# $\beta_2$ -glycoprotein I ( $\beta_2$ -GPI) mRNA is expressed by several cell types involved in antiphospholipid syndrome-related tissue damage

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## SUMMARY

We report here the expression of  $\beta_2$ -GPI mRNA by cell types involved in the pathophysiology of the anti-phospholipid syndrome (APS), i.e. endothelial cells as a target of autoantibodies in the APS, astrocytes and neurones involved in APS of the central nervous system (CNS). Lymphocytes were also included in the study, as it has been demonstrated that patients with systemic lupus erythematosus-associated CNS diseases have serum anti-lymphocyte antibodies cross-reacting with brain antigens, and intrathecally synthesized anti-neurone antibodies. Reverse transcriptase-polymerase chain reaction followed by restriction enzyme digestion of the product obtained demonstrated the presence of  $\beta_2$ -GPI mRNA in all cell types here tested, cultured both in presence and absence of fetal calf serum. In both culture conditions, the same cell types were immunoreactive to an anti- $\beta_2$ -GPI MoAb, as determined by indirect immunofluorescence technique. Taken together, these results indicate a direct cell synthesis of  $\beta_2$ -GPI, suggesting an antigenic function of  $\beta_2$ -GPI in the APS, including the CNS disease that occurs in this syndrome.

Keywords  $\beta_2$ -glycoprotein I anti-phospholipid syndrome endothelium central nervous system

## INTRODUCTION

 $\beta_2$ -GPI is an apolipoprotein involved in lipid metabolism and classified among members of the complement control protein superfamily [1]. The  $\beta_2$ -GPI sequence, determined by protein and cDNA analysis, is strongly homologous among species [2,3]. The attention of the immunologists has recently focused on  $\beta_2$ -GPI because of its requirement for the formation of the antigenic epitopes of some anti-phospholipid autoantibodies (aPL) [4-6]. In fact, clinical and experimental reports suggest that  $\beta_2$ -GPI is one of the main target antigens for aPL in the anti-phospholipid syndrome (APS), either primary (PAPS) or secondary to systemic lupus erythematosus (SLE) [7-10]. APS is characterized by fetal loss and a wide spectrum of clinical manifestations, including neurological symptoms such as focal cerebral and ocular ischaemia, the myelopathy of lupoid sclerosis and Degos' disease, and, less frequently, Guillan-Barré polyradiculoneuritis, migraine, chorea and seizures [11,12]. Because of the frequent association with aPL, the clinical relevance of anti- $\beta_2$ -GPI autoantibodies (a  $\beta_2$ -GPI) in APS has been long debated [13–16]. A direct pathogenic role of the  $a\beta_2$ -GPI in APS has been recently suggested by the identification of these antibodies as a population distinct from

Correspondence: Professor Guido Valesini, Allergologia e Immunologia Clinica III, Policlinico Umberto I, 00161 Roma, Italy. anti-cardiolipin antibodies (aCL) [17] and by a series of clinical reports. Thus,  $a\beta_2$ -GPI have been associated with thrombosis and thrombocytopenia occurring in the APS more significantly than aPL [8,18-21]. Moreover, APS has been recently reported in patients having  $a\beta_2$ -GPI but not aPL. For this reason, to describe this primary variant of the APS, some authors coined the term 'aPL-cofactor syndrome' [22,23]. Interestingly, central nervous system (CNS) involvement is frequent in these conditions. The binding of  $a\beta_2$ -GPI to several cell elements, such as macrophages, apoptotic thymocytes, trophoblast cells, activated platelets and, in particular, endotheliocytes has been previously reported [24-31]. More recently, we demonstrated that  $a\beta_2$ -GPI bind cerebrovascular endothelium, astrocytes and neurones [32]. The  $a\beta_2$ -GPI immunoreactivity indicates that  $\beta_2$ -GPI is present in cell types other than hepatocytes, which represent the main site of synthesis in the organism.

