2179-2192.
261.	Shahid S, <b>Ahmad F</b> , Hassan MI, Islam A (2019). Mixture of Macromolecular Crowding Agents Has a Non-additive Effect on the Stability of Proteins. <i>Appl Biochem Biotechnol.</i> <b>188</b> , 927-941.
262.	Pandey P, Meena NK, Prakash A, Kumar V, Lynn AM, <b>Ahmad F</b> (2019) Characterization of Heterogeneous Intermediate Ensembles on the Guanidinium Chloride-induced Unfolding Pathway of $\beta$ -lactoglobulin, <i>J Biomol Str Dyn.</i> doi: 10.1080/07391102.2019.1593245

263.	Dahiya R, Mohammad T, Roy S, Anwar S, Gupta P, Islam A, <b>Ahmad F</b> , and Hassan MI (2019) Investigation of inhibitory potential of quercetin to the pyruvate dehydrogenase kinase 3: Towards implications in anticancer therapy <i>I J Bio Macromol</i> <b>136</b> , 1076-1085.
264.	Rahman S, Islam A, Hassan MI, Kim J, and Ahmad F (2019) Unfoldness of the Denatured State of Proteins Determines Urea: Methylamine Counteraction in terms of Gibbs Free Energy of Stabilization, <i>I J Biol Macromol</i> <b>132</b> , 666-676.
265.	Amir M, Kumar V, Dohare R, Rehman T, Hussain A, MF Alajmi, EI Seedi HR, Hassan AMH, Islam A, <b>Ahmad F</b> and Hassan MI (2019) Investigating architecture and structure-function relationships in cold shock DNA-binding domain family using structural genomics-based approach. <i>Int J Biol Macromol</i> <b>133</b> , 484-494.
266.	Naiyer A, Islam A, Hassan MI, <b>Ahmad F*</b> , Sundd M (2019) Backbone and side chain <sup>1</sup> H, <sup>15</sup> N and <sup>13</sup> C chemical shift assignments of the molten globule state of L94G mutant of horse cytochrome-c, Biomolecular NMR Assignments. DOI: 10.1007/s12104-019-09917-7
267.	Amir M, Ahmad S, Ahmad S, Kumar V, Mohammad T, Dohare R, Rehman T, MF Alajmi, Hussain A, <b>Ahmad F</b> and Hassan MI (2019) Structural and functional impact of non-synonymous SNPs in the CST complex subunit TEN1: Structural genomics approach. <i>Biosci Rep.</i> 39(5): BSR20190312.
268.	Amir M, Mohammad T, Kumar V, AlAjmi M, Rehman MT, Hussain A, Khan PA, Dohare R, Islam A, <b>Ahmad F</b> , Hassan MI (2019) Structural Analysis and Conformational Dynamics of STN1 Gene Mutations Involved in Coat plus Syndrome. <i>Front Mol Biosci.</i> <a href="https://doi.org/10.3389/fmolb.2019.00041">https://doi.org/10.3389/fmolb.2019.00041</a>
269.	Shamsi A, Mohammad T, Shahwan M, Hassan MI, <b>Ahmad F</b> , Islam A (2019) Unraveling binding mechanism of Alzheimer's drug Rivastigmine tartrate with human transferrin: Molecular docking and multi spectroscopic approach towards neurodegenerative diseases. doi: 10.3390/biom9090495
270.	Shahid S, Hasan S I, <b>Ahmad F</b> , Hasan M I, Islam A (2019) Carbohydrate based Macromolecular Crowding-Induced Stabilization of Proteins: Towards Understanding the Significance of the Size of Crowder, <i>Biomolecules.</i> <a href="https://doi.org/10.3390/biom9090477">https://doi.org/10.3390/biom9090477</a>
271.	<u>Naqvi</u> AAT, <u>Jairajpuri</u> DS, <u>Noman</u> OMA, <u>Hussain</u> A, <u>Islam</u> A, <b>Ahmad F</b> , <u>Alajmi</u> MF, and <u>Hassan</u> MI (2020) Evaluation of pyrazolopyrimidine derivatives as microtubule affinity regulating kinase 4 inhibitors: Towards therapeutic management of Alzheimer's disease <i>J Biomol Stru Dyna</i> DOI: 10.1080/07391102.2019.1666745
272.	Amir M, Ahmad S, Mohammad T, , <u>Jairajpuri</u> DS, Hasan G, Dohare R, Rehman T, MF Alajmi, Hussain A, <b>Ahmad F</b> and Hassan MI (2020) Investigation of conformational dynamics of Tyr89Cys mutation in protection of telomeres 1 gene

