prof. Dr J.P. Abrahams

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- Born 28 May 1961, Oslo
- | Nationality: Dutch
- | married to Alison Sutton
- | 2 daughters (Élise & Gwendolyne)

I am a structural biologist, currently at Basel University (prof.) and the Paul Scherrer Institute (head of LBR). I love seeing important molecular structures of life that nobody has seen before, in ways that nobody has tried before. For this purpose, I develop novel methods in high-resolution bioimaging, which include computation, chemistry and physics. I currently focus my efforts on the analysis of protein 3D nano-crystals.

After my Ph. D. in Leiden (cum laude, 1990), I moved to the LMB in Cambridge, where, working in the groups of Andrew Leslie and John Walker, I solved the structure of the F1-ATPase. I could only solve this structure with the novel approaches in X-ray data collection and crystallographic phasing that I developed specifically for this purpose and that have now become part of mainstream crystallography. Together with Robin Carrell, I solved structures of serpins and with Richard Henderson I worked on a giant image plate scanner. In 1997 I returned to Leiden to become a full professor. Here, I studied serpins, viruses, ribosomal complexes, DNA repair proteins, microtubule complexes, enzymes, and amyloid-formation using X-ray crystallography, EM, NMR, AFM and spectroscopic techniques. Between 1997 and 2012, as a PI, I raised in excess of 36 MEuro in competitive funding for this and other work; the majority of these funds were for joint research projects and joint infrastructure. In this period, four companies were spun out of my research group.

Being inspired by the visionary, multidisciplinary approach at the LMB in Cambridge, I decided upon my move to Leiden that I should devote part of my energy to integrating approaches in bioimaging and establishing a joint, top-end infrastructure for structural biology. Hence, I initiated and directed the Cyttron I & II consortia¹, raised the funds for and established NeCEN², and initiated and directed the Leiden Cell Observatory³. In 2015, I moved to Switzerland as professor at the Basel Biozentrum and head of the Laboratory of Biomolecular Research of the Paul Scherrer Institute, in order to advance biomolecular cell research with the anticipated billion Euro SwissFEL X-ray source and a new electron diffraction instrument that will be built under my direction and guidance.

Current and future research

One of the main challenges in structural biology is that many proteins fail to produce (sufficiently large) 3D crystals required for structure solution. In order to meet this challenge, I am now analysing 3D nanocrystals of proteins using electron diffraction. My recent successes include visualising 2D, phased projection structures of single 100 nm protein crystals with a resolution beyond 2 Å using cryo-EM, the collection of large wedges of continuous 3D rotation diffraction data (up to 35 Deg) of similar nano-crystals with a resolution of up to 3Å and phasing of small organic molecules by direct methods. In order to achieve these results, I implemented quantum area detectors for electron diffraction and developed novel concepts, algorithms and software for analysing diffraction and imaging data. In the near future I anticipate adapting and improving these methods, in close

¹ Public-private partnership of 6 universities and 8 commercial enterprises, budget of 38 MEuro (2003-2015): http://cyttron.org

² Netherlands Centre for Electron Nanoscopy: joint EM centre of the Netherlands at Leiden University: http://www.necen.nl

³ The Leiden University joint facility for bioimaging: http://cellobservatory.leidenuniv.nl

collaboration with scientists at the PSI, to allow determining and refining protein structures *ab initio*. This approach is vital for making good use of the novel ultra-high intensity X-ray sources (XFEL, SwissFEL, new synchrotrons with diffraction limited storage rings). It will allow thorough, low-cost optimisation and characterisation of the marginal, but biologically important samples that are destined for such competitive, top-end X-ray sources. As a spin-off of my effort, a novel approach in atomic refinement of 3D cryo-EM single particle reconstructions of large molecular complexes is emerging from my research group.

However, many of the important molecular structures of life probably will not crystallise at all. In order to characterise these structures, I will expand techniques that I am currently developing to include asymmetric structures. I will focus my efforts on diffraction studies, as in both EM and X-ray imaging, the shot-noise of quanta that did not interact with the sample is much higher than the useful elastic signal. Only by collecting diffraction data, these non-interacting quanta (be they photons or electrons) can be discarded: they end up in the direct beam. In order to retrieve the phases of the diffraction patterns, I envisage using ptychography (lensless imaging by overlapped diffraction) and structured beam diffraction (phase retrieval using prior knowledge of the shape of the diffracted beam). I am convinced that these approaches will provide more accurate structures to a higher resolution than is currently feasible.

When establishing a new method, one is allowed to be a bit opportunistic concerning the biological samples are investigated. But new methods are only worth developing if they provide the tools that biologists need. Through collaborations that I will establish with leading biological labs, including the BioZentrum in Basel, I will ensure the methods I develop, are designed for the analysis of important biological molecular systems that cannot be tackled in any other practical way. Separately, I will continue a project I started a few years ago on disease associated nano-crystallisation of misfolded proteins into amyloid, which is associated with important ailments including Alzheimer's and Parkinson's Disease.

International position

In the 2010 national evaluation of all Chemistry groups of the Netherlands, the Abrahams group was awarded a score placing it amongst the top chemistry groups of the Netherlands (5/5 for quality, 5/5 viability, 4/5 productivity, 4/5 relevance). Concerning the scientific quality of the group, the review committee wrote⁴: "*The Biophysical Structural Chemistry group is pioneering in the development of new methodologies for structural biology and is a top "methods" group worldwide both through the combination of different experimental techniques for structure determination of biological macromolecules and the development of novel software for both EM and X-ray analysis. The research is reflected in publications on extremely challenging experimental systems (the ribosome, F1-ATPase etc). The leadership in the field is also evident in the establishment and/or contributions to the cell observatory, the Cytronl/II and the upcoming national centre for electron microscopy (NeCEN)."*

⁴ <u>http://www.qanu.nl/sites/default/files/bestanden/files/Chemistry_OZ_2011.pdf</u> (p. 127)

Qualifications:	
Full professorship:	1997 (Leiden University)
Ph. D. Diploma:	1991 cum laude (Leiden University)
Doktoraal Diploma:	1985 cum laude (Leiden University)
Kandidaats Diploma:	1984 (Leiden University, the Netherlands)
European Baccalaureate:	1979 (European School, Bergen, the Netherlands)

Employment :

2015-now:	Head of the Laboratory of Biomolecular Research at the Paul Scherrer Institute
	(Switzerland)
	Full professor at the Biozentrum, Basel University (Switzerland)
	Full professor at the Institute of Biology Leiden, Leiden University (The Netherlands)
1997-2015:	Full professor at the Leiden Institute of Chemistry, Leiden University (The
	Netherlands)
1990-1997:	Post-doc, then staff scientist at MRC Laboratory of Molecular Biology, Cambridge UK
1985-1990:	Netherlands Scientific Organisation (Leiden University)

Grants & Prizes (37.5 MEuro in total):

