# The PD-1/PD-L1 axis contributes to T-cell dysfunction in chronic lymphocytic leukemia

Davide Brusa,<sup>1</sup> Sara Serra,<sup>1</sup> Marta Coscia,<sup>2</sup> Davide Rossi,<sup>3</sup> Giovanni D'Arena,<sup>4</sup> Luca Laurenti,<sup>5</sup> Ozren Jaksic,<sup>6</sup> Giorgio Fedele,<sup>7</sup> Giorgio Inghirami,<sup>8</sup> Gianluca Gaidano,<sup>3</sup> Fabio Malavasi,<sup>9</sup> and Silvia Deaglio<sup>1</sup>

<sup>1</sup>Human Genetics Foundation (HuGeF) and Department of Medical Sciences, University of Turin, Turin, Italy; <sup>2</sup>Division of Hematology, Laboratory of Hematology Oncology, Center of Experimental Research and Medical Studies, San Giovanni Battista University Hospital, Turin, Italy; <sup>3</sup>Division of Hematology, Department of Translational Medicine, Amedeo Avogadro University of Eastern Piedmont, Novara, Italy; <sup>4</sup>Department of Onco-Hematology, IRCCS "Centro di Riferimento Oncologico della Basilicata," Rionero in Vulture, Italy; <sup>5</sup>Institute of Hematology, Catholic University of the Sacred Heart, Rome, Italy; <sup>6</sup>Department of Hematology, Dubrava University Hospital, Zagreb, Croatia; <sup>7</sup>Department of Infectious, Parasitic and Immunomediated Diseases, Istituto Superiore di Sanità, Rome, Italy; <sup>8</sup>Department of Molecular Biotechnology and Health Sciences, Center of Experimental Research and Medical Studies; and <sup>9</sup>Laboratory of Immunogenetics, Department of Medical Sciences, University of Turin, Turin, Italy

©2013 Ferrata Storti Foundation. This is an open-access paper. doi:10.3324/haematol.2012.077537

#### **Online Supplementary Design and Methods**

#### **B-cell purification**

PBMC were obtained by Ficoll-Hypaque (GE Healthcare, Milan, Italy) centrifugation. B lymphocytes were purified by negative selection by using anti-CD3, -CD14, and -CD16 antibodies and sheep anti-mouse Dynal magnetic beads (Invitrogen, Monza, Italy), with a purity constantly over 95%.

#### Immunohistochemistry and immunofluorescence microscopy

Sections were deparaffinized and endogenous peroxidase activity was blocked. Epitope retrieval was performed in 0.01M citrate buffer, pH 6.0 (40 min, 98°C). Antibodies used were goat polyclonal anti-PD-1 (2.5 µg/mL) and anti-CD23 (10 µg/mL) both from R&D Systems, rabbit polyclonal anti-PD-L1 (3 µg/mL, Novus Biologicals, Segrate, Italy) and anti-CD2 (1.3 µg/mL, Sigma, Milan, Italy), mouse monoclonal anti-CD3 (1:200, clone PS1) and anti-CD4 (1:10, clone 4B12) both from Novocastra (Milan, Italy). Goat HRP-polymer detection system (Biocare Medical, Milan, Italy), anti-rabbit HRP-conjugated Abs and 3,3'diaminobenzidine (EnVision™ System, Dako, Milan, Italy) were used to visualize the reaction. Slides were analyzed using a DMI 3000 B optical microscope (Leica Microsystems), equipped with a DCF 310 FX digital camera (Leica Microsystems). Quantification of staining intensity was performed using the LAS Version 3.8 software (Leica Microsystems) by comparing the percentage of PD-L1+ areas in 20x magnification images of PC to other parts of the same slide. At least 6 independent images of PC versus 6 of other areas were considered for each sample. For immunofluorescence, tissue sections were incubated with the following secondary Abs: AlexaFluor-633-conjugated goat antimouse IgG (Life Technologies), DyLight-488-conjugated bovine anti-goat IgG and DyLight-594-conjugated donkey anti-rabbit IgG (both from Jackson Immunoresearch, West Grove, PA, USA). Samples were counterstained with 4,6-diamidino-2-phenylindole and mounted in Slow-Fade Gold reagent (both from Life Technologies).

Immunofluorescent slides were analyzed using a TCS SP5 laser scanning confocal microscope equipped with 4 lasers (Leica

Microsystems); images were acquired with LAS AF Version Lite 2.4 software (Leica Microsystems) and processed with Photoshop (Adobe Systems, San Jose, CA, USA). Staining quantification was performed by comparing PD-L1 mean pixel intensity in 63x magnification images of CD3high versus CD3low areas. At least 10 independent images of the different areas per slide were analyzed using ImageJ software (National Institutes of Health. Available from: http://rsbweb.nih.gov/ij/).

