Noninvasive Tracking of Hematopoietic Stem Cells in a Bone Marrow Transplant Model

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Received: 20 February 2020; Accepted: 03 April 2020; Published: 10 April 2020

Abstract: The hematopoietic stem cell engraftment depends on adequate cell numbers, their homing, and the subsequent short and long-term engraftment of these cells in the niche. We performed a systematic review of the methods employed to track hematopoietic reconstitution using molecular imaging. We searched articles indexed, published prior to January 2020, in PubMed, Cochrane, and Scopus with the following keyword sequences: (Hematopoietic Stem Cell OR Hematopoietic Progenitor Cell) AND (Tracking OR Homing) AND (Transplantation). Of 2191 articles identified, only 21 articles were included in this review, after screening and eligibility assessment. The cell source was in the majority of bone marrow from mice (43%), followed by the umbilical cord from humans (33%). The labeling agent had the follow distribution between the selected studies: 14% nanoparticle, 29% radioisotope, 19% fluorophore, 19% luciferase, and 19% animal transgenic. The type of graft used in the studies was 57% allogeneic, 38% xenogeneic, and 5% autologous, being the HSC receptor: 57% mice, 9% rat, 19% fish, 5% for dog, porcine and salamander. The imaging technique used in the HSC tracking had the following distribution between studies: Positron emission tomography/single-photon emission computed tomography 29%, bioluminescence 33%, fluorescence 19%, magnetic resonance imaging 14%, and near-infrared fluorescence imaging 5%. The efficiency of the graft was evaluated in 61% of the selected studies, and before one month of implantation, the cell renewal was very low (less than 20%), but after three months, the efficiency was more than 50%, mainly in the allogeneic graft. In conclusion, our review showed an increase in using noninvasive imaging techniques in HSC tracking using the bone marrow transplant model. However, successful transplantation depends on the formation of engraftment, and the functionality of cells after the graft, aspects that are poorly explored and that have high relevance for clinical analysis.

Keywords: hematopoietic stem cell; nanoparticle; homing; tracking; near-infrared fluorescence image; magnetic resonance image; bioluminescence; molecular imaging; noninvasive imaging
1. Introduction

Studies from the early 1950s established that total body irradiation in animal models causes death from hemorrhage and infection, indicating that the hematopoietic system is primarily affected [1]. However, it was also shown that transplantation of genetically identical (i.e., syngeneic) bone marrow cells rescues these animals from death induced by irradiation [1].

Later on, Edward Donnal Thomas and colleagues pioneered the application of the results from these early animal studies for the treatment of leukemia in humans. The approach used here was to kill leukemic cells by high-dose irradiation, followed by restoration of the hematopoietic system with bone marrow transplantation [2]. These early findings provided the rationale for using hematopoietic stem cell transplantation (HSCT) as the first stem cell-based therapy for the treatment of a wide plethora of hematopoietic disorders.

According to a comprehensive report from the Worldwide Network for Bone Marrow Transplantation (WBMT), by the end of 2012, more than one million patients had undergone HSCT [3]. The vast majority of HSCT transplantation procedures were used to treat malignant disorders (87%), most of them leukemias (72%), followed by lymphoproliferative disorders (14.7%) and solid tumors (0.6%) [3]. It is noteworthy that HSCT also cures several genetic diseases, such as severe combined immunodeficiency, Wiskott–Aldrich syndrome, thalassemia, and sickle-cell anemia [4].

The dissemination of HSCT as a therapeutic modality is closely linked to the identification and typing of the major histocompatibility complex (also termed human leukocyte antigens (HLA)) in the early 1960s. As a consequence of these discoveries, allogeneic transplantation of HSCs between HLA-matched individuals became feasible. Indeed, almost half of HSCT procedures are allogeneic according to the latest global survey [3].