	associated with familial melanoma. <i>J Biomol Stru Dyna.</i> <a href="https://doi.org/10.1080/07391102.2019.1705186">https://doi.org/10.1080/07391102.2019.1705186</a>
273.	Wahiduzzaman, Hassan, M. I., Islam, A., and <b>Ahmad, F.</b> (2020) Urea Stress: Myo-inositol's efficacy to counteract destabilization of TIM-beta-globin complex by urea is as good as that of the methylamine. <i>Int J Biol Macromol 151</i> , 1108-1115.
274.	Shamsi, A., Al Shahwan, M., Ahamad, S., Hassan, M. I., <b>Ahmad, F.</b> , and Islam, A. (2020) Spectroscopic, calorimetric and molecular docking insight into the interaction of Alzheimer's drug donepezil with human transferrin: implications of Alzheimer's drug. <i>J Biomol Struct Dyn 38</i> , 1094-1102.
275.	*Amir, M., Ahmad, S., Ahamad, S., Kumar, V., Mohammad, T., Dohare, R., Alajmi, M. F., Rehman, T., Hussain, A., Islam, A., <b>Ahmad, F.</b> , and Hassan, M. I. (2020) Impact of Gln94Glu mutation on the structure and function of protection of telomere 1, a cause of cutaneous familial melanoma. <i>J Biomol Struct Dyn 38</i> , 1514-1524.
276.	Naiyer, A., Islam, A., Hassan, M. I., <b>Ahmad, F.</b> , and Sundd, M. (2020) Backbone and side chain (1)H, (15)N and (13)C chemical shift assignments of the molten globule state of L94G mutant of horse cytochrome-c. <i>Biomol NMR Assign 14</i> , 37-44
277.	Ghosh, S., Shahid, S., Raina, N., <b>Ahmad, F.</b> , Hassan, M. I., and Islam, A. (2020) Molecular and macromolecular crowding-induced stabilization of proteins: Effect of dextran and its building block alone and their mixtures on stability and structure of lysozyme. <i>Int J Biol Macromol 150</i> , 1238-1248.
278.	Kumar, B., Mohammad, T., Amaduddin, Hussain, A., Islam, A., <b>Ahmad, F.</b> , Alajmi, M. F., Singh, S., Pandey, K. C., Hassan, M. I., and Abid, M. (2020) Targeting metacaspase-3 from Plasmodium falciparum towards antimalarial therapy: A combined approach of in-silico and in-vitro investigation. <i>J Biomol Struct Dyn</i> , J Biomol Struct Dyn. 2021 Feb;39(2):421-430. doi: 10.1080/07391102.2019.1711194.
279.	*Gupta, P., Khan, F. I., Ambreen, D., Lai, D., Alajmi, M. F., Hussain, A., Islam, A., <b>Ahmad, F.</b> , and Hassan, M. I. (2020) Investigation of guanidinium chloride-induced unfolding pathway of sphingosine kinase 1. <i>Int J Biol Macromol 147</i> , 177-186.
280.	Parray, Z. A., <b>Ahmad, F.</b> , Alajmi, M. F., Hussain, A., Hassan, M. I., and Islam, A. (2020) Formation of molten globule state in horse heart cytochrome c under physiological conditions: Importance of soft interactions and spectroscopic approach in crowded milieu. <i>Int J Biol Macromol 148</i> , 192-200.
281.	Shamsi, A., Mohammad, T., Anwar, S., Alajmi, M. F., Hussain, A., Hassan, M. I., <b>Ahmad, F.</b> , and Islam, A. (2020) Probing the interaction of Rivastigmine Tartrate, an important Alzheimer's drug, with serum albumin: Attempting treatment of Alzheimer's disease. <i>Int J Biol Macromol 148</i> , 533-542.

