

Memory Sequence

Andreas Gogol-Döring,Knut Reinert

The Cognition of Sequences Snehlata Jaswal,2018-01-24 It is impossible to perceive the innumerable stimuli impinging on our senses, all at once. Out of the myriad stimuli, external and internal, a few are selected for further processing; and even among these, we try to put each in some sort of relation with the others, to be able to make some sense about them all. Time, of course, is an elementary dimension we use to organize our experiences. Thus, the perception of sequences is basic to human cognition. Nevertheless, research addressing sequences is rather sparse. Partly, this is due to difficulty in designing experiments in this area due to huge individual differences. Then, there is the assumption that temporal order has more to do with memory than perception. Another problem is that sequences seem endemic to the auditory world. So much so that some researchers have suggested that sound provides the ‘auditory scaffolding’ for sequencing behavior. Little wonder that research studies addressing sequences in modalities other than audition are extremely rare. This research topic aimed to gather a holistic picture of sequencing behaviour among humans by collecting snapshots of the current research on the topic of sequencing. We particularly sought contributions which addressed sequences beyond the auditory modality. The single unifying criteria for these diverse contributions was that they shed new light on previously unexplored empirical relationships and/or provoked new lines of research with incisive ideas regarding sequencing behavior. Seasoned researchers contributed their views on perception, memory, and production of sequences.

Sequence Learning Ron Sun,C.Lee Giles,2003-06-29 Sequential behavior is essential to intelligence in general and a fundamental part of human activities, ranging from reasoning to language, and from everyday skills to complex problem solving. Sequence learning is an important component of learning in many tasks and application fields: planning, reasoning, robotics natural language processing, speech recognition, adaptive control, time series prediction, financial engineering, DNA sequencing, and so on. This book presents coherently integrated chapters by leading authorities and assesses the state of the art in sequence learning by introducing essential models and algorithms and by examining a variety of applications. The book offers topical sections on sequence clustering and learning with Markov models, sequence prediction and recognition with neural networks, sequence discovery with symbolic methods, sequential decision making, biologically inspired sequence learning models.

Advances in Genomic Sequence Analysis and Pattern Discovery ,

Narrative memory and the impact of trauma on individuals with reference to one short sequence from “Memento” Michael Schmid,2007-01-12 Seminar paper from the year 2004 in the subject American Studies - Culture and Applied Geography, grade: 1,3, Free University of Berlin (John F. Kennedy Institut Berlin), course: HS American Cultural Memory: Trauma, Collective Imagery and the Politics of Remembering, language: English, abstract: The text by Bessel Van der Kolk and Onno Van der Hart “The Intrusive Past” provides an overview of the work and achievement of Jean – Martin Charcot’s and Pierre Janet’s study about how the mind processes memories and the effects of traumatic memories on consciousness. With the following text, I will present a couple of central aspects of Janet’s study and the phenomena of dissociation and the reconstruction of the past through narrative memory and project them onto one short sequence from “Memento” (2001) to further support my argument. The main point of this text is to illustrate how narrative memory reshapes the past in a variety of ways and that the main character in “Memento”, who has lived through a traumatic experience, creates and recreates his past through the means of a combination of the already mentioned dissociation and narrative memory. Janet considered “the memory system as the central organizing apparatus of the mind, which categorizes and integrates all aspects of experience and automatically integrates them into ever – enlarging and flexible meaning schemes.” He differentiates between the subconscious automatic integration of familiar and expectable experiences into existing meaning schemes and the difficult integration of frightening and novel experiences, which might either totally resist integration or be remembered extremely vivid. The subconscious integration of memories occurs because they fit easily into the meaning scheme, they do not pose a threat or form a contradiction to the already existing beliefs, values and meanings of the world. Whereas the automatic integration of new information happens without conscious attention, the narrative memory is something very deliberate and conscious. Narrative memory is not the act of remembering something that happened in the past but an act of recreating the past, of changing the memory. Janet explains this phenomena as mental constructs, “which people use to make sense out of experience.” This suggests that the individual’s existing meaning schemes may be entirely unable to integrate a specific terrifying experience, which causes the memory to be stored differently, and therefore might not be available for the act of remembering.

