

REVIEW

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The role of puberty and adolescence in the pathobiology of pediatric multiple sclerosis

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Abstract

Multiple sclerosis (MS) is increasingly recognized in the paediatric age. In a smaller, but well-established, proportion of paediatric MS patients [20% of total paediatric MS cases: 0.2% to 0.7% of the total MS patients] the onset of disease is before 10 years of age [pre-pubescent (*childhood*) MS]; in the majority [80%] of paediatric MS patients, however [1.7% to 5.6% of the total MS population], the onset of disease is between 10 and 18 years [post-pubertal (*juvenile*) MS]. Notably, while pre-pubertal MS occurs almost equally in both genders (female/male ratio = 0.9:1; reverting to 0.4–0.6/1 in pre-school MS children) the female/male ratio rises to 2.2/3:1 in the post-pubertal age. Interestingly, precocious puberty has been associated to: (a) a higher risk of developing MS; and (b) a more severe disease course. In addition to that, males are more susceptible to MS (and manifest more neurodegeneration) than females the latter being however more inflammatory than males; pregnancy however reduces MS relapses. All the above findings led to the suggestion of an underlying female sex hormonal involvement in the pathophysiology of MS vs. a protective role of male sex hormones. Epigenetic perspectives indicate that the interplay between genetic background, environmental triggers and neuroendocrine changes, typically occurring around the time of *adolescence*, could all play a combined role in initiating and/or promoting MS with onset in the paediatric age including many of the most frequent disease-associated risk factors (e.g., overweight/obesity, low vitamin D levels, reduced sunlight exposure, Epstein-Barr virus infection). According to this proposed *complex multifactorial model*, susceptibility to MS may be thus acquired during pre-pubertal age and children have probably to wait until the adolescence to manifest their first clinical signs/symptoms.

Keywords: Paediatric multiple sclerosis, Childhood multiple sclerosis, Early-onset multiple sclerosis, Puberty, Hormones, Pathophysiology, Leptin, PI3K, Demyelination

Background

The *World Health Organisation (WHO) - Multiple Sclerosis International Federation* reported that the interquartile range for signs/symptom onset in MS is between 25.3 and 31.8 years, placing the average age of MS onset at 29.2 years [1]. However, *late-onset* cases have been well documented [2] and the occurrence of MS at the other end of the spectrum of life (i.e., < age 18 years: *childhood MS*) is now well established (1.7% to 10% of total MS patients) [3–19]. A small, but well-established subgroup of paediatric MS cases is younger than (or had the onset of symptoms before) 10 years of age (pre-

pubescent MS: 0.2% to 0.7% of total MS cases) [7, 18] including children with onset of disease in pre-school years [17] and (exceptionally) during early infancy (i.e., < age 2 years) [17, 19]. The mean annual incidence rates for childhood/paediatric MS is at 0.1/100,000–0.9/100,000 [3–16] whilst annual incidence figures for *pre-pubescent* onset MS are at 0.09/100,000 [7, 17–19]. While pre-pubescent MS occurs almost equally in both genders (female/male ratio = 0.9:1; reverting to 0.4/0.6/1 in pre-school MS children, as it occurs in acute disseminated encephalomyelitis - ADEM) [20] the female/male ratio rises to 2.2/3:1 in the post-pubertal age (“*juvenile MS*”: i.e., MS with onset between age 10–18 years) [7, 12, 17–19, 21, 22].

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In the present review, we summarize the gender effects on inflammatory and neurodegenerative processes in MS and the relationship between pubertal hormonal and/or neuroendocrine changes and the risk of paediatric MS.

Pathophysiology of MS and the rationale for disease-modifying therapies

The hallmark of MS is the demyelinated “*MS plaque*” that is unique and different from that seen in other inflammatory diseases and consists of a well-demarcated hypocellular area characterised by the loss of myelin, the formation of astrocytic scars, and the presence of inflammatory mononuclear cell infiltrates, typically concentrated in perivascular, particularly perivenular, cuffs [23–25]. These infiltrates, which are mainly composed of a mixture of innate (CNS-resident) and adaptive (CNS-infiltrating) components of the immune system [24], include [among the *innate* effectors] monocytes/macrophages, dendritic cells, reactive microglial cells, astrocytes, and mast cells, and [among the *adaptive* effectors] autoreactive lymphocyte T cells, B lymphocytes, and plasma cells plus minor additional components (e.g., ependymal cells), which after their migration into the central nervous system (CNS), incite a pro-inflammatory reaction, resulting in local tissue injury, which consists in blood brain barrier (BBB: another innate immune component) leakage, destruction of myelin sheaths, oligodendrocytes damage, and cell death, as well as axonal damage and loss, leading in turn to the glial scar (i.e., to the “*MS plaque*”, as seen at imaging and histopathology) [23].