282.	Anwar, S., Kar, R. K., Haque, M. A., Dahiya, R., Gupta, P., Islam, A., <b>Ahmad, F.</b> , and Hassan, M. I. (2020) Effect of pH on the structure and function of pyruvate dehydrogenase kinase 3: Combined spectroscopic and MD simulation studies. <i>Int J Biol Macromol</i> <b>147</b> , 768-777.
283.	Mohammad, T., Siddiqui, S., Shamsi, A., Alajmi, M. F., Hussain, A., Islam, A., <b>Ahmad, F.</b> , and Hassan, M. I. (2020) Virtual Screening Approach to Identify High-Affinity Inhibitors of Serum and Glucocorticoid-Regulated Kinase 1 among Bioactive Natural Products: Combined Molecular Docking and Simulation Studies. <i>Molecules</i> <b>25</b> , 823.
284.	Nasreen, K., Parray, Z. A., Ahamad, S., <b>Ahmad, F.</b> , Ahmed, A., Freeh Alamery, S., Hussain, T., Hassan, M. I., and Islam, A. (2020) Interactions Under Crowding Milieu: Chemical-Induced Denaturation of Myoglobin is Determined by the Extent of Heme Dissociation on Interaction with Crowders. <i>Biomolecules</i> <b>10</b> , 490.
285.	Khan, S. H., Prakash, A., Pandey, P., Islam, A., Hassan, M. I., Lynn, A. M., and <b>Ahmad, F.</b> (2020) Effects of natural mutations (L94I and L94V) on the stability and mechanism of folding of horse cytochrome c: A combined in vitro and molecular dynamics simulations approach. <i>Int J Biol Macromol</i> <b>159</b> , 976-985.
286.	Anwar, S., Shamsi, A., Kar, R. K., Queen, A., Islam, A., <b>Ahmad, F.</b> , and Hassan, M. I. (2020) Structural and biochemical investigation of MARK4 inhibitory potential of cholic acid: Towards therapeutic implications in neurodegenerative diseases. <i>Int J Biol Macromol</i> <b>161</b> , 596-604.
287.	Parray ZA, Hassan MI, <b>Ahmad F</b> , Islam A. (2020) Amphiphilic nature of polyethylene glycols and their role in medical research. <i>Polymer Testing</i> <b>82</b> :106316.
288.	Parray, Z. A., <b>Ahmad, F.</b> , Hassan, M. I., Hasan, I., and Islam, A. (2020) Effects of Ethylene Glycol on the Structure and Stability of Myoglobin Using Spectroscopic, Interaction, and In Silico Approaches: Monomer Is Different from Those of Its Polymers. <i>ACS Omega</i> <b>5</b> , 13840-13850.
289.	Fatima S, Mohammad T, Jairajpuri DS, Samim M, <b>Ahmad F</b> , Alajmi F, Hussain A, Rehman MT, Hassan MI (2020) Identification and evaluation of glutathione conjugate gamma-L-glutamyl-L-cysteine for improved drug-delivery to the brain. <i>J Biomol Struct Dyn</i> . 2020 Aug;38(12):3610-3620. doi: 10.1080/07391102.2019.1664937
290.	Anwar, S., Shamsi, A., Shahbaaz, M., Queen, A., Khan, P., Hasan, G. M., Islam, A., Alajmi, M. F., Hussain, A., <b>Ahmad, F.</b> , and Hassan, M. I. (2020) Rosmarinic Acid Exhibits Anticancer Effects via MARK4 Inhibition. <i>Sci Rep</i> <b>10</b> , 10300.
291.	Amir M, Mohammad T, Dohare R, Islam A, <b>Ahmad F</b> and Hassan MI (2020) Structure, function and therapeutic implications of OB-fold proteins: A lesson from past to present. <i>Briefings in Functional Genomics</i> , elaa008. 19, 377-389.

