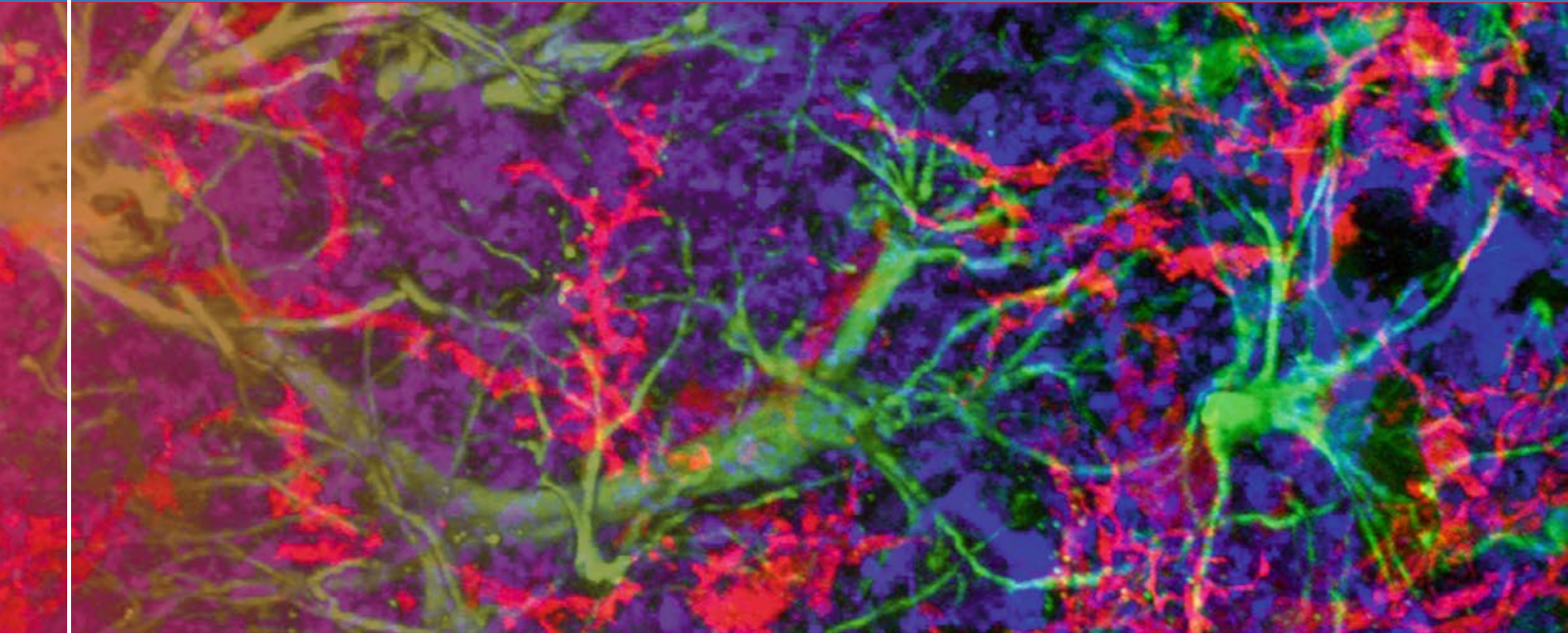


Tsuneya Ikezu
Howard E. Gendelman *Editors*



Neuroimmune Pharmacology

Second Edition

Serge Przedborski · Eliezer Masliah · Marco Cosentino
Associate Editors

 Springer

Neuroimmune Pharmacology

Tsuneya Ikezu • Howard E. Gendelman
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ISBN 978-3-319-44020-0 ISBN 978-3-319-44022-4 (eBook)
DOI 10.1007/978-3-319-44022-4

Library of Congress Control Number: 2016954435

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Printed on acid-free paper

This Springer imprint is published by Springer Nature
The registered company is Springer International Publishing AG
The registered company address is: Gewerbestrasse 11, 6330 Cham, Switzerland

Preface

In the past three decades, enormous strides have been made in our understanding of the relationships between inflammation, immune responses, and degenerative human diseases. Neuroinflammation has grown from a tiny beginning to its current status as the largest field of brain research. Its growth continues to be rapid so this dominance will continue to expand. The field commenced with the obscure identification of activated microglia in the brains of Alzheimer's disease cases. A grant application of ours to the Canadian government to explore the ramifications of this finding was turned down with the terse comment: "This hypothesis is ridiculous!"

To grasp the dimensions of growth in this field, and to gain a perspective of future opportunities, today's neuroscientists only need to read this volume from cover to cover. The developing information has mostly appeared in specialty journals that have dealt only with isolated aspects of these tightly related fields. As a result, contemporary scientists have had a difficult time finding sources, even in review articles, that provide an integrated picture. This updated volume, by assembling chapters that demonstrate the relationship between these historically separated fields, overcome that difficulty. There are 56 chapters which cover a broad spectrum of topics on immunology of the nervous system. Included are diseases that result from immunological dysfunction, current therapeutic approaches, and prospects for the future. Overall, it integrates cutting-edge neuroscience, immunology, pharmacology, neurogenetics, neurogenesis, gene therapy, adjuvant therapy, nanomedicine, pharmacogenetics, biomarkers, proteomics, and magnetic resonance imaging. It is a rich harvest and readers will gain a perspective that has not previously been so readily available. Exposure to such a wealth of ideas is bound to inspire readers to undertake new and productive research initiatives.

The modern era of research into neuroinflammation and its relationship to neurodegenerative diseases began in the 1960s with the elaboration by Ralph van Furth of the monocyte phagocytic system. He injected labeled monocytes into animals and followed their migration and maturation into resident phagocytes in all body tissues. This provided a closure between Metchnikoff's 1882 discovery of mesodermal attack cells in starfish larvae, which he named phagocytes, and del Rio Hortega's 1919 discovery of phagocytic mesodermal cells entering the brain, which he named microglia. Hortega's results had always been questioned, and for more than two further decades, the controversy continued as to whether microglia were truly phagocytes of mesodermal origin or were merely typical brain cells of epidermal origin. Resolving the controversy required development of the techniques of immunohistochemistry and monoclonal antibody production. These tools for exploring brain biochemistry at the cellular level opened new vistas for understanding brain functioning and the pathogenesis of human disease. Using these tools, our laboratory and that of Joseph Rogers in Sun City at that time demonstrated that HLA-DR was strongly expressed on activated microglia. The identification of HLA-DR, a well-known leukocyte marker displayed by antigen-presenting cells, on these cells vindicated both Hortega and van Furth. The way was paved for many productive investigations exploring the properties of microglial cells and their relationship to inflammation and immune responses. This example of a conjunction between a fundamental concept and technical advances to establish its validity has been repeated many times since, as the chapters in this volume illustrate.

For a time, the concept that the brain is immunologically privileged held sway among neuroscientists. This was based on a narrow view that only the invasion of brain by lymphocytes could be taken as evidence of an inflammatory response. But immunohistochemistry, coupled with newly developed molecular biological techniques, revealed that a spectrum of inflammatory mediators, including many known to cause tissue damage, was produced within the brain by resident brain cells. These discoveries required entirely new interpretations as to the nature of neuroinflammation and its relationship to immune responses. The innate immune system, operating at the local level in brain, has clearly proved to be the first line of defense. Indeed, the basic discoveries from studying the response of the brain in a variety of neurological diseases are causing a reevaluation of a number of peripheral degenerative disorders where innate immune responses, which had previously been ignored, have been shown to play a critical role in their pathogenesis. In other words, those studying the brain are providing immunologists with revolutionary new concepts regarding classical peripheral diseases. The insights of this volume need to be interpreted in this broader context.

Major neurologic disorders and details of their pathobiology are presented as individual chapters. They involve disorders where innate immune responses predominate, as in Alzheimer's disease, to others such as multiple sclerosis, where adaptive immune responses predominate, and others which seem to involve both. We have suggested that diseases involving self-damage generated by innate immune responses be defined as autotoxic to differentiate them from classical autoimmune diseases where self-damage is generated by adaptive immune responses. The common theme, however, is the involvement of microglia as the effector cells.

Genetics is well covered. It is a rapidly moving field. The methodology for linking familial disease to DNA mutations commenced in the late 1970s through identification of restriction fragment-linked polymorphisms. By 1983, when James Gusella and his colleagues demonstrated a linkage of the G8 fragment to Huntington disease, only about 18 markers were known. Now over 1,500,000 single-nucleotide polymorphisms have been localized so that every centimorgan of the human genome can be explored. This advance has been coupled with rapid methods for sequencing DNA. The report on genetics must be regarded as the tiny tip of a giant iceberg where much below the surface will soon be revealed.

The ultimate objective of neuroscientists studying human disease is to find more effective treatments. Part 3 covers the pharmacology of existing drugs, as well as describing approaches now in clinical evaluation, and those still at the bench level. Some of these include concepts that depart from established therapeutic approaches, giving the reader much food for thought.

There is an important chapter on the new field of biomarkers. Biomarkers have established that neurodegenerative disorders such as Alzheimer's disease commence at least a decade before clinical signs develop. A window of opportunity is opened up where early anti-inflammatory intervention can arrest disease progression and abort disease development. This can be combined with brain imaging to measure objectively the effects of therapeutic agents in diseases where progressive brain degeneration occurs.

In summary, this is a volume not to be put on the shelf as a reference text, but to be read cover to cover by aspiring neuroscientists.

Introducing Neuroimmune Pharmacology

Neuroscience, immunology, and pharmacology are each of and by themselves broad disciplines that, without argument, impact upon a large component of what we come to know as biomedical science (Elenkov et al. 2000; Gendelman 2002; Holzer et al. 2015; McGeer and McGeer 2004). Each, by themselves and even more so when put together, is multidisciplinary and require, for the student, both a broad knowledge and deep understanding of molecular and cellular biology, neuroimmunity, the functional blood–brain barrier, neurochemistry, neuroinfectious and neurodegenerative disorders, cancer, and neurodevelopment. For many, it is considered a branch of immunology, but that is a start point as the field bridges investigations of drug action and development with nervous system biology and disease pathogenesis. To those engaged in this field, linking of the disciplines is ever more challenging as when they are joined, they come interactive. The bridges between disciplines are what we now call multidisciplinary science and require another level of insight. However, we posit that each need be first understood as a single entity. To this end, we forged chapters that cover the structure, function, and biology of each of the fields independent of one another. Then step-by-step they are each combined one with the other to form the basis of our engagement with the environment, to disease and a means to restore homeostasis by protective immune and pharmacologic means. Indeed, drugs that include immune modulators can certainly influence organ function, aging, and tissue homeostasis and improve clinical outcomes.

Indeed, a special feature of “humankind” among other species is the presence of an extraordinary complex immune system that can be used to protect against a plethora of harmful microbial pathogens, including viruses, bacteria, and parasites, as well as abnormal cells and proteins (Petrazyi 2002; Obermeier et al. 2016). This underlies the complexity of the human genome which encodes expansive immune-related genes not found in lower species (Hughes 2002). When the immune system is compromised, disease occurs and often does with ferocity; a wide range of clinical manifestations ensue that follows as a consequence of neurodegenerative, psychiatric, cancer, and infectious diseases, or those elicited by the immune system’s attack on itself. The latter is commonly referred to as “destructive” autoimmunity (Christen and von Herrath 2004). Interestingly enough, the immune system may sometimes be an impediment to therapy. Indeed, modulating its function is required for long-term and successful organ transplantation (Samaniego et al. 2006; Horst et al. 2016). On balance, modulation of the immune system can affect “neuroprotective” responses for certain diseases (Anderson et al. 2014).

Like the immune system, the nervous system contains surveillance functions and also possesses a number of functional roles that include mentation, movement, reasoning, sensation, vision, hearing, learning, breathing, and most behaviors. The nervous system includes defined tissue structures such as the brain, spinal cord, and peripheral nerves. On the cellular level, it includes networks of nerve cells with a variety of functional activities; complex networks and communications; supportive and regulatory cells, called glia; and a protective barrier that precludes the entry of a variety of macromolecules, cells, and proteins. It also possesses connections throughout the body that permits it function. Neuroscience is the discipline used to explore each of the nervous system regions and cells that include their networks and modes of communication in health and disease. As humans, we have ~100 billion neurons that are each

functional units contained within the nervous system. Molecular and biochemical studies along with cell and animal systems were each used alone and together to explore and define neural biology. The task of neurosciences is to better understand the brain's function in the context of ontogeny, organism development, and aberrations during disease. Neuroscience is an interdisciplinary field and evolved as such during the past quarter-century. It includes neurobiology, neurochemistry, neurophysiology, mathematics, psychology, computer neuroscience, and learning and behavior. It also included the field of immunology (Sehgal and Berger 2000). In a historical context, the brain, for a long time, was considered an immune-privileged organ, meaning that its protective shield or barrier, commonly termed the blood–brain barrier, served to protect, defend, and as a consequence exclude ingress of toxins, cells, and pathogens (Streilein 1993; Becher et al. 2000). This has undergone a reassessment of purpose (Louveau 2015). Nonetheless, this is balanced by the fact that resident inflammatory cells do exist inside the brain and are capable of producing robust immune responses. In recent years, we have come to accept that it is the mononuclear phagocytes (MP; perivascular macrophage and microglia) that are disease perpetrators, while the astrocyte serves as “supportive” and “homeostatic” in nurturing neurons and in protection against the ravages of disease (Gendelman 2002; Simard and Nedergaard 2004; Trendelenburg and Dirnagl 2005). The neuron in this neuroimmune model is the passive recipient of the battles that rage between the MP and the astrocyte. Findings that have emerged over the past half-decade have challenged this model. We now know that dependent upon environmental cues and disease, microglia, astrocytes, and other neural cell elements including endothelial cells and oligodendrocytes possess immunoregulatory functions. We also know that microglia and astrocytes dependent upon the environment and stimuli can be supportive, destructive, or both. Even more importantly, neurons can secrete immunoregulatory factors and engage directly into cell–cell–environmental cue stimulations. To make the system perhaps even more complex, local neuroimmune processes can result in the recruitment of T cells and enticement of the adaptive immune response, significantly affecting disease outcomes (Olson and Gendelman 2016). All in all, things appear more complex than once thought even in the past decade.