2017:	SNI NanoArgovia "Applicability of 3D Electron Diffraction in the Pharmaceutical Industry" (co-applicant, 100 kCHF)
2016:	SNF R'Equip grant "Detector Direct electron detector for cryo-EM single particle
	analysis, electron tomography and protein nanocrystallography" (co-applicant, 325 kCHF)
	NanoArgovia "Hybrid Pixel Detectors for Electron Diffraction of Nano-Samples" (co- applicant, 150 kCHF)
2015:	SNF grant "SwissFEDI – a free electron diffraction instrument for nano-diffraction of biological specimens" (741 kCHF)
	SNI grant "A programmable e- beam shaper for diffractive imaging of biological structures at Å resolution" (together with S Tsuijno). Project P1505 (270 CHF)
2014:	TA grant for developing novel electron detectors for structural biology (775 kEuro)
2014:	STW Valorization Grant stage I for See-e ⁻ s, valorizing a novel detector for electron diffraction (25 kEuro)
2013:	LIC: PhD grant (200 kEuro): Max Clabbers
2012:	CSC Grant for Yaowang Li for PhD research in my group (160 kEuro)
2011:	Joint Research PhD grant LUMC/LWN (150 kEuro): Eric van Genderen
2010:	EFRO (6.2 MEuro for a Titan Krios Microscope)
2010:	NWO Groot (6.2 MEuro for a second Titan Krios Microscope)
2010:	FES2009: (10 MEuro for CyttronII, including 1.2 MEuro for own research group)
2009:	NWO-CW Top grant (720 kEuro)
2006:	EU FP6 Max-Inf (30 kEuro for travel, organising workshop)
2005:	STW (600 kEuro)
2004:	BSIK (8.8 MEuro for Cyttron project, including about 1.1 MEuro for own research group).
2003:	NWO (Molecuul tot Cel) (700 kEuro);
2001:	Softlink: FOM & NWO-CW (350 kEuro);

2000:	Avantium Technologies BV(100 kEuro), Leadd BV (50 kEuro), Crystallics BV (200
	kEuro), KeyDP BV (70 kEuro), Isotis (50 kEuro)
1999:	Swiss National Foundation (50 kEuro)
1999:	STW (400 kEuro), Senter (170 kEuro);
1998:	NWO-CW (jonge chemici) (150 kEuro), NWO-MW (NWO-middelgroot) (400 kEuro)
1997:	European Union FP5 (120 kEuro)
1991-1992:	EMBO-fellowship (at LMB, Cambridge)
1990-1991:	Royal Society & Royal Dutch Academy of Sciences (at MRC Laboratory of Molecular
	Biology, Cambridge, UK)
1990:	Shell Travel Award (5000 Euro)
1989:	FEBS (Århus, Denmark; short term)
1987, 1988:	EMBO (Bayreuth, Germany; short term)

Supervised PhD students (PhD obtained):

Yaowang Li (2017, promotor)

Eric van Genderen (2015, promotor)

Jinghui Luo (2014, promotor)

Igor Nederlof (2013, promotor)

Willem-Jan Waterreus (2013, promotor)

Elisabeth Meulenbroek (2012, promotor)

Jan de Sonneville (2011, promotor)

Hanna Lindfors (2010, promotor)

Sandra Zovko (2010, promotor)

Daniel de Geus (2009, promotor)

Linhua Jiang (2009, promotor)

Pavol Skubak (2008, promotor)

Dilyana Georgieva (2008, promotor)

Ying-hui Zhang (2006, promotor)

Ineke Bos (2005, co-promotor)

Ellen Thomassen (2005, promotor)

Marike van Roon (2004, promotor)

Rutger Leliveld (2003, promotor)

Lei Jin (MRC LMB, 1997 supervisor)

Peter Elliott (MRC LMB, 1996 supervisor)

Richard Skinner (MRC LMB, 1995 supervisor)

Currently supervised PhD students (projected PhD date):

Max Clabbers (2018), Inayatullah Mohammed (2018), Jonas Heidler (2020), Poojah Thakkar (2020), Liu Bingrun (2020), Julian Wennmacher (2020)

Stimulation of talent

Dr Linhua Jiang (PhD student in my group 2005-2009) and Dr Zunfeng Liu (postdoc in my group 2007-2012) both became full professor in 2013.

Science Management Main:

2015-now 2003-now	Head of the Laboratory of Biomolecular Research at the Paul Scherrer Institute Initiator & director of CyttronI & CyttronII, two consecutive consortia of Universities and commercial enterprises (18 MEuro & 20M Euro budgets resp.)
Other:	
2013	Board member NeCEN
2009-12	Director & initiator of NeCEN
2008-12	Director & initiator of Leiden Cell Observatory
2011	Chair of the VICI committee STW
2010	Invited member of EU FP7 advisory committee on "4D Biology for health and disease"
2006-10	Member of WECO (science advisory board of the Leiden University Science Faculty)
2008	Invited member of EU KP7 advisory committee on "Detection, Diagnosis and Monitoring"
2005:	Invited member of MAX-INF2 (continuation of MAX-INF), invited member of SRS DUBBLE beam line allocation committee, invited member and rapporteur of EU KP7 advisory committee on Molecular testing and bioimaging
2002:	Invited member of SLS protein crystallography beam line allocation committee
2001:	co-founder of KeyDP BV – a company for high throughput crystallography and bio- informatics; Elected member of Biological Chemistry policy advisory committee (BAC) of the Dutch NWO chemistry council; Invited member of the National Dutch genomics committee, advising and prioritising national programme in fundamental genomics research; Invited member of MAX-INF: EU funded infrastructure Network (protein crystallography)
2000:	Elected member of the board of the Dutch NWO Protein Community; co-founder of Crystallics BV – a company for high throughput polymorph screening
1999:	Elected member of the board of the Dutch Society for Biochemistry & Molecular Biology;