Thus, the migration and/or activation of (innate and adaptive) pro-inflammatory cells into the CNS represent a key stage in the natural history of MS (but what initiates this event still remains unclear) [23]. From a pathophysiologic viewpoint MS appears to be caused by a contact in early childhood with a pathogen coupled with other individual susceptibility factors (e.g., genetic, racial and demographic background), which can elicit their reactivation, triggering innate mechanisms of defence as *toll-like receptors* (TLRs: membrane-spanning, non-catalytic receptors expressed on sentinel cells - e.g., macrophages or dendritic cells - recognizing structurally conserved molecules derived from microbes), that signalizes downstream through its adapter protein *MyD88* (myeloid differentiation primary response 88), and the phosphorylated/degraded protein *IKB* which permits translocation of *NF-KB* (nuclear factor kappa-light-chain enhancer of activated B cells: a protein complex, which controls DNA transcription, cytokine production and cell survival) and the transcription of pro-inflammatory cytokines such as IL-6, TNF, IL-1, IL-12, E-selectin, MCP-1, and IL-8. TLR through IRF7 (Interferon regulatory factor 7) gives the signal to the transcription of IFN α/β (i.e., the cytokines used for communication between cells to trigger the protective

defences of the immune system). Another important signal is given by NOD receptors (nucleotide-binding oligomerization domain: i.e., a cytoplasmic pattern recognition receptor, which regulates the innate system and cooperates with TLRs) activated also by potassium efflux-inducing agents such as ATP and TLR stimulation. Additional signalling is provided by PAMS/PAMP (pathogen-associated molecule patterns), toxins, danger or stress, whose triggering induce the *inflammasome* (i.e., a cytoplasmic multiprotein oligomer) via NLRP (NOD-like receptor protein) that form a complex with ASC (apoptosis-associated speck-like protein containing a CARD: caspase recruitment domain) and caspase-1 (i.e., the interleukin-1 converting enzyme, which converts the IL precursors into mature active IL proteins), activating IL-1b, a major factor inducing inflammation, autophagy and cell death, particularly necrosis [23].

All the above pro-inflammatory soluble factors activate microglia and endothelial cells [i.e., innate effectors], up-regulating expression of adhesion molecules (e.g., E-selectin), facilitating the migration of T cells into the CNS. Matrix metalloproteinases (MMP) degrades BBB enhancing further migration of autoreactive T cells and macrophages via chemokines (CX3CL-1). The Th1 response evoked via IL-12 and IFN- γ further activates macrophages that in turn do so to T cells CD8+. Th2 response via IL-6 mainly stimulates maturation of B cells and production of autoantibodies. Cytotoxic damage to the oligodendrocyte mediates myelin loss and exposure of the axon to reactive oxygen species, slowing or blocking action potentials and the production of neurological manifestations.

There are intents to remyelinate these lesions via OPCs (oligodendrocyte precursor cells), but neuronal factors such as LINGO-1 (Leucine rich repeat and immunoglobulin-like domain-containing protein 1: a protein important for protein-protein interactions, which regulates/modulates neuronal differentiation and growth, regulation of axon guidance and regeneration processes) or TLR2 inhibit their migration [23–25].

Based on these premises, over the last two decades a dozen different preparations of immunomodulatory/immunosuppressive agents, targeting the above CNS autoimmune mechanisms, have been developed, showing beneficial effects in patients with MS and have been approved as first- or second-line disease-modifying therapies (DMTs), including [24, 26]: (a) [*first-line DMTs*] interferon- β (IFN- β 1a and 1b), glatiramer acetate (GA), dimethyl fumarate (DMF), and teriflunomide; and (b) [*second-line DMTs*] mitoxantrone, fingolimod (a small molecule antagonist against SIP and SIP-receptors inhibiting immune cell trafficking), natalizumab (an alpha-4 integrin blocker of immune cell trafficking/migration), alemtuzumab (an anti-CD52 cell-depleting monoclonal

antibody), daclizumab (a blocker of the interleukin 2R α chain), and ocrelizumab (an anti-CD20 cell-depleting monoclonal antibody) [23, 26].

Although these therapies are able to modulate the immune adaptive response, they do not inhibit innate immune cells (e.g., microglial cells, macrophages, and dendritic cells) that participate in the progression of MS. In addition to that, some of these strategies, with their indiscriminate targeting of both pathogenic and protective immune cells, might have side effects. Several new drugs are imminently emerging including strategies targeting the innate immune system [e.g., inhibition of tyrosine kinase, inhibition of NF κ B, scavengers for active oxygen species and nitric oxide, or pharmacological interference with their production], or targeting the inflammasome [23].