These three disciplines and the complexities inherent in each academic field is perhaps the most multidisciplinary, serving to bring scientists and clinicians together with knowledge of neurobiology, immunology, pharmacology, biochemistry, cellular and molecular biology, virology, genetics, gene therapy, medicinal chemistry, nanomedicine, proteomics, pathology, and physiology. Even more than immunology and neuroscience, pharmacology integrates a broad knowledge in scientific disciplines, enabling the pharmacologist a unique perspective to tackle drug-, hormone-, immune-, and chemical-related pathways as they affect human health and behavior. Drug actions and therapeutic developments form the basis of such discoveries, but the central understanding of how they act provides vision for further research to improve human well-being and health.

This textbook is unique in scope by serving to investigate the intersection of this new discipline. Neuropharmacologists study drug actions including neurochemical disorders underlying a broad range of diseases such as psychiatric disorders (e.g., schizophrenia and depression) and neurodegenerative diseases (e.g., Alzheimer's and Parkinson's diseases). Drugs can also be used to examine neurophysiological or neurobiochemical changes as they affect brain, behavior, movement, and mental status. Immunopharmacology seeks to control the immune response in the treatment and prevention of disease. Research does include immunosuppressant agents used in organ transplant as well as developing agents that affect bone marrow function and cell differentiation in cancer therapies.

What then defines the field of *neuroimmune pharmacology*? Is it simply a field that intersects the three disciplines of neuroscience, immunology, and pharmacology in seeking to better define the epidemiology, prevention, and treatment of immune disorders of the nervous systems (Fig. 1)? Are these disorders limited in their scope in affecting behavior, cognition, motor, and sensory symptoms, or do they also involve developmental and degenerative disorders? It is clear that the immune system is linked, in whole or in part, to diseases that develop

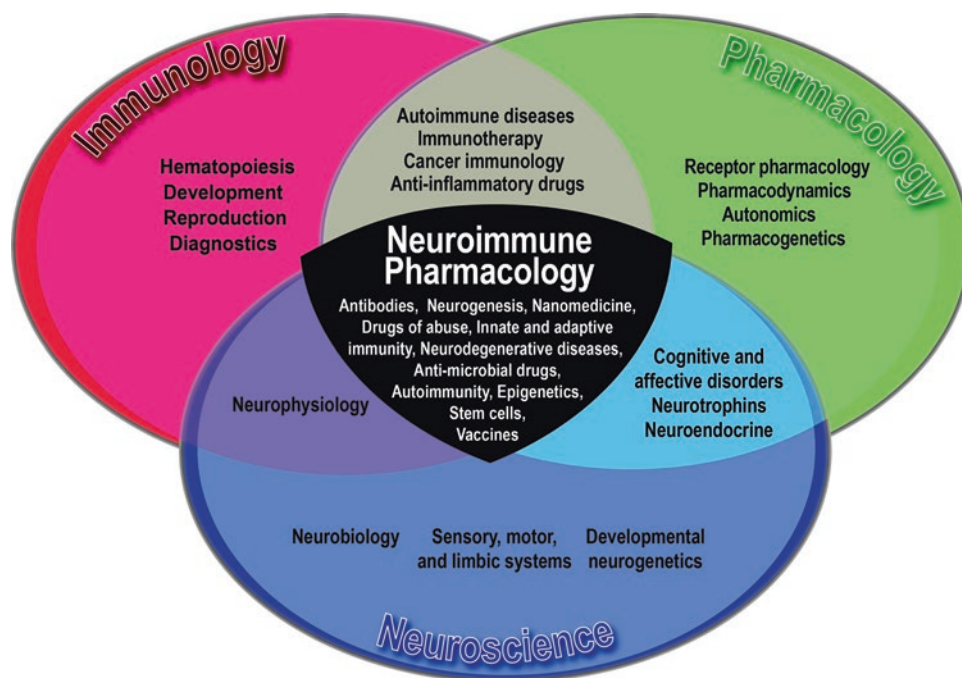


Fig. 1 This Venn diagram pictorially represents the fields of immunology, pharmacology and neuroscience with common elements overlapping into the discipline of neuroimmune pharmacology. The blackened area depicts the diseases and research areas covered in this book and integral to the field of study

as a consequence of genetic abnormalities and a broad range of environmental cues (including microbial infections and abused drugs) and to toxins. So, where does neuroimmune pharmacology find its niche? These can occur, in part, as a consequence of neuropeptides, neurotransmitters, cytokines, chemokines, and abused drugs. Like much in science, we are left with more questions than answers. In the end, we seek avenues for translational research and better understanding of disease mechanisms. Diseases are together linked to microbial agents, by inflammatory processes, by emergence of cancerous cells or tumors, by stress, by environmental cues, and by genetic disturbances. No matter the cause, harnessing the immune processes for pharmacological benefit will, at days end, provide “real” solutions to positively affect some of the most significant and feared disorders of our century.

What do we seek to accomplish by editing such a textbook? First, we would be remiss in not acknowledging the pivotal discoveries made by others when research fields intersect. These include the discovery and characterization of the guanosine triphosphate (GTP) binding and prion proteins (Gilman 1995; Rodbell 1995; Prusiner 1998, 2001), neurotransmission and memory functions (Carlsson 2001; Greengard 2001; Kandel 2001), and odorant receptors (Buck 2000; Axel 2005). We posit that new discoveries can and will be made through the intersections of neuroscience, immunology, and pharmacology and as such sought to define it for the student. The notion that inflammation contributes in significant manner to neurodegeneration and significantly beyond autoimmune diseases is brought front and center and demonstrated without ambiguity for multiple sclerosis, peripheral neuropathies, Alzheimer’s and Parkinson’s disease, and amyotrophic lateral sclerosis as well as for microglial infections of the nervous system including NeuroAIDS where microglial activation is central to disease processes (Hooten et al. 2015; Appel et al. 1995; Toyka and Gold 2003; McGeer and McGeer 2004; Ercolini and Miller 2005; Gendelman 2002; Gendelman and Mosley 2015). Perhaps most importantly, we have laid the groundwork for how the immune system can be harnessed either through its modulation, through altering blood–brain barrier integrity and function, or by drug-delivery strategies that target the brain. This second edition will add on to the success

of the first edition with additional chapters on emerging topics, including but not limited to enteric nervous system, microbiota, innate immunity signaling, exosomes, stress granules, microRNA, autism spectrum disorders, traumatic brain injury, biomarkers, macromolecular therapeutics, and “omics” pharmacology.

No doubt this textbook is an expansive read for the student and scholar alike. To this end, we are humbled by its realization and even more so during this second edition. These words lay only the beginnings to what we believe will be a significant future footprint into the integration between neuroscience, immunology, and pharmacology.

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Acknowledgments

These comprehensive volumes would not have been possible without the tireless, dedicated, and determined effort of one very special person, Reed Felderman. Reed is skillful, resourceful, and enthusiastic. Any task offered was never too minor and never left unattended. The linkages between editors, authors, administrators, and publishers flowed seamlessly. He quickly rose to the task of managing editor with aplomb, and it was his attention to detail that was singularly responsible for the success of this text.

Gregory Baer and his staff at Springer USA are already trusted friends, colleagues, and publishing visionaries. Their leadership together with Ms. Robin Taylor and our administrative staff at our own University of Nebraska Medical Center and Boston University School of Medicine has and continues to be a foundation for all we've done and for this textbook, for our *Journal of Neuroimmune Pharmacology* and beyond. We are lucky to have them as partners in this endeavor and so appreciate being able to continue to work with such talented individuals who are so dedicated to the pursuit of excellence.

To our leaders and visionaries, Jeffrey Gold, Harold Maurer, Donald Leuenberger, Dele Davies, and Deb Thomas, a simple thank you is clearly inadequate for all you have done to ensure the success of our mission and so many of our endeavors. To Carol Swarts, a friend, a colleague, a confidant, and a soul mate whose gifts of time and person ensure our sustained academic successes: Carol, you have no parallel. To Susan Leeman, the pioneer of molecular biology in neuroinflammation, our mentor and a friend who always navigate our neuroinflammation research. Your contribution to science is immeasurable. Thank you all for keeping the fire burning very bright for all we do. To Harriet Singer, Frani Blumkin and Jim Roubal who have been steadfast in support for so many years. Your faith in our research and unquestioned faith strengthens us each and every day.

To our chapter contributors and reviewers whose expertise, knowledge, intellect, and dedications proved invaluable in completing the tasks.

To Seiko Ikezu and Bonnie Bloch, our partners in life, life journeys, and always best friends.

To Yohei and Michiko Ikezu and Soffia Gendelman, our parents, navigators, and role models.

To Yumiko Aoyama, for her continuous support of family and friendship, a simple thank you is nearly inadequate.

To our children and grandchildren who are all our life treasures: Clark and David Ikezu and Sierra, Jason and Sacha Tobias and Adam and Jen Wolf-Gendelman and Lesley Gendelman and Emma Ehrenkranz.

We salute *the Journal of Neuroimmune Pharmacology* that provided and continues to be the source of inspiration for this work. To a special friend, Joel Alperson, for the gift of his ears and to our patients, students, mentors, and colleagues who inspire and guide us always, thank you all so much.

Tsuneya Ikezu
Howard E. Gendelman

Contents

Part I Immunology of the Nervous System

1	Innate and Adaptive Immunity in Health and Disease	3
	Howard E. Gendelman and Eliezer Masliah	
2	The Blood-Brain Barriers	5
	William A. Banks	
3	Regulation of Nervous System Function by Circumventricular Organs	25
	Emily A.E. Black, Nicole M. Cancelliere, and Alastair V. Ferguson	
4	Anterior Chamber and Retina (Structure, Function and Immunology)	39
	William Rhoades, Leila Kump, and Eyal Margalit	
5	The Vertebrate Retina	55
	Wallace B. Thoreson	
6	Hippocampus, Spatial Memory and Neuroimmunomodulation	69
	Huangui Xiong, Jingdong Zhang, and Jianuo Liu	
7	Anatomical Networks: Structure and Function of the Nervous System	81
	Eliezer Masliah	
8	Immune Sensors and Effectors of Health and Disease	93
	Manmeet K. Mamik and Christopher Power	
9	LRRK2	107
	Darcie A. Cook and Malú G. Tansey	
10	Astrocytes, Oligodendrocytes and Schwann Cells	117
	Malabendu Jana and Kalipada Pahan	
11	Overview of Mononuclear Phagocytes	141
	Mary G. Banoub and Howard E. Gendelman	
12	Macrophages, Microglia and Dendritic Cell Function	153
	James Hilaire and Howard E. Gendelman	
13	Microglial Biology and Physiology	167
	Oleg Butovsky, Charlotte Madore, and Howard Weiner	
14	Human Lymphocyte Biology and Its Application to Humanized Mice	201
	Larisa Y. Poluektova	

15 Stem Cells and Neurogenesis for Brain Development, Degeneration and Therapy	217
Justin Peer, Hainan Zhang, Hui Peng, Krysten Vance, Yunlong Huang, and Jialin C. Zheng	
16 Innate Immunity Signaling	245
Tsuneya Ikezu	
17 Cytokines and Chemokines.....	261
Yunlong Huang and Jialin Zheng	
18 Growth and Neurotrophic Factors in HIV-Associated Neurocognitive Disorders	285
Palsamy Periyasamy, Ming-Lei Guo, and Shilpa Buch	
19 RNA Binding Proteins in Health and Disease	299
Tara E. Vanderweyde and Benjamin Wolozin	
20 Exosomes and Neuroregulation	313
Denise A. Cobb and Howard E. Gendelman	
21 MicroRNA Implications in Neurodegenerative Disorders	329
Amrita Datta Chaudhuri and Sowmya V. Yelamanchili	
Part II Immunology of Neurodegenerative, Neuroinflammatory, Neuroinfectious and Neuropsychiatric Disorders	
22 Neurodegeneration.....	345
Serge Przedborski	
23 Multiple Sclerosis	355
Irene Falk and Steven Jacobson	
24 Guillain-Barré Syndrome, Chronic Inflammatory Demyelinating Polyradiculoneuropathy, and Axonal Degeneration and Regeneration.....	365
Ralf Gold and Klaus V. Toyka	
25 Guillain-Barré Syndrome and Acute Neuropathy	373
Helmar C. Lehmann and Kazim A. Sheikh	
26 Autoimmunity	395
Marco Cosentino, Natasa Kustrimovic, and Franca Marino	
27 HIV-Associated Neurocognitive Disorders	407
Howard Fox and Phillip Purnell	
28 Neuroimmunomodulation of Human T-Lymphotropic Virus Type I/II Infection	421
Akinari Yamano, Yoshihisa Yamano, and Steven Jacobson	
29 Viral Encephalitis.....	437
Clinton Jones and Eric M. Scholar	
30 Alzheimer’s Disease	451
Tsuneya Ikezu	
31 Parkinson’s Disease.....	477
John Loike, Vernice Jackson-Lewis, and Serge Przedborski	

32 Amyotrophic Lateral Sclerosis	493
Ericka P. Simpson and Stanley H. Appel	
33 Huntington's Disease	505
Adam Labadorf, Andrew G. Hoss, and Richard H. Myers	
34 Prion Diseases	519
Qingzhong Kong and Richard A. Bessen	
35 Glaucoma	533
Shane J. Havens, Deepta A. Ghate, and Vikas Gulati	
36 Ocular Manifestations of Systemic Autoimmune Diseases	553
Aniruddha Agarwal, Yasir J. Sepah, and Quan Dong Nguyen	
37 Neurogenesis and Brain Repair	575
Tomomi Kiyota	
38 Chronic Traumatic Encephalopathy	599
Anumantha Kanthasamy, Vellareddy Anantharam, Huajun Jin, Shivani Ghaisas, Gary Zenitsky, and Arthi Kanthasamy	
39 The Neuroimmune System in Psychiatric Disorders	621
Jonna M. Leyrer-Jackson, Gregory K. DeKrey, and Mark P. Thomas	
40 Autism Spectrum Disorders	643
Theoharis C. Theoharides and Irene Tsilioni	
41 Drugs of Abuse	661
Toby K. Eisenstein and Thomas J. Rogers	
Part III Therapies and Diagnostics	
42 Therapeutic Strategies in Neurodegenerative Diseases	681
Kristi M. Anderson and R. Lee Mosley	
43 Immunomodulatory Therapy for Multiple Sclerosis	713
Irene Cortese and Avindra Nath	
44 Therapeutic Considerations in HIV-Associated Neurocognitive Disorders	737
Stephanie A. Cross and Dennis L. Kolson	
45 Immunotherapy for Alzheimer's Disease	753
Tsuneya Ikezu	
46 Immunotherapies for Movement Disorders: Parkinson's Disease and Amyotrophic Lateral Sclerosis	767
Charles Schutt, Howard E. Gendelman, and R. Lee Mosley	
47 Regenerative and Repair Strategies for the Central Nervous System	799
Donald S. Sakaguchi	
48 New Generation of Adjuvants for Protection Against Disease and to Combat Bioterrorism	819
Sam D. Sanderson, Joseph A. Vetro, and Bala Vamsi Krishna Karuturi	
49 Medicinal Chemistry and Brain Drug Penetrance	831
James Hilaire and Howard E. Gendelman	
50 Polymer Nanomaterials for Drug Delivery Across the Blood Brain Barrier	847
Alexander V. Kabanov and Elena V. Batrakov	