292.	Amir, M., Mohammad, T., Prasad, K., Mustafa Hasan, G., Kuma, V., Dohare, R., Islam, A., <b>Ahmad, F.</b> and Hassan, I. (2020) Virtual high-throughput screening of natural compounds in-search of potential inhibitors for protection of telomeres 1 (POT1). <i>J Biomol Stru Dyn</i> <b>38</b> , 4625-4634. doi: 10.1080/07391102.2019.1682052.
293.	Amir, M., Khan, P., Queen, A., Dohare, R., Alajmi, M. F., Hussain, A., Islam, A., <b>Ahmad, F.</b> , and Hassan, M. I. (2020) Structural Features of Nucleoprotein CST/Shelterin Complex Involved in the Telomere Maintenance and Its Association with Disease Mutations. <i>Cells</i> <b>9</b> , 359.
294.	Khan S, Kaushik R, Sharma M, Raina N, <b>Ahmad F</b> , Islam A. (2020) Molecular basis of pathogenic parasitic infections: insights from parasite kinome. <i>Front Biosci</i> <b>25</b> ,1488-1509.
295.	Amir, M., Alam A, Ishrat R, Alajmi, M. F., Hussain, A., Rehman, M. T., Islam, A., <b>Ahmad, F.</b> , Hassan, M.I. and Dohare, R. (2020) A Systems View of the Genome Guardians: Mapping the Signaling Circuitry Underlying Oligonucleotide/Oligosaccharide-Binding (OB) Fold Proteins. <i>OMICS: A Journal of Integrative Biology</i> . 4 OMICS. 2020 Sep;24(9):518-530. doi: 10.1089/omi.2020.0072
296.	Gupta, P., Khan, S., Fakhr, Z., Hussain, A., Alajmi, M. F., Rehman, M. T., Islam, A., <b>Ahmad, F.</b> , and Hassan, M. I. (2020) Identification of Potential inhibitors of Calcium/calmodulin-dependent Protein Kinase IV from Bioactive Phytoconstituents. <i>Oxidative Medicine and Cellular Longevity</i> , doi.org/10.1155/2020/2094635
297.	Almary, K.A., Srivastava, S., Shahbaaz, M., Gupta, P., Khan, P., Syed, S. B., Azum, N., Asiri, A. M., Islam, A., <b>Ahmad, F.</b> , and Hassan, M. I. (2020) Unravelling the unfolding pathway of human Fas-activated serine/threonine kinase by urea-induced denaturation. <i>J Biomol Struct Dyn</i> , 1-10. doi: 10.1080/07391102.2020.1790423
298.	Shamsi, A, Mohammad, T, Anwar, S, Hassan, MI, <b>Ahmad, F</b> , Hasan, I, and Islam, A (2020) Biophysical insight into implications of PEG-400 on alpha-crystallin structure: Multi spectroscopic and microscopic approach, <i>ACS Omega</i> <b>5</b> , 19210–19216. Guptadoi: 10.1021/acsomega.0c02648
299.	Raina, N., Singh, A.K., Hassan, M.I., <b>Ahmad, F.</b> , and Islam, A. (2020) Concentration Dependent Effect of Ethylene Glycol on the Structure and Stability of Holo $\alpha$ -Lactalbumin: Characterization of Intermediate State amidst Soft Interactions, <i>Int J Biol Macromol</i> <b>164</b> , 2151-2168.
300.	Shamsi A, Anwar S, Shahbaaz M, Mohammad T, Al-Ajmi MF, Hussain A, Hassan MI, <b>Ahmad F</b> , and Islam A (2020) Evaluation of binding of Rosmarinic acid with human transferrin and its impact on protein structure: Targeting polyphenolic acid-induced protection of neurodegenerative disorders, <i>Oxidative Medicine and Cellular Longevity</i> , doi.org/10.1155/2020/1245875