Allogeneic HSCT comes with the risk of developing a serious immune reaction termed graft versus host disease (GVHD), in which alloreactive donor T cells attack the recipient’s tissues [5]. GVHD is the primary immune barrier to allogeneic HSCT efficacy and is the second cause of death in patients that undergo this procedure, falling behind only mortality caused by the primary disease [6]. For autologous HSCT, on the other hand, the main factor limiting its efficacy is graft failure. Graft failure is a rare complication of HSCT and may be caused by several factors, such as a low dose of injected HSCs, old HSC donors, bone marrow fibrosis in the recipient, storage techniques affecting HSC integrity, and pre-HSCT treatment with chemotherapy and/or irradiation [7].

Studying the dynamics of HSC engraftment and of its progeny is of paramount importance to define the mechanistic basis of hematopoietic reconstitution and the complications of HSCT. For a long time, most of the data regarding the engraftment and expansion of HSCs in small animals were obtained from post-mortem analysis of hematopoietic organs. With the development of imaging hardware, molecular imaging, and imaging reporters, HSCs could be tagged with radioisotopes, fluorophores, contrast agents, reporter constructs, ligands, probes, or cell transduction by reporter genes that codified proteins, such as GFP and luciferase, allowing migration homing and tracking upon transplantation [8–12]. These advances have provided both temporal and spatial information of experimental HSCT that no other technique could provide. In addition, molecular imaging involves a set of noninvasive techniques that allow a serial analysis of the same individual, thereby significantly decreasing the number of animals used for experimentation [13–15].

To study the dynamics of hematopoietic reconstitution, in which cells localize deep inside hematopoietic organs and other tissues, different modalities of molecular imaging are used for HSC tracking and engraftment, such as (i) optical imaging represented by bioluminescence imaging (BLI) [9,11,16–20] and fluorescence imaging (FLI) [21–25], (ii) nuclear imaging represented by positron emission tomography (PET) [26–28] and single-photon emission computed tomography (SPECT) [29–31], and (iii) magnetic resonance imaging (MRI) [32–34].

BLI generally provides high sensitivity with a high signal-to-noise ratio when compared to fluorescent imaging [35]; moreover, its tissue penetrability in higher than that of fluorescence imaging [14,36], and has a high throughput for imaging of small animals [9,15], as also a wide temporal detection window (0 days to 1 year)[9], despite its use being limited to preclinical studies. In vivo BLI enables real-time monitoring of gene expression and cell fate through visual
representation of the bioluminescence generated by oxidation of specific substrates by luciferase enzymes, providing a dynamic profile of engraftment and proliferation in live recipient animals [9,37]. Thus, cells genetically engineered to express luciferase can be tracked after substrate injection using highly sensitive charged-coupled device (CCD) cameras to detect light [36]. On the other hand, FLI can be obtained using fluorescent probes, which, when in the visible length, makes it possible to detect by intravital microscopy techniques, and when the probes emit a fluorescence signal in the wavelength between 650 and 900 nm (infrared spectrum), it is possible to acquire images of deeper tissues compared to conventional fluorescence imaging [38,39]. Thus, the near-infrared fluorescence (NIRF) imaging has a temporal resolution in the order of s to min, a spatial resolution of 2 to 3 mm with a depth of penetration less than 1 cm, low sensibility, no use of ionizing radiation, and low cost. The NIRF disadvantages are the same as that of the BLI modality (low spatial resolution and low tissue penetration), and also as that of autofluorescence [40,41]. Most of the studies used this technique for early HSC engraftment evaluation in the mouse models, but mainly fishes [21–25].

The nuclear image modalities, PET and SPECT are extremely valuable for investigating molecular processes in vivo, which are based on the administration and detection of radioisotopes. However, the HSCT tracking time depends on the radioisotope half-life, resulting in a small temporal window in signal detection. These techniques have good sensitivity, high penetration without depth limit, and can be translated into clinical practice, but they have a low spatial resolution, require the use of ionizing radiation, and have a high cost [14,42,43].

MRI provides excellent anatomical resolution, in vivo cell tracking and graft-host with 3-dimensional information, high penetration depth, does not require the use of ionizing radiation, and is easy to translate into clinical application, although this modality is limited by low sensitivity and high instrumentation cost, as well the tracer dilution upon cell division. MRI is now emerging and rapidly expanding in all fields. In addition, it has the advantages of safety, high spatial and temporal resolution, and direct applicability to cell tracking in clinical studies. MRI specificity can be increased by the use of the contrast agents with magnetic character (paramagnetic and superparamagnetic) [44–48].