51	Macromolecular Therapeutics: Development and Delivery Engineering	869
	Gang Zhao, Xin Wei, and Dong Wang	
52	Application to Gene Therapy and Vaccination	885
	Xiaomin Su, William J. Bowers, Michelle C. Janelsins, and Howard J. Federoff	
53	Introduction to Imaging in the Neurosciences	907
	Michael D. Boska and Matthew L. White	
54	Proteomics and Genomics in Neuroimmunological Disorders	941
	Maire Rose Donnelly, Wojciech Rozek, and Pawel S. Ciborowski	
55	Pharmacogenomics of Neurodegenerative Diseases: Roles in Personalized Medicines	959
	Ruby E. Evande, Rinku Dutta, Chalet Tan, Jean L. Grem, and Ram I. Mahato	
56	Control of Neuroinflammation for Therapeutic Gain	971
	Howard E. Gendelman and Eric J. Benner	
	Glossary	979
	Index.....	1007

Abbreviations

¹ H	Proton
¹ H-MRSI	Proton magnetic resonance spectroscopic imaging
2D SDS-PAGE	Two-dimensional polyacrylamide gel electrophoresis
3'UTR	3'-Untranslated region
3HK	3-Hydroxykynurenine
5-ASA	5-Aminosalicylic acid
5-HIAA	5-Hydroxyindole acetic acid
5-HT	5-Hydroxy tryptophan
5-HT	5-Hydroxytryptamine
6-OHDA	6-Hydroxydopamine
8-OHdG	8-Hydroxy-2'-deoxyguanosine
γc	Common γ-chain
AAAD	Aromatic L-amino acid decarboxylase
AAV	Adeno-associated virus
Ab	Antibody
Aβ	Amyloid-β
ABD	Adamantiades-Behçet's disease
ABP	Actin-binding protein
AC	Anterior chamber
ACAID	Anterior chamber-associated immune deviation
αCamKII	α-Calcium/calmodulin-dependent protein kinase II
ACh	Acetylcholine
AChE	Acetylcholinesterase
ACTH	Adrenocorticotrophic hormone (pro-opiomelanocortin; POMC)
AD	Alzheimer's disease
ADHD	Attention deficit hypersensitivity disorder
ADI	Acceleration/deceleration injury
ADAM	A disintegrin and metalloprotease
ADCC	Antibody-dependent cellular cytotoxicity
ADEM	Acute disseminated encephalomyelitis
AF	Activation factor
Ag	Antigens
AGE	Advanced glycation end products
AGM	Aorta-gonad mesonephros
AHSCT	Autologous hematopoietic stem cell transplantation
AICA	Anterior inferior cerebellar artery
AID	Activation-induced (cytidine) deaminase
AIDP	Acute inflammatory demyelinating polyradiculoneuropathy
AIDS	Acquired immunodeficiency syndrome
AIF	Apoptosis inducing factor
AIR	Autoimmune retinopathy
AIRE	Autoimmune regulator

AIS	Anterior chamber-associated immune deviation (ACAID)-inducing signal
ALR	AIM2-like receptors
ALS	Amyotrophic lateral sclerosis
AMAN	Acute motor axonal neuropathy
AML	Acute myeloid leukemia
AMN	Adrenomyeloneuropathy
AMPA	α -Amino-3-hydroxy-5-methylisoxazole-4-propionic acid
AMSAN	Acute motor-sensory axonal neuropathy
ANG	Angiotensin II
ANI	Asymptomatic neurocognitive impairment
ANS	Autonomic nervous system
AP	Area postrema
AP-1	Activating protein-1
APAF 1	Apoptotic peptidase activating factor 1
APC	Antigen-presenting cells
aPKC	Atypical protein kinase C
aPL	Antiphospholipid antibody
APO1	Apoptosis antigen 1 (Fas/CD95)
apoE	Apolipoprotein E
APP	Amyloid precursor protein
APPs	Acute phase proteins
AQP4	Aquaporin-4
ARE	AU-rich response element
ARMD	Age-related macular degeneration
ART	Antiretroviral therapy
ASD	Autism spectrum disorders
ASL	Arterial spin-labeled
ASTIN	Acute stroke therapy by inhibition of neutrophils
AT	Adoptive transfer
ATL	Adult T cell leukemia
ATON	Atacicept in optic neuritis
ATP	Adenosine triphosphate
AVE	Anterior visceral endoderm
AVG	Anti-viral granule
AVP	Arginine vasopressin
AV3V	Anteroventral third ventricular
AZT	3'-Azidothymidine/zidovudine
BACE	Beta-site amyloid precursor protein cleaving enzyme
BBB	Blood-brain barrier
BBMEC	Bovine brain microvessel endothelial cells
BCR	B-cell receptor
BCRP	Breast cancer resistance protein
BCSFB	Blood-CSF barrier
BD	Bipolar disorder
BDNF	Brain-derived neurotrophic factor
bFGF	Basic fibroblast growth factor (FGF2)
bHLH	Basic helix-loop-helix
BHV-1	Bovine herpesvirus-1
BLBP	Brain lipid binding protein
BLIMP-1	B lymphocyte-induced maturation protein-1
BLV	Bovine leukemia virus
BM	Bone marrow
BMDM	Bone marrow-derived macrophages

BMP	Bone morphogenetic protein
BMVEC	Brain microvascular endothelial cell
BP	Biological processes
BrdU	Bromodeoxyuridine
BRMs	Biological response modifiers
BSE	Bovine spongiform encephalopathy
BTLA	B and T lymphocyte attenuator
C/EBP	CCAAT box/enhancer binding protein
C3d	Complement C3 fragment d
C4d	Complement C4 fragment d
CA	Cornu ammonis
CA II	Carbonic anhydrase II
CaMKII	Calcium/calmodulin-dependent protein kinase II
CAMs	Cell adhesion molecules
CAPS	Cryopyrin-associated periodic syndromes
CAR	Cancer-associated retinopathy
CARD	Caspase recruitment domain
CARD15	NOD2/caspase recruitment domain 15
CB	Cannabinoid
CB1	Cannabinoid receptor 1
CB2	Cannabinoid receptor 2
CBA	Cytokine bead arrays
CBF	Cerebral blood flow
CBF1	C promoter binding factor 1
CBP	cAMP-response element binding protein (CREB)-binding protein
CBV	Cerebral blood volume
CCI	Controlled cortical impact
CCK	Cholecystokinin
CCT	Central corneal thickness
CD	Cluster of differentiation
CD11b	Complement component 3 receptor 3 subunit (integrin alpha M; ITGAM)
CD40L	CD40 ligand (TNFSF5)
CDP	Common DC progenitor
CDR	Complementarity-determining region
CDV	Canine distemper virus
CEP	Carboxyethylpyrrole
CFT	2- β -Carbomethoxy-3 β -(4-fluorophenyl) tropane
cGMP	Cyclic guanosine 5'-monophosphate
CGRP	Calcitonin gene-related peptide
CHAT	Choline acetyltransferase
CHN	Congenital hypomyelinating neuropathy
Cho	Choline
CIDP	Chronic inflammatory demyelinating polyradiculoneuropathy
CINC1	Cytokine-induced neutrophil chemoattractant-1
CIS	Clinically isolated syndrome
CJD	Creutzfeldt-Jakob disease
CLA	Cutaneous leukocyte antigen
CLL	Chronic lymphocytic leukemia
CLP	Common lymphoid progenitor
CMAP	Compound motor action potential
CMC	Critical micelle concentration
CMP	Common myeloid precursor
CMT	Charcot-Marie-Tooth (disease)

CMV	Cytomegalovirus
CNG	cGMP-gated
CNPase	Cyclic nucleotide 3' phosphohydrolase
CNS	Central nervous system
CNTF	Ciliary neurotrophic factor
CO	Cytochrome oxidase
COMT	Catechol- <i>O</i> -methyltransferase
Con A	Concanavalin A
COP-1	Copolymer-1
COX	Cyclooxygenase (prostaglandin-endoperoxide synthase; PTGS)
CP	Choroid plexus
CR	Complement receptor
CRalBP	Cellular retinal binding protein (retinaldehyde binding protein 1; RLBP1)
CRD	Carbohydrate-recognition domains
CRE	cAMP-responsive element
Cre	Creatine
CREB	cAMP-response element binding protein
CRH	Corticotrophin-releasing hormone
CRID	Cytokine release inhibitory drugs
CRP	C-reactive protein
CRPM	Collapsing response mediator protein
CRVO	Central retinal vein occlusion
CSF	Cerebrospinal fluid
CSF-1R	Colony-stimulating factor receptor
CSPG	Chondroitin sulfate proteoglycans
CT	Computed tomography
CTA	Computed tomographic angiography
CTE	Chronic traumatic encephalopathy
cTEC	Cortical thymic epithelial cell (TEC)
CTL	Cytotoxic T lymphocyte
CTLA-4	Cytotoxic T lymphocyte antigen-4
CTP	Circulating T-cell progenitors
CVF	Cobra venom factor
CVO	Circumventricular organ
Cx	Connexin
cyto c	Cytochrome c
D1R	Type-1 family of dopamine receptors
D9-THC	D9-tetrahydrocannabinol
DA	Daniel's strain of Theiler's virus
DA	Dopamine
DAF	Decay-accelerating factor
DAG	Diacylglycerol
DAMP	Danger-associated molecular patterns
DAT	Dopamine transporter (solute carrier family 6A3; SLC6A3)
D β H	Dopamine- β -hydroxylase
DC	Dendritic cells
DCX	Doublecortin
ddC	Dideoxycytidine
ddI	Dideoxyinosine
DG	Dentate gyrus
dGTP	Deoxyguanosine triphosphate
DHA	Docosahexaenoic acid
Dhh	Desert hedgehog

DHP	Dihydropyridine
DIGE	Difference gel electrophoresis
DIRA	Deficiency of IL-1 receptor antagonist
DISC	Death-inducing signaling complex
DM	Diabetes mellitus
DN	Double/dominant negative
DNA	Deoxyribonucleic acids
Doc2	Double C2 protein
DOR	Delta-opioid receptor
Dox	Doxorubicin
Dox	Doxycycline
DR	Dopamine receptors
DRPLA	Dentatorubral-pallidoluyian atrophy
DSI	Depolarization induced suppression of inhibition
DSPN	Distal sensory peripheral neuropathy
DSS	Dejerine-Sottas syndrome
DTH	Delayed-type hypersensitivity
DTI	Diffusion tensor imaging
DTR	Diphtheria toxin receptor
DWI	Diffusion weighted imaging
E2F	Early-region-2 transcription factor
EAE	Experimental allergic/autoimmune encephalomyelitis
EAN	Experimental allergic/autoimmune neuritis
EAU	Experimental autoimmune uveitis
EBV	Epstein-Barr virus
ECT	Electroconvulsive therapy
EDSS	Expanded disability status scale
EEG	Electroencephalography
EGC	Embryonic germ cell
EGF	Epidermal growth factor
EGFR	Epidermal growth factor receptor
EIAV	Equine infectious anemia virus
ELAVL4	Embryonic lethal, abnormal vision, Drosophila-like 4
ELP	Early lymphoid progenitors
eLTP	Early long-term potentiation
ELVIS	Extravasation through Leaky Vasculature and subsequent Inflammatory cell-mediated Sequestration
EMG	Electromyography
EMP	Erythromyeloid progenitors
EndoG	Endonuclease G
EOAD	Early-onset Alzheimer's disease
EPI	Echo-planar imaging
EPR	Enhanced permeability and retention
EPS	Extrapyramidal symptoms
EPSC	Excitatory postsynaptic currents
EPSP	Excitatory postsynaptic potential
ER	Endoplasmic reticulum
ERK2	Extracellular signal-regulated kinase 2
ESC	Embryonic stem cell
ESCRT	Endosomal-sorting complex required for transport
ESI-MS/MS	Electrospray ionization-mass spectrometry/mass spectrometry
ET	Endothelin
ETP	Early T lineage progenitor

EV	Extracellular vesicles
FA	Fractional anisotropy
FAD	Familial Alzheimer's disease
fALS	Familial amyotrophic lateral sclerosis
FasL	Fas ligand (FASLG)
FcγR-1	Fc receptor IgG, high affinity-1
fCJD	Familial Creutzfeldt-Jakob disease
FcR	Fc receptor
FDA	Food and Drug Administration
FDC	Follicular dendritic cell
FDG	Fluorodeoxyglucose
FDOPA	6-[(18)F]fluoro-L-dopa
FFI	Fatal familial insomnia
FGF	Fibroblast growth factor
FID	Free induction decay
FIRE	Febrile infection-related epilepsy syndrome
FIV	Feline immunodeficiency virus
FKN	Fractalkine (CX3CL1)
fMRI	Functional magnetic resonance imaging
Foxp3	Forkhead box P3 transcription factor
FPI	Fluid percussion injury
FR	Folate receptor
FRC	Fibroblastic reticular cell
FRS2	Fibroblast growth factor receptor substrate 2
FS	Fisher syndrome
FSH	Follicle-stimulating hormone
FTD	Frontotemporal dementia
FT-ICR	Fourier transformed ion cyclotron resonance mass spectrometry
FTLD	Frontotemporal lobar dementia
FUS	Fused in sarcoma
Fz/PCP	Frizzled/planar cell polarity
GA	Glatiramer acetate
GABA	Gamma-aminobutyric acid
GAD	Glutamate decarboxylase
GalC	Galactocerebroside
GALT	Gut-associated lymphoid tissue
GAP	GTPase activating protein
GAPDH	Glyceraldehyde 3 phosphate dehydrogenase
GBM	Glioblastoma multiforme
GBS	Guillain-Barré syndrome
GC	Germinal center
GCDC	Germinal center dendritic cell
GCL	Ganglion cell layer
GEF	Guanine nucleotide exchange factor
GL	Granular cell layer
GC-MS	Gas chromatography combined with mass spectrometry
G-CSF	Granulocyte colony-stimulating factor
Gd	Gadolinium
GDF	Growth and differentiation factor
GDNF	Glial-derived neurotrophic factor
GDP	Guanosine diphosphate
GEF	GDP-GTP exchange factor
GFAP	Glial fibrillary acidic protein

