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## Multicenter study on recent portal venous system thrombosis associated with cytomegalovirus disease.

Chloé de Broucker, Aurélie Plessier, Isabelle Ollivier-Hourmand, Sebastien Dharancy, Christophe Bureau, Jean-Paul Cervoni, Philippe Sogni, Odile Goria, Olivier Corcos, Riccardo Sartoris, et al.

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2 **with cytomegalovirus disease**  
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18

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31 current study are not publicly available but are available from the corresponding  
32 author upon reasonable request.

33

34 **Authors contributions:**

1 PER, DV and CDB designed the study. CDB and PER wrote the manuscript. AP,  
2 IOH, SD, CDB, JPC, PS, OG, OC, KZ, and AP collected patients' data. NFH and YY  
3 provided virological insight. All authors read and critically revised the manuscript.  
4

1 **Abstract:** 269 words.

2 **Background and aims:** Recent non-malignant non-cirrhotic portal venous system  
3 thrombosis (PVT) is a rare condition. Among risk factors for PVT, cytomegalovirus  
4 (CMV) disease is usually listed based on few reported cases. The aim of this study  
5 was to determine characteristics and outcome of patients with PVT associated with  
6 CMV disease.

7 **Methods:** We conducted a French multicenter retrospective study comparing  
8 patients with recent PVT and CMV disease ("CMV positive"; n = 23) with patients with  
9 recent PVT for whom CMV testing was negative ("CMV negative"; n = 53) or  
10 unavailable ("CMV unknown"; n = 297).

11 **Results:** As compared with patients from the "CMV negative" and "CMV unknown"  
12 groups, patients from the "CMV positive" group were younger, had more frequently  
13 fever, higher heart rate, higher lymphocyte count and higher serum ALT levels ( $p \leq$   
14 0.01 for all). Prevalence of immunosuppression did not differ between the 3 groups  
15 (4%, 4% and 6%, respectively). Extension of PVT was similar between the 3 groups.  
16 Thirteen out of 23 "CMV positive" patients had another risk factor for thrombosis.  
17 Besides CMV disease, number of risk factors for thrombosis was similar between the  
18 3 groups. Heterozygous prothrombin gene mutation was more frequent in "CMV  
19 positive" patients (22%) than in the "CMV negative" (4%,  $p = 0.01$ ) and "CMV  
20 unknown" (8%,  $p = 0.03$ ) groups. Recanalization rate was not influenced by CMV  
21 status.

22 **Conclusions:** In patients with recent PVT, features of mononucleosis syndrome  
23 should raise suspicion of CMV disease. CMV disease does not influence thrombosis  
24 extension nor recanalization. More than half "CMV positive" patients have another  
25 risk factor for thrombosis, with a particular link with prothrombin gene mutation.

1 **Lay summary:** Patients with CMV-associated portal venous system thrombosis have  
2 similar thrombosis extension and evolution as patients without CMV disease. They  
3 more frequently have prothrombin gene mutation, suggesting a synergy between  
4 these two entities to promote thrombosis.

5

6

7

## 1 **Introduction**

2 Recent non-malignant non-cirrhotic extrahepatic portal venous system thrombosis  
3 (PVT) is characterized by new occurrence of a thrombus in the main portal vein  
4 and/or its right or left branches and/or splenic or mesenteric veins [1]. The incidence  
5 of PVT is estimated at 0.7 per 100.000 per year [2]. Recent PVT can lead to intestinal  
6 infarction in 2 to 20% of cases with an estimated mortality of 20% at 30 days [3]. The  
7 causes for PVT include inherited thrombophilia (protein C or S or antithrombin  
8 deficiency; factor V or factor II gene mutation), acquired thrombophilia  
9 (antiphospholipid antibodies, myeloproliferative neoplasms, paroxysmal nocturnal  
10 hemoglobinuria), hormonal factors, as well as local and systemic inflammation [4,5].  
11 Human cytomegalovirus (CMV) infection is very frequent, usually without overt  
12 symptoms. Anti-CMV IgG, representing past infection, are found in 50-65% of adults  
13 in developed countries, and in more than 90% in developing countries [6]. After  
14 primary infection, CMV establishes a latent infection from which intermittent  
15 reactivation can occur, as with other *Herpesviridae* [7,8]. Reinfection with new  
16 strains is also possible. CMV infection is defined by the evidence of CMV (plasma or  
17 organ-specific PCR) with or without symptoms, whereas CMV disease is defined by  
18 CMV infection with organ injury or clinical symptoms suggestive of the disease.  
19 Clinical manifestations depend on patient immunity. In immunocompromised patients  
20 and newborns, organ injury is more common [9]. In immunocompetent patients, viral  
21 replication is frequently asymptomatic, although CMV disease is possible [10]. CMV  
22 infection has been associated with indirect effects, such as increased all-causes  
23 mortality, increased risk of cardiovascular disease and increased risk of deep vein  
24 thrombosis and pulmonary embolism [11–16].

1 Recent PVT associated with CMV infection has only been described in few case  
2 reports so that the specificities of this association are unknown [17,18]. The aim of  
3 this retrospective multicenter study was to describe the characteristics, associated  
4 causes and outcome of patients with CMV-associated recent PVT.