Disease-modifying therapies in pediatric MS

No medication currently approved for adults with (relapsing-remitting) MS has completed testing for pediatric MS in randomized placebo-controlled trials, although several pediatric MS trials have recently been launched [27]. Use of DMTs in pediatric MS remains off-label in many countries, especially in patients younger than 12 years; nevertheless, these medications are widely used. At present, IFN- β and GA continue to be the standard first-line treatments for pediatric patients with MS, as supported by observational studies and experts' consensus guidelines [26, 27]. Trials are on-going evaluating the clinical outcome of pediatric patients with MS treated with fingolimod, dimethyl fumarate, and teriflunomide [27].

Ages at presentation of MS in childhood and the "true" pre-pubertal threshold

Currently, MS in the paediatric age group is divided into two main groups according to the age at presentation of first signs/symptoms [3–19]:

- (1) *Childhood MS* (when the first acute demyelinating event occurs prior to age 12 years);
- (2) *Juvenile MS* (when onset of disease ranges from 12 to 18 years);

A separate group defines (3) *adult MS*, when disease onset is after age of 18 years [1, 2].

The cut-off period up to 12 years to define childhood MS was chosen by most Authors in their studies because this period was (and still is) considered as the pre- or early pubertal period [Tanner stages (i.e., Breast, Genitalia, Pubic hair) I or II]. A restricted number of Authors have proposed, in their studies, a lower cut-off for defining "true" childhood MS at 10 years of age [12, 17, 19]; this (lowered) period better reflects the biological pre-

pubertal period irrespective to gender [Tanner stages (i.e., Breast, Genitalia, Pubic hair) I vs. I or II: Tanner stage I represents the *true* pre-pubertal stage]. By lowering the cut-off period down to 10 years one could be surer: (a) to exclude early pubertal children in analysis of paediatric MS cases, thus avoiding inclusion of MS patients already targeted by the postulated effects of pubertal sex hormones on predisposed tissues [e.g., bone marrow, thymus, central nervous system]; and (b) to limit multiple viral exposures as by age 10 years most children (e.g., in Italy) have usually completed their vaccination schedule of mandatory and recommended vaccines [17, 19].

Pre-pubescent onset MS is characterised by peculiar clinical, laboratory and imaging features and outcome [17, 19, 28], including inversion of sex ratios, low to null family history for MS, preponderance of atypical manifestations at onset (e.g., hemiparesis, seizures, lethargy, brainstem signs/symptoms or cerebellar ataxia), polyfocal presentation, highest relapse number/year and fastest recovery time, more severe neurological deficits at relapses with more completely or near-completely recovery, ADEM/leukodystrophy-like MRI patterns at onset vs. typical MS MRI patterns attained years after the first attacks, a worse outcome in the earliest onsets (i.e., < 2 years of age) vs. a better outcome (as compared to post-pubertal MS) in onsets at toddler ages.

Age- and gender-related peculiarities of pediatric MS vs. similar disorders

A peculiar female responsiveness to environmental triggers is noted across many disease models and is usually attributed to the need, in the female gender, to make repeated, rapid and consistent physiologic accommodations to pregnancy. In female adolescents with MS, a number of genetic, non-genetic and lifestyle factors have possibly sexually dimorphic effects on MS disease predisposition and on its clinical course [29, 30].

Similarly to what occurs in MS, the so-called *pseudotumour cerebri syndrome (PTCS)* is a neurological disorder, which, within childhood, mostly affects post-pubertal females, who often are overweight. PTCS is a condition of unclear aetiology, characterised by increased intracranial pressure (ICP) without any radiographic evidence of brain tissue abnormalities, and with normal chemical and cytological cerebrospinal fluid (CSF) composition [31–33]. Multiple causes have been taken into consideration in the pathophysiology and aetiology of PTCS [32, 33] including obesity, endocrine abnormalities (e.g., hyperaldosteronism, Cushing syndrome, hyperandrogenism, Addison disease), kidney disease (e.g., nephrotic syndrome), systemic disease (e.g., systemic lupus erythematosus, Guillain-Barré syndrome, antiphospholipid antibody syndrome, polycystic ovary syndrome - PCOS, Behcet disease, familial

Mediterranean fever), medications (e.g., recombinant growth hormone therapy, tetracycline, steroids, mycophenolate mofetil, vitamins A and D, cytarabine, and cyclosporine A), viral infections (e.g., chickenpox, measles, reactivation of varicella infection) and changes in CSF volume and in cerebral CSF hemodynamic (increased cerebral blood volume, increased cerebrospinal fluid production, decreased cerebrospinal fluid resorption or venous flow abnormalities); PTCS has been also observed in members of the same family presenting in either an autosomal dominant or recessive manner. A recently proposed *unifying (neuroendocrine) hypothesis* inferred that [32] multiple neuroendocrine interactions (e.g., cortisol, aldosterone, progesterone) could influence the activation of the mineralocorticoid receptor (MR) in the choroid plexus epithelial cells, which in turn stimulates (via a nuclear pathway) the ATPase/Na⁺/K⁺ pump leading to raised intracranial CSF production [25]. Even though it typically affects both genders and all age groups, the post-pubertal PTCS typically occurs in overweight girls/women during their reproductive age [34]. Notably, the overall incidence of PTCS is estimated to be 0,9/100,000 rising to 19/100,000 in overweight women [34, 35].

