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Notes:

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Roberta Morosetti^{*†}, Massimiliano Mirabella^{**§}, Carla Gliubizzi^{**}, Aldobrando Broccolini^{*}, Luciana De Angelis[¶], Enrico Tagliafico^{||}, Maurilio Sampaolesi^{**}, Teresa Gidaro^{*}, Manuela Papacci^{*}, Enrica Roncaglia^{||}, Sergio Rutella^{††}, Stefano Ferrari^{||}, Pietro Attilio Tonali^{**}, Enzo Ricci^{**}, and Giulio Cossu^{**†§§}

^{*}Department of Neurosciences and [†]Interdisciplinary Laboratory for Stem Cell Research and Cellular Therapy, Catholic University, Largo A. Gemelli 8, 00168 Rome, Italy; [‡]Fondazione Don Carlo Gnocchi, 00194 Rome, Italy; ^{**}Institute of Cell Biology and Tissue Engineering, San Raffaele Biomedical Science Park, 00128 Rome, Italy; [¶]Department of Histology and Embryology, University "La Sapienza," 00161 Rome, Italy; ^{||}Department of Biomedical Sciences, University of Modena and Reggio Emilia, 41100 Modena, Italy; ^{**}Stem Cell Research Institute, San Raffaele Hospital, 20132 Milan, Italy; ^{††}Institute of Hematology, Catholic University, 00168 Rome, Italy; and ^{§§}Department of Biology, University of Milan, 20133 Milan, Italy

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Inflammatory myopathies (IM) are acquired diseases of skeletal muscle comprising dermatomyositis (DM), polymyositis (PM), and inclusion-body myositis (IBM). Immunosuppressive therapies, usually beneficial for DM and PM, are poorly effective in IBM. We report the isolation and characterization of mesoangioblasts, vessel-associated stem cells, from diagnostic muscle biopsies of IM. The number of cells isolated, proliferation rate and lifespan, markers expression, and ability to differentiate into smooth muscle do not differ among normal and IM mesoangioblasts. At variance with normal, DM and PM mesoangioblasts, cells isolated from IBM, fail to differentiate into skeletal myotubes. These data correlate with lack in connective tissue of IBM muscle of alkaline phosphatase (ALP)-positive cells, conversely dramatically increased in PM and DM. A myogenic inhibitory basic helix–loop–helix factor B3 is highly expressed in IBM mesoangioblasts. Indeed, silencing this gene or overexpressing MyoD rescues the myogenic defect of IBM mesoangioblasts, opening novel cell-based therapeutic strategies for this crippling disorder.

The idiopathic inflammatory myopathies (IM), characterized by mononuclear cells infiltration of skeletal muscle, are the largest group of acquired muscle diseases and encompass three major forms: dermatomyositis (DM), polymyositis (PM), and inclusion-body myositis (IBM) (1). Causes of DM, PM, and IBM are unknown, but an autoimmune pathogenesis is supported by marked up-regulation of cytokines and adhesion molecules, evidence of a T cell-mediated myocytotoxicity in PM and IBM and of a complement-mediated microangiopathy in DM (2). Current immunotherapies are usually effective in DM and PM patients, whereas IBM, the most frequent myopathy in elderly patients, responds poorly or not at all to immunosuppressive therapies and its course steadily progresses to severe disability. In IBM muscle, the presence of degenerative features, such as vacuolated fibers containing amyloid and amyloid-related proteins (3), reflects a complex pathogenesis involving misfolded and unfolded proteins and increased oxidative stress in the context of a cellular "aged" milieu acting in concert with chronic inflammation (4). Regeneration and repair of muscle fibers are fundamental processes accounting for rebuilding muscle integrity and gradual recovery of muscle strength in IM after suppression of mononuclear cells infiltration. Satellite cell-dependent regeneration occurs also in IBM muscle wherein multiple metabolic pathways normally involved in muscle development are activated (5, 6). However, in IBM, despite the activation of potentially repairing mechanisms, regeneration is inefficient.

Mesoangioblasts are vessel-associated stem cells, firstly isolated from dorsal aorta of mouse embryos (7), able to differentiate into a variety of mesoderm tissues including skeletal, cardiac and smooth muscle (8, 9). When delivered intraarterially, mesoangioblasts

restore to a significant extent muscle morphology and function in a mouse model of muscular dystrophy (10).

Because mesoangioblasts express numerous receptors for inflammatory cytokines, we assumed that the human counterpart of murine mesoangioblasts should be recruited in high numbers during muscle inflammation.

Here, we describe the isolation and functional characterization of pericyte-derived adult mesoangioblasts (herein simply called mesoangioblasts) from diagnostic muscle biopsies of IM patients and show that IBM mesoangioblasts fail to differentiate into skeletal muscle. This differentiation block can be corrected *in vitro* by transient expression of MyoD, making these cells potential attractive candidates for cellular therapy of this disabling disease.