Molecular imaging techniques vary in terms of their characteristics of detecting the corresponding signal (specificity, sensitivity, and depth), but technological advances in the development of new multifunctional contrast agents have allowed the use of combined imaging techniques, providing more information and complementing the results obtained, thus culminating in greater clarity of the location of the cells, in addition to allowing co-registration [46,49,50].

In addition, luciferase can be coupled to nanoparticles and fluorescent proteins to create multimodality reporters or brighter probes [51]. One interesting example is the process of generating bioluminescence resonance energy transfer (BRET) upon the fusion between Renilla luciferase and the fluorescent protein Venus. The resulting structure is an auto-illuminated fluorescent probe that allows subcellular imaging at high resolution and, at the same time, displays enhanced in vivo brightness when compared to Renilla luciferase alone [52].

The potential of molecular imaging to reveal complex biological processes is virtually endless. These applications dramatically enhance the amount of information that can be easily obtained from refined animal models. Therefore, this review aimed at performing a systematic review of the methods employed to track hematopoietic reconstitution using molecular imaging. We hope that this effort will serve as a practical and comprehensive guide to help researchers to refine their in vivo models of HSCT by incorporating molecular imaging strategies.

2. Materials and Methods

2.1. Search Strategy

This systematic review follows the Preferred Reporting Items for Systematic Reviews and Meta-Analyses (PRISMA) guidelines [53]. We searched articles indexed, published prior to January 2020, in PubMed, Cochrane, and Scopus. The following selected criteria of interest, keyword sequences
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((Hematopoietic Stem Cell OR Hematopoietic Progenitor Cell) AND (Tracking OR Homing) AND (Transplantation)), and boolean operators (DecS/MeSH) were used:

SCOPUS: ((TITLE-ABS-KEY (tracking) OR TITLE-ABS-KEY (homing))) AND ((TITLE-ABS-KEY (“Hematopoietic Stem Cell”) OR TITLE-ABS-KEY (“Hematopoietic Progenitor Cells”) OR TITLE-ABS-KEY (“Hematopoietic Cells”) OR TITLE-ABS-KEY (“Hematopoietic Progenitor Cell”) OR TITLE-ABS-KEY (“Hematopoietic Progenitor Cells”)) AND (TITLE-ABS-KEY (transplantation)))

PubMed: (((“Hematopoietic Stem Cell”[Title/Abstract]) OR “Hematopoietic Stem Cells”[Title/Abstract]) OR “Hematopoietic Progenitor Cell”[Title/Abstract]) OR “Hematopoietic Progenitor Cells”[Title/Abstract]) AND ((tracking[Title/Abstract]) OR homing[Title/Abstract]) AND transplantation[Title/Abstract]

Cochrane: “hematopoietic stem cell” in Title Abstract Keyword OR “hematopoietic progenitor cell” in Title Abstract Keyword AND tracking in Title Abstract Keyword OR homing in Title Abstract Keyword AND “transplantation” in Title Abstract Keyword — (Word variations have been searched)

2.2. Inclusion Criteria

Eligibility criteria were established a priori. This review included only original articles written in English, published between 2000 and 2020, that had used (i) in vivo models of bone marrow transplantations, (ii) tracking agents to analyze the hematopoietic stem cells labeled with these agents, and (iii) noninvasive techniques to allow the in vivo homing of hematopoietic stem cells in the bone marrow transplantation model.

2.3. Exclusion Criteria

Reasons for excluding studies were as follows: (i) reviews, (ii) clinical articles, (iii) book chapters, (iv) protocols, (v) editorials/expert opinions, (vi) letters/communications, (vii) publications in languages other than English, (viii) did not analyze stem cell homing with noninvasive techniques, (ix) indexed articles published in more than one database (duplicates), (x) only used invasive techniques to analyze hematopoietic stem cell homing.

