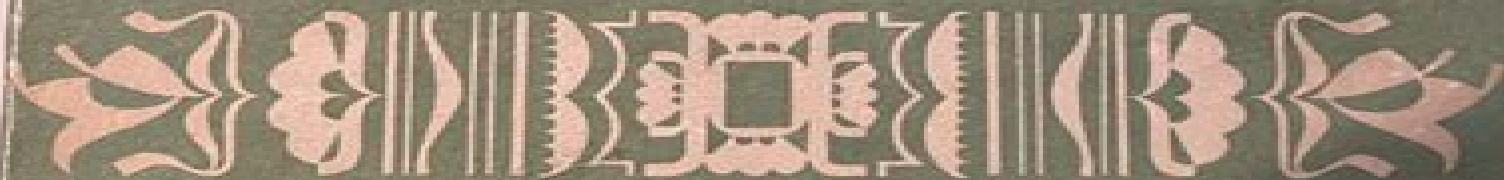


MAN

IN
STRUCTURE
AND
FUNCTION



FRITZ KAHN, M.D.

Man In Structure Function

**J.P.C. de Bruin, M.A. Corner, M.G.P.
Feenstra, C.G. Van Eden, H.B.M. Uylings**



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Man in Structure & [and] Function Fritz Kahn,1946

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Structure Function in Man Stanley W. Jacob,1982-01

Structure and Function in Man Stanley Wallace Jacob,Clarice

Ashworth Francone,Walter J. Lossow,1978

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The Trigeminal System in Man

Kamen G. Usunoff,Enrico Marani,Jaap H.R. Schoen,2012-12-06

This monograph offers a comprehensive review of present knowledge of the structure and connections of the trigeminal nuclei in humans and compares it to laboratory animal findings The authors provide cytoarchitectural data from their own research and trace trigeminal pathways in human material by means of the Nauta technique In humans the trigeminal nuclear complex includes the motor nucleus the principal sensory pontine nucleus the spinal nucleus subdivided into oral interpolar and caudal nuclei and the mesencephalic nucleus and several small nuclei The supratrigeminal nucleus as described in various mammals is not defined in the human brain The primary afferents to all subdivisions of the trigeminal nuclear complex in humans appear to be entirely ipsilateral Some of the extratrigeminal primary afferents described in experimental animals are also present in the human brain and the nucleus ovalis receives primary and possibly secondary afferents from the trigeminal systems A significant difference between the human trigeminal system and the subprimate species is seen in the monosynaptic cortical projection to the motor trigeminal nucleus

On protein structure, function and modularity from an evolutionary perspective Robert Pilstål,2018-05-23 We are compounded entities given life by a complex molecular machinery When studying these molecules we have to make sense of a diverse set of dynamical nanostructures with vast and intricate patterns of interactions Protein polymers is one of the major groups of building blocks of such nanostructures which fold up into more or less distinct three dimensional structures Due to their shape dynamics and chemical properties proteins are able to perform a plethora of specific functions essential to all known cellular lifeforms The connection between protein sequence translated into protein structure and in the continuation into protein function is well accepted but poorly understood Malfunction in the process of protein folding is known to be implicated in natural aging cancer and degenerative diseases such as Alzheimer s Protein folds are described hierarchically by structural ontologies such as SCOP CATH and Pfam all which has yet to succeed in deciphering the natural language of protein function These paradigmatic views centered on protein structure fail to describe more mutable entities such as intrinsically disordered proteins IDPs which lack a clear defined structure As of 2012 about two thirds of cancer patients was predicted to survive past 5 years of diagnosis Despite this about a third do not survive and numerous of successfully treated patients suffer from secondary conditions due to chemotherapy surgery and the like In order to handle cancer more efficiently we have to better understand the underlying molecular mechanisms Elusive to standard methods of investigation IDPs have a central role in pathology dysfunction in IDPs are key factors in cellular system failures such as cancer as many IDPs are hub regulators for major cell functions These IDPs

carry short conserved functional boxes that are not described by known ontologies which suggests the existence of a smaller entity. In an investigation of a pair of such boxes of c-MYC a plausible structural model of its interacting with Pin1 emerged but such a model still leaves the observer with a puzzle of understanding the actual function of that interaction. If the protein is represented as a graph and modeled as the interaction patterns instead of as a structural entity another picture emerges. As a graph there is a parallel from that of the boxes of IDPs to that of sectors of allosterically connected residues and the theory of foldons and folding units. Such a description is also useful in deciphering the implications of specific mutations. In order to render a functional description feasible for both structured and disordered proteins there is a need of a model separate from form and structure. Realized as protein primes patterns of interaction which has a specific function that can be defined as prime interactions and context. With function defined as interactions it might be possible that the discussion of proteins and their mechanisms is thereby simplified to the point rendering protein structural determination merely supplementary to understanding protein function.