Invited lectures & organised conferences, 2002 onwards

12-02-16 Ringberg meeting on Structural Biology with FELs, invited speaker

- 13-01-16 Biozentrum Symposium, keynote speaker
- 16-11-15 Cyttron II symposium, Naturalis, Leiden, conference organiser & chair
- 24-09-15 Engelberg (Switzerland), RegioMeeting 2015 (invited speaker)
- 16-09-15 Vercelli (Italy), XLIV Annual Meeting of the AIC (keynote lecturer)
- 26-08-15 Rovinc (Croatia), EMC15, Hybrid Sessions (chair)
- 25-08-15 Rovinc (Croatia), EMC15, electron diffraction (invited speaker)
- 14-04-15 Bad Ramsach (Switzerland), Basel FP Structural Biology Meetings, invited
- 24-12-14 I broke my foot, was home-bound on doctor's order for 3 months and had to cancel invited lectures at the Ringberg meeting on "Recent progress on soft- and hard-X-ray FEL applications in structural biology aswell as developments in electron microscopy/crystallography and synchrotron based microcrystallography", and at the Gordon Conference on "Polyamines"
- 10-11-14 PSI photon science workshop, invited speaker
- 03-11-14 Lorentz Workshop Life Science & Industry, organiser
- 16-10-14 Basel University, speaker at Symposium 'Nano-Diffraction of Biological Specimens'
- 08-10-14 110th International Titisee Conference on "Structure, forces and dynamics of macromolecular complexes", Titisee, Black Forest, Germany, invited speaker
- 11-09-14 Basel University, invited seminar
- 18-09-14 ICCBM 15, Hamburg 'Macromolecular crystallization for novel radiation sources', session 'Complementary Methods', invited speaker
- 24-06-14 Gordon Conference '3DEM: a rising star in structural biology', Girona, Spain, invited speaker
- 28-03-14 Cyttron II symposium, Naturalis, Leiden, conference organiser & chair
- 16-01-14 Paul Scherrer Institute, Villigen, Switzerland, invited speaker
- 03-10-13 Co-organiser, host and speaker at the Lorentz workshop 'Life Science and Industry'
- 28-06-13 Changzhou University invited speaker
- 22-03-13 Leiden: CyttronII symposium invited speaker, chair
- 16-06-12 Stockholm: Electron crystallography school: new methods invited speaker
- 08-06-12 Hamburg: CFEL seminar, keynote speaker
- 16-03-12 Leiden: CyttronII symposium, keynote speaker
- 05-01-12 Warwick: CCP4 study weekend, invited speaker

- 14-09-10 Dublin: ICCBM13, invited speaker Berlin: Facets of Electron Crystallography, invited speaker 07-07-10 16-03-10 Brussels: "4D Biology for health and disease", invited speaker 05-02-10 LUMC: LUMC meets TUDelft, invited speaker Geneva: Cleveringa lecture 07-05-09 Italian Crystallographic Association (AIC) Congress : Methods & Crystallography (in 22-09-09 Salerno, Italia) (keynote lecture) 25th European Crystallography Meeting (in Istanbul, Turkey) (keynote lecture) 19-08-09 Seminar Swiss Light Source, Villigen, Switserland, invited speaker 15-05-09 07-05-09 SGC seminar (Oxford), invited speaker 21-02-08 Tiling 3D Rotation Space, Lunteren, invited speaker LIFE symposium, keynote lecture 06-03-08 Max-Inf2/Lorentz Center workshop New algorith in macromolecular crystallography 17-05-08 and electron microsocpy, invited speaker 02-11-07 Proteins Killing Tumour Cells (Lorentz workshop), invited speaker 13-06-07 Advances in EM, Valencia, keynote lecture 18-01-05 This Weeks Discoveries - Univ. Leiden, invited speaker 31-01-05
- XFEL meeting, Woudschoten, invited speaker
- 17-02-05 Fysisch Genootschap, keynote lecture
- 23-03-05 Hybrid Methods, Granlibakken, California - invited speaker

Leiden: Reedijk symposium, invited speaker

Erice: Electron Crystallography, invited speaker

18-04-05 Diligentia, keynote lecture

28-10-11

01-06-11

- 25-08-05 ECM – Antwerp, invited speaker
- Structural Biology of Molecular Motors, Inst. Lorentz, invited speaker 27-10-05
- 14-06-04 ETH Structural Biology Symposium, Zürich, invited speaker
- 29-09-04 ELBA Max Planck Forum (Porto Conte, Sardinia) - invited speaker
- Studium Generale, Univ. Leiden, keynote lecture 08-12-04
- 16-12-03 Mathematics in Biology, Wagningen, invited speaker
- 03-05-02 Genomics Momentum, keynote lecture
- 18-02-02 Orleans, SOLEIL, invited speaker
- 18-04-02 IOP symposium, Wageningen, invited speaker
- 04-01-02 CCP4 study weekend, York, invited speaker

Refereed publications

(142 papers/book(chapter)s, 88 as first or last author, total number of citations until August 2016:

- ~8000, H-factor: 34 (ResearchGate))
- Yin Q, Liu Z, Laroche FJF, Zhou X, Shao N, Lin B, Wang R, Yuan N, Ding J, Abrahams JP [1] (2017) "A Novel Capturing Method for Quantification of Extra-Cellular Nanovesicles" J. Nanosci. Nanotech. 17, 908-913
- Su J, Wang H, Wu K, Liu ZS, Yin Q, Wang R, Lv Wei, Yin S, Liu Z, Abrahams JP (2017) [2] "Neutravidin-Mediated Extraction of Isolated Small Diameter Single Walled Carbon Nanotubes for Bio-Recognition" J. Nanosci. Nanotech. 17, 3588-3596
- Clabbers MTB, van Genderen E, Wan W, Wiegers EL, Gruene T, Abrahams JP (2017) [3] "Protein structure determination by electron diffraction using a single three-dimensional nanocrystal" Acta Crystallogr. D (in the press)
- Wang R, Boleij M, Yin Q, Galjart N, Lin B, Yuan N, Zhou X, Tan M, Ding J, Liu Z, Abrahams [4] JP (2017) "Purification of Biotinylated Proteins Using Single Walled Carbon Nanotube-Streptavidin Complexes", J. Nanoscience and Nanotechnology 17, 926-931
- [5] Abrahams JP (2016) "Electron nanodiffraction for structural biology" Acta Crystallogr. A72(a1), s6
- [6] Tiiman A, Luo J, Wallin C, Olsson L, Lindgren J, Jarvet J, Roos P, Sholts SB, Rahimipour S, Abrahams JP, Erikson Karlström A, Gräslund A, Warmländer SKTS (2016) "Specific Binding of Cu(II) lons to Amyloid-Beta Peptides Bound to Aggregation-Inhibiting Molecules or SDS Micelles Creates Complexes that Generate Radical Oxygen Species"J. Alzheimer's Disease 54, 971-982
- Luo J, Wärmländer SKTS, Gräslund A, Abrahams JP (2016) "Cross-interactions between the [7] Alzheimer disease amyloid-ß peptide and other amyloid proteins: A further aspect of the amyloid cascade hypothesis" J. Biol. Chem. 291, 1485-1493
- [8] Wallin, C., Kulkarni Y.S., Abelein, A., Jarvet, J., Liao, Q., Strodel, B., Olsson, L., Luo, J.,

Abrahams, J. P., Sholts, S. B., Roos, P. M., Kamerlin, S.C., Gräslund, A., and Wärmländer, S. K. T. S. (2016) Characterization of Mn(II) ion binding to the amyloid- β peptide in Alzheimer's disease. J. Trace Elem. Med. Biol. 10.1016/j.jtemb.2016.03.009