2.4. Data Extraction, Data Collection, and Risk of Bias Assessment

In this review, seven of the authors (F.A.O.; M.P.N.; I.S.F.; J.B.M.; F.A.; G.N.A.R. and L.F.G.) independently and randomly selected (in pairs), revised, and evaluated the titles and abstracts of the publications identified by the search strategy in the databases cited above, and all potentially relevant publications were retrieved in full. These same reviewers evaluated the full text articles to decide whether the eligibility criteria were met. Discrepancies in study selection and data extraction between the two reviewers were discussed with a third reviewer and resolved.

F.A.O.; M.P.N.; J.M.F. and L.E.B.S searched for in vivo models of bone marrow transplantations; F.A.O.; I.S.F.; J.B.M., G.N.A.R. and L.P.N. searched for tracking agents to analyze the hematopoietic stem cells labeled with these agents; F.A.O.; M.P.N.; I.S.F.; F.A., and L.F.G. searched for noninvasive techniques to allow the in vivo homing of hematopoietic stem cells in the bone marrow transplantation model. The analysis process and table plots were carried out by a full consensus of peers, respecting the distribution above. In cases of disagreement, a third, independent author decided to add or subtract data. The final inclusion of studies into the systematic review was by agreement of all reviewers.

2.5. Data Analysis

All results were described and presented using the percentage distribution for all variables analyzed in the tables.
3. Results

3.1. Selection Process of the Articles Identified According to the PRISMA Guidelines

We searched publications between January 2000 and January 2020, indexed in PubMed, Scopus, and Cochrane Library, and a total of 2191 articles were identified. Of the 50 articles identified in Cochrane Library, none were included because, out of the articles, 42 were reviews, 5 were only protocols, and 3 were clinical articles. Of the 447 articles identified in Pubmed, 114 were excluded after screening (82 reviews, 24 publications before 2000, and 8 publications in other languages) and 325 articles were excluded after assessing eligibility (91 reported no data about HSC, 99 reported no data about animal models, and 135 reported no data about noninvasive imaging); thus, only 8 articles were included from this database. Of the 1694 articles identified in Scopus, after screening, 771 articles were excluded (416 reviews, 106 publications before 2000, 50 publications in other languages, and 199 duplicated in Pubmed search), and after assessing eligibility, 910 articles were excluded (287 reported no data about HSC and 623 reported no data about noninvasive imaging); thus, only 13 articles were included from this database. In total, only 21 nonduplicate full text articles were included in this review [9,11,16–34], as depicted in Figure 1.

Figure 1. Flowchart corresponding to the stages of the PRISMA guidelines [54] of the article screening process for inclusion in this review. Abbreviations: HSC, hematopoietic stem cell; SPECT, single-photon emission computed tomography; PET, positron emission tomography; MRI, magnetic resonance imaging; FLI, fluorescence; BLI, bioluminescence.

3.2. Extraction and Isolation of Hematopoietic Stem Cells

The hematopoietic stem cell (HSC) extraction and isolation process used in the selected studies is described in detail in Table 1. Of the 21 selected studies, nine (43%) [16,18–20,23,25,28,32,34] studies used mice as a cell donor for bone marrow transplant (BMT), seven studies (33%) used humans as cell donors [9,11,21,24,26,27,33], two studies (10%) used rat cells [29,30], and the remaining studies used other cell donors, such as fish [17], axolotl [22], and dogs [31], corresponding 5% for each study. Regarding the source of the cells, the majority of studies (53%) [16,18–20,23,25,28–32] used bone marrow; however, the study by Ushiki [25] used bone marrow and spleen cells, and the second main
source of cells was the umbilical cord used in five studies (24%) [9,11,24,27,33], followed by peripheral blood, which was used in two studies (10%) [21,26]. Furthermore, the study by Lopes [22] used liver and spleen cells, the study by Astuti [17] collected the renal marrow cells from zebrafish, and the study by Sweeney [34] obtained cells from embryonic cells. Of eleven studies that used rodents as cell donors, the age of the animals was, on average, 8 weeks [18,20,23,25,30], ranging from 6 [18] to 10 [20] weeks, and the use of male rodents [29–32] was more prevalent than of female rodents [18], though two studies [23,28] used both genders. In the study by Lange [31], which used dogs, the animals were all male of ~3 years of age, and of the studies that used humans as cell donors, only the study by Massolo [29] reported the age range and gender of the volunteers (aged 18–40 and both genders).