Människan byggs upp av celler de i sin tur består av mindre beständsdelar livets molekyler. Dessa fungerar som mekaniska byggstenar liksom maskiner och robotar som sliter vid fabriken band envar utförandes en absolut nödvändig funktion för cellens och hela kroppens fortsatta överlevnad. De av livets molekyler som beskrivs centralt i den här avhandlingen är proteiner vilka i sin tur består utav en lång kedja med olika typer av länkar som liksom garn lindar upp sig i ett nystan av en mer eller mindre bestämd struktur som avgör dess roll och funktion i cellen. Intrinsiskt ordnade proteiner IDP ger emot denna enkla skildring de är proteiner som saknar struktur och beter sig mer liksom spaghetti i vatten än maskin. IDP är inte funktionella och bör spela centrala roller i cellens maskineri exempelvis oncoproteinet c-Myc som agerar gaspedal för cellen för att c-Mycs funktion leder till att cellerna löper amok delar sig hejdlöst och vi får cancer. Man har upptäckt att c-Myc har en ombytlig struktur vi inte kan se studier av punktvisa förändringar mutationer i kedjan av byggstenar hos c-Myc visar att många länkar har viktiga roller i funktionen. Detta ger oss bättre förståelse om cancer men samtidigt är laboratoriearbetet både komplicerat och dyrt här kan evolutionen hjälpa oss och avslöja hemligheterna snabbare. Molekylär evolution studeras genom att beräkna variation i proteinkedjan mellan beslötade arter som finns lagrade i databaser detta visar snabbt via nätverksanalys och grafteori vilka delar av proteinet som är centrala och kopplade till varandra av nödvändighet för artens överlevnad. På så vis hjälper evolutionen oss att förstå proteinfunktioner via modeller baserade på proteinernas interaktioner snarare än deras struktur. Samma modeller kan nyttjas för att förstå dynamiska förlopp och skillnader mellan normala och patologiska varianter av proteiner mutationer kan uppstå i vår arvsmassa som kan leda till sjukdom. Genom analys av proteinernas kopplingsnätverk i grafmodellerna kan man bättre förstå vilka mutationer som är farligare än andra. Dessutom har det visat sig att en sådan representation kan ge bättre förståelse för den normala funktionen hos ett protein än vad en proteinstruktur kan. Här introduceras även konceptet proteinprimär vilket är en abstrakt representation av proteiner centrerad på deras interaktiva mönster snarare än på partikulär form och struktur. Det är en förhoppning att en sådan

representation skall f renkla diskussionen anbelangande proteinfunktion s till den grad att strukturbest mmelse av proteiner som r en mycket kostsam och tidskr vande process till viss m n kan anses vara sekund r i betydelse j mf rt med funktionellt modellerande baserat p evolution ra data extraherade ur v ra sekvensdatabaser

The Prefrontal Cortex: Its Structure, Function and Pathology J.P.C. de Bruin,M.A. Corner,M.G.P. Feenstra,C.G. Van Eden,H.B.M. Uylings,1991-03-05 Thanks to a resurgence of interest and a recent proliferation of research techniques much new and illuminating data has emerged during the last decade relating to the prefrontal cortex particularly in primates and rodents In view of this progress the 16th International Summer School of Brain Research was held in Amsterdam The Netherlands from 28 August to 1 September 1989 devoted to the topic of The Prefrontal Cortex Its Structure Function and Pathology The edited proceedings embodied in this 85th volume of Progress in Brain Research fall into three sections the first of which following two introductory chapters discusses the present knowledge of the organization of prefrontal cortical systems In the second section developmental and plasticity aspects in rodent and human cortex are considered whilst the third section deals extensively with the functional aspects characteristic for the prefrontal cortex in primates rats and rabbits The last section reviews several topics on dysfunction of prefrontal cortex in rat and man including a historical review on psychosurgery