301.	Parray ZA, <b>Ahmad F</b> , Hassan MI, and Islam A (2020) Conformational changes in cytochrome c directed by ethylene glycol accompanying complex formation: Protein-solvent preferential interaction or/and kosmotropic effect, <b>Spectrochimica Acta Part A: Molecular and Biomolecular Spectroscopy</b> <b>242</b> , doi.org/10.1016/j.saa.2020.118788
302.	Bashir S, Shamsi A, <b>Ahmad F</b> , Hassan MI, Kamal M, and Islam A (2020) Biophysical elucidation of fibrillation inhibition by sugar osmolytes in $\alpha$ -lactalbumin: Multispectroscopic and molecular docking approaches, <b>ACS Omega</b> <b>5</b> , 26871-26882.
303.	Shahid S, Hasan I, <b>Ahmad F</b> , Hassan MI, and Islam A (2020) Effect of Size of Crowder on Protein Stabilization, <i>in Recent Res Adva Biol</i> , Vol 3, Chapter 10. DOI: 10.9734/bpi/rwab/v3.
304.	Naiyer A, Khan B, Islam A, Hassan MI, Sundd M, <b>Ahmad F</b> (2020) Heme-iron ligand (M80-Fe) in Cytochrome c is Destabilizing: Combined in vitro and in silico approaches to monitor changes in structure, stability and dynamics of the protein on mutation, <b>J Biomol Struct Dyn</b> 2:1-18. doi: 10.1080/07391102.2020.1853607
305.	Nasreen, K., Parray, Z.A., Shamsi, A., <b>Ahmad, F.</b> , Ahmed, A., Malik, A., Lakhrm, N.A., Hassan, M.I. and Islam, I.J.B.M. (2021) Crowding Milleu stabilizes apo-myoglobin against chemical-induced denaturation: Dominance of hardcore repulsions in the heme devoid protein. <b>Int J Biol Macromol.</b> <b>81</b> , 552-560. doi: 10.1016/j.ijbiomac.2021.03.089.
306.	Naiyer A, Khan B, Hussain A, Islam A, Alajmi MF, Hassan MI, Sundd M, and <b>Ahmad F</b> (2021) Stability of uniformly labeled ( <sup>13</sup> C and <sup>15</sup> N) cytochrome c and its L94G mutant, <b>Sci Rep.</b> 2021 Mar 24;11(1):6804. doi: 10.1038/s41598-021-86332-w.
307.	Ahanger I, Parray Z, <b>Ahmad F</b> , Hassan MI, Islam, A, Sharma, and A (2021) Heparin Accelerates the Protein Aggregation via the Downhill Polymerization Mechanism: Multi-Spectroscopic Studies to Delineate the Implications on Proteinopathies, <b>ACS Omega.</b> 2021 Jan 12;6(3):2328-2339. doi: 10.1021/acsomega.0c05638.
308.	Shamsi A, Mohammad T, Anwar S, Nasreen K, Hassan MI, <b>Ahmad F</b> , and Islam A (2021) Insight into the binding of PEG-400 with eye protein alpha-crystallin: Multi spectroscopic and computational approach: Possible therapeutics targeting eye diseases, <b>J Biomol Struct Dyn.</b> 2020 Dec 11:1-11. doi: 10.1080/07391102.2020.1858964.
309.	Parray Z, <b>Ahmad F</b> , Hassan MI, Ahmed A, Almajhdi FN, Malik A, Hussain T and Islam A (2021) Structural refolding and thermal stability of myoglobin in the presence of mixture of crowders: Importance of stabilizing and destabilizing forces in the cellular condition, <b>Molecules</b> <b>10</b> ; <b>26</b> (9):2807. doi: 10.3390/molecules26092807.