The main method used for harvesting stem cells from animals in the selected studies was flushing (29%) [18,29,30,32], followed by maceration (21%) [22,23,28]. The other studies reported the methods of differentiation from embryonic cells [34] and femoral bone marrow aspiration in live animals [31]. Of the studies that used humans as cell donors, only one study [26] reported the apheresis method for harvesting stem cells.

Concerning the medium and supplementation used during the harvesting of stem cells, two studies [29,30] used Dulbecco’s modified Eagle medium (DMEM) supplemented with 10% fetal bovine serum (FBS) and heparin. Three other studies [17,22,23] used Phosphate-buffered saline (PBS), and the study by Parada-Kusz [23] supplemented the PBS with 0.5% bovine serum albumin (BSA)/FBS and 2 mM EDTA (Ethylenediaminetetraacetic acid). Furthermore, the study by Lopez [22] used 0.75× PBS supplemented with 5% BFS (APBS, axolotl PBS); the study by Lin [18] used minimum essential medium Eagle alpha modification (α-MEM) supplemented with 10% heparin; the study by Sweeny [34] used StemPro34 medium with a hematopoietic cytokine selection; and in the study by Niemeyer [33], the harvesting of stem cells was performed with citrate buffer.

For isolating mononuclear cells from the pool of cells collected, most of the selected studies (33%) [11,22,24,29–31,33] used the Ficoll density gradient method, although the study by Ushiki [25] used Lympholyt, and the study by Hamilton [21] used Percoll for HSC isolation. The phenotypic characterization identified the primitive lineage in all of the studies that used the isolated cells from humans as CD34 positive (+) [9,11,21,24,27,33] and CD38 negative (−) [9], as well as a subpopulation of CD34(+) and CD133 positive (+) [26]. The same primitive lineage was reported in three studies from rat-isolated cells as CD90 positive (+) [29,30], and dog-isolated cells as CD34(+) [31]. However, in the studies that used cells from mice, the phenotypic characterizations identified the primitive lineage as an early form of mouse hematopoietic stem cells, such as KSL cells [19,32], (c-kit positive (+) [23,34], Sca-1 positive (+) [34], Lin negative (−) [20]), as well as CD45(+) [23,34], CD11b(+) [23], and CD41(+) [34]. The study by Lange [31] identified the isolated cells as the primitive lineage of CD34(+). The cell lines of interest were isolated by sorting in six studies (29%) [20,23,25,26,29,33], using only Magnetic-activated cell sorting (MACS). Another three studies (14%) [22,29,30] used Fluorescence-activated cell sorting (FACS), and a further six studies (29%) [9,11,19,21,24,32] (29%) used both techniques (FACS and MACS). Only four studies [11,24,27,33] reported HSC-isolated cell purity superior to 95%, and one study [9] reported purity between 80% and 90%.
Table 1. Extraction and isolation of hematopoietic stem cells.

