

Expression of Early Growth Response Gene-2 and Regulated Cytokines Correlates with Recovery from Guillain–Barré Syndrome

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Guillain–Barré syndrome (GBS) is an immune-mediated peripheral neuropathy. The goal of this research was the identification of biomarkers associated with recovery from GBS. In this study, we compared the transcriptome of PBMCs from a GBS patient and her healthy twin to discover possible correlates of disease progression and recovery. The study was then extended using GBS and spinal cord injury unrelated patients with similar medications and healthy individuals. The early growth response gene-2 (EGR2) was upregulated in GBS patients during disease recovery. The results provided evidence for the implication of EGR2 in GBS and suggested a role for EGR2 in the regulation of IL-17, IL-22, IL-28A, and TNF- β cytokines in GBS patients. These results identified biomarkers associated with GBS recovery and suggested that EGR2 overexpression has a pivotal role in the downregulation of cytokines implicated in the pathophysiology of this acute neuropathy. *The Journal of Immunology*, 2016, 196: 1102–1107.

Guillain–Barré syndrome (GBS) is an immune-mediated peripheral neuropathy involving both the myelin sheath and axons that affects the peripheral nervous system (1). It has been identified as the main cause of the acute neuromuscular paralysis worldwide, with an annual incidence ranging from 0.81 to 1.89 cases per 100,000 people (2). GBS is characterized by rapidly evolving ascending weakness, mild sensory loss, and hyporeflexia or areflexia, progressing to a nadir over up to 4 wk

(2). Besides the classic presentation of ascending paralysis in demyelinating GBS, clinical variants are based on the types of nerve fibers involved (motor and/or sensory, cranial or autonomic), predominant mode of fiber injury (demyelinating versus axonal), and alterations in consciousness. Consequently, different subtypes of GBS such as acute inflammatory demyelinating polyneuropathy, Miller Fisher syndrome, acute motor axonal neuropathy (AMAN), acute motor sensory axonal neuropathy (AMSAN), acute pan-autonomic neuropathy, and Bickerstaff's brainstem encephalitis have been described previously (1–3).

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Received for publication September 29, 2015. Accepted for publication November 21, 2015.

This work was supported partially by European Union Framework Program 7 Antigen Project 278976. L.M.-H. was supported by a fellowship from the University of Castilla La Mancha (UCLM). M.V. was supported by the research plan of the UCLM.

E.D.-P. and J.d.l.F. conceived the experiments; E.D.-P., L.M.-H., E.P., A.G.-F., M.V., R.T., F.R.G., V.V.d.S., R.R., and I.G.F.d.M. performed the experiments; and J.d.l.F., E.D.-P., and E.P. wrote the paper. All authors approved the paper in its present form.

The RNA sequencing data presented in this article have been submitted to the National Center for Biotechnology Information's Gene Expression Omnibus (<http://www.ncbi.nlm.nih.gov/geo/query/acc.cgi?acc=GSE72748>) under accession number GSE72748.

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The online version of this article contains supplemental material.

Abbreviations used in this article: AMAN, acute motor axonal neuropathy; AMSAN, acute motor sensory axonal neuropathy; Ct, cycle threshold; GBS, Guillain–Barré syndrome; HCV, hepatitis C virus; PGK1, phosphoglycerate kinase 1; SCI, spinal cord injury.

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www.jimmunol.org/cgi/doi/10.4049/jimmunol.1502100