Paediatric PTCS is known to occur in association with a broad variety of conditions, especially obesity and endocrine derangements (e.g., cortisol deficiency or excess, hyperandrogenism, hyperaldosteronism) [32, 36]. Although pre-pubertal PTCS can occur in both genders and ages, post-pubertal PTCS is usually recorded in women during their reproductive age [34, 35]: in this respect, it has been previously proposed that the proneness of some women to develop PTCS could be linked to an estrogenic gynecoid (pear-shaped) fat distribution [34]. Adipose tissue contains aromatase, which may be a link between obesity and PTCS. Aromatase, which catalyses the production of oestrogens from plasma androstenedione, is more prevalent in the fat of the buttock regions (reflecting the typical female fat distribution) vs. the abdominal (visceral) regions [34–36]. Of note, the reports of the onset of PTCS in postmenopausal women following the initiation of hormone replacement therapy further support the notion of an oestrogen involvement in the pathophysiology of this condition [31–33, 37, 38].

The PTCS neuroendocrine pathophysiology [32] cannot be applied to MS, as the mechanism underlying the rise of CSF pressure cannot be compared to the process of demyelination and unlikely involves an autoimmune aetiology [39]. Nonetheless, higher values of ICP have been recently documented in the paediatric MS population [31, 32], thus reflecting the fact that both these conditions (PTCS and MS) could share similar precipitating factors (e.g., obesity, female sex hormones) on a background of alike clinical and anthropometric features. A tenable hypothesis of common

possible trigger(s) underlying both diseases could be represented by the putative involvement of *Leptin*, which seem to be centrally involved either in PTCS and in MS pathophysiology [7, 32] (Fig. 1).

The role of *gender* factors in paediatric MS

The sex discrepancy (with a female preponderance) in MS is evident only in individuals who manifest disease symptoms after puberty [1, 7, 17, 19], implicating a likely role of female sex hormones in initiating and/or promoting the disease [7] and of post-pubertal male (high) levels of testosterone in protecting from the disease [40]. The above notion is supported by a number of clinical and laboratory evidences: (a) men with MS present at an older age, concurrent with the start of the age-related decline in testosterone levels; (b) a decrease of androgen levels in MS adult males is associated with a more severe disease course and a faster progression to disability; and (c) testosterone administration may ameliorate the clinical course of MS in males [41, 42].

Oestrogens (17 β -estradiol-E2- and estriol-E3), progesterone and testosterone may provide anti-inflammatory and neuroprotective effects on induction and effector phases of experimental allergic encephalomyelitis (EAE) [29, 30]. Anti-inflammatory effects appear mainly mediated by oestrogen nuclear receptors alpha (ER α) and beta (ER β) expressed by regulatory CD4+ CD25+ T cells (Treg), regulatory B (Breg) cells and dendritic cells and may be abrogated in the absence of B cells and the co-inhibitory receptor, Programmed Death-1 (PD-1) on CD4+ Foxp3+ Treg cells. E2 protective effects on EAE seem to be mediated by binding to the membrane G-protein-coupled receptor 30(GPR30). Testosterone may work through androgen receptors or after its conversion to oestrogen through ERs, or GPR30. Androgens may induce remyelination in cuprizone-induced CNS demyelination by acting on neural androgen receptors. Experimental studies also showed that androgens exert a protective role against the development of EAE, the animal model of MS [30]. Additionally, therapeutic trials with dihydrotestosterone (DHT) in castrated animals ameliorate both symptoms and inflammation [29, 30, 40].

Some neuroprotective effects of oestrogens in EAE are mediated by ER α expressed on astrocytes: ER β ligands can prevent demyelination and stimulate remyelination and ER β treatment can affect microglia with protective effects in CNS inflammation. Progesterone appears to affect axonal protection and remyelination, and testosterone can restore synaptic transmission deficits in the hippocampus.