| Ref. | Year | Donor | Source of Cells | Donor Age | Donor Gender | Harvest | Medium and Supplementation | Separation | Immunophenotypic Characterization | Technique for Sorting | Purity (%) |
|------|------|-------|-----------------|-----------|--------------|---------|---------------------------|------------|-------------------------------|---------------------|-----------|
| Parada-Kusz et al. [23] | 2018 | Mice (C57BL/6 or CB6F1; B6-Ubi:GFP) | BM | 6-8 w | M-F | Maceration | PBS (0.5% BSA/FBS, 2 mM EDTA) | NR | CD45+; c-kit+; CD11b+ | MACS | NR |
| Sweeney et al. [34] | 2018 | Mice (129/SvJ) | ESC | NR | NR | Differentiation | StemPro34 with cytokines | NR | CD41+; CD45+; Sca1+; c-Kit+ | NR | NR |
| Asiedu et al. [29] | 2017 | Mice (C57BL/6 CD45.2) | BM | NR | M-F | NR | NR | NR | NR | NR | NR |
| Lange et al. [31] | 2017 | Dog (Beagle) | BM | 1-3 y | M | Aspiration | PBS | NR | NR | NR | NR |
| Astuti et al. [17] | 2017 | Fish (zebrafish ubi:Luc) | WKM | NR | NR | Maceration | PBS | NR | NR | NR | NR |
| Saia et al. [20] | 2016 | Mice (Ubi-Luc2Kl) | BM | 8-10 w | NR | NR | NR | Lin- | MACS | NR | NR |
| Lopez et al. [22] | 2014 | Axolotl (GFP+; nucCherryRed+) | SP; LV | NR | NR | Maceration | APBS (5% FBS) | Ficoll | NR | FACS | NR |
| Sambucetti et al. [30] | 2013 | Rat (Lewis) | BM | 7 w | M | Flushing | DMEN (10% FBS, UFH) | Ficoll | CD90+ | FACS | NR |
| Lin et al. [18] | 2011 | Mice (Balb/c) | BM | 6-8 w | F | Flushing | α-MEM (10% UFH) | NR | NR | NR | NR |
| Andrade et al. [16] | 2011 | Mice (FVB H-2Kq) | BM | NR | NR | NR | NR | NR | NR | NR | NR |
| Bengtsson et al. [32] | 2011 | Mice (TgCAGDsRed) | BM | NR | M | Flushing | NR | NR | lin- c-kit+ Sca1+ | MACS; FACS | NR |
| Ohmori et al. [19] | 2010 | Mice (C57BL/6 Ly5.1) | BM | NR | NR | NR | NR | NR | lin- c-kit+ Sca1+ | MACS; FACS | NR |
| Ushiki et al. [25] | 2010 | Mice (Balb/c nu/nu) | BM; SP | 7-9 w | M | Flushing | DMEN (10% FBS, UFH) | Ficoll | CD90+ | FACS | NR |
| Massolo et al. [29] | 2010 | Rat (Lewis) | BM | 7 w | M | Flushing | DMEN (10% FBS, UFH) | Ficoll | CD90+ | FACS | NR |
| Hamilton et al. [21] | 2018 | Human | PB | 18-40 y | M-F | NR | NR | Percoll | CD34+ | MACS; FACS | NR |
| Faivre et al. [27] | 2016 | Human | CB | NR | NR | NR | NR | Ficoll | CD34+ | MACS | 94.6 |
| Staal et al. [24] | 2016 | Human | CB | NR | NR | NR | NR | Ficoll | CD34+ | MACS; FACS | 95 |
| Pantin et al. [26] | 2015 | Human | PB | NR | NR | Aspiration | PBS | NR | CD133+ | MACS | NR |
| Niemeyer et al. [33] | 2010 | Human | CB | NR | NR | NA | Citrate buffer | Ficoll | CD34+ | MACS | >95 |
| Steiner et al. [11] | 2008 | Human | CB | NR | NR | NR | NR | Ficoll | CD34+ | MACS; FACS | >95 |
| Wang et al. [9] | 2003 | Human | CB | NR | NR | NR | NR | CD34+; CD38-; CD11b+ | MACS; FACS | 80-90 |

Abbreviations—Ref.: reference; BM: bone marrow; ESC: embryonic stem cell; WKM: whole kidney marrow; SP: spleen; LV: liver; PB: peripheral blood; CB: cord blood; w: weeks; y: years; M: male; F: female; PBS: phosphate-buffered saline; APBS: 0.753x phosphate-buffered saline; DMEN: Dulbecco’s modified Eagle’s medium; α-MEM: minimum essential medium Eagle—alpha modification; FBS: fetal bovine serum; BSA: bovine serum albumin; EDTA: ethylenediaminetetraacetic acid; UFH: unfractionated heparin; Lin: lineage; FACS: fluorescence-activated cell sorting; MACS: magnetic-activated cell sorting; NR: not reported.
3.3. Lentiviral Transduction of Hematopoietic Stem Cells