5

## 6 **Patients and methods**

### 7 **Inclusion criteria**

8 This retrospective study included three groups of patients with recent PVT.

9 The group of patients with recent PVT and CMV disease, referred to as “CMV  
10 positive” group, included all patients with CMV-associated recent PVT diagnosed  
11 between January 2000 and December 2019 in one of the centers of the French  
12 network for Vascular Liver Diseases. Diagnosis of CMV disease was based on  
13 laboratory tests performed within 3 months before or after diagnosis of recent PVT.  
14 Details are presented in Supplementary Methods [19] [20] [21].

15 The group of patients with recent PVT without CMV disease, referred to as “CMV  
16 negative” group, included all patients with recent PVT, diagnosed between January  
17 2014 and December 2019, at the French Reference Center for Vascular Liver  
18 Diseases (Hôpital Beaujon, Clichy) and tested negative for CMV. Absence of CMV  
19 disease was based on undetectable anti-CMV IgM and/or undetectable plasma CMV  
20 DNA, within 3 months before or after PVT diagnosis.

21 The group of patients with recent PVT untested for CMV disease, referred to as  
22 “CMV unknown” group, included patients with a diagnosis of recent PVT between  
23 January 2004 and December 2019 in one of the centers of the French network for  
24 Vascular Liver Diseases, without available CMV viral load or serology within 3  
25 months before or after diagnosis of PVT.



1 The study was performed in accordance with the ethical guidelines of the 1975  
2 Declaration of Helsinki and was approved by the institutional review board (CPP Ile  
3 de France IV, Paris; France). Informed consent was obtained from all patients  
4 included in the study.

5

#### 6 *Liver surface nodularity (LSN) quantification*

7 LSN quantification was performed on portal venous phase computed tomography  
8 (CT) images using semiautomated CT software (LSN Software, version 0.88; Liver  
9 Nodularity llc) by an abdominal radiologist (RS) blinded to clinical data, using a  
10 method explained by De Vos and al. in [22] and detailed in the Supplementary  
11 Methods. The optimal cutoff value of 2.5 was chosen, based on previously published  
12 data, as reliably differentiating presence or absence of advanced fibrosis [23–26].

13

#### 14 *Definitions*

15 Diagnostic criteria for recent PVT included imaging evidence of solid material in one  
16 or more segment of the portal venous system (portal trunk, left or right portal branch,  
17 splenic vein, superior or inferior mesenteric vein) on a CT-scan or a magnetic  
18 resonance imaging, associated with signs of a recent thrombosis: hyperdense  
19 thrombus on unenhanced CT phase and/or recent abdominal pain and/or systemic  
20 inflammatory response syndrome at diagnosis. Date of diagnosis of PVT was the  
21 date of the first imaging procedure fulfilling PVT diagnostic criteria. Patients having  
22 one of the following conditions at PVT diagnosis were not included in the study:  
23 cirrhosis, portal cavernoma, variceal bleeding, hepatic or biliary malignancies.

1 In patients from the “CMV positive” and “CMV negative” groups, absence of cirrhosis  
2 was ascertained using either the results of a liver biopsy, or the association of at  
3 least 2 out of the 3 following criteria: LSN < 2.5; no cause for cirrhosis; liver stiffness  
4 measurement using Fibroscan® < 10 kPa (Supplementary Figure 1). In patients from  
5 the “CMV unknown” group, the absence of cirrhosis was based on the opinion of the  
6 practitioner in charge of the patient.

7 Other definitions are presented in Supplementary Methods.

8

#### 9 *Investigations for risk factors for thrombosis*

10 Investigations for risk factors for thrombosis are detailed in Supplementary Methods  
11 [27].

12

#### 13 *Statistical analysis*

14 Quantitative variables were expressed as median (interquartile ranges) and were  
15 compared using the Mann-Whitney test. Qualitative variables were expressed as  
16 absolute and relative (percentage) frequencies and compared using the Chi-square  
17 or the Fisher's test, as appropriate. To reduce the risk of bias, we performed  
18 sensitivity analyses consisting in comparing “CMV positive” patients with “CMV  
19 negative” and “CMV unknown” patients, matched 1:1 and 1:4, respectively, using a  
20 propensity score. Covariables included in the propensity score model were selected  
21 based on their known associations with PVT development, namely age and body  
22 mass index (BMI) [28]. The model was then used to obtain matches using the  
23 nearest-neighbor matching method, with a maximal difference of propensity score of  
24 0.05 [29].

1 We analyzed variables associated with complete recanalization of portal venous  
2 system thrombosis using Cox regression univariate analysis. Variables achieving a p-  
3 value below 5% by univariate analysis and with less than 5% of missing data were  
4 included in a Cox regression multivariate analysis.[28][27][29] Duration of follow-up  
5 used for these Cox regression models was the time period between PVT diagnosis  
6 and the first CT-scan or MRI showing a complete recanalization of the portal venous  
7 system, or -in the absence of recanalization- the last imaging procedure performed  
8 within 24 months after PVT diagnosis, or death if it occurred within 24 months after  
9 PVT diagnosis. Cumulative incidence of complete recanalization of the portal venous  
10 system was assessed using the Kaplan-Meier method and compared using the log-  
11 rank test.

