

# Neurology<sup>®</sup>

## Table of Contents

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### IN FOCUS

- 647 Spotlight on the February 25 Issue
- 690 This Week's *Neurology*<sup>®</sup> Podcast





### SPECIAL EDITORIAL



- 648 Call for papers: *Neurology: Neuroimmunology & Neuroinflammation*, a new *Neurology* journal  
*R.M. Ransohoff*

### EDITORIALS

- 650 Refractory status epilepticus: What to put down: The anesthetics or the patient?  
*N.B. Fountain and J.E. Fugate*
- 652 Antiepileptic drug nonadherence in children with epilepsy: Outcomes and potential intervention  
*J. Wagner*
- 654 The pill times 2: What every woman with multiple sclerosis should know  
*A.M. Langer-Gould*

### ARTICLES



- 656 Anesthetic drugs in status epilepticus: Risk or rescue? A 6-year cohort study  
*R. Sutter, S. Marsch, P. Fuhr, P.W. Kaplan, and S. Rüegg*  
  Editorial, p. 650
- 665 Ketogenic diet for adults in super-refractory status epilepticus  
*K.T. Thakur, J.C. Probasco, S.E. Hocker, K. Roehl, B. Henry, E.H. Kossoff, P.W. Kaplan, R.G. Geocadin, A.L. Hartman, A. Venkatesan, and M.C. Cervenka*  

- 669 Comment: Should we induce ketosis in super-refractory status epilepticus?  
*M. Gruenthal*
- 671 Early pediatric antiepileptic drug nonadherence is related to lower long-term seizure freedom  
*A.C. Modi, J.R. Rausch, and T.A. Glauser*  
 Editorial, p. 652

- 674 Pregnancy outcomes in the clinical development program of fingolimod in multiple sclerosis  
*G. Karlsson, G. Francis, G. Koren, P. Heining, X. Zhang, J.A. Cohen, L. Kappos, and W. Collins*  
Editorial, p. 654
- 681 Naive CD4 T-cell activation identifies MS patients having rapid transition to progressive MS  
*E. Zastepa, L. Fitz-Gerald, M. Hallett, J. Antel, A. Bar-Or, S. Baranzini, Y. Lapierre, and D.G. Haegert*  

- 691 APOE  $\epsilon$ 4 worsens hippocampal CA1 apical neuropil atrophy and episodic memory  
*G.A. Kerchner, D. Berdnik, J.C. Shen, J.D. Bernstein, M.C. Fenesy, G.K. Deutsch, T. Wyss-Coray, and B.K. Rutt*
- 698 Prevalence of cortical superficial siderosis in a memory clinic population  
*H.I. Zonneveld, J.D.C. Goos, M.P. Wattjes, N.D. Prins, P. Scheltens, W.M. van der Flier, J.P.A. Kuijjer, M. Muller, and F. Barkhof*  

- 705 Evaluation of SLC20A2 mutations that cause idiopathic basal ganglia calcification in Japan  
*M. Yamada, M. Tanaka, M. Takagi, S. Kobayashi, Y. Taguchi, S. Takashima, K. Tanaka, T. Touge, H. Hatsuta, S. Murayama, Y. Hayashi, M. Kaneko, H. Ishiura, J. Mitsui, N. Atsuta, G. Sobue, N. Shimozawa, T. Inuzuka, S. Tsuji, and I. Hozumi*

### GLOBAL PERSPECTIVES

- 713 Institute of Medicine's Global Forum on Innovation in Health Professional Education  
*P. Cuff, A. Meleis, and J. Cohen*

### SPECIAL ARTICLE

- 716 Summary of evidence-based guideline update: Prevention of stroke in nonvalvular atrial fibrillation: Report of the Guideline Development Subcommittee of the American Academy of Neurology  
*A. Culebras, S.R. Messé, S. Chaturvedi, C.S. Kase, and G. Gronseth*  
 

CONTINUED

# Neurology<sup>®</sup>

## Table of Contents

*continued*



### VIEWS & REVIEWS

- 725 Confidentiality in preclinical Alzheimer disease studies: When research and medical records meet  
*J.J. Arias and J. Karlawish*

### CLINICAL/SCIENTIFIC NOTES

- 730 Corticobasal syndrome associated with antiphospholipid syndrome without cerebral infarction  
*D.-W. Lee, S.-W. Eum, C.O. Moon, H.-I. Ma, and Y.J. Kim*


### REFLECTIONS: NEUROLOGY AND THE HUMANITIES


- e62 Sabertruth  
*D.S. Barron*

### NEUROIMAGES

- 732 Bilateral jugular paragangliomas: A rare cause of raised intracranial pressure  
*M.J. Thurtell, P.A. Kirby, and M. Wall*

### RESIDENT & FELLOW SECTION

- e63 Mystery Case: Eyelid myoclonia with absences in an adult patient  
*Y. Hannawi, S.S. Satpute, and A. Maheshwari*  

- e65 Clinical Reasoning: A 49-year-old man with fever and proximal weakness of his arms  
*C. Seliger, W. Schulte-Mattler, U. Bogdahn, and M. Uhl*

- e70 Teaching Video *NeuroImages*: Orofacial dyskinesia and oral ulceration due to involuntary biting in neuroacanthocytosis  
*I.K. Gooneratne, P.N. Weeratunga, and R. Gamage*  


### PATIENT PAGE

- e71 Taking epilepsy medications consistently: Why it makes a difference  
*S. Karceski*

### WRITECLICK: EDITOR'S CHOICE

- 734 When is a global health program global?  
*J.H. Chin; Author Response: J.A. Aarli, O. Abramsky*
- 734 Education Research: Changing practice: Residents' adoption of the atraumatic lumbar puncture needle  
*T.K.F. Chow; Author Response: C.E. Tung*

### EDITORIAL EXPRESSION OF CONCERN

- 735 Structural and functional brain connectivity in presymptomatic familial frontotemporal dementia

### DEPARTMENT

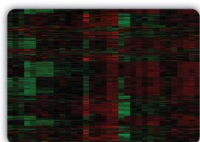
- A37 In the Next Issue of *Neurology*<sup>®</sup>

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**Cover image:** Unsupervised hierarchical clustering (Ward clustering method and Euclidean distance metric) of 1,439 genes having the greatest variance across all samples segregated healthy controls from patients with SPMS and identified 2 SPMS subgroups (SP-1 and SP-2). See page 681.

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