One of the noninvasive techniques used for HSC tracking after bone marrow transplantation is bioluminescence imaging. This technique depends on HSCs expressing the luciferase enzyme that, added to its substrate under ideal conditions, can emit light. This technique was used in 7 out of the 21 studies [9,11,16–20] for HSC migration homing and tracking analysis (Table 2). Three studies [16,17,20] used genetically modified animals that expressed the luciferase enzyme, and the other studies [9,11,18,19] performed luciferase transduction in hematopoietic stem cells to express the bioluminescence signal. The transduction process was performed with different vectors; for instance, the study by Lin [18] used a lower multiplicity of infection (MOI) ranging from 0.5 to 1, and the study by Ohmori [19] used an MOI of 20. Thus, already, the cell dose ranges from $3 \times 10^5$ to $8 \times 10^5$ per well. For improving the transduction efficiency, three of four studies used the transfection agent during the process, such as polybrene (8 µg/mL) [18,19] and retronectin (100 µg/mL) [11], and the culture medium and supplementation varied among studies, as well as the cytokines used in the transduction process. The time of incubation was mainly 24 h, with the exception of the study by Lim [18], which used 48 h, and the study by Wang [9], which used two cycles of 24 h with an interval of 8 h. Luciferase expression evaluation was performed mainly, in three studies [11,18,19], by the flow cytometry technique (FCT); however, the study by Lin [18] also used the polymerase chain reaction (PCR) and colony forming unity (CFU) techniques, and the study by Wang [9] conducted only immunohistochemistry using monoclonal antiluciferase antibody.

3.4. Labeling Strategies and Techniques of Hematopoietic Stem Cells with Tracers

The labeling of HSC with tracers is one way to analyze cell tracking by other noninvasive image techniques. Of the 21 selected studies in this review, in 13 studies (62%) [21,23–34], the HSCs were submitted to the labeling process using various types of tracers, such as radioisotope agents used for PET/SPECT imaging in six studies (46%) [26–31], fluorophore agents used in FLI imaging in four studies (31%) [21,23–25], and magnetic nanoparticles as a contrast agent for MRI exams in three studies (23%) [32–34]. Regardless of the tracers used for labeling, this process allows HSC tracking and engraftment evaluation.

3.4.1. Labeling of HSCs with Radioisotopes/Radiopharmaceuticals

HSC labeling with radioisotopes was reported in six studies [26–31], and all of the details of this process are described in Table 3. The common radioisotope used in three of the selected studies (50%) [29–31] was the 99mTc-exametazime (99mTc), which has a half-life of 6.03 h and it is associated with hexamethylene propyleneamine oxime (HMPAO) molecules (Ceretec, GE Healthcare). Regarding the labeling process, two of these studies [29,30] incubated $2 \times 10^6$ cells with 37 MBq 99mTc-HMPAO for 30 min and the yield was 15 ± 3% after labeling, with an activity to administration of 5.55 MBq; the study by Lange [31] has already labeled with the range of 550–583 MBq 99mTc-HMPAO, and after labeling, the radioisotope activity was between 669–1350 MBq.

Another two studies [26,28] used the zirconium-89 (89Zr) radioisotope obtained by Cyclotron, and one study [28] added an oxine molecule to the radioisotope, which has a half-life of 78.4 h, but the labeling process differed between studies. The study by Asiedu [28] incubated $10^6$ cells with 0.01–5.55 MBq 89Zr-oxine for 20 min, resulting in a cell-associated radioactivity of 0.0036–1.7 MBq and a yield of 26–30%; the study by Pantin [26] already incubated $2 \times 10^6$ cells with 0.37 MBq 89Zr for 30 min, and after labeling, the cell activity was 0.28 MBq. The study by Faivre [27] used [18F] fluorodeoxyglucose (18F-FDG) obtained by Cyclotron, which has a half-life of 1.83 h, and labeling was performed with a radioisotope activity ranging from 301.8 to 945.9 MBq after labeling of 5–10 MBq, with a purity of 94.6 ± 6%.
Table 2. Luciferase transduction in hematopoietic stem cells.