310.	Kumar B, Mohammad T, Amaduddin, Hussain A, Islam A, <b>Ahmad F</b> , Alajmi MF, Singh S, Pandey KC, Hassan MI, Abid M. (2021) Targeting metacaspase-3 from Plasmodium falciparum towards antimalarial therapy: A combined approach of in-silico and in-vitro investigation, <i>J Biomol Struct Dyn</i> . 2021 Feb;39(2):421-430. doi: 10.1080/07391102.2019.1711194.
311.	Anwar S, Khan S, Shamsi A, Anjum F, Shafie A, Islam A, <b>Ahmad F</b> , and Hassan MI (2021) Structure-based investigation of MAP/microtubule affinity-regulating kinase 4 inhibitory potential of Naringenin for therapeutic management of cancer and neurodegenerative diseases. <i>J Cell Biochem</i> . 2021 Oct;122(10):1445-1459. doi: 10.1002/jcb.30022.
312.	Parray ZA, Naqvi AAT, Ahmad F, Hassan MI, and Islam A (2021) Characterization of different intermediate states in myoglobin induced by polyethylene glycol: a process of spontaneous molecular self-organization foresees the energy landscape theory via in vitro and in silico approaches, <i>J Mol Liquids</i> 342, 117502.
313.	Rahban M, Stanek A, Hooshmand A, Khamineh Y, Ahi S, Kazim SN, <b>Ahmad F</b> , Muronetz V, Abousenna MS, Zolghadri S, and Saboury AA (2021), Infection of human cells by SARS CoV-2 and molecular overview on gastrointestinal, neurological and hepatic problems in COVID-19 patients. <i>J. Clin. Med.</i> , 10(21), 4802; <a href="https://doi.org/10.3390/jcm10214802">https://doi.org/10.3390/jcm10214802</a>
314.	Waseem R, Shamsi A, Mohammad T, Alhumaydhi FA, Kazim SN, Hassan MI, <b>Ahmad F</b> , Islam A. (2021) Multispectroscopic and Molecular Docking Insight into Elucidating the Interaction of Irisin with Rivastigmine Tartrate: A Combinational Therapy Approach to Fight Alzheimer's Disease, <i>ACS Omega</i> 6, 7910-7921. doi: 10.1021/acsomega.1c00517.
315.	Amir M, Ahamad S, Mohammad T, Jairajpuri DS, Hasan GM, Dohare R, Islam A, Ahmad F, Hassan MI (2021). Investigation of conformational dynamics of Tyr89Cys mutation in protection of telomeres 1 gene associated with familial melanoma. <i>J Biomol Struct Dyn</i> . 39(1):35-44. doi: 10.1080/07391102.2019.1705186.
316.	Ahanger IA, Bashir S, Parray ZA, Alajmi MF, Hussain A, <b>Ahmad F</b> , Hassan MI, Islam A, Sharma A (2021). Rationalizing the Role of Monosodium Glutamate in the Protein Aggregation Through Biophysical Approaches: Potential Impact on Neurodegeneration. <i>Front Neurosci</i> . 5:636454. doi: 10.3389/fnins.2021.636454.
317.	Wahiduzzaman, Kumar V, Anjum F, Shafie A, Elsbali AM, Islam A, <b>Ahmad F</b> , Hassan MI (2021) Delineating the Aggregation-Prone Hotspot Regions (Peptides) in the Human Cu/Zn Superoxide Dismutase 1. <i>ACS Omega</i> . 6(49):33985-33994. doi: 10.1021/acsomega.1c05321.
318.	Bashir S, Ahanger IA, Shamsi A, Alajmi MF, Hussain A, Choudhry H, <b>Ahmad F</b> , Hassan MI, Islam A (2021). Trehalose Restrains the Fibril Load towards $\alpha$ -Lactalbumin Aggregation and Halts Fibrillation in a Concentration-Dependent Manner. <i>Biomolecules</i> 11(3):414. doi: 10.3390/biom11030414.