Sex hormones play a pivotal role in the human *immune system*, regulating antigen presentation, cytokine gene expression, lymphocyte activation and autoimmune

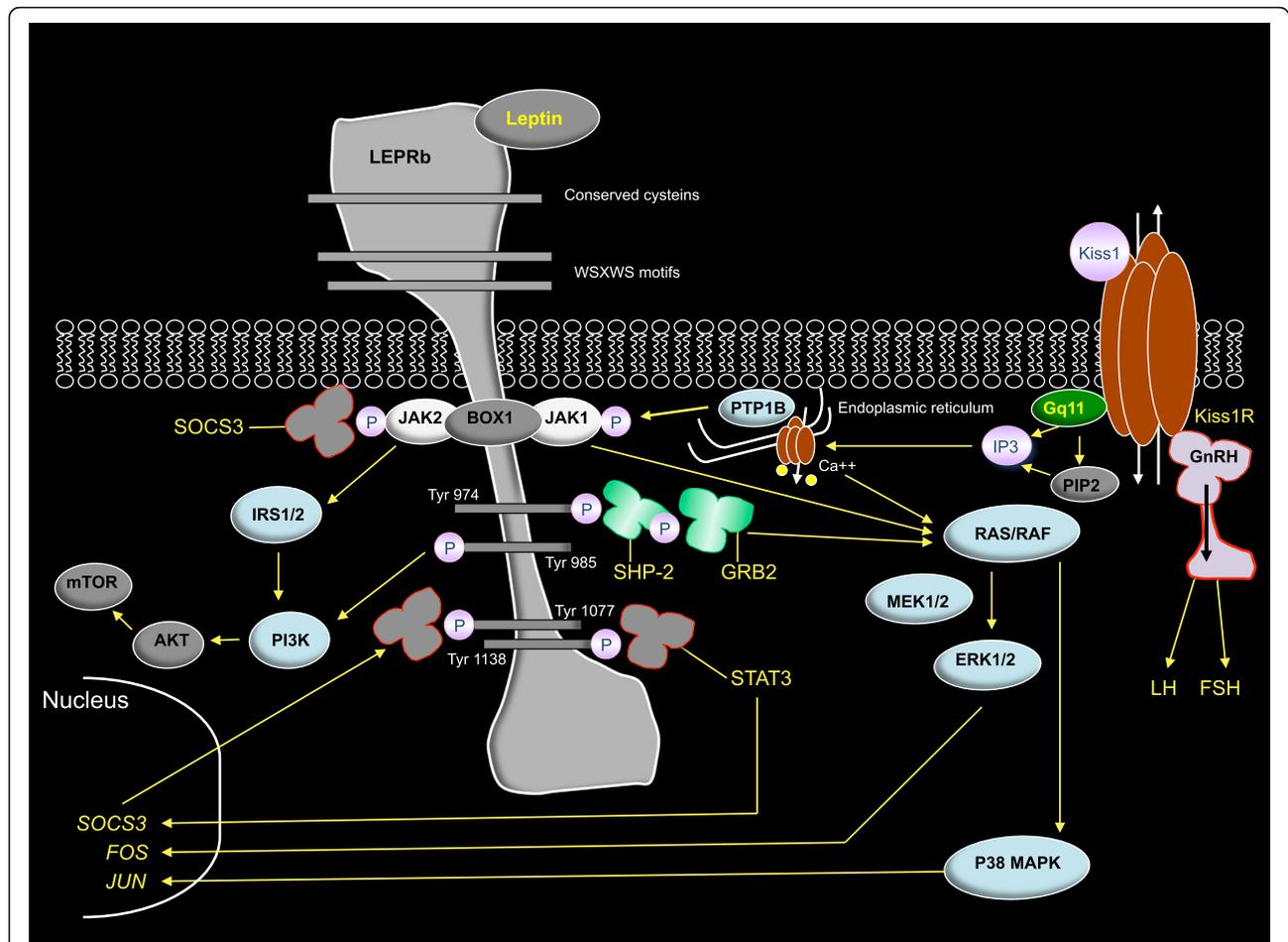


Fig. 1 Mechanisms of Leptin signalling in immune and neuroendocrine cells. Leptin binds to one of its receptors, LEPRb, activating JAK2 by auto-phosphorylation or cross-phosphorylation, and phosphorylates tyrosine residues in the receptor's cytoplasmic domain. Four of the phosphorylated residues [974, 985, 1077, 1138] function as docking sites for cytoplasmic adaptors for STAT factors, particularly STAT3, which dimerizes translocating into the nucleus, where it induces expression of *SOCS3*, *FOS* and *JUN* genes. *SOCS3* participates in a feedback loop that inhibits Leptin signalling by binding to phosphorylated tyrosines. *SHP-2* is recruited to Tyr985 and Tyr974 and activates ERK1/2 and p38 MAPK pathways through the adaptor protein GRB2, ultimately inducing *FOS* and *JUN* gene expression [*FOS* and *JUN* encode for fos and jun proto-oncogene proteins, which form heterodimers (C-fos:c-jun) resulting in the formation of AP-1 (Activator Protein-1) complex, which binds DNA at AP-1 specific sites at the promoter and enhancer regions of target genes and converts extracellular signals into changes of gene expression]. *PTP-1B* is localized on the surface of the *endoplasmic reticulum*, and is involved in negative regulation of LEPRb signalling through dephosphorylation of JAK2 after internalization of the LEPRb complex; the endoplasmic reticulum is also the site of action (via Ca^{++}) of the IP3-PIP2-mediated pathway of the Kiss1 protein, which in turn modulates GnRH secretion and ultimately LH and FSH secretion [neuroendocrine cells are hereby represented as if they were inside the membrane for practical purposes: in the real pathways the Kiss1 protein binds to the Kiss1 receptor (R), which is expressed on the membrane surface of both immune and neuroendocrine cells: the latter cells promote secretion of GnRH, which in turn stimulate secretion of LH and FSH]. JAK2 can also induce phosphorylation of the IRS1 and 2 proteins, which are responsible for PI3K/AKT and mTOR pathway activation