GFP	Green fluorescent protein
GI	Gastrointestinal
GKAP	Guanylate kinase-associated protein
GLC1A	Chromosome 1 open-angle glaucoma gene
Gln	Glutamine
GLP-1	Glucagon-like peptide-1
GM-CSF	Granulocyte-macrophage colony-stimulating factor
GMP	Granulocyte monocyte precursor
GnRH	Gonadotropin-releasing hormone
GO	Gene ontology
GO	Graves' ophthalmopathy
GPCR	G-protein-coupled receptor
GPI	Glycosylphosphatidylinositol
GR	Glucocorticoid receptor
GRIP	Glucocorticoid receptor-interacting protein
GRO- α	Growth-related oncogene alpha
GSH	Glutathione
GSS	Gerstmann-Straussler-Scheinker disease
GSTO1	Glutathione <i>s</i> -transferase omega-1
GT	G-protein transducing
GTP	Guanosine triphosphate
GUCY	Guanylate cyclase
GWAS	Meta-genome-wide association study
HAART	Highly active antiretroviral therapy
HAD	HIV-associated dementia
HAM/TSP	HTLV-I associated myelopathy/tropical spastic paraparesis
HAND	HIV-associated neurocognitive disorder
HAT	Histone acetyltransferase
HAV	Hepatitis A virus
HBsAg	Hepatitis B surface antigen
HCV	Hepatitis C virus
HCMV	Human cytomegalovirus
HD	Huntington's disease
HDAC	Histone deacetylase
HDLS	Hereditary diffuse leukoencephalopathy with spheroids
HES	Hairy and enhancer of split homolog
HEV	High endothelial venule
HFS	High frequency stimulation
Hh	Hedgehog
HHH	Hypervolemic-hemodilution and hypertensive
HHV-6	Human herpes virus-6
HIV-1	Human immunodeficiency virus type 1
HIVE	Human immunodeficiency virus encephalitis
HLA	Human leukocyte antigen
HLH	Helix-loop-helix
HMG-CoA	3-Hydroxy-3-methylglutaryl coenzyme A
HNE	4-Hydroxy-2-nonenal
hnRNP-A1	Heterogeneous nuclear ribonuclear protein-A1
HPA	Hypothalamic-pituitary-adrenal
HPMA	<i>N</i> -(2-Hydroxypropyl) methacrylamide
HRP	Horseradish peroxidase
HSC	Hematopoietic stem cell
HSC70	Heat shock cognate protein 70

Hsp	Heat shock protein
HSV	Herpes simplex virus
HSVE	Herpes simplex virus-mediated encephalitis
HT	Huntington's disease
HTLV	Human T-cell lymphotropic virus type
HTRA2	High temperature requirement serine protease 2
Htt	Huntingtin
HUVEC	Human umbilical vein endothelial cells
HveA	Herpesvirus entry mediator A
I-1	Regulatory protein inhibitor-1
IBS	Inflammatory bowel syndrome
ICAM	Intracellular adhesion molecule
ICAT	Isotope-coded affinity tags
ICE	IL-1 β -converting enzyme
ICGA	Indocyanine green angiography
ICH	Intracerebral hemorrhage
iCJD	Iatrogenic Creutzfeldt-Jakob disease
ICOS	Inducible co-stimulatory molecule
ICV	Intra-cerebro-ventricular
Id	Inhibitor of differentiation
IDE	Insulin degrading enzyme
IDO	Indoleamine 2,3-dioxygenase
IE	Immediate early
IF	Intermediate filament
IFN	Interferon
Ig	Immunoglobulin
IGF	Insulin-like growth factor
IgG	Immunoglobulin G
IGIV	Immunoglobulin intravenous therapy
Ihh	Indian hedgehog
IIDD	Idiopathic inflammatory demyelinating disease
I κ B	Inhibitory kappa B
IKK	I κ B kinase
IL	Interleukin
IL1RA	IL-1 receptor antagonist
ILBD	Incidental Lewy body disease
ILK	Integrin-linked kinase
ILM	Inner limiting membrane
ILV	Intraluminal vesicles
IM	Intramuscular
IMAC	Immobilized metal affinity chromatography
IMPDH	Inosine monophosphate dehydrogenase
iNKR	Inhibitory natural killer cell receptor
INL	Inner nuclear layer
INO	Internuclear ophthalmoplegia
iNOS	Inducible nitric oxide synthase (NOS2A)
IOP	Intraocular pressure
IP	Interferon-inducible protein
IPC	Intermediate progenitor cells
InsP3R	Inositol 1,4,5-triphosphate receptor
IPL	Inner plexiform layer
IRAK	IL-1 receptor-associated protein kinase
IRBP	Interphotoreceptor retinoid-binding protein

IRF	Interferon regulatory factor
ISCOM	Immunostimulating complexes
ISH	In situ hybridization
IT15	Interesting transcript 15
ITAM	Immunoreceptor tyrosine-based activation motif
ITGAM	Integrin, alpha M
ITR	Inverted terminal repeat
IVDU	Intravenous drug use
IVIg	Intravenous immunoglobulin
JAK	Janus kinase
JEV	Japanese encephalitis virus
JNK	c-Jun N-terminal kinase
KLH	Keyhole limpet hemocyanin
KO	Knockout
KOR	Kappa opioid peptide receptor
KYN	Kynurenine
KYNA	Kynurenic acid
LAK	Lymphokine-activated killer (cell)
L-AP4	L-2-Amino-4-phosphonobutyric acid
LAT	Latency-associated transcript
LAMP	Lysosome-associated membrane protein
LB	Lewy bodies
LC	Locus coeruleus
LCA	Leukocyte common antigen
LC-FTICR MS	Liquid chromatography Fourier transform ion cyclotron resonance mass
LC-MS	Liquid chromatography combined with mass spectrometry
LC-UV-SPE-NMR	Liquid chromatography, UV detection, solid phase extraction, and nuclear magnetic resonance
LD	Linkage disequilibrium
LDL	Low density lipoprotein
LFA-1	Leukocyte function-associated antigen-1 (integrin beta 2; ITGB2)
LGN	Lateral geniculate nucleus
LH	Luteinizing hormone
LIF	Leukemia inhibitory factor
Lingo-1	Leucine-rich repeat and Ig domain containing Nogo receptor-interacting protein-1
LMN	Lower motor neuron
LOAD	Late-onset Alzheimer's disease
LOS	Lipooligosaccharide
LPA	Lysophosphatidic acid
LPBN	Lateral parabrachial nucleus of the pons
LPS	Lipopolysaccharide
LRP-1	Lipoprotein receptor-related protein-1
LRR	Leucine-rich repeat
LRRK2	Leucine-rich repeat kinase 2
LT	Lymphotoxins
LTD	Long-term depression
LTNP	Long-term nonprogressors
LTP	Long-term potentiation
LTR	Long-terminal repeat
LT-bR	Lymphotoxin beta receptor
MCAO	Middle cerebral arterial occlusion
MCI	Mild cognitive impairment

MG	Myasthenia gravis
MHV	Mouse hepatitis virus
mI	Myoinositol
MIF	Migration inhibitory factor
Mint1	Munc-18 interacting protein 1
MIP	Macrophage inflammatory protein
MJO	Machado-Joseph disease
MME	Membrane metalloendopeptidase (neprilysin)
MMP	Matrix metalloproteinase
MMSE	Mini-Mental State Examination
MNGC	Multinucleated giant cell
MnPO	Median preoptic nucleus
Mn-SOD	Manganese superoxide dismutase
MOAT	Multispecific organic anion transporter
MOBP	Myelin-associated/oligodendrocyte basic protein
MOG	Myelin oligodendrocyte glycoprotein
MOI	Multiplicity of infection
MOR	mu opioid receptor
MOSP	Myelin/oligodendrocyte-specific protein
MP	Mononuclear phagocytes
MPA	Mycophenolic acid
MPL	Monophosphoryl lipid A
MPO	Myeloperoxidase
MPO	Medial preoptic area
MAC	Membrane attack complex
MPP	Multipotent progenitors
MPP+	1-Methyl-4-phenylpyridinium
MAdCAM-1	Mucosal addressin cell adhesion molecule-1
MAG	Myelin-associated glycoprotein
MAML	Mammalian mastermind-like
MAO	Monoamino-oxidase
MAP	Microtubule-associated protein
MAPK	Mitogen-activated protein kinases
Mash1	Mammalian achaete-scute homologue 1
MBGI	Myelin-based growth inhibitor
MBP	Myelin basic protein
MC	Mast cell
MC-1R	Melanocortin-1 receptor
MCMD	Minor cognitive motor disorder
MCP	Membrane cofactor protein (CD46)
MCP-1	Monocyte chemoattractant protein-1 (CCL2)
M-CSF	Macrophage colony-stimulating factor (CSF1)
MD	Major depression
MDA	Malondialdehyde
MDD	Major depressive disorder
MDM	Monocyte-derived macrophages
MDP	Muramyl-dipeptide
MDP	Monocytes and dendritic cell progenitor
MDR	Multidrug resistant
MDSC	Myeloid-derived suppressor cells
ME	Median eminence
MEG	Magnetoencephalography
MEPP	Miniature end-plate potential

MFS	Miller Fisher syndrome
MHC	Major histocompatibility complex
MHC-II	Class II major histocompatibility complex
MHPG	Methoxy-hydroxy-phenylethanolamine
MLR	Mixed lymphocyte reaction
MLV	Murine leukemia virus
MND	Mild neurocognitive disorder
MP	Mononuclear phagocyte
MPMV	Mason-Pfizer monkey virus
MPTP	1-Methyl-4-phenyl-1,2,3,6-tetrahydropyridine
MR	Mineralocorticoid receptors
MRA	Magnetic resonance angiography
MRI	Magnetic resonance imaging
MRP	Multidrug resistance protein
MRS	Magnetic resonance spectroscopy
MRSI	Magnetic resonance spectroscopic imaging
MS	Multiple sclerosis
MSA	Multisystem atrophy
MSCs	Myelinating Schwann cells
MSH	Melanocyte-stimulating hormone
MSN	Medium spiny neuron
mSOD1	Mutant Cu ²⁺ /Zn ²⁺ superoxide dismutase 1
MSRV	Multiple sclerosis retrovirus
MT	Magnetization transfer
Mtb	Mycobacterium tuberculosis
MTI	Magnetization transfer imaging
mTEC	Medullary thymic epithelial cell
mTOR	Mammalian target of rapamycin
MTR	Magnetization transfer ratio
MUC1	Mucin type 1 glycoprotein
MuLV	Murine leukemia virus
Munc-18	Mammalian homologue of unc-18
MV	Microvesicles
MVB	Multivesicular bodies
MVE	Murray Valley encephalitis virus
MW	Molecular weight
MZ	Marginal zone
NAA	<i>N</i> -Acetyl-aspartate
NAC	<i>N</i> -Acetyl cysteine
NADPH	Nicotinamide adenine dinucleotide phosphate
NB	Nucleotide-binding domain
NCAM	Neural cell adhesion molecule
NCC	Neural crest cell
NE	Norepinephrine
NEP	Neutral endopeptidase metalloendopeptidase
NET	Norepinephrine transporter
NeuN	Neuronal nuclei
NF	Neurofilament
NF-kB	Nuclear factor-k-B
NFAT	Nuclear factor of activated T lymphocytes
NFL	Nerve fiber layer
NFT	Neurofibrillary tangles
Ng-CAM	Neuronal-glia cell adhesion molecule (L1/NILE)

NGF	Nerve growth factor
NgR	Nogo-66 receptor
NICD	Notch intracellular domain
NK	Natural killer (cells)
NKT	Natural killer T (cells)
NLR	Nod-like receptors
NMDA	<i>N</i> -methyl- <i>D</i> -aspartate
NMDAR	<i>N</i> -methyl- <i>D</i> -aspartate receptors
NMJ	Neuromuscular junction
NMO	Neuromyelitis optica
NMR	Nuclear magnetic resonance
NMSCs	Nonmyelinating Schwann cells
nNOS	Neuronal nitric oxide synthase
NNRTIs	Nonnucleoside analogue reverse transcriptase inhibitors
NO	Nitric oxide
NOD	Nucleotide oligomerization domain
NOD	Non-obese diabetic mice
NOS	Nitric oxide synthase
NOT	Nucleus of the optic tract
NP	Nanoparticle
NPC	Neural progenitor cell
NPY	Neuropeptide Y
NPZ-8	Neuropsychological Z score for 8 tests
NR	Nuclear receptor
NRG	Neuregulin
NRL	Nuclear receptor ligand
NRTI	Nucleoside analogue reverse transcriptase inhibitors
NSAID	Nonsteroidal anti-inflammatory drug
NSC	Neural stem cell
NSE	Neuron specific enolase
NSF	<i>N</i> -ethylmaleimide sensitive factor
NT	3-Nitrotyrosine
NTF	Neurotrophin
NVU	Neurovascular unit
OB	Olfactory bulb
OCB	Oligoclonal band
OCD	Obsessive compulsive disorder
ODN	Oligonucleotides
OE	Olfactory epithelium
OHT	Ocular hypertension
OL	Oligodendrocyte
OLM	Outer limiting membrane
OMgp	Oligodendrocyte-myelin glycoprotein
OMP	Olfactory marker protein
ONH	Optic nerve head
ONL	Outer nuclear layer
OP	Oligodendrocyte progenitors
OPC	Oligodendrocyte progenitor cell
OPCA	Olivopontocerebellar atrophy
OPL	Outer plexiform layer
ORF	Open reading frame
ORN	Olfactory response neuron
OSP	Oligodendrocyte-specific protein