The immunoreactivity by  $a\beta_2$ -GPI has been particularly investigated in the endothelial cells because of their relevant involvement in APS pathophysiology. Whether  $a\beta_2$ -GPI immunoreactivity of endotheliocytes is due to the uptake of extracellular  $\beta_2$ -GPI molecules (i.e. of serum origin) or to direct intracellular synthesis of the glycoprotein is still a matter of debate. Some authors described the disappearance of  $a\beta_2$ -GPI immunoreactivity in endotheliocytes grown in serum-free medium [31], whereas

others reported its lasting presence even after serum depletion [33-36]. However, the direct evidence of  $\beta_2$ -GPI synthesis by cell types involved in APS pathophysiology is provided by the demonstration of  $\beta_2$ -GPI mRNA in human fetal astrocytes as well as in human cells of intestine and placenta, as determined by reverse transcriptase-polymerase chain reaction (RT-PCR) [37-39]. In the present study, we performed RT-PCR to determine  $\beta_2$ -GPI mRNA expression by endothelial cells, a known target of autoantibodies in APS, astrocytes and neurones, being cell populations involved in the CNS disease of the APS. We also investigated lymphocytes because this cell type has been a known target of a wide spectrum of antibodies detected in sera of SLE patients, frequently with evidence of CNS disease. These antibodies share some degree of cross-reactivity with neuronal and glial antigens [40]. Moreover, intrathecal synthesis of anti-neurone antibodies has been recently demonstrated in SLE patients with CNS disease [41].

## MATERIALS AND METHODS

Cells

The following human cells were used: LAN5 (neuroblastoma line), human umbilical vein endothelial cells (HUVEC), two established human glial cell lines T67 and T70, respectively derived from a III WHO gemistocytic astrocytoma and from a glioblastoma, as previously described [32,42] (kindly provided by Professor G. Lauro, Department of Biology, III University of Rome, Italy), HEpGL2 hepatoma line used as positive control, according to Averna et al. and Chamley et al. [38,39] (kindly provided by R. Nicotra, Istituto Regina Elena, Roma, Italy), and normal skin fibroblasts as negative controls. Cell lines were cultured in their usual medium supplemented by 5-20% fetal calf serum (FCS) and gentamycin (0.05 mg/ml) (GIBCO BRL, Paisley, UK). After 3 days, one sample of each cell line was repeatedly washed in PBS pH7.3 to remove the culture medium and then analysed; a second sample was cultured for a further 3 days in FCS-free medium before analysis. Peripheral blood lymphocytes from three healthy donors were separated by Ficoll (Nycomed, Oslo, Norway) gradient centrifugation followed by hypotonic lysis of the erythrocytes and washed three times in PBS pH 7.3.

### Detection of $\beta_2$ -GPI mRNA by RT-PCR

Total RNAs, extracted from  $5-10 \times 10^6$  cells of each line by Ultraspec RNA isolation system (Biotecx, Houston, TX) according to the manufacturer's instructions, were treated with DNase I RNase-free (GIBCO BRL) and then converted to first-strand cDNA copies by random primers of 4 µg of total RNA with Super Script H<sup>-</sup>RNase RT, as suggested by the supplier (GIBCO BRL). Oligonucleotide primers designed for PCR amplification of the human  $\beta_2$ -GPI and  $\beta$ -actin mRNAs were checked by Genebank. Based on the coding sequences [2,43], the following primers were used:  $\beta_2$ -GPI-F 5' – TCTGCCATGCCAAGTTGTAAAG – 3' (784–805);  $\beta_2$ -GPI-R 5' – CATCGGATGCATCAGTTTTCCA – 3' (1045–1024);  $\beta$ actin-F 5' – AAGAGAGGCATCCTCACCCT – 3' (222–241);  $\beta$ actin-R 5' – TACATGGCTGGGGTGTTGAA – 3' (439–420) [44].