12 All tests were bilateral and performed with a first-species risk of 0.05. Statistical  
13 analyses were performed using SPSS version 22.0 software (SPSS Inc., Chicago,  
14 IL). CTAT form is available in the online Supplement.

15

## 16 **Results**

17

### 18 **Study population**

#### 19 *“CMV positive” group*

20 Twenty-three patients were included in the “CMV positive” group (Clichy, n=16;  
21 Caen, n=2; Lille, n=2; Besançon, n=1; Paris Cochin, n=1; Toulouse, n=1)  
22 (Supplementary Figure 2). Their virological characteristics are summarized in  
23 Supplementary Table 1.

24 Fifteen patients had a confirmed CMV disease: 13 had confirmed primary infection  
25 (11 had positive anti-CMV IgM antibodies with low avidity IgG; 2 had  
26 seroconversion); 1 immunocompromised patient (Patient 1) had colitis and plasma

1 CMV DNA at 4.61 LogUI/mL; and 1 (Patient 13) had colitis, colon biopsies with  
2 detectable CMV DNA and histological lesions compatible with CMV disease. Eight  
3 patients had probable CMV disease, based on positive anti-CMV IgM at diagnosis,  
4 but unavailable avidity. Plasma CMV DNA was available in 3 of them and was  
5 detectable in all cases. Out of the 8 patients with probable CMV disease, neutrophil  
6 to lymphocyte ratio was below 1 in 6 patients, and Downey cells were observed in 4  
7 patients (including one with neutrophil to lymphocyte ratio above 1). These  
8 proportions were similar to that observed in patients with confirmed CMV disease: 5  
9 out of 13 had neutrophil to lymphocyte ratio below 1 (unavailable in 2) and 4 out of 11  
10 had Downey cells (unavailable in 4). A recent EBV infection was ruled out by positive  
11 anti-EBNA IgG in all 6 patients with a probable CMV disease tested. Supplementary  
12 results and Supplementary Table 2 detail characteristics of PVT including features  
13 attesting recentness of the thrombus.

14

#### 15 *“CMV negative” and “CMV unknown” groups*

16 In 53 patients with recent PVT, CMV disease could be ruled out (“CMV negative”  
17 group): 48 had undetectable anti-CMV IgM antibodies and 23 had undetectable  
18 plasma CMV DNA. In 297 patients with recent PVT, neither CMV serology or nor viral  
19 load at the diagnosis of PVT was available, so that these patients were included in  
20 the “CMV unknown” group.

21

#### 22 **Characteristics at diagnosis of PVT**

23 Characteristics of the patients are presented in Table 1 and Supplementary Table 3.  
24 None of the patients from the “CMV positive” and “CMV negative” and 3 patients of  
25 the “CMV unknown” group were receiving anticoagulation at the time of PVT

1 diagnosis. Patients with CMV disease were younger at the time of PVT diagnosis  
2 than patient from “CMV negative” and “CMV unknown” groups. “CMV positive”  
3 patients had more commonly signs of viral infection including tachycardia, fever and  
4 elevated transaminases and lymphocytes than patients from the control groups.  
5 Similar results were obtained when restricting the “CMV unknown group” to patients  
6 with available liver stiffness measurement (Supplementary Table 4). Similar results  
7 were also obtained when matching, using a propensity score, “CMV positive” patients  
8 with “CMV negative” and “CMV unknown” patients (Supplementary Table 5). There  
9 was no difference in site or extension of PVT nor in rate of immunosuppression  
10 between “CMV positive” patients and patients from the two control groups.

11

## 12 **Risk factors for thrombosis**

13 Heterozygous prothrombin G20210A gene mutation was 3 to 5-fold more frequent in  
14 patients from the “CMV positive” group than in patients from the two control groups  
15 (Table 2, Figure 1). Anti-cardiolipin antibodies were more commonly present at the  
16 time of PVT diagnosis in the “CMV positive” group, but these antibodies disappeared  
17 in all but one patient, 3 months after CMV disease. There was no difference in other  
18 risk factors for thrombosis. Besides CMV disease, the number of risk factors for  
19 thrombosis was similar between the 3 groups. Similar results were obtained when  
20 restricting the analysis to patients in the “CMV unknown group” with available liver  
21 stiffness measurement (Supplementary Table 4). Similar results were also obtained  
22 when matching, using a propensity score, “CMV positive” patients with “CMV  
23 negative” and “CMV unknown” patients, (Supplementary Table 5).