After labeling with carboxyfluorescein diacetate N-succinimidyl ester (CFSE, 5  $\mu$ M final concentration, Invitrogen), purified CLL cells or peripheral blood normal B lymphocytes were incubated (2.5x10 $^{5}$ /well) in 96-well plates in the presence of CpG ODN2006K (1  $\mu$ g/mL; TibMolBiol, Genoa, Italy) and recombinant human IL-2 (100 IU/mL; R&D Systems). The proliferative response was measured after five days by staining with anti-CD19-PE and anti-PD-L1-PE-Cy7 mAbs (eBioscience). PBMC were similarly labeled and incubated with anti-CD3 (0.5  $\mu$ g/mL) and anti-CD28 (0.5  $\mu$ g/mL, both soluble) mAbs for 3-5 days. Cells were stained with anti-CD8-PE, -CD4-PE-Cy5 and -PD-1-PE-Cy7 before being analyzed by flow cytometry.

#### **ELISpot** assay

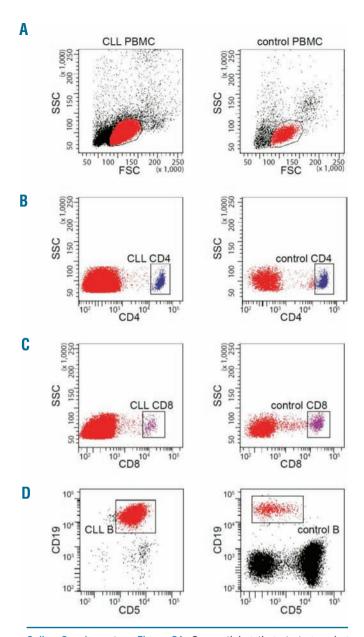
Nitrocellulose membrane 96-well microtiter (Multiscreen, Millipore, Milan, Italy) were coated with an anti-IFN-γ Ab (Biolegend). 2.5 x106 PBMC from CLL patients or control donors were cultured for five days with a mitogenic combination of anti-CD3 and anti-CD28 antibodies (see above). Soluble PD-L1 (from R&D System; 10 µg/mL) or blocking anti-PD-1 and anti-PD-L1 antibodies (both from eBiosciences; final concentration 5 µg/mL) were used to interfere with the PD-1/PD-L1 axis. After five days cells were recovered and seeded (2.5x10<sup>5</sup>) in triplicate wells in the presence of the indicated treatments. After 24 h, cells were lysed with distilled water and a biotinylated anti-IFN-y mAb was added to the wells, followed by streptavidin-horse radish peroxidase-conjugated (both from Biolegend). The substrate solution (3-amino-9-ethylcarbazole, Sigma) was added for 30 min and the reaction stopped by rinsing with tap water. Red spots (indicative of reactive lymphocytes) were detected with the AID Elispot-Reader (Bioline Amplimedical, Milan, Italy).

### Online Supplementary Table S1. Characteristics of CLL patients and of controls.

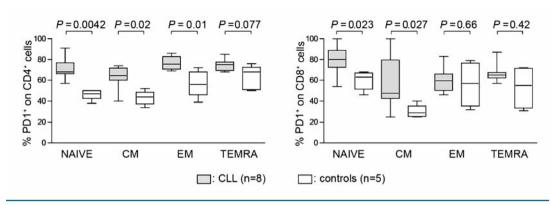
| oontrois.  |  |
|--|--|
| Variable   | N. (%)   |
| Controls<br>Mean age (years)<br>Sex (males)  | 33<br>65<br>18 (54)  |
| Patients<br>Mean age (years)<br>Sex (males)  | 117<br>63<br>60 (51)   |
| Binet stage at diagnosis, n. 98<br>A<br>B/C  | 76 (78)<br>22 (22)   |
| Treatment, n. 94<br>Treated<br>Fludarabine-based<br>Campath-based<br>Other                                     | 38 (40)<br>20 (53)<br>3 (8)<br>15 (39)                       |
| IgVH gene mutation status, n. 90<br>UM (<2%)   | 35 (39)  |
| Genomic aberrations, n. 76 deletion 11q/17p deletion 11q only deletion 17p only deletion 13q trisomy 12 normal | 12 (16)<br>4 (5)<br>11 (14)<br>35 (46)<br>18 (24)<br>20 (26) |