- [9] Luo J, Wärmländer SKTS, Gräslund A, Abrahams JP (2016) "Reciprocal molecular interactions between the Aβ peptide linked to Alzheimer's Disease and insulin linked to Diabetes Mellitus type II" ACS Chem Neurosci. DOI: 10.1021/acschemneuro.5b00325
- [10] Van Genderen E, Li, YW, Nederlof I, Abrahams JP (2016) "Lattice filter for processing image data of 3D protein nano-crystals" Acta Crystallogr. D72, 34-39
- [11] van Genderen E, Clabbers MTB, Das PP, Stewart A, Nederlof I, Barentsen KC, Portillo Q, Pannu NS, Nicolopoulos S, Gruene T, Abrahams JP (2016) "Ab initio structure determination of nanocrystals of organic pharmaceutical compounds by electron diffraction at room temperature using a Timepix quantum area direct electron detector" Acta Crystallogr. A72, 236-242, doi: 10.1107/S2053273315022500
- [12] Abrahams JP, Van Genderen E, Nederlof I, Li, YW, (2015) "Electron diffraction and imaging of 3D nanocrystals of pharmaceuticals, peptides and proteins" Acta Crystallogr. A71, S103
- [13] Clabbers MTB, Van Genderen E, Nederlof I, Abrahams JP (2015) "Electron crystallography of 3D nano-crystals" Acta Crystallogr. A71, S405
- [14] Afanasyev P, Ravelli RBG, Matadeen R, De Carlo S, van Duinen G, Alewijnse B, Peters PJ, Abrahams JP, Portugal RV, Schatz M, van Heel M (2015) "A posteriori correction of camera characteristics from large image data sets" Scientific Reports **5**, doi:10.1038/srep10317
- [15] Luo J, Wärmländer SKTS, Gräslund A, Abrahams JP "Alzheimer Peptides Aggregate into Transient Nanoglobules That Nucleate Fibrils" (2014) Biochemistry 53, 6302-6308
- [16] Luo J, Wärmländer SKTS, Gräslund A, Abrahams JP "Non-chaperone Proteins Can Inhibit Aggregation and Cytotoxicity of Alzheimer Amyloid-β Peptide" (2014) Biomacromolecules, DOI: 10.1021/bm401874j
- [17] Luo J, Mohammed I, Wärmländer SKTS, Hiruma Y, Gräslund A, Abrahams JP "Endogenous Polyamines Reduce the Toxicity of Soluble Aβ Peptide Aggregates Associated with Alzheimer's Disease" (2014) Biomacromolecules, DOI: 10.1021/bm401874j
- [18] Luo J, Wärmländer SKTS, Yu CH, Muhammad K, Gräslund A; Abrahams JP (2014) "The Aβ peptide forms non-amyloid fibrils in the presence of carbon nanotubes" Nanoscale 6, 6720-6726
- [19] Luo, J, Abrahams JP (2014) "Cyclic peptides as inhibitors of amyloid fibrillation" Chemistry Eur. J 20, 2410-2419
- [20] Abelein A, Abrahams JP, Danielsson J, Gräslund A, Jarvet J, Luo J, Tiiman A, Wärmländer SKTS (2014) "The hairpin conformation of the Amyloid β peptide is a common structural motif along the aggregation pathway" J. Biol. Inorg. Chem. 19, 623-624
- [21] Ten Bruggencate F, Laroche, F, Zhang, Y, Song G, Yin S, Abrahams JP, Liu Z (2013) "Visualizing the localization of transfection complexes during graphene nanoparticle-based transfection" J. Mater. Chem. B 1, 6353-6358
- [22] Luo J, Otero JM, Yu C-H, Wärmländer SKTS, Gräslund A, Overhand M, Abrahams JP (2013) "Inhibiting and reversing Aβ peptide (1-40) fibrillation by gramicidin S and engineered analogues" Chemistry 19, 17338-17348
- [23] Luo J, Wärmländer SKTS, Gräslund A, Abrahams JP (2013) "Human lysozyme inhibits the in vitro aggregation of Aβ peptide, which in vivo is associated with Alzheimer's disease" Chem. Comm 49, 6507-6509
- [24] Liu ZF, Voskamp P, Zhang Y, Chu FQ, Abrahams JP (2013) "Capture of unstable protein complex on the streptavidin-coated single-walled carbon nanotubes" Journal of Nanoparticle Research 15
- [25] Nederlof I, van Genderen E, Li YW, Abrahams JP (2013) "A Medipix quantum area detector allows rotation electron diffraction data collection from submicrometre three-dimensional protein crystals" Acta Crystallogr D69 1223-1230
- [26] Nederlof I, Li YW, van Heel M, Abrahams JP (2013) "Imaging protein three-dimensional nanocrystals with cryo-EM" Acta Crystallogr. D69, 852-859
- [27] Luo JH, Yu CH, Yu Hx, Borstnar R, Kamerlin SCL, Graslund A, Abrahams JP, Warmlander SKTS (2013) "Cellular Polyamines Promote Amyloid-Beta Peptide Fibrillation and Modulate the Aggregation Pathways" Biophys J 104 398A-398A
- [28] Luo JH, Zwier R, Abrahams JP (2013) "An efficient nanolitre-volume multi-channel device for highly viscous materials used in membrane protein crystallization" Appl. Crystallogr. 46, 829-831
- [29] Luo JH, Yu CH, Yu Hx, Borstnar R, Kamerlin SCL, Gräslund A, Abrahams JP, Wärmländer SKTS (2013) "Cellular Polyamines Promote Amyloid-Beta Peptide Fibrillation and Modulate the Aggregation Pathways" Neuroscience 4, 454-462
- [30] Liu ZF, Galli F, Waterreus WJ, Meulenbroek E, Koning RI, Lamers GEM, Olsthoorn RCL,

Pannu NS, Oosterkamp TH, Koster AJ, Dame RT, Abrahams JP (2012) "Single-Walled Carbon Nanotubes as Scaffolds to Concentrate DNA for the Study of DNA-Protein Interactions" Chem. Phys. Chem 13, 1569-1575