OVA	Ovalbumin
OVLT	Organum vasculosum of the lamina terminalis
PO	Myelin protein zero
p75NTR	p75 neurotrophin receptor (nerve growth factor receptor; NGFR)
PACAP	Pituitary adenylate cyclase-activating polypeptide
PACT	Protein activator of the interferon-induced protein kinase
PAF	Platelet-activating factor
PAG	Periaqueductal gray
PAMP	Pathogen-associated molecular pattern
PANDAS	Pediatric autoimmune neuropsychiatric disorders associated with Streptococcus
PANSS	Positive and negative syndrome scale
PARP	Poly(ADP-ribose) polymerase
PASAT	Paced Auditory Serial Addition Test
PBBS	Peripheral benzodiazepine binding sites
PBL	Peripheral blood lymphocyte
PBMC	Peripheral blood mononuclear cell
PBR	Peripheral benzodiazepine receptor
PCP	Phencyclidine
PCR	Polymerase chain reaction
PD	Parkinson's disease
pDC	Plasmacytoid dendritic cells
PD1	Program death-1
PDE	Phosphodiesterase
PDGF	Platelet-derived growth factor
PDTC	Pyrrolidine dithiocarbamate
PE	Plasma exchange
Pe	Periventricular
PEG	Polyethylene glycol
PEI	Polyethyleneimine
PENK	Proenkephalin
PERG	Pattern electroretinogram
PET	Positron emission tomography
PFS	Periodic fever syndromes
PG	Prostaglandin
Pgp	P-glycoprotein
PHF	Paired helical filament
PI	Phosphatidylinositol
PI3K	Phosphatidylinositol-3-kinase
PICA	Posterior inferior cerebellar artery
PICK1	Protein interacting with C kinase 1
PKA	cAMP-dependent protein kinase
PKG	Protein kinase G
PLGA	Poly(D,L-lactide-co-glycolide)
PLOSL	Polycystic lipomembranous osteodysplasia with sclerosing leukoencephalopathy
PLP	Proteolipid protein
PMCA	Plasma membrane bound Ca ²⁺ -ATPase
PMD	Pelizaeus-Merzbacher disease
PML	Progressive multifocal leukoencephalopathy
PMN	Polymorphonuclear (leukocyte)
PMP22	Peripheral myelin protein 22
PNS	Peripheral nervous system
POAG	Primary open-angle glaucoma
polyQ	Polyglutamine

POMC	Pro-opiomelanocortin
POU3F2	POU class 3 homeobox 2
PP	Protein phosphatase
PPAR	Peroxisome proliferator activated receptor
PPF	Paired-pulse facilitation
PPG	Poly(propylene glycol)
PP-MS	Primary progressive multiple sclerosis
PR	Photoreceptor
PrPc	Cellular prion protein
PrPres	Protease resistance prion
PrPsc	Disease-associated prion protein
PRR	Pattern recognition receptor
PS	Presenilin (PSEN)
PSA-NCAM	Poly-sialylated form of the neural cell adhesion molecule
PSCs	Perisynaptic Schwann cells
PSD	Postsynaptic density
PSP	Progressive supranuclear palsy
PSW	Periodic sharp wave
Ptc	Patched, a hedgehog receptor
PTM	Post-translational modifications
PTP	Post-tetanic potentiation
PTSD	Post-traumatic stress disorder
PTZ	Pentylentetrazol
PVL	Periventricular leukomalacia
PVM	Perivascular macrophages
PVN	Paraventricular nucleus
PYD	Pyrin domain
PYY	Peptide YY
RA	Rheumatoid arthritis
Rag	Recombination-activating gene
RAGE	Receptor for advanced glycation end product
RANTES	Regulated upon activation normal T-cell expressed and secreted (CCL5)
RAS	Renin-angiotensin system
Rb	Retinoblastoma
RBP	RNA-binding proteins
REM	Rapid eye movement
RER	Rough endoplasmic reticulum
RF	Radiofrequency
RFLPs	Restriction fragment length polymorphisms
RGC	Retinal ganglion cell
RISC	RNA-induced silencing complex
RIG1	Retinoic acid-inducible gene 1
RIM	Rab3-interacting molecule
RIP	Receptor-interacting protein
RMS	Rostral migratory stream
RNAi	RNA interference
RNI	Reactive nitrogen intermediates
RNP	Ribonucleoprotein
RNS	Reactive nitrogen species
ROCK	Rho kinase
ROHHAD	Rapid-onset obesity, hypoventilation, hypothalamic dysfunction, and autonomic dysregulation
ROI	Reactive oxygen intermediate

ROR γ	Retinoic-acid-receptor-related orphan receptor- γ
ROS	Reactive oxygen species
RP	Relapsing polychondritis
RPE	Retinal pigment epithelial (cells)
RRM	RNA recognition motifs
RRMS	Relapsing and remitting multiple sclerosis
RSV	Rous sarcoma virus
RTK	Receptor tyrosine kinase
rt-PA	Recombinant tissue plasminogen activator
RT-PCR	Reverse transcription polymerase chain reaction
RyR	Ryanodine receptor
sALS	Sporadic amyotrophic lateral sclerosis
SAP	Synapse-associated protein
SAPAP	SAP-associated protein (discs, large homolog-associated protein-1; DLGAP1)
SAPK	Stress-activated protein kinase (JNK, MAPK8)
sAPP	Secreted β -amyloid precursor protein
SBMA	Spinobulbar muscular atrophy
SCs	Schwann cells
SC	Superior colliculus
SCA-3	Spinocerebellar ataxia-3
scFv	Single-chain Fv antibodies
SCI	Spinal cord injury
SCID	Severe combined immunodeficiency
sCJD	Sporadic Creutzfeldt-Jakob disease
SCPs	Schwann cell precursor
sCrry	Soluble complement receptor-related protein y
SDF-1	Stromal cell-derived factor 1 (CXCL12)
SEC	Sinus endothelial cell
SELDI-TOF	Surface enhanced laser desorption ionization time-of-flight
SER	Smooth endoplasmic reticulum
SERCA	Sarco(endo)plasmic reticulum Ca ²⁺ -ATPase
SERT	Serotonin transporter
sFI	Sporadic fatal insomnia
SFO	Subfornical organ
SG	Stress granule
SGLPG	Sulfated glucuronyl lactosaminyl paragloboside
SGZ	Subgranular zone
Shh	Sonic hedgehog
sIg	Surface immunoglobulin
sIL-2R	Soluble IL-2 receptor
SITA	Swedish interactive thresholding algorithm
SIV	Simian immunodeficiency virus
SIVE	Simian immunodeficiency virus encephalitis
SLE	Systemic lupus erythematosus
SMA	Spinal muscular atrophy
SMAC	Second mitochondrial-derived activator of caspase
SMase	Sphingomyelinase
SMN	Survival motor neuron gene
SN	Substantia nigra
SNAP	Sensory nerve action potential
SNAP-25	Synaptosome-associated protein of 25,000 daltons
SNARE	NSF attachment receptor
SNpc	Substantia nigra pars compacta

SNPs	Single-nucleotide polymorphisms
SNS	Sympathetic nervous system
SOCS	Suppressors of cytokine signaling
SOD1	Superoxide dismutase 1
SON	Supraoptic nuclei
SP1	Specificity protein 1
SPARC	Secreted protein acidic and rich in cysteine
SPECT	Single photon emission computed tomography
SPG-II	Spastic paraplegia type II
SR-A	Scavenger receptor type A
SRBCs	Sheep red blood cells
SREBP	Sterol regulatory element-binding protein
SRF	Serum response factor
SSRI	Selective serotonin reuptake inhibitors
STAT	Signal transducers and activators of transcription
STP	Short-term potentiation
SV5	Simian virus 5
SVZ	Subventricular zone
SWAP	Short wavelength automated perimetry
SWATH	Sequential window acquisition of all theoretical mass spectra
SWI	Susceptibility weighted imaging
SYN	α -Synuclein
SZ	Schizophrenia
T	Tesla
Tat	Trans-activator of transcription
TBE	Tick-borne encephalitis virus
TBI	Traumatic brain injury
TBP	TATA-binding protein
TCA	Tricarboxylic acid
TCR	T-cell receptor
TCV	T-cell vaccination
TDO	Tryptophan 2,3-dioxygenase
TE	Echo time
TEC	Thymic epithelial cell
TES	Traumatic encephalopathy syndrome
Teff	T effector cells
TF	Transcription factor
TG	Trigeminal ganglia
TGF	Transforming growth factor
TH	Thyroid hormone
TH	Tyrosine hydroxylase
Th1	T helper type 1 cell
Th2	T helper type 2 cell
TIA	Transient ischemic attack
TIR	Toll/IL-1 receptor
TJ	Tight junction
TLR	Toll-like receptor
TMEV	Theiler's mouse encephalomyelitis virus
TMT	Trimethyltin
TMZ	Temozolomide
TN	Terminal nuclei
TNF	Tumor necrosis factor
TNFR	Tumor necrosis factor receptor

TOR1	Target of rapamycin 1
TRAF	TNF-receptor mediated factor
TRAIL	TNF-related apoptosis-inducing ligand (TNFSF10)
TRANCE	TNF-related activation-induced cytokine (TNFSF11)
TRANCER	TRANCE receptor (TNFRSF11A)
TRE	Tax responsive element
Treg	T regulatory cells
TREM	Triggering receptor expressed on myeloid cells
Trk	Receptor tyrosine kinase
TRP	Transient receptor potential
TRPC	Transient receptor potential canonical channels
TSA	Tissue-specific antigen
TSE	Transmissible spongiform encephalopathies
TSP	Thrombospondins
TULP-1	Tubby-lie protein 1
Tyk2	Protein tyrosine kinase 2
UACA	Uveal autoantigen with coiled domains and ankyrin repeats
UMN	Upper motor neuron
V1	Primary visual cortex
VAMP	Vesicle-associated membrane protein
VCAM	Vascular cell adhesion molecule
VEGF	Vascular endothelial growth factor
VEP	Visual evoked potential
VIP	Vasoactive intestinal peptide
VMAT-2	Vesicular monoamine transporter-2 (solute carrier family 18; SLC18A2)
VZV	Varicella-zoster virus
WDM	Welander distal myopathy
WKAH	Wistar-King-Aptekman-Hokudai
WNV	West Nile virus
X-ALD	X-adrenoleukodystrophy
XIAP	X-linked inhibitor of apoptosis protein
ZO-1	Zonula occludens-1 (TJP1)

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Part I

Immunology of the Nervous System

Howard E. Gendelman and Eliezer Masliah

Abstract

Neuroinflammatory processes play a significant role in health and disease of the nervous system. These regulate development, maintenance and sustenance of brain cells and their connections. Linked to aging, epidemiologic, animal, human, and therapeutic studies all support the presence of a neuroinflammatory cascade in disease. This is highlighted by the neurotoxic potential of microglia. In steady state, microglia serve to protect the nervous system by acting as debris scavengers, killers of microbial pathogens, and regulators of innate and adaptive immune responses. In neurodegenerative diseases, activated microglia affect neuronal injury and death through production of glutamate, pro-inflammatory factors, reactive oxygen species, quinolinic acid amongst others and by mobilization of adaptive immune responses and cell chemotaxis leading to transendothelial migration of immunocytes across the blood-brain barrier and perpetuation of neural damage. As disease progresses, inflammatory secretions engage neighboring glial cells, including astrocytes and endothelial cells, resulting in a vicious cycle of autocrine and paracrine amplification of inflammation perpetuating tissue injury. Such pathogenic processes contribute to neurodegeneration. Research from others and our own laboratories seek to harness such inflammatory processes with the singular goal of developing therapeutic interventions that positively affect the tempo and progression of human disease.

Key words

Alzheimer's disease • Microglia • Neurodegenerative disorders • Neuroinflammatory processes • Parkinson's disease

Neuroinflammatory processes play a significant role in health and disease of the nervous system. These regulate development, maintenance and sustenance of brain cells and their connections. Linked to aging, epidemiologic, animal, human, and therapeutic studies all support the presence of a neuroin-

flammatory cascade in disease. This is highlighted by the neurotoxic potential of microglia. In steady state, microglia serve to protect the nervous system by acting as debris scavengers, killers of microbial pathogens, and regulators of innate and adaptive immune responses. In neurodegenerative diseases, activated microglia affect neuronal injury and death through production of glutamate, pro-inflammatory factors, reactive oxygen species, quinolinic acid amongst others and by mobilization of adaptive immune responses and cell chemotaxis leading to transendothelial migration of immunocytes across the blood-brain barrier and perpetuation of neural damage. As disease progresses, inflammatory secretions engage neighboring glial cells, including astrocytes and endothelial cells, resulting in a vicious cycle of autocrine and

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paracrine amplification of inflammation perpetuating tissue injury. Such pathogenic processes contribute to neurodegeneration. Research from others and our own laboratories seek to harness such inflammatory processes with the singular goal of developing therapeutic interventions that positively affect the tempo and progression of human disease.

As the life expectancy of the human population continues to increase, the possibility of developing neuroinflammatory and neurodegenerative diseases have increased considerably during the past 50 years. Of the neurodegenerative disorders Alzheimer's disease continued to be the leading cause of dementia in the aging population. Traditionally, neurodegenerative disorders have been defined as conditions where there is selective loss of neurons within specific region of the brain accompanied by astrogliosis. However, in the past 20 years, we have learned that the pathological process leading to the dysfunction of selected circuitries in the brain initiates with damage to the synapses rather than with the loss of neurons. In fact, neuronal loss is a late event that is probably preceded by damage to axons and dendrites followed by shrinkage of the neuronal cell body and abnormal accumulation of filamentous proteins.

Therefore, the revised concept of neurodegeneration suggests that neuronal injury initiates at the synaptic junction and propagates throughout selected circuitries leading to neuronal dysfunction which resolves in the classical clinical symptoms characteristic to each of the neurodegenerative disorders (Hashimoto and Masliah 2003). So for example in Alzheimer's disease early damage to the synapses between the entorhinal cortex and the molecular layer of the dentate gyrus (perforant pathway) resolves in the short term memory deficits characteristic of this dementing disorder. Later on disconnection of the cortico-cortico fibers in the frontal, parietal, and temporal cortex resolved in more severe memory deficits, and alterations in executive functions and abstraction. Degeneration of connections between the nucleus basalis of Meynert and the neocortex resolves in attention and memory deficits that usually associated with loss of cholinergic neurons. Other circuitries and neuronal populations are also affected in Alzheimer's disease illustrating the complexity of these disorders and the fact

that the concept of single population is affected is limited. That is the case with several other disorders including Parkinson's disease where degeneration is not limited to the dopaminergic system, but also involves the limbic system, the raphe nucleus, the insula, and other systems.

In response to the injury neurons produce adhesions molecules and trophic factors that recruit astroglial and microglial cells to participate in the process of repair of the damage. In addition the microvasculature and other glial systems might also participate in the process. Thus, neurodegeneration is accompanied by astrogliosis, microgliosis, and microvascular remodeling. While astroglial cells initially produce trophic factors and cytokines that aid in the tissue repair, eventually these factors could amplify the inflammatory response, increase vascular permeability and result in microglial activation, which in turn might lead to the production of more proinflammatory cytokines and chemokines. A critical balance between the repair and proinflammatory factors often determines the future rate and progression of the degenerative process.