Computer Analysis of Sequence Data, Part I Annette M. Griffin,Hugh G. Griffin,2008-02-02 DNA sequencing has become increasingly efficient over the years, resulting in an enormous increase in the amount of data generated. In recent years, the focus of sequencing has shifted, from being the endpoint of a project, to being a starting point. This is especially true for such major initiatives as the human genome project, where vast tracts of DNA of unknown function are sequenced. This sheer volume of available data makes advanced computer methods essential to analysis, and a familiarity with computers and sequence analysis software a vital requirement for the researcher involved with DNA sequencing. Even for nonsequencers, a familiarity with sequence analysis software can be important. For instance, gene sequences already present in the databases can be extremely useful in the design of cloning and genetic manipulation experiments. This two-part work on Analysis of Data is designed to be a practical aid to the researcher who uses computers for the acquisition, storage, or analysis of nucleic acid (and/or protein) sequences. Each chapter is written such that a competent scientist with basic computer literacy can carry out the procedure successfully at the first attempt by simply following the detailed practical instructions that have been described by the author. A Notes section, which is included at the end of each chapter, provides advice on overcoming the common problems and pitfalls sometimes encountered by users of the sequence analysis software. Software packages for both the mainframe and personal computers are described.

Sequence Kenneth Rogers, Jr.,2012-03-05 Andrea Remus and Thomas Charon are trapped inside the Pandora Complex, a phenomena which causes thoughts to travel from one memory to the next as they are downloaded into the Pandora system. Each download causes Andrea and Thomas to feel as though they are reliving their memories for the first time. As they increase contact with Pandora, their minds become trapped in the past. Will Andrea and Thomas regain their full memories? What dangers will they encounter as their intense visits to the past cause a stunning affect? The riveting Sequence questions our reality, while taking the reader on a remarkable journey into the mind.

Next Generation Sequencing Technologies and Challenges in Sequence Assembly Sara El-Metwally,Osama M. Ouda,Mohamed Helmy,2014-04-19 The introduction of Next Generation Sequencing (NGS) technologies resulted in a major transformation in the way scientists extract genetic information from biological systems, revealing limitless insight about the genome, transcriptome and epigenome of any species. However, with NGS, came its own challenges that require continuous development in the sequencing technologies and bioinformatics analysis of the resultant raw data and assembly of the full length genome and transcriptome. Such developments lead to outstanding improvements of the performance and coverage of sequencing and improved quality for the assembled sequences, nevertheless, challenges such as sequencing errors, expensive processing and memory usage for assembly and

sequencer specific errors remains major challenges in the field. This book aims to provide brief overviews the NGS field with special focus on the challenges facing the NGS field, including information on different experimental platforms, assembly algorithms and software tools, assembly error correction approaches and the correlated challenges.

Temporal sequence in the perception of speech Warren H. Fay,2018-11-05 No detailed description available for Temporal sequence in the perception of speech.

Biological Sequence Analysis Using the SeqAn C++ Library Andreas Gogol-Döring,Knut Reinert,2009-11-11 An Easy-to-Use Research Tool for Algorithm Testing and Development Before the SeqAn project, there was clearly a lack of available implementations in sequence analysis, even for standard tasks. Implementations of needed algorithmic components were either unavailable or hard to access in third-party monolithic software products. Addressing these concerns, the developers of SeqAn created a comprehensive, easy-to-use, open source C++ library of efficient algorithms and data structures for the analysis of biological sequences. Written by the founders of this project, Biological Sequence Analysis Using the SeqAn C++ Library covers the SeqAn library, its documentation, and the supporting infrastructure. The first part of the book describes the general library design. It introduces biological sequence analysis problems, discusses the benefit of using software libraries, summarizes the design principles and goals of SeqAn, details the main programming techniques used in SeqAn, and demonstrates the application of these techniques in various examples. Focusing on the components provided by SeqAn, the second part explores basic functionality, sequence data structures, alignments, pattern and motif searching, string indices, and graphs. The last part illustrates applications of SeqAn to genome alignment, consensus sequence in assembly projects, suffix array construction, and more. This handy book describes a user-friendly library of efficient data types and algorithms for sequence analysis in computational biology. SeqAn enables not only the implementation of new algorithms, but also the sound analysis and comparison of existing algorithms. Visit SeqAn for more information.