One quarter of cDNA synthesis reaction volume was combined for PCR amplification in a 100- $\mu$ l final volume containing each primer and Taq polymerase (GIBCO BRL). PCR was performed for either 35 ( $\beta_2$ -GPI) or 25 ( $\beta$ -actin) cycles, each cycle consisting of denaturation at 94°C (45 s), annealing at 60°C (30 s), extension at 72°C (30 s), after predenaturation at 95°C (2 min), and final extension at 72°C (10 min). RT-PCR products (15  $\mu$ l) were

electrophoresed on 2% agarose gels in TAE buffer. To rule out the possibility of amplification of contaminating genomic DNA, RNA samples treated with DNase were submitted to PCR amplification without RT. The optical density (OD) measurements of the lanes on the agarose gels were performed by means of a computerassisted image analysis system (MCID, Imaging Research, St Catherine's, Ontario, Canada) using dedicated software. The system was calibrated using a set of OD standards (Kodak, Rochester, NY). Non-linear distance calibration was performed using the molecular length marker (50-bp DNA ladder; GIBCO BRL). Background subtraction was automatically performed before analysis. In order to compare the results obtained from each cell line cultured in presence or absence of FCS, a semiquantitative approach was used by comparing in the same cell line samples the OD value of  $\beta_2$ -GPI lane to that of  $\beta$ -actin. The RT-PCR products were digested with the restriction enzyme Alu I (GIBCO BRL) and electrophoresed on 2% agarose gels in TAE buffer.

## Detection of $\beta_2$ -GPI by indirect immunofluorescence

Cells were cultured for 3 days on cover slides in medium containing FCS, then repeatedly washed in PBS to remove the culture medium. For each cell line, some samples were immediately fixed in PBS 4% formaldehyde (v/v) (1 h at room temperature); other samples were grown for another 3 days in FCS-free medium before fixation. The presence of  $\beta_2$ -GPI was detected by indirect immunofluorescence (IIF) using the antibody anti-human  $\beta_2$ -GPI affinity-purified mouse monoclonal ( $a\beta_2$ -GPI MoAb 1A4, isotype IgG2, 1:10 diluted in PBS) prepared as previously reported [17,32,45]. The immune reaction was revealed by goat antimouse IgG conjugated with FITC (GAM-FITC; Sigma, St Louis, MO; 30 min at room temperature) and observed under a fluorescence microscope with an FITC outfit (Leitz, Wetzlar, Germany), as described in our previous study [32]. The  $a\beta_2$ -GPI MoAb 1A4 immunoreactivity with the lymphocytes was detected on fresh cells spotted on slides and fixed in PBS 4% formaldehyde (v/v). Negative controls were performed on each cell line replacing the a  $\beta_2$ -GPI MoAb 1A4 with mouse non-immune IgG.

### RESULTS

The expression of mRNA for  $\beta_2$ -GPI and  $\beta$ -actin in the different cell lines cultured in medium containing FCS for 3 days and in fresh lymphocytes is shown in Fig. 1. The  $\beta_2$ -GPI mRNA was identified from the HEpGL2, HUVEC, LAN5, T67, T70 cell lines and from lymphocytes, whereas no detectable signal was obtained from the fibroblasts. The RT-PCR of RNAs resulted in amplification of the expected bands, i.e. 262 bp for  $\beta_2$ -GPI and 218 bp for  $\beta$ actin. Similar results were obtained from RT-PCR of RNAs extracted from the cell lines grown for another 3 days in FCSfree medium (data not shown). When RT-PCR products were digested with Alu I restriction enzyme and subjected to electrophoresis analysis, two bands of the expected length (i.e. 101 and 161 bp) were observed, as shown in Fig. 2. The possibility of amplification of contaminating genomic DNA was excluded, since no products were obtained from the RNA samples subjected to PCR without RT. Moreover, the  $\beta$ -actin-specific primer pairs were selected from two exons separated by one intronic sequence [44]. As shown in Fig. 1, the RT-PCR product of  $\beta$ -actin mRNA (218 bp), but no gene fragment (659 bp) was observed. The immunoreactivity of the  $a\beta_2$ -GPI MoAb 1A4, as detected by IIF,

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was observed with HEpGL2, HUVEC, T67, T70 and LAN5 cell lines both in presence and absence of FCS in the culture medium, and with fresh PBL, but never with the fibroblasts. The  $a\beta_2$ -GPI MoAb 1A4 immunoreactivity with HUVEC, T67 and T70 glial lines, LAN5 neuroblastoma line and PBL in serum-free condition is shown in Fig. 3 (left panels). The fluorescent pattern was homogeneous and localized within the cytoplasm and/or on the cell membrane. The intensity of fluorescence, as well as the pattern, were similar to those obtained in cells cultured in medium containing FCS (data not shown). Technique controls performed on each cell type replacing the  $a\beta_2$ -GPI MoAb 1A4 with non-immune IgG were negative (Fig. 3, right panels).