The understanding of the mechanisms of neurodegeneration and inflammatory response in these neurological conditions has seen a tremendous progress in the past 10 years. It is now recognized that probably small soluble misfolded protein aggregates denominated oligomers are responsible for the injury. So for example in Alzheimer's disease A β protein oligomers might damage the synapses in neocortical regions and the limbic system while in Parkinson's disease α -synuclein oligomers may damage the axons in the striatum, brainstem and cortical regions. While significant progress has been made in understanding the fundamental mechanisms for the neuronal injury, less is known about the reasons for the selective neuronal vulnerability characteristic to these neurological conditions and the role of the innate immune system in this process.

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William A. Banks

Abstract

The blood-brain barriers (BBBs) are mainly located at the levels of the vasculature, choroid plexus, and the circumventricular organs and play multiple roles in neuroimmunology. The ability of the BBBs to separate the blood and its contents from the central nervous system (CNS) is largely responsible for the CNS being an immune-privileged region. However, the BBB then revises this separation in a regulated way by a variety of mechanisms, including the ability to transport cytokines, regulate the entry of immune cells into the brain, and to itself secrete into the blood and into the CNS immunoactive substances. The BBB thereby participates in a number of neuroimmune axes that allow communication between the CNS and the peripheral immune cells. Failure of the highly regulated activities of the BBBs can be both a cause and consequence of immune diseases.

Keywords

Active transport • Adsorptive endocytosis • Adsorptive transcytosis • AIDS • Blood-brain barrier • Brain • Brain endothelial cell • Central nervous system • Cytokine • Endothelin • Facilitated diffusion • Immune cell • Interleukin • Neuroimmune • Neurovascular unit • Opiate • Transmembrane diffusion • Transport • Tumor necrosis factor • Virus

2.1 Introduction

The roles played by the blood-brain barrier (BBB) in neuroimmunology are diverse and ever expanding. Conceptually, the BBB is those processes that restrict, control, or otherwise influence the exchange of substances between the peripheral circulation and the brain interstitial fluid and cerebrospinal fluid (CSF). More concretely, there are multiple BBBs: the barrier formed by monolayers of endothelial cells (the vascular BBB), the barrier formed by ependymal/epithelial cells (the blood-CSF barrier) and a barrier formed by tanycytes interposed between the circumventricular organs and the

adjacent brain tissue. It is the restrictive properties of the BBB that limit and control the trafficking of immune cells and prevent the unrestricted leakage of immune active substances from the blood into the brain that renders the central nervous system (CNS) an immune-privileged area. But the BBB also transports immunoactive substances between the blood and CNS, responds to immunoactive substances secreted into the blood or CNS fluids, and secretes substances into those compartments. These last three processes of transport, responsiveness, and secretion allow the BBB to be in constant cross talk with other cells in both the CNS and periphery in a formation referred to as the neurovascular unit (NVU). The BBB both responds to and influences the CNS and peripheral microenvironment and does so through reacting to immunoactive substances, including cytokines, chemokines, prostaglandins, and nitric oxide. Several consequences arise from this cross talk, including that under physiological conditions the metabolic needs of the brain are met by the BBB. The BBB helps to keep the CNS informed of

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peripheral events including immune events and is central to the formation and functioning of neuroimmune axes. When any of these physiological consequences are violated, disease can arise; and the BBB can be a target, a cause, or a conduit to treatment of those diseases.

2.1.1 Structures and Functions of the BBB

Evidence for an interface between the circulation and the CNS dates back to the end of the nineteenth century (Bradbury 1979). The best known of those early studies were done by a young Paul Erlich who found that some dyes did not stain the brain after their peripheral injection. Erlich concluded erroneously that the lack of staining was because these dyes did not bind to brain tissue. Several decades later, Goldmann, a student of Erlich's, found that these dyes could stain the brain when injected intravenously. Thus, these studies were reinterpreted as evidence in favor of some sort of barrier between the CNS and blood.

The location and nature of that barrier was controversial through much of the twentieth century. Elegant studies by Davson and colleagues identified the barrier at the vascular level. However, alternative opinions were held until Reese and coworkers conducted classic studies with the electron microscope in the late 1960s (Brightman and Reese 1969; Reese and Karnovsky 1967). Previous work had shown no difference between vascular beds of peripheral tissues and the CNS when studied grossly or at the light microscope level. However, Reese and coworkers found numerous differences at the ultrastructural level. These included a much-reduced rate of pinocytosis and an absence of intracellular fenestrations. Currently, the most widely discussed finding is the presence of tight junctions between adjacent endothelial cells. The tight junctions, low rate of pinocytosis, and low number of intracellular fenestrations effectively eliminate intercellular gaps and pores. This, in turn, essentially eliminates the production of a plasma-derived ultrafiltrate and hence the leakage of serum proteins into the brain.

From this single change, the lack of a production of an ultrafiltrate, evolves a large number of consequences for CNS function. Obviously, it is the basis of the restriction of protein access, which first defined the BBB in late nineteenth century. The need for an efficient lymphatic system is eliminated, but the lack of a classic lymphatic system means that the CNS needs other methods to rid itself of the free water and wastes produced by metabolism and the secretions of the choroid plexus. Without production of an ultrafiltrate, the CNS depends on other methods to extract nourishment from the blood. The BBB addresses this need with a large number of selective transporters for substances from electrolytes to regulatory proteins (Davson and Segal 1996b, c). Because the CNS is not equipped to handle an ultrafiltrate, the reintroduction of a

leaky BBB, as with hypertensive crisis, can result in increased intracranial pressure and encephalopathy (Al-Sarraf and Phillip 2003; Johansson 1989; Mayhan and Heistad 1985).

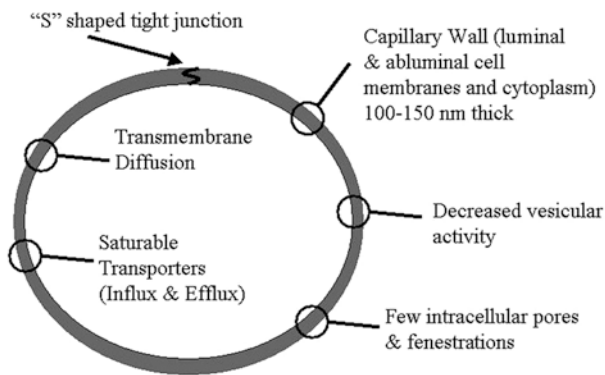
2.1.2 The Various BBBs

The BBB is not a single barrier but several barriers, which are in parallel. This contrasts with the testis-blood barrier, which consists of several barriers in series (Holash et al. 1993; Neaves 1977). The most studied of these barriers are the vascular barrier and the choroid plexus. Often, the terms BBB and blood-cerebrospinal (CSF) fluid barrier are used to refer specifically to the vascular barrier and the barrier formed at the choroid plexus, respectively. The least studied barriers are the barriers formed by tanycytes at the circumventricular organs (CVO) and other specialized neural barriers, such as the blood-retinal barrier (Neuwelt et al. 2008).

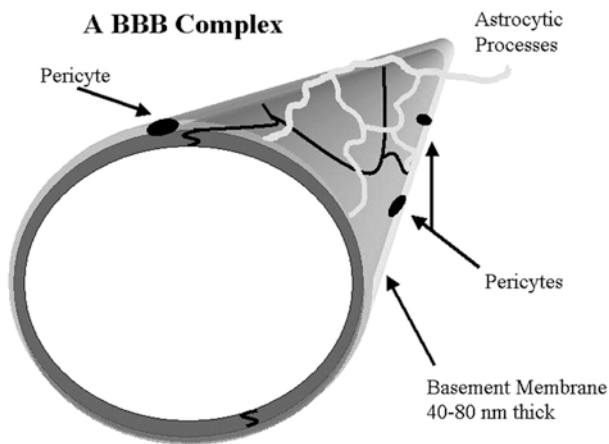
2.1.2.1 Vascular BBB

The vascular BBB occurs because of the modifications, noted by Reese and co-workers, in the endothelial cells that comprise the capillary bed and line the venules and arterioles of the CNS (upper panel Fig. 2.1). It is likely that these three regions are highly specialized. For example, immune cells primarily cross at the venules, and most of the classic transporters are located at the capillaries (Engelhardt and Wolburg 2004). No CNS cell is more than about 40 μm from a capillary. This means that a substance that can cross the vascular BBB can immediately access the entire CNS. Substances that cross the vascular BBB can be either flow-dependent or not dependent on flow rate. A flow-dependent substance is one in which the BBB extracts from the blood nearly the maximal amount possible (Kety 1987). The only way to increase the amount of a flow-dependent substance entering the brain is to increase the flow rate to the brain. Glucose is an example of a flow-dependent substance (Rapoport et al. 1981). A brain region that is particularly active has its increased demand for glucose met by an increase in regional blood flow. In contrast, transport of a cytokine such as tumor necrosis factor- α (TNF- α) is not flow dependent. Only a small percent of the TNF- α in blood is extracted by the brain via the saturable transporter for TNF- α located at the BBB (Gutierrez et al. 1993). Alterations of blood flow within physiological limits do not alter the uptake of TNF- α from blood by brain. However, extreme changes in the rate of blood flow or capillary tortuosity can result in rheological changes, such as the loss of laminar flow. Such alterations likely occur in stroke, AIDS, and Alzheimer's disease (de la Torre and Mussivand 1993; Nelson et al. 1999). This may result in impaired permeation of flow-dependent and non-flow dependent substances.

The Brain Endothelial Cell



A BBB Complex



The Neurovascular Unit

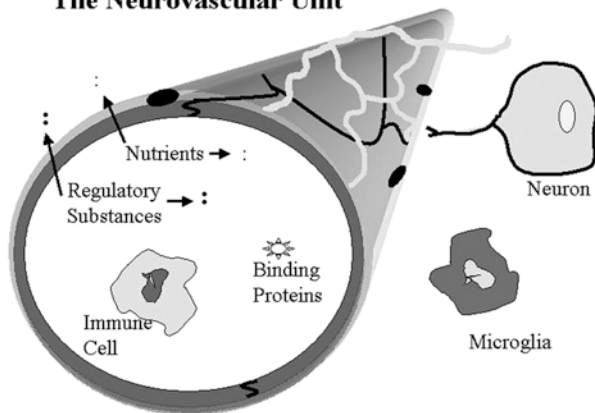


Fig. 2.1 The vascular blood-brain barrier: the *upper panel* illustrates the brain endothelial cell. This is the functional and anatomical site of both barrier function and of saturable and non-saturable mechanisms of passage. The major modifications allowing both barrier function and selective penetration of substances are indicated. The *middle panel* illustrates other cell types and structures important in BBB function. Pericytes are embedded in a basement membrane and astrocytes form a structure over the capillary bed above the pericytes and basement membrane. Both cell types are in paracellular communication with the brain endothelial cells. The *lower panel* illustrates the neurovascular unit, a concept that emphasizes integration of peripheral, BBB, and central interactions

The vascular BBB has regional variations in terms of function and susceptibilities to disease. Huber et al. noted that some regions of the brain suffer larger and earlier disruptions to their barriers during diabetes mellitus than others (Huber 2008; Huber et al. 2006). Many peptides and regulatory substances have unique regional variations in their transport rates across the BBB (Banks et al. 1996; Banks and Kastin 1998). It is assumed that these reflect on brain function. For example, the brain region with the highest rate of transport of leptin is the region of the arcuate nucleus, an area important in leptin-mediated control of feeding (Banks et al. 1996; Schwartz et al. 2000).

2.1.2.2 Choroid Plexus

The choroid plexus are bags composed of monolayers of epithelial cells that project into the ventricles and contain a capillary plexus (Johanson 1988). The capillaries do not have barrier function and so produce an ultrafiltrate that fills the bag. The epithelial cells have tight junctions and so prevent the ultrafiltrate from entering the ventricular space. Unlike the capillaries, the epithelial cells of the choroid plexus have a high rate of vesicular turnover that is responsible for the production of the CSF. However, the CSF is not an ultrafiltrate but a secreted substance. The choroid plexus also has many selective transport systems, some of which are specific to it or are enriched in comparison to the vascular BBB. In some cases, the choroid plexus seems not to be a system complementary to the vascular BBB, but one that is contrary to it. For example, efflux of amyloid beta peptide by the vascular BBB is impaired with aging but is increased with aging at the choroid plexus (Pascale et al. 2011). Thus, the choroid plexus should not be thought of as a secondary barrier but as an independent system with its own unique characteristics.

2.1.2.3 Barriers at Circumventricular Organs

The CNS of mammals contains seven regions of the brain where the vasculature does not fully participate in a BBB (Gross et al. 1987). These regions have at least one side that faces a ventricle and so are termed circumventricular organs (CVOs). Together, they comprise about 0.5% of the brain by weight. Their capillaries allow the production of an ultrafiltrate and so their cells are in more intimate contact with the circulation. They are known to play vital roles as sensing organs for critical peripheral events; for example, they act as emetic centers and are important in blood pressure modulation (Johnson and Deckwerth 1993; Ferguson 1991). They can relay their signals to the rest of the brain by neurons that project from them to distant brain regions or project to them from other brain regions. However, the mixing of their interstitial fluids with that of adjacent brain tissue or CSF has been shown to be limited in most studies (Peruzzo et al. 2000; Plotkin et al. 1996; Rethelyi 1984). Diffusion through brain tissue is poor, and this alone would tend to produce a

limit to mixing within a few hundred microns of the CVO (Cserr and Berman 1978; de Lange et al. 1995). However, the major factor preventing leakage of substances from the CVO into the adjacent CSF and brain tissue is a physical barrier to diffusion. The epithelial cells that line the ventricles form tight junctions when they are next to CVOs, thus limiting CVO-to-CSF diffusion. A functional barrier formed by bands of tanyocytes also exists for the diffusion of substances from the CVO to the adjacent brain region (Peruzzo et al. 2000; Plotkin et al. 1996; Rethelyi 1984). These tanyocytes themselves can project to the cells and vessels within the CVO and can respond dynamically to relay information to adjacent regions of the brain that have BBBs (Langlet et al. 2013). A recent review by Rodriguez (Rodriguez et al. 2010) has explored the various aspects of the CVO barriers.