Analysis of Single Sequence of Words Retrieved from Long Term Memory Tarow Indow,1993

Supervised Sequence Labelling with Recurrent Neural Networks Alex Graves,2012-02-06 Supervised sequence labelling is a vital area of machine learning, encompassing tasks such as speech, handwriting and gesture recognition, protein secondary structure prediction and part-of-speech tagging. Recurrent neural networks are powerful sequence learning tools—robust to input noise and distortion, able to exploit long-range contextual information—that would seem ideally suited to such problems. However their role in large-scale sequence labelling systems has so far been auxiliary. The goal of this book is a complete framework for classifying and transcribing sequential data with recurrent neural networks only. Three main innovations are introduced in order to realise this goal. Firstly, the connectionist temporal classification output layer allows the framework to be trained with unsegmented target sequences, such as phoneme-level speech transcriptions; this is in contrast to previous connectionist approaches, which were dependent on error-prone prior segmentation. Secondly, multidimensional recurrent neural networks extend the framework in a natural way to data with more than one spatio-temporal dimension, such as images and videos. Thirdly, the use of hierarchical subsampling makes it feasible to apply the framework to very large or high resolution sequences, such as raw audio or video. Experimental validation is provided by state-of-the-art results in speech and handwriting recognition.

Data Structures and Algorithms in Java Michael T. Goodrich,Roberto Tamassia,Michael H. Goldwasser,2014-01-28 The design and analysis of efficient data structures has long been recognized as a key component of the Computer Science curriculum. Goodrich, Tomassia and Goldwasser's approach to this classic topic is based on the object-oriented paradigm as the framework of choice for the design of data structures. For each ADT presented in the text, the authors provide an associated Java interface. Concrete data structures realizing the ADTs are provided as Java classes implementing the interfaces. The Java code implementing fundamental data structures in this book is organized in a single Java package, net.datastructures. This package forms a coherent library of data structures and algorithms in Java specifically designed for educational purposes in a way that is complimentary with the Java Collections Framework.

Attributes of Memory (Psychology Revivals) Peter Herriot,2013-11-05 First published in 1974, Attributes of Memory rejected the prevalent stress on the structure of memory. It suggests that the view of memory as a sequence of stores through which information passes is mistaken. Instead, the author emphasizes the coding process of memory by which the nominal stimulus, the stimulus as presented, is transformed into the functional stimulus, the stimulus as coded. Dr Herriot proposes that there are many different forms of coding, and that efficiency of recall or recognition performance is a function of the nature of coding employed. He suggests that the subject's linguistic system is the most frequently employed linguistic device; that is, that the underlying attributes and rules of language are used automatically when material is verbal. Since the basic function of language is to communicate meaning, those forms of coding which are meaningful in nature are most effective in memory. The book cites a great deal of experimental evidence, including many studies of the time. As well as stating a point of view, it should be useful to undergraduate and postgraduate students as a review of the early literature, read in its historical context.

Encyclopedia of the Mind Harold Pashler,2013-01-14 It's hard to conceive of a topic of more broad and personal interest than the study of the mind. In addition to its traditional investigation by the disciplines of psychology, psychiatry, and neuroscience, the mind has also been a focus of study in the fields of philosophy, economics, anthropology, linguistics, computer science, molecular biology, education, and literature. In all these approaches, there is an almost universal fascination with how the mind works and how it affects our lives and our behavior. Studies of the mind and brain have crossed many exciting thresholds in recent years, and the study of mind now represents a thoroughly cross-disciplinary effort. Researchers from a wide range of disciplines seek answers to such questions as: What is mind? How does it operate? What is consciousness? This encyclopedia brings together scholars from the entire range of mind-related academic disciplines from across the arts and humanities, social sciences, life sciences, and computer science and engineering to explore the multidimensional nature of the human mind.

