



## LETTER TO THE EDITOR

ARTHUR SANIOTIS, MOHAMMED MAAN AL-SALIHI, MACIEJ HENNEBERG

## KNOWLEDGE OF EVOLUTION ASSISTS IN UNDERSTANDING BRAIN VARIATIONS IN NEUROSURGERY

Anatomical structures are considered to be "ideal" types remaining unchanged over extended time. This is far from the truth. For example, neuroanatomical structures such as Circulus Arteriosus Cerebri (CAC) exhibit considerable variation up to 45%, (Qiu et al. 2015, Forgo et al. 2018) with some variants linked to risk for aneurysms and stroke (Tarulli et al. 2014, van Seeters et al. 2015). Why do such variants, among others, exist at all? Biological variation is the source and the result of evolution (Saniotis, Hennenberg 2020).

Neurological pathologies are related to evolutionary origins of brain parts. For example, Pick's disease has a predilection to primate-specific structures (i.e., the frontal and temporal lobes), and spares the primary sensory and motor cortices. The subcortical white matter (of the isocortex) is involved in Krabbe's leukodystrophy but allocortical fibers, such as the

olfactory tracts, fornix, and mammillothalamic tract are spared (Basma *et al.* 2020).

Importantly, even though natural selection has been relaxed in extant *Homo*, neuroanatomical structures are still undergoing microevolution. Certain heritable traits which can be observed now are a consequence of relaxed natural selection (Rűhli, Henneberg 2013). Some evolutionary scientists have speculated that improved living conditions have led to altered embryonic developmental processes leading to an increased prevalence in the median artery, left vertebral artery origin from the aortic arch, tarsal coalitions and accessory renal arteries (Saniotis, Henneberg 2020, Rűhli, Henneberg 2013). Of relevance to neurosurgeons is the rise in spina bifida occulta during the 20<sup>th</sup> century, which reflects a decline in differential mortality and fertility that has provided less chance for natural

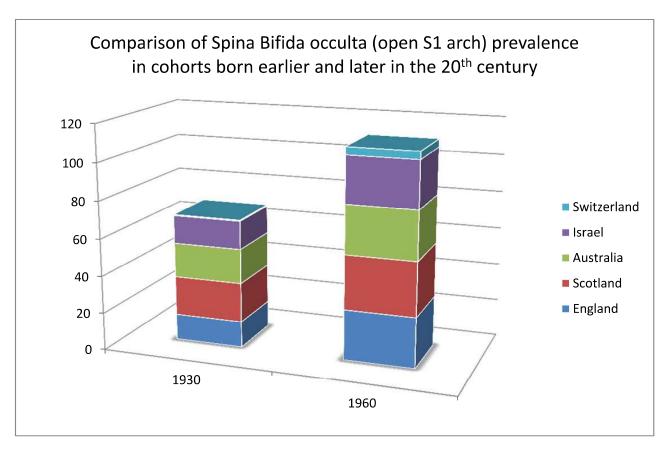


FIGURE 1: The prevalence of the spina bifida occulta (open S1 arch) increased in the 20<sup>th</sup> century. Data from Henneberg 2011.

selection to remove harmful mutations from the human gene pool (Saniotis *et al.* 2020).

Tumors are effects of evolution since neoplastic processes involve mutations. Tumors tend to grow in specific regions of the brain. Such preferential growth may be related to distinct cytoarchitectonic properties of phylogenetic origins. Glial tumors occur in phylogenetically "recent" systems (i.e., the association cortex) or "older constantly active" regions (i.e., in memory, or limbic system).

In a recent study, Hindenes *et al.* (2020) identify a trend in missing segments in CAC. Based on previous studies on increasing anatomical variations, we may speculate that this increase of missing segments in CAC reflects relaxed selection. This should certainly be brought to the attention of neurosurgeons since it identifies changes to brain hemodynamics and their implications for strokes. Therefore, increasing awareness of the evolutionary basis of neuroanatomical structures and their ongoing microevolution may provide

neurosurgeons important information on human development and possible anomalies.