2.1.2.4 Other Specialized Neural Barriers

These barriers include the blood-retinal barrier, blood-spinal cord barrier, and blood-nerve barriers (for a full listing, see Neuwelt et al. (2008)). These barriers generally include tight junctions participating in a vascular barrier but exhibit varying degrees of leakiness. They can also vary markedly in transporter activity from the vascular BBB during both health and in response to disease (Pan et al. 2008a; Pan and Kastin 2003; Prockop et al. 1996).

2.1.3 Concept of the Neurovascular Unit and Comparison to Peripheral Vascular Beds

The endothelial cell is the anatomical location of the barrier aspect of the vascular BBB and of its various saturable transporters (Fig. 2.1). Capillary beds from peripheral tissues have numerous intracellular and intercellular pores and fenestrations and high rates of pinocytosis that account for their leakiness. The brain endothelial cell engages in comparatively little macropinocytosis, has few intracellular pores or fenestrations, and intercellular pores or gaps are eliminated because of tight junctions.

However, the brain endothelial cell does not function in isolation. The abluminal (brain side) of the capillary is encased in a basement membrane 40–80 nm thick. This membrane does not act as a barrier to molecules but may restrict viral-sized particles (Muldoon et al. 1999). It also holds pericytes in close approximation to the endothelial cell (Balabanov and Dore-Duffy 1998). The pericyte is a pluripotent cell and a modulator of BBB function (Dore-Duffy et al. 2000; Deli et al. 2005; Dore-Duffy 2008). It, like astrocytes and microglia, secretes cytokines and other immune active substances both constitutively and when induced (Kovac et al. 2011). In the rat hippocampus, astrocytes project endfeet that surround the capillary in what looks at the ultrastructural level like a

complete covering, albeit without intercellular tight junctions (Mathiisen et al. 2010). Astrocytes and pericytes both secrete substances that induce tight junction formation in endothelial cells (Deli et al. 2005; Daneman et al. 2010). All the major cell types of the NVU (pericytes, astrocytes, microglia, and endothelial cells) secrete a variety of substances, including cytokines, into their local environment (Fabry et al. 1993; Nath et al. 1999; Banks 2014). Pericytes and astrocytes play interrelated but distinct roles in BBB function. For example, pericytes are the primary protectors of the blood-retinal and blood-brain barriers during glycemic stress (Romeo et al. 2002; Nakaoka et al. 2007).

It is clear that immune cell trafficking occurs across the normal BBB (Greenwood et al. 2011). One study found that about one in 5000 intravenously injected lymphocytes resided in the brain at any given time and that uptake was affected by strain and immune activation (Banks et al. 2012). The microglia may be at equilibrium with circulating macrophages/monocytes, although whether monocytes cross the BBB of the adult healthy animal to become microglia seems to still be unresolved (Williams and Hickey 1995). Other immune cells also enter and exit the CNS at unknown rates and frequencies as influenced by yet to be determined factors. Clearly, secretions of prostaglandins, nitric oxide, and cytokines from each of these cells are important for intercellular communication and can influence endothelial cell permeability (Chao et al. 1994; Nath et al. 1999; Shafer and Murphy 1997).

The concept of the neurovascular unit (NVU) emphasizes the interactive role that cells and events within the CNS and in the circulation play on BBB permeability as well as the consequences of the permeability itself. The NVU includes other factors long known to influence the penetration of substances across the BBB, such as degradation, sequestration, and serum protein binding. The encompassing concept of the NVU is particularly useful when considering the next section, the mechanisms of transport across the BBB.

2.2 Mechanisms of Transport Across the BBB

Substances can enter or exit the CNS by a variety of mechanisms. Some of these mechanisms are operational in both the blood-to-brain (influx) and the brain-to-blood (efflux) directions; whereas, others are unidirectional.

2.2.1 Blood to CNS

Saturable and nonsaturable modes predominate influx. Within each of these categories are a diverse number of mechanisms. These different mechanisms tend to favor certain groups or types of substances.

2.2.1.1 Nonsaturable Passage

A hallmark of nonsaturable passage is that the percent of material crossing into the CNS is not affected by the amount of material available for transport. The two main mechanisms of nonsaturable passage are transmembrane or transcellular diffusion and the extracellular pathways. The former is much better studied and its principles are widely applied by industry for the development of CNS drugs; the latter has received much less attention.

Transcellular Diffusion

The most studied nonsaturable mechanism by which small molecules cross the BBB is by transmembrane or transcellular diffusion (Rapoport 1976). The major determinant of passage is the degree to which the substance is lipid soluble. A substance that is too lipid soluble will be unable to repartition into the brain's interstitial fluid and so will become trapped in the cell membranes of the BBB. A ratio of about 10:1 in favor of lipid versus aqueous solubility is near ideal for maximal passage across the BBB. The second most important determinant is molecular weight with passage being favored for smaller molecules. Other physicochemical determinants, such as charge, can occasionally become dominant for specific compounds. Work by Lipinski and colleagues in Caco-2 cells, an immortalized cell line derived from a gastrointestinal cancer, clearly shows that smaller, less charged, more lipid soluble drugs are favored in transmembrane diffusion (Lipinski et al. 1997). Many exogenous substances, including many drugs with CNS activity, enter the brain predominantly by way of transmembrane diffusion. Morphine and ethanol are prime examples of common substances that cross the BBB by this mechanism (Oldendorf 1974).

Although higher molecular weight (MW) is an impediment to transmembrane diffusion at the BBB, there seems to be no absolute molecular weight cut-off. A previous study which had thought to define such an absolute limit had discovered, in retrospect, early evidence for an efflux system (Levin 1980). The largest substance to date noted to have a measurable uptake by brain by transcellular diffusion is cytokine-induced neutrophil chemoattractant-1 (CINC1), with a MW of about 7.8 kDa (Pan and Kastin 2001a). A surprisingly large number of small, lipid soluble compounds cross the BBB at a rate considerably greater or lesser than that predicted by their physicochemical characteristics (Oldendorf 1971, 1974). Binding to serum proteins and efflux systems are major factors decreasing influx, and the presence of a saturable blood-to-brain transporter is a major factor increasing influx.

Extracellular Pathways

Albumin derived from serum is present in small amounts in the CSF, showing that the BBB is not absolute. The amount of protein in CSF, however, is very small, being about 0.5%, or 1/200th, of that in plasma. The CSF is not an ultrafiltrate but a

secreted fluid. This means that the relative and absolute concentrations of proteins, electrolytes, minerals, and other substances can differ tremendously to that of plasma. The extracellular pathways are another avenue by which substances can enter the CNS (Balin et al. 1986; Broadwell 1993). These represent what have sometimes been termed "functional leaks" at discreet areas of the brain, including the large vessels of the pial surface and subarachnoid space, the circumventricular organs, the nasal epithelium, the sensory ganglia of spinal and cranial nerves, and some deep brain regions, such as the nucleus tractus solitarius (Broadwell and Banks 1993).

The amount of a substance that enters the brain by the extracellular pathways is small. However, this route may be therapeutically relevant for compounds which have favorable peripheral pharmacokinetics, such as a long serum half-life and a small volume of distribution (Banks 2004). Antibodies, erythropoietin, and enzymes can access the brain by way of the extracellular pathways (Banks et al. 2004a, 2005a, 2007; Kozłowski et al. 1992; Grubb et al. 2008), and this may underlie their therapeutic benefits when given in high doses (Grubb et al. 2008; Alafaci et al. 2000; Ehrenreich et al. 2002; Erbyraktar et al. 2003; Morgan et al. 2000; Janus et al. 2000; Hock et al. 2003; Farr et al. 2003).

2.2.1.2 Receptor-Mediated and Saturable Transports

Saturable processes represent a diverse group of mechanisms. Included in this group are diapedesis and adsorptive endocytosis/transcytosis that share characteristics with the saturable systems.

Active Transport Versus Facilitated Diffusion

Saturable transporters (Yeagle 1987) can be divided into those which require energy (active transport) and those which do not (facilitated diffusion). Both are dependent on a protein which acts as the transporter, may have co-factors, and be modulated by physiologic and disease processes. Energy requiring systems can be unidirectional; that is, they may have only an influx or efflux component. Non-energy requiring saturable transport (facilitated diffusion) is bidirectional; that is, it transports substances in both directions with net flux being from the side of higher concentration to the side of lower concentration.

Most of the classic saturable transporters at the BBB are facilitated diffusion systems (Kaur et al. 1992). For example, GLUT-1, the transporter for glucose, is a facilitated diffusion transporter. If the level of glucose is artificially raised above that of serum (or if radioactive glucose is introduced into the CNS, but not the serum), efflux of glucose can be shown.

Transcytotic Versus Transmembrane Transport

Saturable transporters can also be categorized based on whether they use pores/channels or vesicles to transport their ligands across the BBB. In the pore system, the molecule

crosses from one side of the cell membrane to the other by passing through a cavity in the transporter protein. The substance is thus transported either into or out of the cytoplasm of the BBB cell; a second set of transporters on the opposing cell membrane completes the transfer across the BBB or the substance can rely on transmembrane diffusion. With vesicular transport, the transported substance adheres to a binding site, usually a glycoprotein. Invagination then produces a vesicle that is then routed to the opposite membrane, and the contents of the vesicle are released from the cell surface. A specificity of transport distinguishes these vesicles from the macropinocytosis whose reduction is a defining characteristic of the BBB (Reese and Karnovsky 1967).

Most small molecules, such as glucose, electrolytes, and amino acids, use pores or channels. Pore systems may be either active or facilitated diffusion systems. Vesicular transporters, on the other hand, are energy requiring and so are characterized by unidirectional transport. The best described of these vesicular dependent systems is receptor-mediated transcytosis and is characterized by clathrin- and transglutaminase-dependence (Davies et al. 1980). However, non-clathrin dependent vesicles, such as podocytes, are also likely active at the BBB.

It is reasonable to assume that very large molecules would be required to use vesicles rather than pores and channels to cross, but the molecular weight at which vesicles would be requisite is not known. It has been proposed that interleukin-2 (IL-2) is transported (Drach et al. 1996) by p-glycoprotein (P-gp). As P-gp is a pore system (Begley 2004), IL-2 would be the largest substance currently known to be transported by a pore system. Peptides much smaller than IL-2 are known to cross by vesicular dependent pathways (Shimura et al. 1991; Terasaki et al. 1992). It is clear, then, that the size of the ligand alone does not dictate the need for vesicular transport.

Diapedesis of Immune Cells

A major shift in thinking about the relation of immune cells to the CNS and BBB has occurred over the last few decades. The CNS was once viewed as separate from the immune system and sterile in terms of immune cell occupancy except under conditions of brain infection. As reviewed above, it is now clear that immune cells patrol the normal CNS, although many important questions remain. For example, a major type of brain cell, the microglia, is known to be derived from peripheral macrophages, although the extent to which the pools of peripheral macrophages and microglia mix in the normal postnatal condition is unknown.

Adsorptive Endo- and Trans-cytosis

Adsorptive endocytosis occurs when a glycoprotein on the brain's endothelial surface binds another glycoprotein in ligand like fashion (Broadwell et al. 1988; Broadwell 1989). This second glycoprotein (the ligand) may be free or attached

to the surface of a virus or immune cell (Mellman et al. 1986). The binding can initiate endocytosis with the subsequent vesicle having several potential fates (Banks and Broadwell 1994). In some cases, the vesicle is routed to lysosomes, the glycoprotein destroyed, and the vesicle rerouted to the endothelial cell surface for discharge of contents. In other cases, the vesicle can be routed to the Golgi complex and endoplasmic reticulum. In other cases still, the vesicle can be discharged at the endothelial cell surface opposite to that of uptake. In this case, the vesicle has crossed the width of the endothelial cell, and hence crossed the BBB, in a transcytotic event. What determines the fate of these vesicles is largely unknown, but at least some vesicles can engage in more than one fate (Broadwell 1993). It may be that binding of a large amount of glycoprotein to the endothelial cell can overwhelm the lysosomal pathway and result in the vesicles being routed to the transcytotic or Golgi complex pathways.

Several principles of adsorptive endocytosis and transcytosis (also termed adsorptive-mediated transcytosis) are clear. Many of the glycoprotein ligands are toxic, and endocytosis may represent a mechanism to rejuvenate or repair the membrane (Raub and Audus 1990; Vorbrodts 1994; Westergren and Johansson 1993). Viruses and other pathogens that can infect or cross the BBB have often co-opted adsorptive endocytosis/transcytosis mechanisms (Marsh 1984; Chou and Dix 1989; Schweighardt and Atwood 2001). These processes may also be related to diapedesis as many of the events of immune cell passage across the BBB resemble these endocytic mechanisms. For example, both LFA-1 (leukocyte function-associated antigen-1) and ICAM (intercellular adhesion molecule), important to immune cell passage across the BBB, are glycoproteins. Although adsorptive endocytosis is in some sense saturable because of a finite amount of any single glycoprotein on a cell surface, it is not easy to demonstrate classical saturable kinetics for this process. In fact, excess glycoprotein can sometimes further stimulate endocytosis and so lead to a paradoxical increase, rather than decrease, in the rate of passage across the BBB (Banks et al. 1997). Glycoprotein distribution on brain endothelial cells is polarized; that is, a glycoprotein may be enriched on either the luminal or abluminal membranes (Vorbrodts 1994; Zambenedetti et al. 1996). The tight junctions act as a "fence" to keep the glycoproteins confined to their respective sides of the endothelial cell (Deli et al. 2005). This means that the movement of a glycoprotein molecule (or a virus whose coat displays that glycoprotein) can be unidirectional as its transcytosis can only be initiated from the side of the brain endothelial cell that contains the ligand's complementary glycoprotein (Villegas and Broadwell 1993; Broadwell 1989). The possession and distribution of glycoproteins similarly dictate which viruses can invade the brain; neurovirulent viruses that invade the brain as free virus (as opposed to entering in Trojan horse fashion inside an infected

immune cell) can do so because they possess a glycoprotein ligand capable of binding to the BBB.

Other molecules besides glycoproteins can also induce adsorptive endocytosis/transcytosis type mechanisms. A classic example is polycationic molecules such as the poly-L-lysines and the protamines. Protamines are peptides of about 30 amino acids that contain an abundance of arginine molecules. They can induce adsorptive transcytosis so vigorously as to result in BBB disruption (Vorbrodt et al. 1995; Hardebo and Kahrstrom 1985). One of the proofs that viruses co-opt adsorptive transcytosis like mechanisms is that they, like protamine, bind to heparins and heparans; indeed, protamine sulfate blocks viruses such as HIV-1 from binding to the BBB (Banks et al. 2004c; Ramos-Kuri et al. 1996; Bobardt et al. 2004). Many of the highly charged penetrating peptides, such as those derived from Tat, and many of the antibodies that target receptor-mediated transporters likely are taken up by adsorptive endocytosis/transcytosis related mechanisms (Niewoehner et al. 2014; Weissmann 1976; Herve et al. 2008).