- [31] Meulenbroek EM, Thomassen EAJ, Pouvreau L, Abrahams JP, Gruppen H, Pannu NS (2012) "Structure of a post-translationally processed heterodimeric double-headed Kunitz-type serine protease inhibitor from potato" Acta Crystallogr. D68, 794-799
- [32] Moldovan G, Sikhrulidze I, Mattheson J, Derbyshire G, Kirkland AI, Abrahams JP (2012) "Characterisation of a counting imaging detector for electron detection in the energy range 10-20 keV" Nucl. Inst. & Meth. In Physics Res 681, 21-24
- [33] Zhou X, Laroche F, Lamers GEM, Torraca V, Voskamp P, Lu T, Chu FQ, Spaink HP, Abrahams JP, Liu ZF (2012) "Ultra-small graphene oxide functionalized with polyethylenimine (PEI) for very efficient gene delivery in cell and zebrafish embryos" Nano Res. 5, 703-709
- [34] Jiang L, Georgieva D, Abrahams JP (2011) "EDIFF: a program for automated unit cel determination of electron diffraction data" J. Appl. Cryst. 44, 1132-1136
- [35] Jiang L, Georgieva D, Nederlof I, Liu ZF, Abrahams JP (2011) "Image Processing and Lattice Determination for Three-Dimensional Nanocrystals" Microscopy and Microanalysis 17, 879-885.
- [36] Sikharulidze I, van Gastel R, Schramm S, Abrahams JP, Poelsema B, Tromp RM, van der Molen SJ (2011) "Low energy electron microscopy imaging using Medipix2 detector" Nucl. Inst. & Meth. In Physics Res A633, S239-S242
- [37] Pannu NS, Waterreus WJ, Skubak P, Sikharulidze I, Abrahams JP (2011) "Advances in the CRANK software suite for experimental phasing" Acta Crystallogr. D67, 331-337
- [38] Georgieva D, Jansen J, Sikharulidze I, Jiang L, Zandbergen HW, Abrahams JP (2011) "Evaluation of Medipix 2 detector for recording electron diffraction data in low dose conditions" J. Instrumentation 6 C01033
- [39] Nicolopoulos S, Rauch EF, Georgieva D, Abrahams JP (2011) "Low-resolution electron crystallography challenges in organic and inorganic crystals with transmission electron microscope (TEM)" Acta Crystallogr. D67, C188
- [40] Nederlof I, Georgieva D, Abrahams JP (2011) "Electron diffraction of submicron threedimensional protein crystals" Acta Crystallogr. D67, C228
- [41] Nederlof I, Hosseini R, Georgieva D, Luo J, Li D, Abrahams JP "A straightforward and robust method for introducing human hair as a nuceant into high throughput crystallization trials" (2011) Crystal Growth & Design 1170-1176
- [42] Abrahams JP (and 27 co-authors) (2011) "4D Biology for Health and Disease workshop report" New Biotechnology 28, 291-293
- [43] Liu ZF, Galli F, Janssen KGH, Jiang LH, van der Linden HJ, de Geus DC, Voskamp P, Kuil ME, Olsthoorn RCL, Oosterkamp TH, Hankemeier T, Abrahams JP (2010) "Stable Singlewalled carbon nanotube-streptavidin complex for biorecognition" J. Phys. Chem. C114, 4345-4352
- [44] Liu ZF, Jiang LH, Galli F, Nederlof I, Olsthoorn RCL, Lamers GEM, Oosterkamp TH, Abrahams JP (2010) "Graphene oxide.streptavidin complex for biorecognition towards affinity purification" Advanced Functional Materials 20, 2857-2865
- [45] Abrahams JP (2010) "The Strong Phase Object Approximation May Allow Extending Crystallographic Phases of Dynamical Electron Diffraction Patterns of 3D Protein Nano-Crystals", Zeitschr. F. Kristallografie 225, 67-76
- [46] Jiang L, Liu Z, Georgieva D, Kuil ME, Abrahams JP (2010) "A Novel Approximation Method of CTF Amplitude Correction for 3D Single Particle Reconstruction" Ultramicroscopy 4, 350-358
- [47] Van Gastel R, Sikharulidze I, Schramm S, Abrahams JP, Poelsema B, Tromp RM, van der Molen SJ (2009) "Medipix detector applied to low energy electron microscopy" Ultramicroscopy 110, 33-35,
- [48] van Haren J, Draegestein K, Keijzer N, Abrahams JP, Grosveld F, Peeters PJ, Moechars D, Galjart N (2009) "Mammalian Navigators are Microtubule Plus-End Tracking Proteins that can Reorganize the Cytoskeleton to Induce Neurite-Like Extensions" Cell Motility and the Cytoskeleton 66, 824-838
- [49] Waterreus WJ, Pannu N, Skubak P, Sikhuralidze I, Abrahams JP, de Graaff RAG (2009) "Recent advances in CRANK" Acta Crystallogr. A65, s160
- [50] Abrahams JP, Jiang LH, Sikharulidze I, Nederloff I, Zandbergen H, Georgieva D (2009) "Macromolecular electron crystallography" Acta Crystallogr. A65, s7
- [51] Jiang L, Georgieva D, Zandbergen HW, Abrahams JP (2009) "Unit-cell determination from randomly oriented electron-diffraction patterns" Acta Crystallogr. D65, 625-632
- [52] Jiang, L., Georgieva, D., IJspeert K., Abrahams, J.P. (2009). An intelligent peak search program for digital electron diffraction images of 3D nano-crystals. Image and Signal processing, CISP'09, Vol. 17-19