## Online Supplementary Table S2. Clinical characteristics of patients undergoing lymph node biopsy.

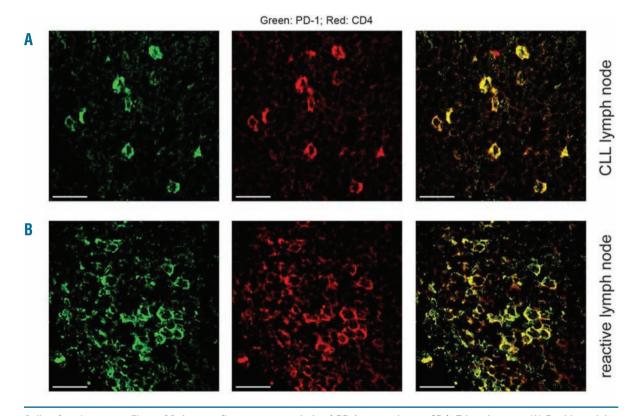
| Variable   | N. (%)   |  |
|--|----------|--|
| CLL lymph nodes<br>Median age at diagnosis (years), n.12 | 20<br>64 |  |
| Binet stage at diagnosis, n. 12                          |          |  |
| A  | 2 (17)   |  |
| B/C  | 10 (83)  |  |
| Treatement, n. 12  |          |  |
| Treated  | 8 (66)   |  |
| Genomic aberrations, n. 8                                |          |  |
| deletion 11q/17p   | 4 (50)   |  |
| deletion 17  | 0 (0)    |  |
| deletion 13  | 0 (0)    |  |
| trisomy 12   | 1 (12.5) |  |
| normal   | 3 (37.5) |  |



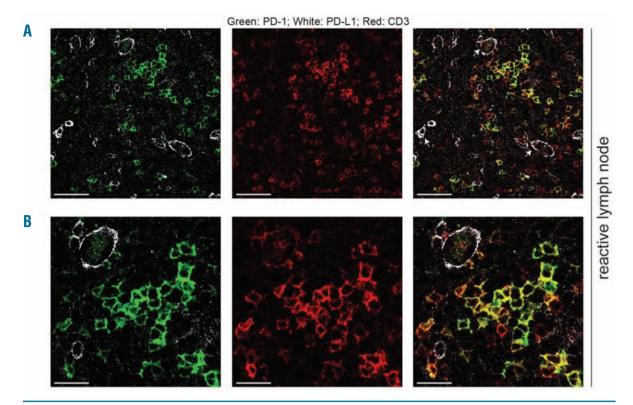
Online Supplementary Figure S1. Sequential gating strategy using multicolor flow cytometry. (A) Lymphocytes from PBMC preparations from CLL patients (left) or age- and sex-matched controls (right) were first identified based on a combination of forward and side scatter (FSC and SSC). CD4\* (B) or CD8\* (C) T-lymphocyte subsets were next gated on the basis of SSC and stained with a specific antibody. Double CD19/CD5 staining (D) was used to gate on CD5\* CLL or CD5- normal B lymphocytes.



Online Supplementary Figure S2. Comparative analysis of PD-1 expression on CD4<sup>+</sup> and CD8<sup>+</sup> T-cell subpopulations in CLL patients and in age- and sex-matched donors. Cumulative data showing the distribution of PD-1 surface levels in CD4<sup>+</sup> (left) or CD8<sup>+</sup> (right) lymphocyte subpopulations in 8 CLL patients (gray box plots) and 5 matched controls (open bars).



Online Supplementary Figure S3. Immunofluorescence analysis of PD-1 expression on CD4<sup>+</sup> T lymphocytes. (A) Double staining of a representative CLL LN section with anti-PD-1 (green) and anti-CD4 (red) indicates that PD-1 expressing cells are CD4<sup>+</sup> T lymphocytes. (B) Double staining of a representative germinal center from a reactive LN tissue section with anti-PD-1 (green) and anti-CD4 (red). Original magnification 63x, zoom factor of 2. Scale bar represents 25  $\mu$ m. Images were acquired using a TCS SP5 laser scanning confocal microscope (Leica Microsystems) with an oil immersion 63x/1.4 objective lenses equipped with the LAS AF software and processed with Adobe Photoshop (Adobe Systems).



Online Supplementary Figure S4. Immunofluorescence analysis of PD-1, PD-L1 and CD3 expression in reactive lymph node samples. (A-B) Triple staining of a representative reactive LN section with anti-PD-1 (green), anti-PD-L1 (white) and anti-CD3 (red), showing that PD-L1 expressing cells are restricted to endothelial cells (white arrows), while PD-1-positive elements are predominantly CD3<sup>+</sup>. Original magnification 63x. Zoom factor of 2 for the images below (B). Scale bars represent 50 µm (A) and 25 µm (B). Immunofluorescence was analyzed using a TCS SP5 laser scanning confocal microscope (Leica Microsystems) with an oil immersion 63x/1.4 objective lenses, images were acquired with the LAS AF software and processed with Adobe Photoshop (Adobe Systems).