#### DISCUSSION

The results of the present study demonstrate  $\beta_2$ -GPI mRNA expression by endothelial cells, astrocytes, neurones and lymphocytes, thus indicating that these cell types synthesize  $\beta_2$ -GPI. The RT-PCR product of total RNA from these cell types resulted in amplification of the expected band for  $\beta_2$ -GPI mRNA, i.e. 262 bp. The same band was also obtained from the HEpGL2 hepatocyte cell line used as positive control for  $\beta_2$ -GPI mRNA expression both in this study and in previous reports [38,39].

After digestion with Alu I restriction enzyme we observed the two expected bands (101 and 161 bp). This confirms that the RT-PCR product resulted from the amplification of the 262 bp sequence belonging to the  $\beta_2$ -GPI mRNA. The results' viability is supported by the observations that no RT-PCR product was obtained from RNA of fibroblasts, used as negative control, and that PCR performed without RT on each RNA excluded the possibility of amplifying the contaminating genomic DNA. This latter possibility was further excluded since we observed the RT-PCR product of  $\beta$ -actin mRNA (218 bp), but no gene fragment (659 bp), as previously demonstrated [44].

The demonstration of  $\beta_2$ -GPI mRNA in endotheliocytes, CNS cells and lymphocytes, by RT-PCR, extends the knowledge that the liver is not the unique site of synthesis of  $\beta_2$ -GPI. In this respect,  $\beta_2$ -GPI mRNA has been previously demonstrated using the same technique in fetal astrocytes, cells of intestine and placenta [37–39]. The question whether  $\beta_2$ -GPI is synthesized by endotheliocytes is particularly relevant, since these cells are targeted by aPL



**Fig. 1.** Shown are reverse transcriptase-polymerase chain reaction (RT-PCR) products of mRNAs for  $\beta_2$ -GPI (left lanes) and for  $\beta$ -actin (right lanes) from HEpGL2 (1), fibroblasts (2), human umbilical vein endothelial cells (HUVEC) (3), T67 (4), T70 (5), LAN5 (6), and lymphocytes (7). The RT-PCR of RNAs results in amplification of the expected bands: 262 bp for  $\beta_2$ -GPI, 218 bp for  $\beta$ -actin. M, 50-bp DNA ladder.

and  $a\beta_2$ -GPI in APS and aPL cofactor syndrome. In view of this, the present results of  $a\beta_2$ -GPI immunoreactivity of endothelial cells cultured for 3 days in FCS-free conditions, and in particular  $a\beta_2$ -GPI immunoreactivity localized within the cytoplasm, provide evidence in favour of the presence of endogenous  $\beta_2$ -GPI in the endothelial cells. These results are consistent with previous reports of the persistence of  $a\beta_2$ -GPI immunoreactivity in endotheliocytes cultured in  $\beta_2$ -GPI-deficient medium [33–36].

Since translocation of extracellular  $\beta_2$ -GPI within the cytoplasm is unknown, the intracellular localization of  $a\beta_2$ -GPI immunoreactivity suggests  $\beta_2$ -GPI synthesis by these cells. The  $\beta_2$ -GPI molecules, synthesized in the cytoplasm, might be successively carried to the cell surface, as routinely occurs for numerous molecules expressed on the cell membrane. Thus, the  $a\beta_2$ -GPI immunoreactivity observed on the cell membrane in serum-free cultured cells, as reported here and in our previous study [32], might be due, at least in part, to this latter mechanism. Other authors [31] reported that  $a\beta_2$ -GPI binding to endothelial cell surface, as determined by ELISA, disappears after 5h of culture in serum-free conditions and is re-established after addition of serum purified  $\beta_2$ -GPI. The lack of a $\beta_2$ -GPI binding in serum-free cultured endotheliocytes reported by these authors [31] might be due to the short period of culture in serum-free medium. It is conceivable that cells require longer than 5 h to synthesize and carry  $\beta_2$ -GPI to the cell surface. However, we found  $\beta_2$ -GPI mRNA in cells cultured both in the absence and presence of serum. Moreover, the  $\beta_2$ -GPI mRNA amount was similar in both culture conditions.