Results

Mesoangioblasts Are Efficiently Isolated from IM Muscle Biopsies.

After 10–15 days of organ culture from biopsies of three normal controls, three DM, three PM, and six IBM, we isolated a population of cells morphologically different from satellite cells. Approximately $3\text{--}4 \times 10^4$ cells could be obtained from each biopsy. From the first passage on, cells were characterized by a triangular, adherent, refractive shape and by a floating/loosely adherent round component, particularly abundant in DM (Fig. 1A). Peculiar cell morphology, phenotypic characteristics, and differentiation potential indicated that our cells were human mesoangioblasts, as recently characterized (A. Dellavalle, M.S., R. Tonlorenzi, E.T., B. Sacchetti, L. Perani, B. G. Galvez, G. Messina, R.M., S. Li, G. Peretti, J. S. Chamberlain, W. E. Wright, Y. Torrente, S.F., P. Bianco, and G.C., unpublished data).

Cells were kept in culture up to 25 population doublings (PD) (25 for control and DM, 19 for IBM, and 20 for PM) with a proliferation rate comparable for DM, IBM, and PM and independent from patients' age. Doubling time from all biopsies was 33.5 ± 2.38 h (Fig. 1B). At both early and late passages, cells kept a diploid karyotype (data not shown). There were no differences in the number of cells isolated from freshly dissected or fresh-frozen muscles at both early

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The authors declare no conflict of interest.

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Abbreviations: ALP, alkaline phosphatase; IM, idiopathic inflammatory myopathies; IBM, inclusion-body myositis; DM, dermatomyositis; PM, polymyositis; SDMC, satellite-derived myogenic cells; *mdx*, mouse muscular dystrophy; bHLH, basic helix–loop–helix; BHLHB3, bHLH domain containing class B3 transcription factor.

[§]To whom correspondence should be addressed. E-mail: mirabella@rm.unicatt.it.

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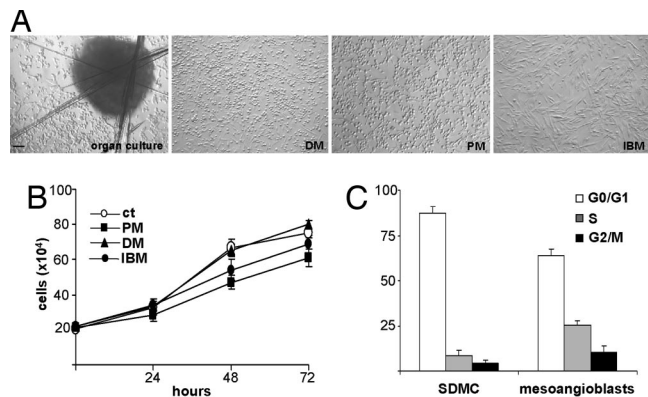


Fig. 1. Cell morphology, growth curve, and cell cycle. (A) From the organ culture on, refractive triangular, adherent, and round loosely adherent/floating cells were observed. (Scale bar: 40 μ m.) (B) Cell growth was assessed after 24 h, 48 h, and 72 h. Viable cells were judged by trypan blue exclusion. Results are expressed as absolute counts. Bars represent mean \pm SD of triplicate samples of one representative experiment of three. (C) Cell cycle distribution of proliferating normal human SDMC and mesoangioblasts from three controls, three DM, three PM, and six IBM (run in duplicate) after 24 h of culture was assessed by propidium iodide and FACS. For each sample, percentage of cells in G₀/G₁, S, or G₂/M phases of cell cycle is indicated. One representative experiment of three is shown.

and late passages, neither phenotypic characteristics of the 12 IM studied were significantly different after 8 and 18 PD (Fig. 2A and Fig. 7, which is published as supporting information on the PNAS web site). Because we isolated on average of $3\text{--}4 \times 10^4$ cells from a single biopsy, the estimated final number of cells after 25 PD is $50\text{--}120 \times 10^{10}$, and the real number that could be obtained before the appearance of senescent cells in significant proportion is between 10 and 20×10^9 cells. This number would be suitable for intraarterial delivery to adult patients, based on a per kg comparison with the mouse model used before (10).

Clonogenic Potential, Cell Cycle, and Phenotypic Characteristics Do Not Differ Among IM Mesoangioblasts. We dissociated mesoangioblasts to single cell suspension and cloned them by limiting dilution: clones appeared in 9.75 ± 3.9 , 8.87 ± 3.1 , and 10.5 ± 4.0 wells for DM, PM, and IBM, respectively, all with the same double morphology of the original cells. By replating the clones at clonal density they were able to give rise to new clones.

