presented by the authors may have revealed leftward tongue retraction due to a right hemilingual weakness. Among the known causes of a geographic tongue, a fungal infestation from poor hygiene is frequent. This relatively immobile weakened tongue may predispose the patients to decreased spontaneous cleansing, which explains hemigeographic tongue from a contralateral cerebral stroke.

Author Response: Montserrat G. Delgado, Sergio Calleja, Oviedo, Spain: We thank Dr. Chang for the interest in our NeuroImage¹ and the valuable clinical contribution. In our daily clinical practice, a hemigeographic tongue is not a frequent epiphenomenon in patients with hemispheric stroke, in spite of the fact that fungal infestation is possible in these medical setting situations (both patients presented hemigeographic tongue while hospitalized due to acute ischemic stroke). After some months, while weakness persisted, the hemigeographic tongue disappeared in

one patient; the second patient was lost to follow-up. As Dr. Chang explains, hemilingual weaknesses may be seen from contralateral cerebral strokes, but this is the first report on this disorder, to our knowledge, which suggests that the mechanism of hemigeographic tongue may be more complex. In this way, a CNS disorder may provoke an alteration of the secretion of certain trophic factors that could be implicated in the trigeminal trophic syndrome pathogenesis.^{2,3}

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CORRECTION

Familial occurrence and heritable connective tissue disorders in cervical artery dissection

In the article “Familial occurrence and heritable connective tissue disorders in cervical artery dissection” by S. Debetto et al.,¹ there is an omission in table 4. The “Gene with mutation” for Patient 1 (ID: I) should have read “COL3A1, NM_000090.3:c.1618G>A (p.Gly540Arg)” rather than “COL3A1, p.Gly540Arg” as originally published. The authors regret the omission. In addition, the authors would like to clarify the following: Patient 2 (ID: II) had a G to A transition (NM_000090.3:c.2212G>A) resulting in a p.Gly738Ser substitution according to the HGVS nomenclature, corresponding to a Gly571Ser substitution in the legacy amino acid numbering for type III collagen. The HGVS codon number is obtained by adding 167 amino acids to the legacy number. The authors regret that only the legacy numbering had previously been mentioned.

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Author disclosures are available upon request (journal@neurology.org).

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