Exploring Latin: Structures, Functions, Meaning Lucie Pultrová,Martina Vaníková,2024-11-18 This two volume work contains a selection of papers first presented at the 22nd International Colloquium on Latin Linguistics held in Prague 2023 The papers address important issues in Latin linguistics with a variety of theoretical and methodological approaches The first volume Word contains texts concerning Latin phonology etymology flexion and derivation and lexical semantics both with respect to individual words and to entire word classes Both diachronic and synchronic perspectives are employed in the discussion of the various issues The second volume Clause and Discourse includes papers dealing with issues of syntax and semantics and with the structure of texts and pragmatic aspects One of the subchapters entitled Conversation and Dialogue contains papers presented at the conference in a separate workshop of the same name linked by a common methodological framework of Conversation Analysis This book provides essential texts for researchers in the field of Latin linguistics and may also be of use to linguists who work primarily with other languages

Man-Machine-Environment System Engineering Shengzhao Long,Balbir S. Dhillon,Long Ye,2024-09-28 From this book reader will learn the best research topics and the latest development trend in MMESE theory and application Man Machine Environment System Engineering MMESE is a scientific study on the design concepts and quantitative analysis of a complex giant system using physiology psychology system engineering computer science environment science management theory education and other related disciplines methods MMESE focuses mainly on the relationship and the optimum combination between Man Machine and Environment The three optimized goals of the MMESE study are safety efficiency and economy Researchers and professionals who study a human centered interdisciplinary subject crossing above disciplines will be mostly benefited from this proceedings In 1981

with direct support from one of the greatest modern Chinese scientists Xuesen Qian Man Machine Environment System Engineering MMESE the integrated and advanced science research topic was established in China by Professor Shengzhao Long Man Machine Environment System Engineering Proceedings of the 24th Conference on MMESE is the academic showcase of latest research papers selected from more than 500 submission in this field in 2024 *Evolution and Man's Place in Nature* Henry Calderwood,1893 **Grasslands, Systems Analysis and Man** A. I. Brey Meyer, George M. van Dyne,1980-03-20 This 1980 book was a synthesis of much of the recent work on the functioning of grassland ecosystems at the time Research Grants Index National Institutes of Health (U.S.). Division of Research Grants,1975 Research Awards Index ,1978 **The Weather and Climate** Shaun Lovejoy, Daniel Schertzer,2013-04-04 A new method of modeling the atmosphere synthesizing data analysis techniques and multifractal statistics for atmospheric researchers and graduate students An Essay on Man and Christian Civilization William Young Craig,1906 **Steroidal Activity in Experimental Animals and Man** Ralph I. Dorfman,2013-10-22 Methods in Hormone Research Volume III Steroidal Activity in Experimental Animals and Man Part A focuses on protection against irradiation damage anti mammary tumor activity body lipids anti aldosterone central depressant action and copulatory reflex activity The selection first offers information on estrogens and anti estrogens including relative potencies in vaginal smear tests vaginal tests and uterine weight tests mechanism of estrogen action and mode action of anti estrogens The book then examines irradiation protection and anti mammary tumor activities in rats and mice The publication surveys steroids and lipid metabolism and the activities of adrenocorticosteroids in experimental animals and man Discussions focus on adrenocortical estrogenic and androgenic and anabolic steroids relative activities of corticosteroids assessed by laboratory and clinical methodology and comparative efficacies of corticosteroids The manuscript then ponders on anti aldosterones steroid central depressants and copulatory reflex response to steroids The selection is a vital source of data for readers interested in steroidal activity in man and animals **Genetic Diversity and Human Behavior** J.N. Spuhler,2017-09-05 Genetic differences in humans like those between individuals of any animal or plant species and those between species are all products of the evolutionary development of the living world These differences with their behavioral consequences can only be understood in the light of evolution Our understanding of evolution however has itself evolved The Darwin Wallace theory of evolution appeared in the nineteenth century Since then development of evolutionary thought has gone through several stages The contributions in this volume describe those stages The Mind of Man Gustav Spiller,1902 The Brain from Ape to Man Frederick Tilney,1928 This work is a classic study of the evolution of the central nervous system in the higher mammals

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