24

## 25 **Evolution of patients according to CMV status**

### 1 *Complete recanalization of the portal venous system*

2 Twenty patients from the “CMV positive” group and 42 patients from the “CMV  
3 negative” group had one or more cross-sectional imaging (CT scan or MRI) available  
4 during follow-up, allowing reliable analysis of PVT recanalization. Median duration  
5 between PVT diagnosis and last cross-sectional imaging was 16 months (3-44) and  
6 13 months (8-24) in patients from the “CMV positive” and “CMV negative” groups,  
7 respectively ( $p=0.789$ ). Number of abdominal cross-sectional imaging (CT scan or  
8 MRI) in the first 24 months was similar between “CMV positive” and “CMV negative”  
9 patients [1 (1-2), vs. 2 (0-2), respectively;  $p = 0.320$ ]. Out of these 20 CMV positive  
10 and 42 CMV negative patients, anticoagulation was initiated at time of PVT diagnosis  
11 in all but one (Patient 1). In the latter patients, total duration of anticoagulation was  
12 17 months (5-54) and 24 months (12-34), respectively ( $p=0.696$ ). Twelve patients  
13 interrupted anticoagulation during follow-up, including 6 of the 20 “CMV positive” and  
14 6 of the 42 “CMV negative” group. During the first 24 months after PVT diagnosis, 10  
15 (50%) patients of the “CMV positive” group and 12 (27%) of the “CMV negative”  
16 group had a complete recanalization of the portal venous system ( $p=0.155$ ).  
17 Cumulative incidence of complete recanalization of the portal venous system at 12  
18 and 24 months of follow-up was 47 and 58% in the “CMV positive” group versus 24  
19 and 50% in the “CMV negative” group (Supplementary Figure 3). We performed a  
20 univariate (Supplementary Table 6) and then a multivariate analysis to identify  
21 variables associated with complete recanalization of PVT at 24 months. As shown in  
22 Table 3, the only variable independently predicting complete recanalization of PVT at  
23 24 months was a lower number of occluded segments at diagnosis. Similar results  
24 were obtained when matching, using a propensity score, “CMV positive” patients with  
25 “CMV negative” patients (Supplementary Table 7). Individual outcome of the patients

1 from the “CMV positive” group, including duration of anticoagulation, is detailed in  
2 Supplementary Figure 4.

3

#### 4 *Extension of PVT*

5 Two patients from the “CMV positive” group developed an extension of PVT. Patient  
6 1 had obliterative portal venopathy and colitis at diagnosis of inferior mesenteric vein  
7 thrombosis. He was initially the only patient not treated with anticoagulation and  
8 developed 2 months later an extension of thrombosis to portal trunk. Patient 5 had at  
9 diagnosis of PVT (involving superior mesenteric vein, splenic vein and portal trunk  
10 thrombosis) no risk factor for thrombosis on top of CMV. Despite anticoagulation, he  
11 developed at month 33 a left portal branch thrombosis. A myeloproliferative  
12 neoplasm was then diagnosed based on detection of *CALR* mutation. In a third  
13 patient (Patient 7), without any risk factor for thrombosis on top of CMV nor any  
14 cause for cirrhosis, recanalization occurred, and anticoagulation was discontinued at  
15 month 22. Six months later, because of a decreasing portal flow velocity,  
16 anticoagulation was resumed, and portal flow velocity normalized. No extension of  
17 PVT was observed in the “CMV negative” group.

18

#### 19 *Portal hypertension related complications*

20 Out of the 15 patients of the “CMV positive” group and the 36 patients of the “CMV  
21 negative” group who underwent gastroscopy during follow-up, 3 (20%) and 12 (33%)  
22 patients had esophageal varices, respectively ( $p = 0.506$ ). Absence of endoscopy  
23 was related to complete recanalization of the portal venous system in 12 patients, to  
24 loss of follow-up in 6 patients, while there was no explanation in 7 patients  
25 (Supplementary Figure 5). No gastro-intestinal bleeding occurred during follow-up.

1 The only portal hypertension related complication was ascites in 2 patients from the  
2 “CMV negative” group at 5 and 47 months after PVT. There were three deaths during  
3 follow-up: one in the “CMV positive” group (Patient 1, 121 months after PVT  
4 diagnosis) and 2 in the “CMV negative” group (at 8 and 313 months after PVT  
5 diagnosis). Causes of death were extra-hepatic malignancies in 2 patients and  
6 unknown in the third one.

7

## 8 **Discussion**

9 Association of recent PVT and CMV disease has long been described. However,  
10 data reported so far (45 patients in 40 articles, summarized in Supplementary Table  
11 8 [30–69]) were too fragmented to have a clear view of the impact of CMV disease  
12 on PVT presentation and outcome. Despite the rarity of this association, thanks to  
13 the French network on vascular liver diseases, we were able to fill this gap in  
14 knowledge. We collected data from 23 well characterized patients with recent PVT  
15 associated with CMV disease. Diagnosis of CMV disease was based on international  
16 guidelines as well as on data review by an expert virologist: 15 patients had  
17 confirmed CMV disease and 8 patients had highly likely CMV disease attested by  
18 detectable plasma anti-CMV IgM as well as in 7 out of these 8 patients either an  
19 elevated lymphocytes/neutrophil ratio or detectable Downey cells. Patients with CMV  
20 disease were compared with two control groups: patients with virological tests ruling  
21 out CMV disease (“CMV negative” group; n=53) and a large group of patients with  
22 unknown CMV status (“CMV unknown” group; n=297) having similar geographic  
23 origin and date of inclusion as patients of the “CMV positive” group. The large  
24 number of patients included in the “CMV unknown” group documents the  
25 unsystematic CMV testing across centers over the study period, which might have



1 induced a bias. Yet, characteristic of the patients of the two control groups were  
2 similar, suggesting that most patients with unknown CMV status did not have a CMV  
3 disease. Moreover, patients from the “CMV positive” and “CMV unknown” group were  
4 included in multiple French centers limiting the risk of bias due to a specific local  
5 recruitment.