2.2.2 CNS to Blood

Traditionally, passage in the brain-to-blood direction (efflux) has been neglected. However, efflux often accounts for the inability of otherwise effective drugs to accumulate in the CNS. Pharmacogenomic studies have suggested that the individual variation in efflux mechanisms may explain why some individuals are less sensitive to the CNS effects of drugs or more sensitive to their toxicities (Loscher and Potschka 2002). Efflux mechanisms are important to the homeostasis of the CNS, ridding the brain of toxins (Taylor 2002). The rate of efflux can be, in addition to synthesis and degradation, an important determinant of the level of a substance produced within the CNS (Chen et al. 1997; Chen and Reichlin 1998; Maness et al. 1998).

2.2.2.1 Nonsaturable

Efflux, like influx, has both saturable and non-saturable mechanisms of entry. Transmembrane diffusion occurs for both influx and efflux. Other mechanisms, such as bulk flow, are unique for efflux.

Transmembrane Diffusion

Many of the principles that govern influx by transmembrane diffusion are also important in efflux. The dramatic role that efflux by transmembrane diffusion can play can be illustrated by comparing the fate of small, lipid soluble molecules to that of a protein after intrathecal administration. Intrathecal application of small, lipid soluble molecules, such as anesthetics, can have a local effect on spinal cord function but have little or no effect on the brain (Bernards 1999). These substances readily cross the brain endothelial

cell by transmembrane diffusion and do this as easily in the brain-to-blood direction as in the blood-to-brain direction. Therefore, they are cleared from the CSF before they are able to reach the brain (McQuay et al. 1989). In contrast, proteins such as leptin and lysosomal enzymes are too large and water soluble to undergo significant transmembrane diffusion (McCarthy et al. 2002; LeBel et al. 1999). Leptin, tetanus antitoxin, and the lysosomal enzyme idursulfase can reach the brain after intrathecal administration in amounts sufficient to produce CNS effects (Calias et al. 2012; McCarthy et al. 2002; Kabura et al. 2006; LeBel et al. 1999).

Efflux by transmembrane diffusion can also contribute to the poor diffusion of substances within brain parenchyma. Diffusion within the interstitial space of the brain is dependent on Brownian motion and the production of metabolic free water as driving forces and so is very slow (Cserr 1984; Cserr and Berman 1978). However, efflux by non-saturable (and saturable) mechanisms can further reduce the distance a substance will ultimately diffuse. For example, the less lipid soluble drug atenolol can diffuse about three times further into brain tissue than can the more lipid soluble drug acetaminophen (de Lange et al. 1993).

Bulk Flow and the Glymphatic System

Bulk flow refers to the reabsorption of CSF into the blood, which occurs at the level of the arachnoid villi (Davson and Segal 1996a) and cribriform plate (Widner et al. 1987; Yamada et al. 1991). Any substance dissolved in CSF will enter the blood by this mechanism (Pollay and Davson 1963; Jones and Robinson 1982). The glymphatic system provides an important mechanism for the mixing of the CSF and brain interstitial fluid, with aquaporin-dependent fluid production at the astrocytes providing the circulant and arteriole pulsations providing the directionality (Iliff et al. 2012). Thus, the glymphatics are important not only for bulk flow, but also for the extracellular pathways, and possibly for the movement of CSF from the spinal cord into the cranium.

The characteristics of this system result in several surprising but important phenomena. For example, CSF drained at the cribriform plate, which is likely the dominant route for CSF drainage at normal CSF pressures (Boulton et al. 1999), can enter into the cervical lymphatic system. This can provide a direct route from the CNS to the cervical lymphatics (Oehmichen et al. 1979), as has been illustrated for gp120, the glycoprotein of the human immunodeficiency virus, HIV-1 (Cashion et al. 1999). This route to the lymphatics may explain why substances injected into the brain can produce a different immune response than when the substance is injected peripherally (Knopf et al. 1995; Cserr and Knopf 1992). Another example is that in some cases, the levels of a substance in blood achieved after injection into the CSF can be sustained longer and at higher levels than after an intravenous bolus (Maness et al. 1998; Chen

et al. 1997; Chen and Reichlin 1998) This is because the central injection acts similarly to an intravenous infusion, slowly delivering drug to the blood. Impairment in lymphatic circulation can result in decreased bulk flow and so may contribute to increased levels of protein and toxins in the CSF (Iliff et al. 2013).

2.2.2.2 Saturable Transport

The last decade has seen a huge increase in the interest of efflux by saturable mechanisms. Just as efflux by transmembrane diffusion can limit diffusion of a substance within the CNS, so can the presence of a saturable efflux transporter (Blasberg 1977). Much of this interest centers on the multi-drug efflux transport systems (Begley 2004), most notably p-glycoprotein (P-gp). However, other efflux transporters for peptides, proteins, endogenous substances, and drugs are known to play important roles in physiology and disease (Martins et al. 1997; Taylor 2002; Drion et al. 1996; Mealey et al. 2001). For example, peptide transport system-1 is a major regulator of brain levels of methionine enkephalin, an endogenous opiate which suppresses voluntary ethanol drinking (Plotkin et al. 1998). Depression and recovery of peptide transport systems-1 with ethanol drinking may relate to alcohol withdrawal seizures (Banks and Kastin 1989, 1994). IL-2 is currently the only cytokine known to be transported by a saturable efflux system (Banks et al. 2004b); some have postulated this transporter may be P-gp. Poor accumulation of protease inhibitors, antibiotics, AZT, anti-cancer drugs, and many other substances occurs because of efflux systems (Glynn and Yazdanian 1998; King et al. 2001; Lee et al. 1998; Loscher and Potschka 2002; Masereeuw et al. 1994; Spector and Lorenzo 1974). P-gp plays a major role in the efflux of intrathecally administered opiate analgesics (Thompson et al. 2000). Brain-to-blood transport of a corticotropin-releasing hormone is sufficient to influence splenic levels of beta-endorphin (Martins et al. 1997). Impaired efflux of amyloid β peptide, the peptide believed to cause Alzheimer's disease, develops with aging in mice which overexpress amyloid precursor protein, thus promoting further accumulation within brain of amyloid β protein (Gherzi-Egea et al. 1996; Banks et al. 2003; Deane et al. 2004). Evidence suggests that impaired transport develops in humans as well and so may be a major mechanism for induction of Alzheimer's disease (Tanzi et al. 2004; Shibata et al. 2000).

2.3 Neuroimmune Interactions

The above discussion of BBB fundamentals is tailored towards understanding the role of the BBB in neuroimmune interactions. Below are specific examples of how the BBB is involved in neuroimmune interactions.

2.3.1 Binding Sites at the BBB: Receptors and Transporters

An important distinction for understanding the function of the BBB is that of receptors vs transporters. The term "receptor" has undergone a transformation of its usage since its introduction in the late nineteenth century when it was first used to denote some physiological function. Eventually, the term receptor was used to denote a physical binding site through which a drug or hormone could exert its effects on a cell. In the 1980s, a distinction was made between "receptor" and "binding site", the former being coupled to intracellular machinery that translated its binding into a cellular effect. Binding sites on the brain endothelial cell can represent transporters, but they can also represent traditional receptors, that is, binding sites coupled to intracellular machinery. For example, brain endothelial cells have both insulin receptors and transporters. As a result, insulin is transported across the BBB to exert effects inside the CNS, but insulin also alters a number of functions of the brain endothelial cell. As examples of the latter, insulin alters the BBB transport of zidovudine (AZT (Ayre et al. 1989), tryptophan (Cangiano et al. 1983), and leptin (Kastin and Akerstrom 2001) and alters brain endothelial cell alkaline phosphatase activity (Catalan et al. 1988). BBB studies have assumed that a binding site represents transporter function and are so designed as to not consider whether receptors as well as transporters may exist at the BBB. However, a great deal of indirect evidence and some direct evidence indicates that the vascular BBB and the choroid plexus probably possess a large variety of receptors that can alter BBB functions. Besides insulin, substances which bind to and alter the function of brain endothelial cells include mu opiate receptor ligands (Baba et al. 1988; Vidal et al. 1998; Chang et al. 2001), cytokines (Ban et al. 1991; Cunningham et al. 1992; van Dam et al. 1996; Vidal et al. 1998; Moser et al. 2004; Khan et al. 2003), leptin (Kastin et al. 2000; Bjorbaek et al. 1998; Hsueh et al. 2013a), acetylcholine (Grammas and Caspers 1991), adrenergics (Walsh et al. 1987; Kalaria and Harik 1989), glutamate (Koenig et al. 1992; Krizbai et al. 1998), and chemokines (Sanders et al. 1998).

2.3.2 Permeability to Cytokines and Related Substances

The BBB is known to transport several cytokines in the blood-to-brain direction. For example, the BBB transports the IL-1s, IL-6, and TNF- α by three separate transport systems. Additionally, nerve growth factor, brain derived neurotrophic factor, interferons, neurotrophins, eotaxin, fibroblast factor 19, and leukemia inhibitory factor (Poduslo

and Curran 1996; Erickson et al. 2014; Hsueh et al. 2013b; Pan et al. 1997b, 1998a, b) are also transported across the BBB. In some cases, the same gene which gives rise to a cytokine's receptor also produces the cytokine's transporter; whereas in other cases, the receptor and transporter are different proteins (Pan and Kastin 2002; Banks et al. 2002). Recently, a BBB transporter for pituitary adenylate cyclase activating polypeptide (PACAP) was found to be the same protein which acts as a neuronal receptor for enterostatin, but not for PACAP, and acts as a lipid transporter in the liver (Martinez et al. 2003; Park et al. 2004; Dogrukol-Ak et al. 2009). In general, BBB transporters occur throughout the CNS, including the spinal cord, although the transport rate across the BBB can vary greatly among CNS regions (McLay et al. 1997; Pan et al. 1997b, 1998b; Banks et al. 1994). Enough cytokine is transported into the brain to affect CNS function. For example, IL-1 α crosses the BBB at the posterior division of the septum where it mediates cognitive impairments (Banks et al. 2001). Similarly, serum TNF- α crosses the BBB to induce CNS release of TNF- α , which in turn can induce apoptosis in the substantia nigra (Qin et al. 2007).

The cytokine transporters are not static but respond to physiological and pathological events. The transport rates of IL-1 and TNF- α each show diurnal variations (Pan et al. 2002; Banks et al. 1998b). The transport rate of TNF- α is altered in animals with experimental allergic encephalomyelitis (EAE), spinal cord injury, or blunt trauma to the brain (Pan et al. 1996, 1997a, 2003b; Pan and Kastin 2001b; Pearce et al. 2003).

2.3.3 Permeability to Other Neuroimmune Substances

Other substances with neuroimmune actions are handled by the BBB in a variety of ways. Monoamines are largely excluded by the BBB (Hardebo and Owman 1990; Kalaria et al. 1987), and opiates and opiate peptides as a rule enter the brain by transmembrane diffusion but are transported by saturable systems in the brain-to-blood direction (King et al. 2001; Banks and Kastin 1990; Elferink and Zadina 2001). Pituitary adenylate cyclase activating peptide, a member of the VIP/secretin/PACAP family, has immune functions (Arimura 1992). Transport of its two major forms across the BBB is complex, involving both brain-to-blood and blood-to-brain components (Banks et al. 1993). Its blood-to-brain transport is altered with CNS injury (Somogyvari-Vigh et al. 2000). Some of the other immune active substances whose passage across the BBB has been investigated are melanocyte stimulating hormone (Martins et al. 1996; Wilson et al. 1984), corticotrophin releasing hormone (Martins et al. 1996), and enkephalins (Banks et al. 1986; Elferink and Zadina 2001).

2.3.4 Permeability to Immune Cells

As discussed above, immune cells cross the BBB by the highly regulated process of diapedesis. The mechanism by which immune cells cross the BBB has also been greatly clarified by recent work. Two major assumptions about how immune cells would enter the CNS has not withstood investigation. The first assumption was that immune cells would enter the CNS by leaking across a disrupted BBB. However, disruptions to the BBB are usually mediated by increased vesicular activity in the endothelial cells (Vorbrot et al. 1995; Lossinsky et al. 1983; Mayhan and Heistad 1985). These vesicles of 100 nm or so could not accommodate the passage of an immune cell 10,000 nm in diameter. Even in diseases where there is both increased immune cell trafficking into the CNS and a disrupted BBB, there is often a mismatch between the site of immune cell entry and BBB disruption (Engelhardt and Wolburg 2004).

The second major assumption is that immune cells would cross between opposing endothelial cells taking the "paracellular route." However, evidence suggests that many immune cells favor a transcellular route and that cells can cross both the vascular BBB and the choroid plexus (Kivisakk et al. 2003; Wolburg et al. 2005). In brief, immune cells tunnel through venular endothelial cells leaving the intercellular tight junctions intact (Engelhardt 2008; Wolburg et al. 2005). This tunneling process is complex and is initiated when LFA-1 on an immune cell binds to ICAM on the brain endothelial cell. Other paracellular messengers, which likely include cytokines, are then released (Male 1995; Persidsky et al. 1997). Protrusions and invaginations of the endothelial cell and protrusions of the immune cell occur, with the immune cell possibly using the tight junction as an initial anchoring site (Lossinsky et al. 1991). Other ligands which have been postulated to play a role in this transcytotic process include PECAM, VE-cadherin, members of the JAM family and CD99 (Engelhardt and Wolburg 2004). Some plasma inevitably accompanies the passage of the immune cells, which can give the appearance of a disrupted BBB (Greenwood et al. 1995; Avison et al. 2004; Persidsky et al. 2000).

The immune processes that induce an immune cell to cross the BBB are complex. Quan has shown that injection of interleukin-1 into brain tissue induces immune cell trafficking, but that such induction can be blocked by injection of lipopolysaccharide (LPS) into the periphery (Ching et al. 2005, 2006; Quan et al. 1994). Although trafficking is a product of a complex interaction between the immune cell and the BBB, the degree to which trafficking occurs can reside primarily with the immune cell or the BBB and genetics or immune events can shift that dominance (Banks et al. 2012). Such events may underlie eufinflammation, the phenomenon by which subclinical activation of the innate immune system renders it increasingly resistant to such activation by increasingly stronger stimuli (Tarr et al. 2014).