The Cognitive Neuroscience of Memory Amanda Parker,Timothy J. Bussey,Edward L. Wilding,2005-08-18 Recent advances in techniques available to memory researchers have led to a rapid expansion in the field of cognitive neuroscience of memory. This book provides accessible coverage of four key areas of recent advance, including research on functional imaging, electrophysiological and lesion studies, and developments from the computational modelling approach. The first section reviews functional imaging studies in humans, with particular emphasis on how imaging methods have clarified the cortical areas involved in memory formation and retrieval. The second section describes electrophysiological and lesion research in monkeys, where lesion and disconnection studies are rapidly adding to our knowledge of both information processing and modulatory aspects of memory formation. In the third section, electrophysiological and lesion studies in rats are reviewed allowing for a detailed study of the role of novelty and exploration in memory formation. The final section reviews current research in computational modelling which has allowed the development of new theoretical and experimental approaches to the study of memory encoding and retrieval. This volume draws together the current developments in each field, allowing the synthesis of ideas and providing converging evidence from a range of sources. It will be a useful resource for both advanced undergraduate and postgraduate students of psychology, as well as researchers in the field and anyone with an interest in cognitive neuroscience.

Narrative Memory and the Impact of Trauma on Individuals with Reference to One Short Sequence from Memento Michael Schmid,2007-11 Seminar paper from the year 2004 in the subject American Studies - Culture and Applied Geography, grade: 1,3, Free University of Berlin (John F. Kennedy Institut Berlin), course: HS American Cultural Memory: Trauma, Collective Imagery and the Politics of Remembering, 3 entries in the bibliography, language: English, abstract: The text by Bessel Van der Kolk and Onno Van der Hart The Intrusive Past provides an overview of the work and achievement of Jean - Martin Charcots and Pierre Janet's study about how the mind processes memories and the effects of traumatic memories on consciousness. With the following text, I will present a couple of central aspects of Janet's study and the phenomena of dissociation and the reconstruction of the past through narrative memory and project them onto one short sequence from Memento (2001) to further support my argument. The main point of this text is to illustrate how narrative memory reshapes the past in a variety of ways and that the main character in Memento, who has lived through a traumatic experience, creates and recreates his past through the means of a combination of the already mentioned dissociation and narrative memory. Janet considered the memory system as the central organizing apparatus of the mind, which categorizes and integrates all aspects of experience and automatically integrates them into ever - enlarging and flexible meaning schemes. He differentiates between the subconscious automatic integration of familiar and expectable experiences into existing meaning schemes and the difficult integration of frightening and novel experiences, which might either totally resist integration or be remembered extremely vivid. The subconscious integration of memories occurs because they fit easily into the meaning scheme, they do not pose a threat or form a contradiction to the already existing beliefs, values and meanings of the world. Whereas the automatic integration of new information happens without consci

Foundations of Human Memory Michael Jacob Kahana,2014-05-01 Foundations of Human Memory provides an introduction to the scientific study of human memory with an emphasis on both the major theories of memory and the laboratory studies that have been used to test those theories and inspire their further development. Written with the undergraduate student in mind, the text assumes no specific background in the subject, but a general familiarity with scientific method and quantitative approaches to the treatment of data. Foundations of human memory is organized around the major empirical paradigms used to study memory in the laboratory and the theories used to explain data obtained using those paradigms. The text begins with a focus on memory for individual items, building up to memory for associations between items, and finally to memory for entire sequences of items and the problem of memory search. Several major theories of memory are considered in detail, including strength theory, summed-similarity theory, neural network based theories, retrieved-context theory, and theories based on the division of memory into separate short-term and long-term storage systems. The text emphasizes basic research over applied problems, but brings in real-world examples and neuroscientific evidence as appropriate.