6

7 The first major finding of this study was that CMV disease does not influence initial  
8 extension nor outcome of recent PVT. Indeed, we observed that the number of  
9 segments occluded in the portal venous system was not different between patients  
10 with CMV disease and patients of the two control groups. Moreover, cumulative  
11 incidence of complete recanalization was similar between patients of the “CMV  
12 positive” and “CMV negative” groups, with figures in line with those previously  
13 reported in a prospective European multicentric study [5]. The number of completely  
14 occluded segments at PVT diagnosis was the only variable independently associated  
15 with a lower incidence of recanalization. This information was lacking in the literature  
16 and one could have thought that an acute event, like a CMV disease, would have  
17 been associated with a better outcome of PVT. Our data do not allow us to draw  
18 conclusions with regard to anticoagulation initiation since only 1 out of 23 CMV  
19 positive patients did not receive anticoagulation at PVT diagnosis, nor on  
20 anticoagulation duration since anticoagulation was interrupted only in 6 of these  
21 patients. Yet, analysis of individual cases suggest that caution is needed when  
22 considering discontinuation of anticoagulation since one patient without any  
23 additional risk factor for thrombosis besides CMV infection had a decreasing portal  
24 flow velocity following anticoagulation interruption that normalized after  
25 anticoagulation was resumed. Our results did not allow us to test the effect of anti-

1 CMV antiviral treatment as only 3 patients received such treatment and they all had a  
2 severe presentation or extended thrombosis. Literature available regarding antiviral  
3 treatment for thrombosis in other vascular beds in patients with CMV disease is also  
4 limited and does thus not allow extrapolations [13–16,70].

5  
6 The second major finding of this study is that more than half of the patients with  
7 CMV-associated PVT had another risk factor for thrombosis. Number of thrombosis  
8 risk factors for thrombosis, regardless of the presence of CMV disease, was not  
9 different between the three groups. This suggests that CMV disease is not a strong  
10 risk factor for PVT and may rather be a trigger for PVT in susceptible patients. This  
11 view is reinforced by the rarity of the association of CMV disease with PVT,  
12 contrasting with the high incidence of CMV infection in the general population (1%  
13 per year in young adults approximately) [20]. As a practical consequence, diagnosing  
14 CMV disease in a patient with recent PVT does not deter from performing a  
15 comprehensive screening for risk factor for thrombosis.

16  
17 The third major finding of this study is the strong link between CMV-associated  
18 recent PVT and prothrombin G20210A gene mutation (Figure 1). Indeed, in our  
19 study, prothrombin gene mutation was detected in 22% of the patients with CMV-  
20 associated recent PVT vs. 4 and 8% in the two control groups. Detailed analysis of  
21 available literature supports our findings, since 5 out of the 25 patients (20%)  
22 reported with CMV-associated recent PVT and available data had prothrombin gene  
23 mutation, vs. 6% in all PVT patients in recent studies [28,71,72]. The prevalence of  
24 prothrombin G20210A gene mutation in general western European population is  
25 around 2% (Figure 1) [73]. This association could be explained by the synergy

1 between prothrombin gene mutation and CMV to promote thrombin generation:  
2 prothrombin G20210A gene mutation is associated with increased plasma  
3 prothrombin levels and dysthrombinemia with unstable prothrombin, hence more  
4 easily activated [74]; CMV surface contains procoagulant phospholipids allowing  
5 assembly of prothrombinase enzyme complex, and thus favors production of  
6 thrombin [75–77]. This effect was observed *in vitro* with infected cells and viral  
7 particles. Another hypothesis for CMV prothrombotic effect is the transient presence  
8 of antiphospholipid antibodies secondary to infection of endothelial cell by CMV,  
9 observed at diagnosis of PVT in our study and in the literature [78,79]. CMV disease  
10 was not associated with antiphospholipid syndrome, as presence of antiphospholipid  
11 antibodies was similar in the 3 groups at 12 weeks, as described in the literature.

12

13 In conclusion, CMV disease can be associated with recent PVT, but without influence  
14 on thrombosis extension, localization nor recanalization. Accordingly, diagnosis of  
15 CMV disease should not influence clinical decisions on PVT management. Other risk  
16 factors for thrombosis are often present so that identification of CMV disease does  
17 not obviate the need for a complete work-up for risks factors for thrombosis. In  
18 particular, a special link exists between prothrombin gene mutation and CMV  
19 disease.

