Supplementary Figure Legend

**FigureS1. Comparative transcriptomic profile of MS tetramer negative memory T cells.** A) Most enriched Biological Processes by genes differentially regulated between MS tetramer negative and HD tetramer negative memory T cells; B) Network showing the interconnection among the most enriched Biological Processes enriched by genes differentially regulated between MS tetramer negative and HD tetramer negative memory T cells; C) Hierarchical Clustering Analysis of the transcriptomic profile of memory T cells in MS patients and healthy controls