## **REFERENCES**

BASMA J., GULEY N., MICHAEL L., M. II, ARNAUTOVIC K., BOOP F., SORENSON J., 2020: The evolutionary development of the brain as it pertains to neurosurgery. *Cureus* 12,1: e6748 https://doi.org/10.7759/cureus.6748

FORGO B., TARNOKI A. D., TARNOKI D. L., KOVACS D. T., SZALONTAI L., PERSELY A., HERNYES A., SZILY M., LITTVAY L., MEDDA E., SZABO A., KOZAK L. R., RUDAS G., SAS A., SEPSI M., KOSTYAL L., OLAH C., 2018: Are the variants of the Circle of Willis determined by genetic or environmental factors? Results of a twin study and review of the literature. *Twin Res Hum Genet* 21, 5: 384–93. https://doi.org/10.1017/thg.2018.50

HENNEBERG M., 2011: Changing frequencies of anatomical variants may pose clinical problems. In: 6<sup>th</sup> APICA & 13<sup>th</sup> PIN-PAAI Conference "The Future of Anatomy" GrahaBIK-

- IPTEKDOK Faculty of Medicine, Airlangga University Surabaya. Pp. 69-74.
- HINDENES L. B., HÅBERG A. K., JOHNSEN L. H., MATHIESEN E.B., ROBBEN D., VANGBERG T. R., 2020: Variations in the Circle of Willis in a large population sample using 3D TOF angiography: The Tromsø Study. *PLoS ONE* 15, 11: e0241373.
  - https://doi.org/10.1371/journal.pone.0241373
- QIU C., ZHANG Y., XUE C., JIANG S., ZHANG W., 2015: MRA study on variation of the circle of Willis in healthy Chinese male adults. *Biomed Res* 2015: 976340. https://doi: 10.1155/2015/976340
- RŰHLI F. J., HENNEBERG M., 2013: New perspectives on evolutionary medicine: the relevance of microevolution for human health and disease. *BMC Med* 11: 115. https://doi.org/10.1186/1741-7015-11-115
- SANIOTIS A., HENNEBERG M., 2020: Anatomical variations and evolution: re-evaluating their Importance for Surgeons. *ANZ Journal of Surgery* 91, 5: 837–840. https://doi: 10.1111/ans.16429
- SANIOTIS A., HENNEBERG M., MOHAMMADI K., 2020: Genetic load and morphological changes to extant humans. *Journal of Biosocial Science* 53,4: 639-642. https://doi:10.1017/S0021932020000413
- TARULLI E., SNEADE M., CLARKE A., MOLYNEUX A. J., FOX A. J., 2014: Effects of Circle of Willis anatomic variations on angiographic and clinical outcomes of coiled anterior communicating artery aneurysms. *Am J Neuroradiol* 35,8: 1551–1555. https://doi: 10.3174/ajnr.A3991
- VAN SEETERS T., HENDRIKSE J., BIESSELS G. J., VELTHUIS B. K., MALI W. P., KAPPELLE L. J. VAN DR GRAAF Y., SMART STUDY GROUP 2015: Completeness of the Circle of Willis and risk of ischemic stroke in patients without cerebrovascular disease. *Neuroradiology* 57, 12: 1247–1251. https://doi.org/10.1007/s00234-015-1589-2

- Arthur Saniotis<sup>1, 2, \*</sup> E-mail: arthur.saniotis@hirszfeld.pl Mohammed Maan Al-Salihi<sup>3</sup> Maciej Henneberg<sup>2,4</sup>
- <sup>1</sup> Department of Anthropology, Ludwik Hirszfeld Institute of Immunology and Experimental Therapy, Polish Academy of Sciences, Wroclaw, Poland
- <sup>2</sup> Biological Anthropology and Comparative Anatomy Research Unit, School of Biomedicine, University of Adelaide, Adelaide, SA, Australia.
- <sup>3</sup> College of Medicine, University of Baghdad, Baghdad, Iraq
- <sup>4</sup> Institute of Evolutionary Medicine, The University of Zürich, Zürich, Switzerland

<sup>\*</sup>Corresponding author.