20

21

22 **Abbreviation list:**

23 APLS, antiphospholipid syndrome; CMV: cytomegalovirus; HCV, hepatitis C virus;  
24 HIV, Human immunodeficiency virus; Ig, Immunoglobulin; OPV, obliterative portal

1 venopathy; PNH, paroxysmal nocturnal hemoglobinuria; PVT, Portal venous system  
2 thrombosis

3

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5

**1 REFERENCES:**

2 Author names in bold designate shared co-first authorship.

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1 **TABLES:**2 **Table 1: Clinical and laboratory characteristics of patients with recent portal**  
3 **venous system thrombosis, according to CMV status**

	N	“CMV positive” group n=23	N	“CMV negative” group n=53	<i>p</i> value pos. vs. neg.	N	“CMV unknown” group n=297	<i>p</i> value pos. vs unk.	<i>p</i> value neg. vs. unk.
<b>Comorbidity</b>									
Gender (female)	23	8 (35)	53	21 (40)	0.799	297	117 (39)	0.825	1.000
Age (years)	23	36 (31–47)	53	51 (38–62)	<b>0.001</b>	297	47 (36-59)	<b>0.002</b>	0.339
BMI (kg/m <sup>2</sup> )	23	28 (26–32)	47	28 (23–32)	0.824	239	26 (23-30)	<b>0.034</b>	<b>0.033</b>
Obesity (BMI > 30 kg/m <sup>2</sup> )	23	8 (35)	48	19 (40)	0.797	238	58 (25)	0.314	<b>0.033</b>
Alcohol consumption (%) *	23	1 (4)	53	1 (2)	0.516	249	13 (5)	1.000	0.477
Immunosuppression	23	1 (4)	53	2 (4)	1.000	297	19 (6)	1.000	0.752
Positive anti-HCV antibodies	21	0	52	0	-	218	5 (2)	1.000	0.587
Positive HBs antigen	21	0	52	1 (2)	-	218	3 (1)	1.000	0.247
Diabetes	23	4 (17)	53	6 (11)	0.479	231	12 (5)	<b>0.045</b>	0.117
Arterial hypertension	23	2 (9)	52	14 (27)	0.125	231	28 (12)	1.000	<b>0.016</b>
Liver stiffness measurement using Fibroscan®	8	6 (4-7)	31	5 (5-7)	0.875	143	5 (4-7)	0.824	0.718
<b>Clinical characteristics at diagnosis</b>									
No symptoms at diagnosis	23	0	53	8 (15)	0.097	297	39 (17)	<b>0.031</b>	0.840
Duration of symptoms	23		45		0.413	198		0.431	0.356
< 1 week		10 (44)		24 (45)			98 (50)		
1 week- 1 month		11 (48)		18 (34)			66 (33)		
1 – 6 months		2 (9)		1 (2)			19 (10)		
> 6 months		0		2 (4)			15 (8)		
Body temperature > 38.5°C	23	11 (48)	52	10 (19)	<b>0.014</b>	286	57 (20)	<b>0.006</b>	1.000
Abdominal pain	23	20 (87)	53	45 (85)	1.000	280	221 (79)	0.434	0.357
Heart rate (bpm)	19	105 (88–107)	49	80 (70–97)	<b>&lt;0.001</b>	160	76 (70-88)	<b>&lt;0.001</b>	0.196
<b>Laboratory characteristics at diagnosis</b>									
Leukocytes count (G/L)	23	7.6 (6.6–10.9)	52	8.1 (5.2–10.8)	0.374	281	7 (5.5-10.4)	0.117	0.644
Neutrophils (G/L)	22	3.4 (2.5–5.9)	52	4.6 (3.3–7.8)	0.100	274	4.1 (2.8-7.1)	0.261	0.217
Eosinophils (G/L)	21	0.1 (0.0-0.3)	52	0.1 (0.0-0.2)	0.392	274	0.1 (0.1-0.2)	0.694	0.059
Lymphocytes (G/L)	21	3.1 (2.4–4.9)	52	1.6 (1.2–2.2)	<b>&lt;0.001</b>	271	1.7 (1.2-2.3)	<b>&lt;0.001</b>	0.549
Platelets count (G/L)	23	221 (157–288)	51	276 (202–348)	<b>0.019</b>	281	257 (188-330)	<b>0.044</b>	0.409
Prothrombin time (%)	20	82 (75–97)	51	85 (75–96)	0.720	278	87 (73-100)	0.702	0.646
Serum ALT (UI/L)	22	99 (55-204)	52	30 (19-46)	<b>&lt;0.001</b>	279	43 (26-66)	<b>&lt;0.001</b>	0.005

Serum albumin (g/L)	23	34 (31–36)	50	33 (30–38)	0.820	273	37 (33-42)	<b>0.010</b>	<b>&lt;0.001</b>
Serum bilirubin (μmol/L)	22	9 (7–11)	52	12 (8–16)	0.067	276	10 (7-16)	0.151	0.496
Serum ferritin (μg/L)	19	573 (261–1154)	44	237 (83–508)	<b>0.007</b>	189	168 (51-382)	<b>&lt;0.001</b>	0.085
Serum CRP (mg/L)	22	76 (22–152)	49	51 (8–162)	0.723	212	30 (5-100)	<b>0.024</b>	<b>0.044</b>
Triglyceride (mmol/L)	16	1.81 (1.21-2.18)	44	1.1 (0.8-1.5)	<b>0.005</b>	200	1.08 (0.74-1.54)	<b>0.001</b>	0.504

1

2 Data are expressed as median (range) or absolute value (percentage) and were compared using the  
3 Mann-Whitney test for quantitative variables, the Chi-square or Fisher's test for qualitative variables.  
4 *p-values* were calculated between "CMV positive" (pos.), "CMV negative" (neg.) and "CMV unknown"  
5 (unk.) groups.