- [53] Pannu, N.S., Ravelli, R.B.G., Abrahams, JP (2009) "The Max-Inf2/Lorentz Center workshop on New algorithms in macromolecular crystallography and electron microscopy" Acta Crystallogr. D65, 623-624.
- [54] Jiang LH, Schaffitzel C, Bingel-Erlenmeyer R, Ban N, Korber P, Koning RI, de Geus DC, Plaisier JR, Abrahams JP (2009) "Recycling of Aborted Ribosomal 50S Subunit-Nascent Chain-tRNA Complexes by the Heat Shock Protein Hsp15" J. Mol. Biol. 386, 1357-1367
- [55] de Geus DC, Thomassen EAJ, Hagedoorn PL, Pannu NS, van Duijn E, Abrahams JP (2009) "Crystal Structure of Chlorite Dismutase, a Detoxifying Enzyme Producing Molecular Oxygen" J. Mol. Biol. 387 192-206
- [56] de Geus DC, van Roon AMM, Thomassen EAJ, Hokke CH, Deelder AM, Abrahams JP (2009) "Characterization of a diagnostic Fab fragment binding trimeric Lewis X" Proteins, Struct. Func. & Bioinf. 76, 439-447
- [57] Meulenbroek EM, Paspaleva K, Thomassen EAJ, Abrahams JP, Goosen N, Pannu NS (2009) "Involvement of a carboxylated lysine in UV damage endonuclease" Protein Science 18, 549-558
- [58] Zovko S, Abrahams JP, Koster AJ, Galjart N, Mommaas AM (2008) "Microtubule plus-end conformations and dynamics in the periphery of interphase mouse fibroblasts" Mol. Biol. Cell 19, 3138-3146
- [59] de Geus DC, Thomassen EAJ, van der Feltz CL, Abrahams JP (2008) "Cloning, expression, purification, crystallization and preliminary X-ray diffraction analysis of chlorite dismutase: a detoxifying enzyme producing molecular oxygen" Acta Crystallogr. F64, 730-732
- [60] Abrahams JP, Georgieva D, Jiang LH, Zandbergen H (2008) "Prospects for structure solution by electron diffraction of three-dimensional protein crystals" Acta Crystallogr. A64, C75-C76
- [61] Paspaleva K, Thomassen E, Pannu NS, Iwai S, Moolenaar GF, Goosen N, Abrahams JP (2007) "Crystal Structure of the DNA Repair Enzyme Ultraviolet Damage Endonuclease" Structure 15, 1-9
- [62] Georgieva DG, Kuil ME, Oosterkamp TH, Zandbergen HW, Abrahams JP (2007) "Heterogeneous nucleation of three-dimensional protein nanocrystals" Acta Cryst. D63 564-570
- [63] Pannu NS, Skubak P, Sikhuralidze I, Abrahams JP, de Graaff RAG (2007) "Recent advances in the CRANK automated structure solution suite" Acta Crystallogr. A63, s116
- [64] Plaisier JR, Jiang L, Abrahams JP (2007) "Cyclops: New modular software suite for cryo-EM" J. Struct. Biol. 157, 19-27
- [65] Georgieva DG, Abrahams JP, Zandbergen HW, Nikolopoulos S, Boulahya K (2007) "Solving Ab-initio Protein and Nanostructures with Precession Electron Diffraction" Microscopy and Microanalysis 13, 952-953
- [66] Schaffitzel C, Oswald M, Berger I, Ishikawa T, Abrahams JP, Koerten HK, Koning RI, Ban N (2006) "Structure of the E-coli signal recognition particle bound to a translating ribosome" Nature 448, 1076-1076
- [67] Kuil ME, Abrahams JP, Marijnissen JCM (2006) "Nano-dispensing by electrospray for biotechnology" Biotechnol. J. 1, 969-975
- [68] Thomassen EAJ, Dekking EHA, Thompson A, Franken KL, Sanal O, Abrahams JP, van Tol MJD, Koning F (2006) "The impact of single amino acid substitutions in CD3 gamma on the CD3 epsilon gamma interaction and T-cell receptor-CD3 complex formation" Human Immunology 67, 579-588
- [69] Schaffitzel C, Oswald M, Berger I, Ishikawa T, Abrahams JP, Koerten HK, Koning RI, Ban N (2006) "Structure of the E-coli signal recognition particle bound to a translating ribosome" Nature 444, 503-506
- [70] Schmauder R, van Rijn R, Abrahams JP, Kuil ME, Schmidt T (2005) "FCS in non-ideal solutions" Biophys. J 88, 655A
- [71] Thomassen EAJ, van Veen HA, van Berkel PHC, Nuijens JH, Abrahams JP (2005) 'The protein structure of recombinant human lactoferrin produced in the milk of transgenic cows closely matches the structure of human milk-derived lactoferrin' TRANSGENIC RESEARCH 14, 397-405
- [72] Rohn JL, Zhang Y-H, Leliveld SR, Danen-van Oorschot AAAM, Henriquez NV, Abrahams JP and Noteborn MHM (2005) Relevance of apoptin's integrity for its functional behaviour. Journal of Virology 79, 1337-1338.
- [73] Zhang Y-H, Rohn JL, Abrahams JP and Noteborn MHM (2004) Underlying molecular mechanisms of tumorigenesis (2)-Unbalance between proliferation and cell death. Journal of Medicine Molecular Biology 1, 195-206.
- [74] Ness SR, de Graaff RAG, Abrahams JP, Pannu NS (2004) "CRANK: New methods for automated macromolecular crystal structure solution" Structure 12, 1753-1761
- [75] Van Roon AM, Bink HH, Plaisier JR, Pleij CW, Abrahams JP and Pannu NS (2004) "Crystal

structure of an empty capsid of turnip yellow mosaic virus" J. of Mol. Biol. 341, 1205-1214.

- [76] Thomassen EA, Pouvreau L, Gruppen H and Abrahams JP (2004) Crystallization and preliminary X-ray crystallographic studies on a Kunito type potato serine protease inhibitor. Acta Crystallogr. D60, 1464-1466.
- [77] Van Roon AM, Pannu NS, De Vrind JP, Van der Marel GA, Van Boom JH, Hokke CH, Deelder AM and Abrahams JP (2004) Structure of an anti-Lewis X Fab fragment in complex with its Lewis X antigen. Structure 12, 1227-1236.
- [78] Plaisier JR, Koning RI, Koerten HK, van Heel M, Abrahams JP (2004) "TYSON: Robust searching, sorting, and selecting of single particles in electron micrographs" J. Struct. Biol. 145, 76-83
- [79] Bos IG, Lubbers YT, Eldering E, Abrahams JP and Hack CE (2004). Effect of reactive site loop elongation on the inhibitory activity of C1-inhibitor. Biochimica Biophysica Acta 1699, 139-144.
- [80] Leliveld SR, Dame RT, Rohn JL, Noteborn MHM and Abrahams JP (2004) "Apoptin's functional N- and C-termini independently bind DNA" FEBS Letters 557, 155-158.
- [81] Hilge M, Siegal G, Vuister GW, Güntert P, Gloor SM, Abrahams JP. (2003) ATP-induced conformational changes of the nucleotide-binding domain of Na,K-ATPase. Nature Struct. Biol. 10, 468-474.
- [82] Danen-van Oorschot AAAM, Zhang YH, Leliveld SR, Rohn JL, Seelen MCMJ, Bolk MW, van Zon A, Erkeland SJ, Abrahams JP, Mumberg D, Noteborn MHM. (2003) "Importance of nuclear localization of apoptin for tumor-specific induction of apoptosis" Journal of Biological Chemistry 278, 27729-27736.
- [83] Leliveld SR, Zhang Y-H, Rohn JL, Noteborn MHM, Abrahams JP. (2003) "Apoptin Induces Tumor-specific Apoptosis as a Globular Multimer" J. Biol. Chem. 278, 9042-9051.
- [84] Zhang YH, Leliveld SR, Kooistra K, Molenaar C, Rohn JL, Tanke HJ, Abrahams JP, Noteborn MHM. (2003) "Recombinant apoptin multimers kill tumor cells but are nontoxic and epitopeshielded in a normal-cell-specific fashion" Experimental Cell Research 289, 36-46.
- [85] Leliveld SR, Noteborn MHM, Abrahams JP. (2003) "Prevalent conformations and subunit exchange in the biologically active apoptin protein multimer" European Journal of Biochemistry 270, 3619-3627.
- [86] Leliveld SR, Dame RT, Mommaas MA, Koerten HK, Wyman C, Danen-van Oorschot AAAM, Rohn JL, Noteborn MHM, Abrahams JP. (2003) "Apoptin protein multimers form distinct higher-order nucleoprotein complexes with DNA" Nucleic Acids Research 31, 4805-4813.
- [87] Koning R, van den Worm S, Plaisier JR, van Duin J, Abrahams JP, Koerten H (2003) Visualization by Cryo-electron Microscopy of Genomic RNA that Binds to the Protein Capsid Inside Bacteriophage MS2. Journal of Molecular Biology 332, 415-422.
- [88] Thomassen E, Gielen G, Schütz M, Schoehn G, Abrahams JP, Miller S, van Raaij MJ. (2003) The Structure of the Receptor-binding Domain of the Bacteriophage T4 Short Tail Fibre Reveals a Knitted Trimeric Metal-binding Fold. Journal of Molecular Biology 331, 361-373.
- [89] Van Roon AM, Pannu NS, Hokke CH, Deelder AM, Abrahams JP. (2003) "Crystallization and preliminary X-ray analysis of an anti-LewisX Fab fragment with and without its LewisX antigen" Acta Crystallograpica D59, 1306-1309.
- [90] Abrahams JP, Thomassen EAJ (2003) "Mechanism of Thrombin's Enigmatic Sodium Switch Revealed" Structure 11, 363-364.
- [91] Bos IGA, Lubbers YTP, Roem D, Abrahams JP, Hack CE, Eldering E (2003) "The functional integrity of the serpin domain of C1-inhibitor depends on the unique N-terminal domain, as revealed by a pathological mutant" Jour. .Biol. Chem. 278, 29463-29470
- [92] McCoy AJ, Pei XY, Skinner R, Abrahams JP, Carrell RW (2003) Structure of betaantithrombin and the effect of glycosylation on antithrombin's heparin affinity and activity. Journal of Molecular Biology 326, 823-833.
- [93] Plaisier JR, Koning RI, Koerten HK, van Roon AM, Thomassen EAJ, Kuil ME, Hendrix J, Broennimann C, Pannu NS, Abrahams JP. (2003) "Area detectors in structural biology" Nuclear Instrumental Methods Physical Research A509, 274-282.
- [94] Bodenstaff ER, Hoedemaeker FJ, Kuil ME, de Vrind HPM and Abrahams JP. (2002) The prospects of protein nanocrystallography. Acta Crystallograpica D58, 1901-1906.
- [95] Hoedemaeker, P.J., Visschers, R.W., Alting, A.C., de Kruif, C.G., Kuil, M.E. & Abrahams, J.P. (2002) "A novel pH dependent dimerisation motif in β-lactoglobulin from pig (Sus scrofa)" Acta Crystallogr. 58D, 480-486
- [96] Kuil, M.E., Bodenstaff, R., Hoedemaeker, P.J. & Abrahams, J.P. (2002) "Protein nanocrystallogenesis" Enzyme Microb. Tech. 30 262-265
- [97] Schmauder R, Schmidt T, Abrahams JP, Kuil ME (2002) "Screening crystallisation conditions using fluorescence correlation spectroscopy" Acta Crystallogr. 58D, 1536-1541
- [98] Bos IGA, Hack CE, Abrahams JP (2002) "Structural and functional aspects of C1-inhibitor"