These results provide evidence that endothelial cells constitutively express  $\beta_2$ -GPI mRNA, since the absence of serum  $\beta_2$ -GPI does not affect, at least for a 3-day period,  $\beta_2$ -GPI mRNA transcription in the endothelial cells. The same result was obtained on astrocytes and neurones. The role of  $\beta_2$ -GPI in the growth and long-term survival of endothelial cells has been demonstrated by purification and characterization of an endothelial cell viabilitymaintaining factor from fetal bovine serum identified as  $\beta_2$ -GPI [46]. Therefore,  $\beta_2$ -GPI synthesis by cells in serum-free medium



**Fig. 2.** Shown are the fragments obtained by digestion with the Alu I restriction enzyme of the reverse transcriptase-polymerase chain reaction (RT-PCR) products of  $\beta_2$ -GPI mRNA from HEpGL2 (a), human umbilical vein endothelial cells (HUVEC) (b), T67 (c), T70 (d), LAN5 (e), lymphocytes (f). M, 50-bp DNA ladder. The two fragments are of the expected molecular length (101 and 161 bp), confirming the specificity of the primers used.

 $\beta_2$ -GPI mRNA expression by astrocytes, neurones, endotheliocytes and lymphocytes (a)



**Fig. 3.** Shown is the immunoreactivity by the  $a\beta_2$ -GPI MoAb 1A4 with endothelial cells (human umbilical vein endothelial cells (HUVEC)) (a) astrocytoma line T67 (b), glioma line T70 (c), neuroblastoma line LAN5 (d), all cultured in serum-depleted medium for 3 days, and lymphocytes (e). The  $a\beta_2$ -GPI binding was revealed by indirect immunofluorescence using a goat anti-mouse conjugated with FITC (left panels). Control techniques were performed on each cell type replacing the  $a\beta_2$ -GPI with non-immune IgG (right panels). Calibration bar = 25  $\mu$ m. © 1999 Blackwell Science Ltd, *Clinical and Experimental Immunology*, **115**:214–219

might represent a mechanism of protection from the absence of extracellular  $\beta_2$ -GPI. In other words, the cell synthesis of  $\beta_2$ -GPI might assure endothelial cell survival even in disadvantageous conditions.

The demonstration of  $\beta_2$ -GPI synthesis by lymphocytes suggests that cells of the immune system are another potential target of  $a\beta_2$ -GPI. With regard to previous reports that anti-lymphocyte antibodies detected in serum of SLE patients with neurological disease cross-react with brain tissue antigens [47,48], the  $\beta_2$ -GPI expression by lymphocytes appears of particular interest. In vitro studies demonstrated that anti-lymphocyte antibodies are cytotoxic for neurones and astrocytes [48]. These antibodies, detected also in cerebrospinal fluid of SLE patients with evidence of intrathecal synthesis, were shown to correlate with CNS lupus disease activity [41,49]. The demonstration that astrocytes and neurones synthesize  $\beta_2$ -GPI suggests the direct antigenic function of this molecule within the CNS and, consequently, the putative role of APSassociated autoantibodies in the CNS damage.  $a\beta_2$ -GPI might contribute to the CNS pathologies by interaction with brain cytotypes, besides the interaction with cerebral vessel endothelium [32].

Further studies are advised on a larger variety of cell types and tissues in order to check the specificity of  $\beta_2$ -GPI expression. However, the present results clearly indicate that  $\beta_2$ -GPI is expressed by different cell types that are supposed to be involved in immune-mediated tissue lesions in APS.

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