6 Abbreviations: ALT, alanine transaminase; BMI, body mass index; CRP, C-reactive protein; HCV,  
7 hepatitis C virus.

8 \* Alcohol consumption  $\geq$  140 g per week.



1 **Table 2: Risk factors for thrombosis identified at diagnosis of recent portal**  
 2 **venous system thrombosis, according to CMV status**

	N	“CMV positive” group n=23	N	“CMV negative” group n=53	<i>p</i> value pos. vs. neg.	N	“CMV unknown” group n=297	<i>p</i> value pos. vs. unk.	<i>p</i> value neg. vs. unk.
Factor V Leiden	23	0	51	2 (4)	1.000	285	18 (6)	0.628	0.750
Prothrombin gene mutation	23	5 (22)	51	2 (4)	<b>0.010</b>	286	22 (8)	<b>0.033</b>	0.222
Protein C deficiency	23	3 (13)	53	7 (13)	1.000	189	15 (8)	0,423	0.278
Protein S deficiency	23	2 (8)	53	3 (6)	1.000	187	12 (6)	0,332	0.728
Antithrombin deficiency	21	1 (5)	53	6 (11)	0.665	192	10 (5)	1,000	0.122
Myeloproliferative neoplasm	23	1 (4)	52	9 (17)	0.264	287	31 (11)	0.713	0.239
JAK2 <sup>v617f</sup> mutation	21	0	50	7 (14)	0.180	282	27 (10)	0.235	0.319
Antiphospholipid syndrome	22	1 (4)	51	1 (2)	0.515	277	17 (6)	1.000	0.327
Lupus anticoagulant	21	5 (24)	48	4 (8)	0.119	273	31 (11)	0.155	0.801
Anticardiolipin antibodies	20	6 (30)	49	4 (8)	<b>0.029</b>	251	12 (5)	<b>0.001</b>	0.308
Anti-β2-Gp1 antibodies	20	3 (15)	49	1 (2)	0.070	249	4 (2)	<b>0.010</b>	1.000
PNH	20	0	52	1 (2)	1.000	261	1 (0)	1.000	0.305
Behçet’s disease	23	0	53	1 (2)	1.000	271	0	-	0.164
Oral contraceptives	8	5 (72)	17	4 (24)	0.061	131	60 (46)	0.254	0.118
Other systemic factors*	23	1 (4)	53	0 (0)	0.307	284	9 (3)	0.655	0.906
Local factors	23	2 (9)	53	12 (23)	0.205	297	67 (23)	0.285	1.000
Personal history of thrombosis	23	2 (9)	53	10 (17)	0.327	297	38 (13)	1.000	0.417
1 <sup>st</sup> degree-relative history of thrombosis	23	5 (22)	53	14 (26)	1.000	297	52 (17)	0.805	0.861
Number of risk factors for thrombosis (0 / 1 / 2 / 3 and more)**	23	10/12/1/0	53	18/27/4/4	0.828	297	95/128/55/19	0.634	0.329

3  
 4 Data are expressed as median (range) or absolute value (percentage) and were compared using the  
 5 Mann-Whitney test for quantitative variables, the Chi-square or Fisher’s test for qualitative variables.  
 6 *p-values* were calculated between “CMV positive” (pos.), “CMV negative” (neg.) and “CMV unknown”  
 7 (unk.) groups.

8 Abbreviations: CMV, cytomegalovirus; PNH, paroxysmal nocturnal hemoglobinuria.

9 \* inflammatory bowel disease (n=2), systemic lupus erythematosus (n=3), sarcoidosis (n=1), celiac  
 10 disease (n=1), rheumatoid arthritis (n=1), juvenile idiopathic arthritis (n=1), psoriasis (n=1).

11 \*\* The following risk factors for thrombosis were taken into account: factor V Leiden, prothrombin gene  
 12 mutation, myeloproliferative neoplasm, confirmed antiphospholipid syndrome, PNH, Behçet’s disease,  
 13 oral contraceptive use, systemic disease, local inflammation or surgery, personal or 1<sup>st</sup> degree-relative  
 14 history of thrombosis.

1 **Table 3: Multivariate analysis using Cox regression model of variables**  
2 **associated with complete recanalization of portal venous system thrombosis at**  
3 **24 months in 62 patients with recent PVT and follow-up imaging available (20**  
4 **patients from the “CMV positive” group and 42 from the “CMV negative”**  
5 **group)**

6

<b>Variable</b>	<b>Hazard ratio</b>	<b>95% CI</b>	<b><i>p</i> value</b>
Abdominal pain	0.581	0.188-1.802	0.348
Number of occluded segments of the portal venous system*	0.591	0.403-0.866	<b>0.007</b>
Serum ALT (UI/L)	1.002	0.999-1.004	0.219

7  
8 This analysis included variables associated persistence of portal venous system thrombosis at 24  
9 months by univariate analysis, with *p* value < 0.05 and with available data for more than 95% of the  
10 patients. Regarding imaging features, only number of completely occluded segments was included in  
11 the analysis and not each specific location.