Multiple Biological Sequence Alignment Ken Nguyen,Xuan Guo,Yi Pan,2016-06-10 Covers the fundamentals and techniques of multiple biological sequence alignment and analysis, and shows readers how to choose the appropriate sequence analysis tools for their tasks This book describes the traditional and modern approaches in biological sequence alignment and homology search. This book contains 11 chapters, with Chapter 1 providing basic information on biological sequences. Next, Chapter 2 contains fundamentals in pair-wise sequence alignment, while Chapters 3 and 4 examine popular existing quantitative models and practical clustering techniques that have been used in multiple sequence alignment. Chapter 5 describes, characterizes and relates many multiple sequence alignment models. Chapter 6 describes how traditionally phylogenetic trees have been constructed, and available sequence knowledge bases can be used to improve the accuracy of reconstructing phylogeny trees. Chapter 7 covers the latest methods developed to improve the run-time efficiency of multiple sequence alignment. Next, Chapter 8 covers several popular existing multiple sequence alignment server and services, and Chapter 9 examines several multiple sequence alignment techniques that have been developed to handle short sequences (reads) produced by the Next Generation Sequencing technique (NSG). Chapter 10 describes a Bioinformatics application using multiple sequence alignment of short reads or whole genomes as input. Lastly, Chapter 11 provides a review of RNA and protein secondary structure prediction using the evolution information inferred from multiple sequence alignments. • Covers the full spectrum of the field, from alignment algorithms to scoring methods, practical techniques, and alignment tools and their evaluations • Describes theories and developments of scoring functions and scoring matrices •Examines phylogeny estimation and large-scale homology search Multiple Biological Sequence Alignment: Scoring Functions, Algorithms and Applications is a reference for researchers, engineers, graduate and post-graduate students in bioinformatics, and system biology and molecular biologists. Ken Nguyen, PhD, is an associate professor at Clayton State University, GA, USA. He received his PhD, MSc and BSc degrees in computer science all from Georgia State University. His research interests are in databases, parallel and distribute computing and bioinformatics. He was a Molecular Basis of Disease fellow at Georgia State and is the recipient of the highest graduate honor at Georgia State, the William M. Suttles Graduate Fellowship. Xuan Guo, PhD, is a postdoctoral associate at Oak Ridge National Lab, USA. He received his PhD degree in computer science from Georgia State University in 2015. His research interests are in bioinformatics, machine leaning, and cloud computing. He is an editorial assistant of International Journal of Bioinformatics Research and Applications. Yi Pan, PhD, is a Regents' Professor of Computer Science and an Interim Associate Dean and Chair of Biology at Georgia State University. He received his BE and ME in computer engineering from Tsinghua University in China and his PhD in computer science from the University of Pittsburgh. Dr. Pan's research interests include parallel and distributed computing, optical networks, wireless networks and bioinformatics. He has published more than 180 journal papers with about 60 papers published in various IEEE/ACM journals. He is co-editor along with Albert Y. Zomaya of the Wiley Series in Bioinformatics.

Long Short-Term Memory Networks With Python Jason Brownlee,2017-07-20 The Long Short-Term Memory network, or LSTM for short, is a type of recurrent neural network that achieves state-of-the-art results on challenging prediction problems. In this laser-focused Ebook, finally cut through the math, research papers and patchwork descriptions about LSTMs. Using clear explanations, standard Python libraries and step-by-step tutorial lessons you will discover what LSTMs are, and how to develop a suite of LSTM models to get the most out of the method on your sequence prediction problems.

Space, Time and Number in the Brain Stanislas Dehaene,Elizabeth Brannon,2011-07-13 The study of mathematical cognition and the ways in which the ideas of space, time and number are encoded in brain circuitry has become a fundamental issue for neuroscience. How such encoding differs across cultures and educational level is of further interest in education and neuropsychology. This rapidly expanding field of research is overdue for an interdisciplinary volume such as this, which deals with the neurological and psychological foundations of human numeric capacity. A uniquely integrative work, this volume provides a much needed compilation of primary source material to researchers from basic neuroscience, psychology, developmental science, neuroimaging, neuropsychology and theoretical biology. The first comprehensive and authoritative volume dealing with neurological and psychological foundations of mathematical cognition Uniquely integrative volume at the frontier of a rapidly expanding interdisciplinary field Features outstanding and truly international scholarship, with chapters written by leading experts in a variety of fields

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