Immunobiology, vol. 205, 518-533.

- [99] Hilge M, Perrakis A, Abrahams JP, Winterhalter K, Piontek K, Gloor SM (2001) "Structure elucidation of beta-mannanase: from the electron-density map to the DNA sequence" Acta Crystallogr. 57D, 37-43
- [100] De Graaff, R.A.G., Hilge, M., van der Plas, J.L. & Abrahams, J.P. (2001) "Matrix methods for solving protein sub-structures of chlorine and sulphur from anomalous data" Acta Crystallogr. 57D, 1875-1862
- [101] Leslie AWG, Abrahams JP, Braig K, Lutter R, Menz RI, Orris GL, van Raaij MJ, Walker JE (1999) "The structure of bovine mitochondrial F₁-ATPase: an example of rotary catalysis" Biochem. Soc. Trans. 27, 37-42
- [102] Hoedemaeker, F.J., Siegal, G., Roe S.M., Driscoll, P.C. & Abrahams, J.P. (1999) "Crystal structure of the C-terminal SH2 domain of the p85α regulatory subunit of phosphoinositide 3-kinase: and SH2 domain mimicking its own substrate", J. Mol. Biol.. 292, 763-770
- [103] Elliott, P.R., Abrahams, J.P., Lomas, D.A. (1998) "Wild type α1-antitrypsin is in the canonical conformation" J. Mol. Biol. 275, 419-425
- [104] Skinner, R. Chang, W.-S.W., Jin, L. Pei, X. Huntingdon, J., Abrahams, J.P., Carrell, R.W. & Lomas, D.A. (1998) "Implications for function and therapy of a 2.9 Å structure of binary complexed antithrombin", J.Mol. Biol. 283, 9-14
- [105] Jin, L., Abrahams, J.P., Skinner, R., Petitou, M., Pike, R.N. & Carrell, R.W. (1998) "The anticoagulant activation of antithrombin by heparin" Proc. Natl. Acad. Sci. USA 94, 14683-14688
- [106] Abrahams, J.P. & De Graaff, R.A.G. (1998) "New developments in phase refinement" Curr. Op. Struct. Biol. 8, 601-605
- [107] Wardell, M.R., Skinner, R., Carter, D.C., Twigg, P.D., Abrahams, J.P. (1997) "Improved diffraction of antithrombin crystals grown in space" Acta Crystallogr. 53D, 622-625
- [108] Abrahams, J.P. (1997) "Bias reduction in phase refinement by modified interference functions: introducing the γ-correction" Acta Crystallogr. 53D, 371-376
- [109] Shirakihara, Y, Leslie, A.W.G., Abrahams, J.P., Walker, J.E., Ueda, T., Sekimoto, Y., Kambara, M., Saika, K., Kagawa, Y., Yoshida, M. (1997) "The crystal structure of the nucleotide free a3b3 subcomplex of F1 ATPase from the thermophilic Bacillus PS3 is a symmetric trimer" Structure 5, 825-836
- [110] Carrell, R., Skinner, R., Jin, L., Abrahams, J.P. (1997) "Structural mobility of antithrombin and its modulation by heparin" Thrombosis and Heamostasis 78, 516-519
- [111] Skinner, R., Abrahams, J.P., Whisstock, J.C., Lesk, A.M., Carrell, R.W., Wardell, M.R. (1997) "The 2.6 Å structure of antithrombin indicates a conformational change at the heparin binding site" J. Mol. Biol 226, 601-609
- [112] JP Abrahams (1996) "Electron density modification with Solomon" CCP4 Newsletter 32,
- [113] Abrahams, J.P. & Leslie, A.W.G. (1996) "Methods used in the structure determination of mitochondrial F1 ATPase", Acta Cryst. D52, 30 - 42
- [114] Van Raaij, M.J., Abrahams, J.P., Leslie, A.W.G. & Walker, J.E. (1996) "The structure of bovine F1 ATPase complexed with the antibiotic inhibitor aurovertin B", Proc. Natl. Acad. Sci. USA 93, 6913-6917
- [115] Abrahams JP (1996) "Likelihood-weighted real space restraints for refinement at low resolution" CCP4 Newsletter
- [116] Abrahams, J.P., Buchanan, S.K., van Raaij, M.J., Fearnley, I.M., Leslie, A.W.G. & Walker, J.E. (1996) "The structure of bovine F1 ATPase complexed with the peptide efrapeptin", Proc. Natl. Acad. Sci. USA 93 9420-9424
- [117] Elliott, P.R., Lomas, D.A., Carrell, R.W. & Abrahams, J.P. (1996) "Inhibitory conformation of the reactive loop of α1-antitrypsin", Nature Struct. Biol. 3, 676-681
- [118] Abrahams, J.P., Leslie, A.G.W, Lutter, R. & Walker, J.E. (1994) "Structure at 2.8 Å resolution of F1 ATPase from bovine heart mitochondria", Nature 370, 621-628
- [119] Wardell MR, Abrahams JP, Bruce D, Skinner R, Leslie AGW (1994) "Crystallisation and preliminary X-ray diffraction analysis of two conformations of intact human antithrombin" J. Mol. Biol. 234, 1253-1258
- [120] Lutter, R., Abrahams, J.P., Van Raaij, M.J., Todd, R.J., Lundquist, T., Buchanan, S.R., Leslie, A.G.W. & Walker, J.E. (1993) "Crystallization of F1-ATPase from bovine heart mitochondria", J.Mol. Biol. 229, 787-790
- [121] Abrahams, J.P., Lutter, R., Todd, R.J., Van Raaij, M.J., Leslie, A.G.W. & Walker, J.E. (1993)
 "Inherent asymmetry of F1-ATPase from bovine heart mitochondria at 6.5 Å resolution" EMBO J. 12, 1775-1780
- [122] Abrahams, J.P., Van Raaij, M.J., Bosch, L., Kraal, B., Walters, J.A.L.I. & Wijmenga, S. (1993) "Study by imino proton NMR of Phe-tRNAPhe in complex with elongation factor Tu and GTP" Spectroscopy 11, 37-44