The cell cycle distribution was similar for all mesoangioblasts of 12 IM in three separate experiments (each one conducted in duplicate) (G₀/G₁, $65.5 \pm 6.4\%$; S, $23.7 \pm 4.3\%$; G₂/M, $10.8 \pm 2.9\%$) regardless of the IM type. The pattern of distribution was significantly different ($P \leq 0.01$) from that observed in control proliferating satellite-derived myogenic cells (SDMC) (G₀/G₁, $87.58 \pm 3.6\%$; S, $8.43 \pm 3.0\%$; G₂/M, $4 \pm 1.8\%$) (Fig. 1C). Results were always consistent throughout all experiments. Cells from all IM were strongly positive for CD44 and CD13, positive for CD49b, homogeneously negative for CD34, CD133, CD45 by FACS (Fig. 2A), consistently with what observed in normal human mesoangioblasts (A. Dellavalle, M.S., R. Tonlorenzi, E.T., B. Sacchetti, L. Perani, B. G. Galvez, G. Messina, R.M., S. Li, G. Peretti, J. S. Chamberlain, W. E. Wright, Y. Torrente, S.F., P. Bianco, and G.C., unpublished data). By immunocytochemistry and Western blot, all of the cells were positive for vimentin, weakly positive for α -SMA and desmin, and did not express glial fibrillar acidic protein (GFAP), nestin, β III-tubulin, and MyoD (data not shown). Alkaline phosphatase (ALP) staining was positive in all IM mesoangioblasts, with the highest levels observed in PM and only a weak labeling in IBM (Fig. 2B). Together, these markers identify human adult mesoangioblasts as the *in vitro* progeny of pericytes.

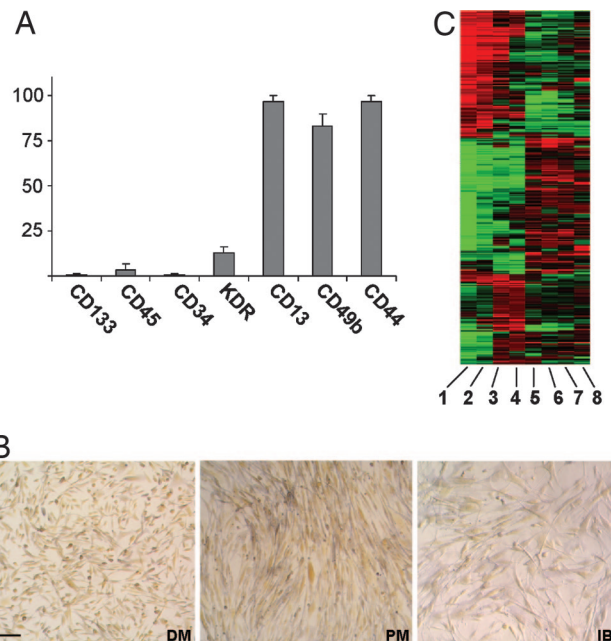


Fig. 2. FACS, immunophenotyping, ALP histochemistry, and gene expression profiling of IM mesoangioblasts. (A) More than 90% of cells from all samples were strongly positive for CD44 and CD13 with high percentage of cells CD49b-positive. None of the markers positive in murine mesoangioblasts were significantly expressed. Bars represent the mean \pm SD of 36 samples from the 12 patients with IM (3 DM, 3 PM, and 6 IBM) (each performed in triplicate). (B) IM mesoangioblasts *in vitro* are all ALP-positive. After simultaneous staining in the same culture conditions, more intensely labeled cells can be observed in PM and to a lesser extent in DM, whereas IBM mesoangioblasts are only weakly positive. (Scale bar: 20 μ m.) (C) Clustering results show two main classes: mesoangioblasts from normal controls (lanes 5–8), mesoangioblasts from DM (lanes 1 and 2), and IBM (lanes 3 and 4). Clustering procedure pairs together DM and IBM replicates.

The Ability to Differentiate into Smooth Muscle Cells (SMCs) and in Osteoblasts Are Similar Among All IM Mesoangioblasts. Murine mesoangioblasts differentiate into mature SMCs upon TGF β treatment (11, 12). Therefore, we exposed mesoangioblasts from all patients to TGF β . Approximately 80% of cells from all biopsies differentiated into strongly positive α -SMA-positive SMC, with no significant difference between the various IM (Fig. 8 which is published as supporting information on the PNAS web site).

Similarly to murine mesoangioblasts (8), human cells responded to BMP2 with a rather low percentage ($\approx 5\%$) differentiating into strongly ALP-positive osteoblast-like cells expressing osteocalcin and osteopontin (data not shown). In contrast, both control and IM mesoangioblasts failed to differentiate into neurons or glia when grown in neural stem cell differentiation media (data not shown).

