

**Figure 1.** Inductors, markers and functions of macrophages subtypes in tissues. All macrophage subtypes may take a pro- or anti-inflammatory phenotype in obesity. MoMFs are a common precursor for subtypes of migrating macrophages in tissue. When damaged, MoMFs can assume a tissue-specific phenotype. Determining the exact origin of macrophages (resident or migratory) is a task in modern science. The plasticity of macrophages between M1 and M2 is also present in the resident forms and is shown by an arrow. The arrow indicates the stimuli, markers and functions that characterize both types of polarization. ABCA1—ATP binding cassette subfamily a member 1; AP—atherosclerosis plaque; AT—adipose tissue; DAMPS—damage-associated molecular patterns; GCLM—glutamate–cysteine ligase modifier subunit; GST—glutathione S-transferase; Hb/Hp—hemoglobin/haptoglobin; HO-1—heme oxygenase 1; IL—Interleukin; KC—Kupffer cells; LCM—liver capsular macrophages; LPM—large peritoneal macrophages; LPS—lipopolysaccharide; MAOA—monoamine oxidase A; MoMF—monocyte-derived macrophage; Mox—oxidized macrophages; MR—mannose receptor;

NAM—neuro-associated macrophages; NE—norepinephrine; Nox2—NADPH oxidase-2; OxPL—oxidized phospholipids; PLIN2—Perilipin 2; PPAR $\gamma$ —peroxisome proliferator-activated receptor gamma; ROS—reactive oxygen species; SAM—sympathetic neuron-associated macrophages; SLC6A2—carrier family 6 member 2; SNS—sympathetic nervous system; Srnx-1—sulfiredoxin-1; TH—tyrosine hydroxylase; TLR—Toll-like receptor; Txnrd1—thioredoxin reductase 1; MMe—metabolically activated macrophages; \*—mouse-specific markers; #—cannot be M1 and M2. This figure has been created by modifying the templates from Servier Medical Art (<https://smart.servier.com>).

## 2. Macrophage Subpopulation Mhem

During the endocytosis of the Hb/Hp complex, heme is released from erythrocytes, which stimulates the transition of macrophages to the Mhem phenotype (**Figure 1**). The main markers for Mhem macrophages are heme oxygenase 1 (HMOX1) and CD163 [5][6]. Mhem macrophages stimulate activating transcription factor (ATF)-1 in various ways, promoting LXR $\beta$  (NR1H2) and HMOX1 expression in humans. This process increases the expression of LXR $\alpha$  and ABCA1, which subsequently increase cholesterol efflux, associated with increased production of IL-10 and apolipoprotein (Apo) E [5][6][7]. Moreover, Mhem have an increased adaptation to plaque hemorrhage. Thus, the M (Hb) and Mhem phenotypes prevent foam cell formation and oxidative stress.

## 3. Macrophage Subpopulation M4

Another type of macrophage, iron-loaded M4 macrophages, is predominantly detected in areas of neovascularization in atherosclerotic plaques (**Figure 1**). After migration through the endothelium, monocytes under the influence of CXCL4 differentiate into M4. The main markers of M4 macrophages are CD68+MR+ [5][6][8] (**Figure 1**).

M4 macrophages are called CXCL4-differentiated macrophages that express the phenotypic markers metalloproteinase 7 (MMP7) and calcium-binding protein S100A8 [9]. At the same time, M4 macrophages do not express CD163 and exhibit low expression of the scavenger receptors CD36 or SR-1, which leads to a failure to induce the expression of the atheroprotective protein HMOX1 when cells are exposed to the Hb/Hp complex [5][6][10]. Thus, M4 macrophages have a pro-atherogenic profile and can be involved in complications of late atherosclerosis, such as acute coronary syndrome and arterial thrombosis. They produce the enzyme MMP12, which can be involved in the degradation of the fibrous coating of the plaque and the plaque destabilization. Furthermore, M4 macrophages express IL-6 and TNF- $\alpha$ , which increase inflammation. However, the fundamental role of M4 cells in atherogenesis is unknown and requires research [5][6].

Using scRNA-seq technology, a new type of macrophage, Trem2hi, has been identified, which is characterized by high expression of Trem2 (triggering receptors expressed on myeloid cells 2), Spp1 (secreted phosphoprotein1), CtSL (cathepsin L) and CD9. The number of Trem2hi macrophages in plaques decreases with a high-fat diet [11]. Trem2 controls the expression of genes associated with energy metabolism and lipid catabolism [12].

Thus, in individuals with obesity, the structure and function of the cardiovascular system adapt to excess body weight. Metabolic disorders, such as obesity, are accompanied by endothelial cell dysfunction and decreased vascular density [13]. The modern paradigm argues that metabolic changes are associated with obesity secondary to endothelial dysfunction. The hypothesis regarding the ability of the endothelium to cause metabolic dysregulation itself must be revised and supplemented.

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