processes [30, 41]. Also, immune central tolerance at the thymus level is strictly dependent on the hormonal status [29, 30, 42]. Elevation of sex steroids during puberty has been, de facto, linked to the typical decline of the thymus, which starts around adolescence; the thymus rejuvenation after ablation of sex steroids further supports this notion [29, 30, 43]. It is unsure whether puberty and its related hormonal changes affect the susceptibility to environmental factors such as infections.

There are gender-related differences in immune response and women have higher levels of immunoglobulin and more vigorous T-cell activation when compared to males [44]. Oestrogens appear to have a controversial role on inflammation in EAE. At lower levels, oestrogens - such as estradiol - may promote inflammation; but at higher levels, oestrogens - such as the pregnancy hormone estriol - may induce a shift in the immune response from a T helper 1 (TH1) response to a T helper 2 (TH2) response, muting inflammation [45]. This

would explain the reason for which disease activity usually decreases during late pregnancy, which is typically characterized by high levels of estradiol and also the beneficial effects of estradiol administration to non-pregnant MS females in improving the disease manifestations [46–48].

Studies in EAE also show that low dose oestrogen therapy may have also profound effects in inhibiting the development of autoimmunity, likely influencing the immune reaction towards a protective anti-inflammatory cytokine response [29, 30]. However, in one of these studies, oestrogen treatment at the onset of active EAE failed to reduce disease severity, a result that is consistent with the hypothesis that naive cells are more sensitive to sex hormones than differentiated effector cells [49].

Of note, post-pubertal EAE female mice develop increased myelin reactive T-cell responses compared to age-matched mice that had been prevented from entering puberty via pre-pubertal ovariectomy surgery [49, 50]. Together, these studies suggest that puberty in females enhances central nervous system (CNS) autoimmune mechanisms, further explaining the female preponderance of MS, at the post-pubertal ages.

Lastly, the role of the female chromosome X on immunity and MS should be also regarded as crucial. This could involve hormones-independent mechanisms, including microRNAs and cytokine genes present on chromosome X [51].

Precocious puberty and the risk of MS

Recent MS studies, further deepened, the (causal) relationship between puberty and the disease: initially, only the peri-pubertal period was regarded as typically associated to a dramatic, female-specific, rise in disease incidence; later studies, however, demonstrated that an earlier occurrence of puberty and menarche was also associated to higher risks of disease onset and a more severe clinical course. One study demonstrated that the age at first symptoms increased by 1.16 years as the age of menarche increased by one year [52]. A further MS collaborative Canadian study showed that females with MS were younger at menarche (i.e., 12.4/12 years vs. 12.6/12 years) compared to controls [53]. An association between earlier age at menarche in females and a more severe disease course has been also recorded [54].

A potential effect of age of puberty and menarche on MS, further strengthens the putative involvement of female sex hormones in disease pathophysiology, due to oestrogen-related changes in CNS and immune system (as outlined above). Thus, an earlier menarche may possibly upset a delicate oestrogen balance, making some susceptible girls prone to develop MS.

However, the question whether younger age at puberty is a real trigger for the disease or a mere trigger factor on a background of multiple genetic and environmental determinants remains unsolved. Additionally, it has also been speculated that earlier menarche is a surrogate for the effect of an MS disease causative factor that influences the risk of MS independently by oestrogens, whilst affecting the age of menarche as a by-product [53].

Of note, puberty onset requires specific changes in the secretion of the pituitary gonadotropins, luteinizing hormone (LH) and follicle-stimulating hormone (FSH), which are dependent on the release of Gonadotropin releasing hormone (GnRH) from the hypothalamus (Fig. 1); thus, timing of puberty is strictly dependent on a specific genetic susceptibility and on environmental conditions that can influence physiological and pathological processes acting on the hypothalamic-pituitary axis, including nutrition, adiposity, bone mass, emotional and psychological factors, light-darkness cycles/melatonin and endocrine disrupting compounds [55].

Interestingly, also low vitamin D levels are usually associated with an earlier age of pubertal onset in the paediatric population [56]. In this context, it is still unclear whether environmental exposures affecting puberty timing also affect the risk of developing MS in an independent manner. Thus, the correlation between precocious timing of puberty and menarche and the risk of onset and/or worsening of MS and the higher prevalence of obesity within the paediatric MS population are overall factors, which reflect a possible underlying endocrinology/metabolic involvement in the pathophysiology of MS.