- [123] JP Abrahams (1993) "Compression of X-ray images" CCP4 Newsletter 28, 3-4
- [124] Abrahams, J.P., Bosch, L., Kraal, B., De Groot, H.J.M., Raap, J. & Lugtenburg, J. (1992) "Magic angle spinning 13C NMR analysis of the complex of elongation factor Tu, GTP and [1-13C]phenylalanyl-tRNAphe", Spectroscopy 10, 1-8
- [125] Batenburg FDH, Bos V, Riethoven JJM, Abrahams JP, Pley CWA (1992) "Porting and optimizsing STAR: a case study of suffering and surfacing" APL 1992:265-274
- [126] Abrahams, J.P., Kraal, B., Clark, B.F.C. & Bosch, L.(1991) "Isolation and stability of ternary complexes of elongation factor Tu, GTP and aminoacyl-tRNA", Nucl. Acids Res. 19, 553-557
- [127] Abrahams, J.P., Acampo, J.J.C, Kraal, B. & Bosch, L. (1991) "The influence of tRNA located at the P-site on the turnover of EF-Tu.GTP on ribosomes", Biochimie 73, 1089-1092
- [128] Abrahams, J.P., Van Raaij, M.J., Ott, G., Kraal, B. & Bosch, L. (1991) "Kirromycin drastically reduces the affinity of E. coli elongation factor Tu for aminoacyl-tRNA", Biochemistry 30, 6705-6710
- [129] Abrahams, J.P., Van den Berg, M., Van Batenburg, F.D.H. & Pleij, C.W.A. (1990) "Prediction of RNA secondary structure, including pseudoknotting, by computer simulation", Nucl. Acids Res. 18, 3035-3044
- [130] Abrahams, J.P., Acampo, J.J.C., Ott, G., Sprinzl, M., De Graaf, J.M., Talens, A. & Kraal, B. (1990) "The interaction between aminoacyl-tRNA and the mutant elongation factors Tu Ar and B0", Biochem. Biophys. Acta 1050, 226-229
- [131] Abrahams, J.P., Kraal, B. & Bosch, L. (1988) "Zone-inteference gel electrophoresis: a new method for studying weak protein - nucleic acid complexes under native equilibrium conditions" Nucl. Acids Res. 16, 10099-10108
- [132] Van Belkum A, Abrahams JP, Pleij CWA, Bosch L (1985) "Five pseudoknots are present at the 204 nulcleotides long 3' noncoding region of tobacco mosaic virus RNA" Nucl. Acids. Res. 13, 7673-7686

Book chapters / books:

- [133] Nederlof I, van Genderen E, Hoedemaeker PJ, Abrahams JP, Georgieva D (2012) "Protein Crystal Growth" in Modern Aspects of Bulk Crystal and Thin Film Preparation, DOI: 10.5772/29424
- [134] Abrahams JP, Georgieva D, Jiang LH, Nederlof I (2012) "Electron diffraction of protein 3D nanocrystals" in U. Kolb (eds.) Uniting Electron Crystallography and Powder Diffraction, NATO Science for Peace and Security Series B: Physics and Biophysics, 389-398
- [135] Sikharulidze I, van Gastel R, Schramm A, Abrahams JP, Poelsema B, Tromp RM, van der Molen, SJ "Improved Imaging in Low Energy Electron Microscopy and Photo Emission Electron Microscopy Using MEDIPIX2 Pixel Detector" Astroparticle, Particle and Space Physics, Detectors and Medical Physics Applications ISBN-10 981-4307-51-3 133-139,
- [136] Kallergi A, Bei Y, Kok P, Abrahams JP, Dijkstra J & Verbeek FF (2009) "Cyttron: a virtualized microscope supporting image integration and knowledge discovery" in: Proteins killing tumour cells (Transworld Research Network), 291-316
- [137] Georgieva D, Jiang LH, Zandbergen H, Nicolopoulos S, Abrahams JP "3D electron diffraction of protein crystals: data collection, cell determination and indexing" in: EMC 2008: Vol. 1 Instrumentation and Methods (Luysberg et al , eds.) Springer-Verlag ISBN 978-3-540-85156-1, 759-760
- [138] Abrahams JP, Plaisier JR, Ness S & Pannu NS (2007) "Phase refinement through density modification" in: "Macromolecular crystallography conventional and high throughput methods" (Oxford Scientific Press), 143-154
- [139] Georgieva D, Abrahams JP & Kuil ME (2006) "Protein Nanocrystallization" in: Advanced techniques in biophysics (Springer), 1-26
- [140] Abrahams JP & Ban, N (2003) "X-ray Crystallographic structure determination of large asymmetric macromolecular assemblies" in: Meth. Enzomology 374 (Elsevier Acad. Press), 163-187
- [141] Abrahams JP (1991) "Enzyme regulation by RNA a study of elongation factor Tu" Leiden University, *PhD Thesis*
- [142] Kraal B, Abrahams JP, Bosch L (1989) "Effects of Kirromycin on the Elongation Factor EF-Tu and its Interactions with GDP or GTP and tRNA. The Application of zone-Interference Gel Electrophoresis, a New Method for the Analysis of Weak Complexes" In The Guanine-Nucleotide Binding Proteins (Bosch L, Kraal B, Parmeggiani A, eds.) ISBN 978-1-4757-2037-2, 121-129