319.	Parray ZA, <b>Ahmad F</b> , Alajmi MF, Hussain A, Hassan MI, Islam A (2021) Interaction of polyethylene glycol with cytochrome c investigated via in vitro and in silico approaches. <i>Sci Rep</i> <b>11</b> (1):6475. doi: 10.1038/s41598-021-85792-4.
320.	Alamry KA, Srivastava S, Shahbaaz M, Khan P, Gupta P, Syed SB, Azum N, Asiri AM, Islam A, <b>Ahmad F</b> , Hassan MI (2020), Unravelling the unfolding pathway of human Fas-activated serine/threonine kinase induced by urea. <i>J Biomol Struct Dyn.</i> <b>39</b> (15):5516-5525. doi: 10.1080/07391102.2020.1790423.
321.	Raina N, Hassan MI, <b>Ahmad F</b> , Islam A, Singh AK (2021). PEG mediated destabilization of holo $\alpha$ -lactalbumin probed by <i>in silico</i> and <i>in vitro</i> studies: deviation from excluded volume effect. <i>J Biomol Struct Dyn.</i> 1-13. doi: 10.1080/07391102.2021.1987990.
322.	Ahanger I, Parray ZA, Nasreen K, <b>Ahmad F</b> , Hassan MI, Islam, A and Sharma, A (2021). Heparin Accelerates the Protein Aggregation via Downhill Polymerization Mechanism: Multi-spectroscopic studies to delineate the implications on proteinopathies. ACS Omega 2021 6 (3), 2328-2339 DOI: 10.1021/acsomega.0c05638.
323.	Waseem R, Shamsi A, Shahbaz M, Khan T, Kazim SN, <b>Ahmad F</b> , Hassan MI, and Islam, A (2021), Effect of pH on the Structure and Stability of Irisin, a Multifunctional Protein: Multispectroscopic and Molecular Dynamics Simulation Approach, <i>J Mol Struct.</i> <b>1252</b> , 132141.
324.	Waseem R, Shamsi A, Mohammad T, Kazim SN, Hassan MI, Chaudhary AA, Al-Zharani M, Rudayni A, <b>Ahmad F</b> , and Islam A (2022), FNDC5/Irisin: Physiology and Pathophysiology. <i>Molecules</i> <b>27</b> , 1118. doi: 10.3390/molecules27031118. PMID: 35164383.
325.	Parray ZA, <b>Ahmad F</b> , Chaudhary AA, Rudayni HA, Al-Zharani M, Hassan MI, Islam A (2022) Size dependent interplay of volume exclusion versus soft interactions: cytochrome c in macromolecular crowded environment, <i>Front Mol Biosci.</i> <b>9</b> , 49683. doi: 10.3389/fmolb.2022.849683.
326.	Rahban M, Zolghadri S, Salehi N, <b>Ahmad F</b> , Haertlé T, Rezaei-Ghaleh N, Sawyer N, and Saboury AA (2022) Thermal stability enhancement: fundamental concepts of protein engineering strategies to manipulate the flexible structure, <i>Int. J. Biol. Macromol.</i> <b>214</b> :642-654. doi: 10.1016/j.ijbiomac.2022.06.154. Epub 2022 Jun 27. PMID: 35772638.
327.	Idrees D, Naqvi AAT, Hassan MI, <b>Ahmad F</b> , and Gourinath S (2022) Insight into the Conformational Transitions of Serine acetyl transferase Isoforms in <i>E. histolytica</i> : Implications for Structural and Functional Balance, <i>ACS Omega</i> <b>7</b> , 24626-24637. DOI: 10.1021/acsomega.2c02467.
328.	<b>Ahmad F</b> (2022) Protein Stability [Determination] problems, <i>Front. Mol. Biosci.</i> <b>5</b> :9:880358. doi: 10.3389/fmolb.2022.880358. eCollection 2022.