Some important MS-associated risk factors (i.e., low vitamin D level, obesity) are also known to be causes of early puberty per se, further supporting the possibility that earlier puberty is a surrogate for the effect of an MS disease causative factor that influences both the risk of MS and an earlier menarche.

Obesity and Leptin: Correlations between metabolism and paediatric MS

Paediatric obesity has been demonstrated in one study to be a risk factor for later development of adult-onset MS in women whilst obesity occurring in adulthood carried out a null risk of developing MS [57]. Another study found that paediatric obesity was independently associated with an increased risk of paediatric onset MS in girls but not in boys: the association between body mass index (BMI) and paediatric MS was strikingly pronounced in extremely obese adolescent girls [7].

Despite these findings, still there are many underexplored and/or not yet fully understood aspects on these relationships. The relative percentages of body fat during the paediatric age are known to be associated with

accelerated sexual maturation and precocious puberty: it is unclear if overweight and obesity may predispose independently to both earlier puberty and MS, or if it happens in a consecutive manner [58, 59]. Obesity is characterized by a low-grade inflammation state and it is known to be associated with a T helper 17 (**Th17**) bias predisposing to autoimmune reactions [60]. Additionally, interactions between obesity and vitamin D status remains unexplored [47].

Adipose tissue is not an inert tissue implied only in energy storage, but can be regarded as a part of an *endocrine organ*, which releases many mediators that in turn may predispose to both puberty and MS; additionally, some of these mediators and/or adipokines released by adipocytes are involved in several inflammatory processes, including tumor necrosis factor alpha (**TNF- α**), interleukin 6 (**IL-6**) and Leptin. The adipokine hormone **Leptin** is an amino-acid cytokine-like protein, which is known to play a crucial role in regulating puberty, especially in females. At central (hypothalamic) level, Leptin facilitates puberty onset likely stimulating the Kisspeptin1 (**Kiss1**) pathways, the upstream regulators of GnRH neurons [55] (Fig. 1).

Besides its metabolic role in promoting puberty onset, Leptin has many additional central and/or peripheral actions, including regulation of both innate and adaptive immunity. In fact Leptin stimulates the secretion of pro-inflammatory cytokines (e.g., IL-6, IL-18) and at the adaptive immune system level, Leptin promotes switch towards pro-inflammatory Th1 immunological responses [61].

Leptin mediates its effects by binding to Leptin receptors (**LepRs**) expressed in the brain and a in wide array of peripheral tissues. Various alternatively spliced isoforms of LepRs have been described, but the long isoform of Leptin receptor (**LepRb**) is primarily responsible for Leptin signalling (Fig. 1). The binding of Leptin to LepRb activates a number of signalling pathways, including AK2/STAT 3 and STAT5, SHP2/MAPK and PI3K/AKT/mTOR [59–61]. Notably, the activation of the phosphatidylinositol-3 kinase (**PI3K**) pathway by Leptin is one of the most studied effects of Leptin signalling in the brain and it has been demonstrated to play crucial roles in several metabolic and energetic processes [61]. A number of studies have demonstrated the relevance of PI3K as an underlying mechanism of Leptin actions in vivo [52]. In rats, peripheral Leptin administration was found to activate PI3K in the brain and pre-treatment with inhibitors of PI3K abolished the anorectic response induced by Leptin [61–63].

Interestingly, the balance of PI3K/AKT pathway is essential for oligodendrocytes survival and axon myelination and *gain of functions* mutations of genes enclosed in this pathway (especially mutations in *PIK3CA* gene) have been linked to various types of overgrowth

syndromes [known as *PIK3-related overgrowth syndromes*, or **PROS**] [64, 65], which are also characterized by diffuse white matter abnormalities and increased signal on T2-weighted images on MRI [66, 67]. Studies that investigate potential interactions between Leptin and the PI3K signalling in MS patients are needed.

One of the principal observations, which indicate that Leptin could represent a key mediator in the pathogenesis of MS, is due to its female-specific rise during the peri-pubertal age. In fact, numerous studies showed that during pubertal age Leptin levels continue to increase in girls but not in boys due to the testosterone-related inhibition on Leptin secretion [63]. Besides its interaction with the PIK/AKT pathway, possibly implicated in early cytodegenerative processes of myelin, the actions of Leptin include a strong influence in both the innate and the adaptive immune system. In the innate system, Leptin stimulate the activation of the monocyte-macrophage lineage and the secretion of pro-inflammatory cytokines; in the adaptive immune system Leptin induces pro-inflammatory Th1 responses [61, 62]. Interestingly, the Leptin deficient mice are resistant to the induction of EAE, and administration of Leptin in this animal model shifts the Th2-type response, characteristic of this animal model, to a Th1-type response [68].