12 Abbreviation list: ALT, alanine aminotransferase; CI, confidence interval.

13 \*The following segments were considered: right portal branch, left portal branch, portal trunk, splenic  
14 vein, superior mesenteric vein.

15 Variables with Hazard ratio > 1 are associated with complete recanalization at 24 months.

1 **FIGURE LEGEND**

2 **Figure 1: Prevalence of prothrombin gene mutation in patients with recent**  
3 **portal venous system thrombosis in our study as well as in the literature,**  
4 **according to CMV status**

5

6 Data were compared using the Fisher's test.

7 <sup>a</sup>Prevalence of prothrombin gene mutation in the general population is based on a study by

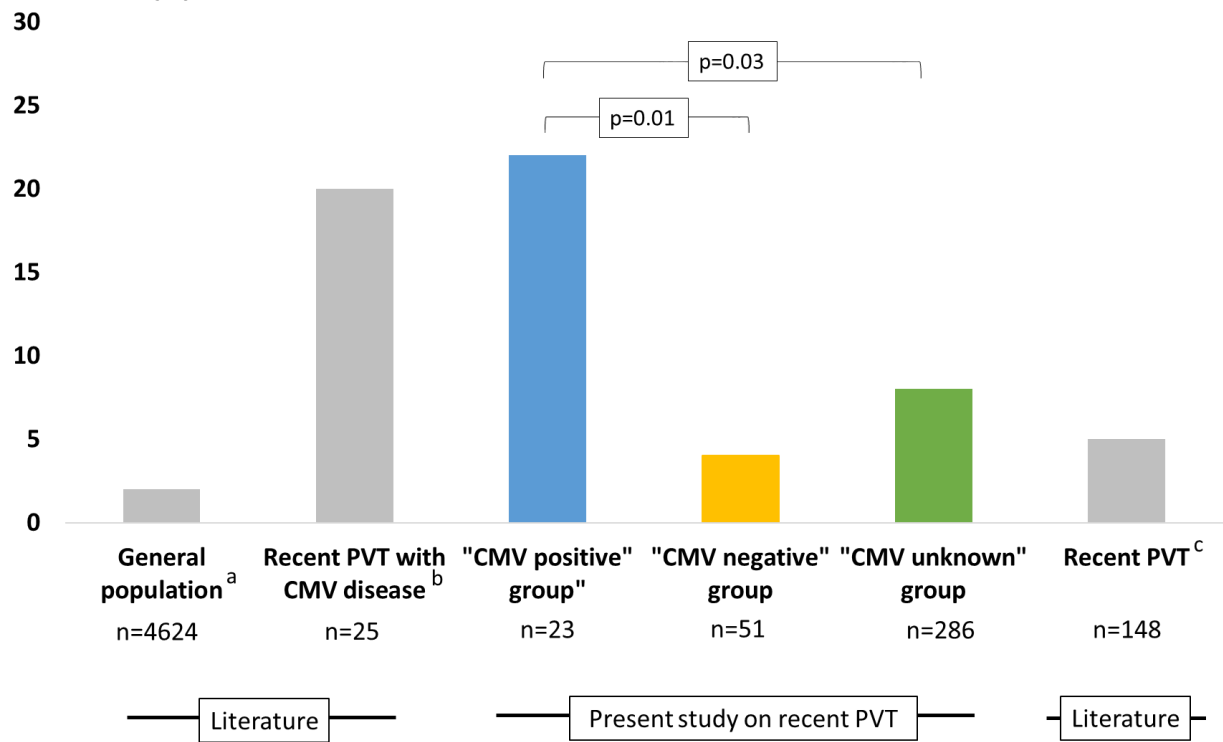
8 Rosendaal and colleagues[73]. <sup>b</sup>Prevalence of prothrombin gene mutation in cases of recent PVT and

9 CMV disease reported so far in the literature is based on studies summarized in Supplementary Table

10 8. <sup>c</sup>Prevalence of prothrombin gene mutation in patients with recent PVT from the literature based on

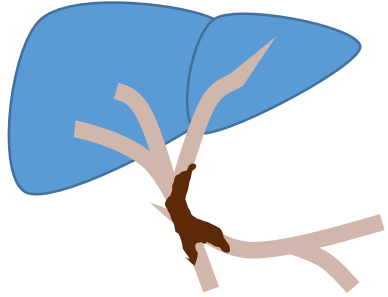
11 2 recent studies on PVT [28,71]

Prevalence (%)



# Recent portal venous system thrombosis (PVT) associated with cytomegalovirus disease

*A multicentric controlled cohort study of the French Network for the vascular liver diseases*



## Recent PVT

	cytomegalovirus disease n = 23	No cytomegalovirus disease n = 53	Unavailable cytomegalovirus status n = 297
	<ul style="list-style-type: none"><li>• Younger</li><li>• More signs of viral infection</li><li>• &gt; 50% patients had another risk factor for thrombosis</li></ul>		
Prothrombin G20210A gene mutation	22%	4%	8%

**No difference regarding localization, extension or recanalization of PVT**