329.	Parray, ZA, Naqvi AAT, Ahanger IA, Shahid M, <b>Ahmad F</b> , Hassan MI, and Islam A (2022), Measuring structural changes in cytochrome c under crowded conditions using in vitro and in silico approaches. <i>Polymers</i> <b>14</b> , 4808-. doi: 10.3390/polym14224808.
330.	Waseem R, Yadav NS, Khan T, <b>Ahmad F</b> , Kazim SN, Hassan MI, Prakash A, and Islam A (2023) Molecular Basis of Structural Stability of Irisin: A Combined Molecular Dynamics Simulation and In vitro Studies for Urea-induced Denaturation. <i>J. Mol. Liquids</i> <b>372</b> , 121120. doi:10.1016/j.molliq.2022.121120.
331.	Ahanger IA, Parray ZA, Raina N, Bashir S, <b>Ahmad F</b> , Hassan MI, Shahid M, Sharma A, and Islam, A (2023), Counteraction of the cetyltrimethylammonium bromide-induced protein aggregation by Heparin: Potential impact on protein aggregation diseases using biophysical approaches. <i>J. Mol. Struc</i> <b>1276</b> , 134714-134730.
332.	Raina N, Singh E Ahanger I, Israil I, Shahid M, Hassan M I, <b>Ahmad F</b> Singh A K and Islam A (2023) Polyethylene Glycol Induced Structural Modulation of Holo alpha-Lactalbumin: In vitro and In vivo Approach towards Protein Stability, <i>J. Mol. Liquids</i> <b>382</b> ,121909. doi: 10.1016/j.molliq.2023.121909.
333.	Rahban M, <b>Ahmad F</b> , Piatyszek MA, Haertlé T, Saso L, Saboury AA (2023) Stabilization challenges and aggregation in protein-based therapeutics in the pharmaceutical industry, <i>RSC Adv.</i> <b>13</b> , 35947-35963. doi: 10.1039/d3ra06476j.
334.	Tanwar S, Qais F A, Naaz F, Rashid N, <b>Ahmad F</b> , and Rehman S ur (2023), Identification of a novel Sorcin isoform with a different C-terminal but intact dimerization property. <i>Sci Rep</i> <b>13</b> ,15262. doi: 10.1038/s41598-023-40913-z.
335.	Juneja P, Rashid N, Qais FA, Tanwar S, Sultan I, <b>Ahmad F</b> , and Rehman S ur (2024) Alternative splicing generates a novel ferroportin isoform with a shorter C-terminal and intact iron and hepcidin binding property, <i>IBMB Life</i> (in press)
336.	
1.	Bashir S, Alam S, Chaudhary AA, Sami N, Khan A, Idris U, Khushnood S, Khan A, <b>Ahmad F</b> , Hassan MI, and Islam A (2021), Neurodegenerative diseases due to Amyloids: An aggregomics perception of proteins in health and pathology, <i>Reveu Neurologique</i> (under preparation)
2.	

3.	
4.	

1. Wahid, M., Bhaskar Singh, S., Srinivasan, A., **Ahmad, F.**, Singh, T. P. and Baig, M. A. (2005). The Sequence of Mitochondrial Cytochrome C From Buffalo, **GenBank Code: DQ176430**.

2. Rehman, M. H., Bhaskar Singh, Srinivasan, A., **Ahmad, F.**, Singh, T. P. and Baig, M. A. (2005). The Sequence of Mitochondrial Cytochrome C From Goat, **GenBank Code: DQ176429**.

3. BMRB ID: 27254

Naiyer A., Islam A., Hassan M. I., Sundd M., and **Ahmad F.** (2019) Solution structure of molten globule state of L94G mutant of horse cytochrome-*c*. [deposition date: 2017/11/15]; [Original release date: 2019/11/08]

4. PDB ID 5ZKV

Naiyer A., Islam A., Hassan M. I., Sundd M., and **Ahmad F.** (2019) Solution structure of molten globule state of L94G mutant of horse cytochrome-*c*.

#### BOOK CHAPTER

1. **Ahmad, F.** (1993), Measuring the Conformational Stability of Enzymes *in the Thermostability of Enzymes* (M.N. Gupta, ed.), pp 95-112, Narosa Publishing House, India.
2. Bashir S, Sami N, Bashir S, **Ahmad F**, Hassan MI, and Islam A (2020) Management of Insulin Through Co-Solute Engineering: A Therapeutic Approach, Editors (BD Singh & T Tripathi), pp 283-315, Springer Nature Singapore Pvt. Ltd.
3. Shahid S, Hasan I, **Ahmad F**, Hassan MI, and Islam A (2020) Effect of Size of Crowder on Protein Stabilization, *in Recent Res Adva Biol*, Vol 3, Chapter 10. DOI: 10.9734/bpi/rrab/v3.
- 4.