Moreover, it has been observed that Leptin is able to maintain environmental conditions that promote loss of immune self-tolerance [69, 70]; in particular, both in vitro and in vivo, leptin can affect the generation, proliferation and responsiveness of t_{reg} cells, a key type of t cells that is involved in the control of immunological tolerance [71].

Thus, the crucial involvement of leptin in initiating and promoting puberty, the observation that it continues to rise in female but not in male adolescents, the increased levels of leptin in the pediatric obese population, the central role of this hormone in regulating inflammatory and autoimmune processes, the demonstration of its necessary role for the induction of the animal model of MS (i.e., EAE), are all convincing evidences of the involvement of leptin in pathogenesis of MS, especially in the post-pubertal pediatric age group.

Conclusions and future directions

In conclusion, within the post-pubertal MS group both the disease prevalence and the female-male ratio are much higher if compared to the pre-pubertal MS group. Furthermore, a more precocious onset of puberty has been associated to both a higher risk of developing MS and an even more severe disease course. Additional support linking puberty with the pathogenesis of MS may be driven from the observations of remarkable similarities between the neuroendocrine mechanisms underlying the onset of puberty and those associated with the

postpartum period [72]. In fact, the hypogonadotrophic state of the postpartum phase resembles the pre-pubertal (hypogonadotrophic) state [73, 74].

Interestingly, the postpartum recovery of gonadotropin release follows a predictable sequence of a preferential rise of FSH followed by LH secretion, a pattern identical to that of the peri-pubertal state [72]. For these reasons, the neuroendocrine changes in the postpartum period have been also known as “*puberty in miniature*” and have been frequently associated to the significantly increased risk of onset and relapse of MS after partum. Thus, it is possible that the biological mechanisms, which are responsible for the development of the clinical manifestations of MS in the pubertal or post-pubertal periods, are also involved in the onset or the reactivation of the disease in the post-partum period.

Thus, recent researches in the field suggest that the neuroendocrine changes, typically occurring around the time of puberty, could play a role in initiating and/or promoting pediatric MS associated with additional genetic/non-genetic (e.g., environmental) factors.

According to this multifactorial model, susceptibility to MS may be thus acquired during a wide window of risk through childhood and most pre-pubertal children acquiring susceptibility to the disease have probably to wait until the “*pubertal switch*” to manifest the clinical symptoms. Further experimental studies are required in future to fully understand the gene-neuroendocrine-immune-environment-lifestyle interactions underlying the molecular pathobiology of pediatric-onset MS.

Abbreviations

AKT: AK (Akr mouse) strain transforming; FOS: Finkel osteogenic sarcoma [Finkel-Biskis-Jinkins murine osteogenic sarcoma virus]; FSH: Follicle stimulating hormone; GnRH: Gonadotropin [LH/FSH] releasing hormone; Gq11: Guanine nucleotide binding protein q 11; GRB2: Growth factor receptor-bound protein 2; IRS: Insulin receptor substrate; JAK2: Janus kinase 2; JUN: Jinkins avian sarcoma virus oncogene; Kiss1: Kissprotein 1; Kiss1R: Kissprotein receptor; LEPRb: Leptin receptor, long form b; LH: Luteinizing hormone; MAPK: Mitogen-activated protein kinase; MEK2: MAPK-extracellular kinase 2; mTOR: Mammalian target of rapamycin; PI3K: Phosphatidylinositol-3-kinase; PIP3: Phosphatidylinositol tri-phosphate; PTP-1B: Tyrosine-protein phosphatase non-receptor type 1; RAF: Rapidly accelerated fibrosarcoma protein; RAS: *Rat sarcoma viral (V-ras)* oncogene homolog; SHP-2: SHP protein tyrosine phosphatase-2; SOCS3: suppressor of cytokine signalling 3; STAT3: transducer and activator of transcription 3; IP3: Inositol 1,4,5-trisphosphate receptor, type 3; ERK1/2: Extracellular-signal regulated kinases 1 and 2

Acknowledgements

We wish to thank Dr. G.H. Tutino (Catania) for his valuable comments and support.

Funding

None

Availability of data and materials

Not applicable

Authors' contributions

VS and MR conceived the review, participated in its design and coordination and wrote the initial draft. VS and GR reviewed the existing literature. VS and

AP drafted the final version. AP along with MR designed and drew Fig. 1. All authors read and approved the final manuscript.

Ethics approval and consent to participate

This was submitted to and approved by the Ethical Committee (Catania 1) based at the AOU “Policlinico-Vittorio Emanuele”, Catania.

Consent for publication

All Authors were informed and gave their consent to publication in MSDD.

Competing interests

The authors declare that they have no competing interests.

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Received: 2 March 2017 Accepted: 10 November 2017

Published online: 22 February 2018

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