Genome-Wide Gene Expression in IM Mesoangioblasts. Proliferating mesoangioblasts from normal and IM muscle were analyzed for gene expression by Affimetrix gene array. As expected, gene expression profile was similar in all samples with only few genes differentially expressed. Clustering results are shown in Fig. 2C. Two main classes were defined: the first included mesoangioblasts from normal muscle, whereas the second consisted of mesoangioblasts from DM and IBM. Interestingly, the clustering procedure paired together DM and IBM replicates.

A summary of the analysis is shown in Table 1, which is published as supporting information on the PNAS web site. In particular, mesoangioblasts from controls and IM (DM and IBM) did not express myogenic factors such as MyoD, or Pax3, Pax7, MEF2C, or MEF2D. As expected for mesoderm cells, mesoangioblasts did not

combined with regenerative cell therapy, given the inflammatory background of IBM muscle. Also rare cases of unresponsive PM and DM would be potentially treatable as well by targeting muscle groups essentials for motor and respiratory functions.

Methods

Patients. Diagnostic muscle biopsies were performed after informed consent at the Neurology Department of Catholic University. We used fresh and fresh-frozen muscles from three normal controls (one fresh, two frozen) (48–84 years of age, average 64 ± 18.33 years of age) and 12 patients with IM: 3 DM (2 fresh, 1 frozen), 3 PM (1 fresh, 2 frozen) (33–75 years of age, average 52.5 ± 9.5 years of age), and 6 sporadic IBM (3 fresh, 3 frozen) (56–75 years of age, average 67.4 ± 18.1 years of age). Diagnosis was based on clinical evaluation and laboratory studies. None of the patients received steroids or immunosuppressive therapy before biopsy. This research was approved by the ethical committee of our institution.

Cell and Organ Cultures. Fragments of intramuscular vessels and surrounding mesenchymal tissue were plated as described (8, 10). Details are provided in *Supporting Methods*, which is published as supporting information on the PNAS web site.

A fragment from the same muscle biopsy was also cultured to obtain primary muscle cultures from satellite cells by using the explantation reexplantation method (18).

Characterization of Human Mesoangioblasts from IM by FACS, Cell-Cycle Analysis, and Growth Curve. Cells (5×10^4) were incubated with FITC-, PE-, or APC-conjugated mAbs directed against AC133/1, CD34, VEGF-RII (KDR), CD45, CD49b, CD44, and CD13. Details are provided in *Supporting Methods*.

In Vitro Differentiation. Skeletal muscle differentiation. Mesoangioblasts were (i) cultured under standard differentiating conditions for SMC (17); (ii) cocultured with a 4-fold excess of C2C12 myoblasts; (iii) cultured in normal human SMC-conditioned medium and then exposed to differentiation medium. At each time point, cells were fixed or harvested for protein extraction. Differentiation assays were performed in all IM samples studied and repeated at least three times for each patient with consistent results. **Smooth muscle, osteoblasts, and neural differentiation.** Differentiation of mesoangioblasts into SMCs, osteoblasts, and neural cells was tested as described (8, 12, 25).

Immunostainings. Immunostainings were performed as described (5, 6). Details are available in *Supporting Methods*.

Double Immunohistochemistry-Histochemical ALP Staining. Immunocytochemistry for MyoD or Pax7 was performed with peroxidase-antiperoxidase followed by histochemical ALP staining on the same unfixed frozen sections of the biopsies used for mesoangioblast isolation.

Gene Expression Profiling and Data Analysis. Proliferating mesoangioblasts from normal and IM muscles were analyzed for gene expression by Affimetrix gene array (26, 27). Details are provided in *Supporting Methods*.

Western Blot Analysis and RT-PCR. Protein expression was analyzed by Western blot according to standard methods. Details on antibodies and primers are available in *Supporting Methods*. Primers and PCR conditions for BHLHB3 have been described (28).

Cell Transduction. Mesoangioblasts from three IBM patients were adenoMyoD-transduced (29), cultured for 24 h in growth medium, and then either shifted to differentiation medium for 7 days or injected *in vivo*. Details are available in *Supporting Methods*.

Intramuscular Transplantation of DM, IBM, and AdenoMyoD-Transduced IBM Mesoangioblasts into Irradiated *scid/mdx* Mice. Mesoangioblasts from DM, IBM, and IBM adenoMyoD-transduced were injected into the right or left TA of six mice (two per group). Details are available in *Supporting Methods*.

BHLHB3 siRNA. Predesigned siRNA directed against human BHLHB3 (Hs BHLHB3 1 and Hs BHLHB2 HP siRNA; Qiagen, Valencia, CA) were transfected into IBM mesoangioblasts. Details are available in *Supporting Methods*.

Statistical Analysis. All data were expressed as mean \pm SD. One-way ANOVA was used to compare differences between groups. Statistical significance was set at $P \leq 0.05$.

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