| Ref. | Tracking agent | Vector | Cell dose | Transfection agent | MOI | Medium Supplemented | Cytokines SDT (ng/mL) | Incubation (hours) | Evaluation |
|------|----------------|--------|-----------|-------------------|-----|---------------------|----------------------|-------------------|------------|
| Lin et al. [18] | FLUC | Lentiviral plasmid, pCSO-re-re-cppt-MCU3-LUC | NR | Polybrene 8 μg/mL | 0.5–1 | DMEN; Iscove’s MEM (20% FBS) | mIL-3: 20; mIL-6: 50; rSCF: 50 | 48 | FCT; CFU; PCR |
| Ohmori et al. [19] | LUC | LentiLox vectors | NR | Polybrene 8 μg/mL | 20 | StemPro 34 SFM | SFC: 100; TPO: 100; IL-6: 100; Flt-3L: 100; sIL-6R: 200 | 24 | FCT |
| Steiner et al. [11] | FLUC | Plasmid pHIV-GFPFFLuc | 3–8 × 10⁵ per well | Retronectin 100 μg/mL | NR | α-MEM (10% FCS; 2mM L-Gln) | NR | Overnight | FCT |
| Wang et al. [9] | FLUC | Lentiviral vector SMPU-R-MNCU3-LUC | 1–10 × 10⁶ per plate; 3–30 × 10³ per well | NR | NR | IL-3: 5; IL-6: 16.5; SCF: 25 | 24 (2 cycles) | IHC |

Abbreviations—Ref.: reference; FLUC: firefly luciferase; LUC: luciferase; MOI: multiplicity of infection; DMEN: Dulbecco’s modified Eagle’s medium; IMEN: Iscove’s modified Eagle’s medium; α-MEM: minimum essential medium Eagle—alpha modification; FBS: fetal bovine serum; FCS: fetal calf serum; SFM: serum-free media; Gln: Glutamine; Cytokines SDT: cytokines stimulation during transduction; mIL: murine interleukin; IL: interleukin; SCF: stem cell factor; TPO: thrombopoietin; Flt-3L: fms-like tyrosine kinase 3 ligand; sIL-6R: soluble IL-6 receptor; FCT: flow cytometry technique; CFU: colony forming unity; PCR: polymerase chain reaction; IHC: Immunohistochemistry; NR: not reported.

Table 3. Labeling of hematopoietic stem cells with radioisotopes/radiopharmaceuticals.

| Ref. | Radioisotope | Radiopharmaceuticals | Half-life (h) | Manufacture | Labeling activity (MBq) | Activity after labeling (MBq) | Cell dose | Incubation (min) | Yield (%) |
|------|--------------|-----------------------|--------------|-------------|-------------------------|-------------------------------|-----------|-----------------|-----------|
| Asiedu et al. [28] | 89Zr | Oxime | 78.4 | Cyclotron | 0.01–5.55 | 0.0036–1.7 | 1 × 10⁶ | 20 | 26–30 |
| Lange et al. [31] | 99mTc | HMPAO | 6.03 | Ceretec, GE Healthcare | 550–583; 669–1350 | NR | NR | NR | NR |
| Faivre et al. [27] | 18F | FDG | 1.83 | Cyclotron | 301.8–945.9 | 5–10 | NR | 30 | 94.6 ± 6 |
| Pantin et al. [26] | 89Zr | NA | 78.4 | Cyclotron | 0.37 | 0.28 | 2 × 10⁶ | 30 | NR |
| Sambuceti et al. [30] | 99mTc | HMPAO | 6.03 | Ceretec, GE Healthcare | 37 | 5.55 | 2 × 10⁶ | 30 | 15 ± 3 |
| Massolo et al. [29] | 99mTc | HMPAO | 6.03 | Ceretec, GE Healthcare | 37 | 5.55 | 2 × 10⁶ | 30 | 15 ± 3 |

Abbreviations—Ref.: reference; 89Zr: radioactivity zirconium isotope; 99mTc: radioactivity technetium metastable isotope; 18F-FDG: 18Fluoride -fluorodeoxyglucose; HMPAO: hexamethypropyleneamine oxime; GE: General Electric; min: minute; h: hour; MBq: Mega Becquerel; NR: not reported; NA: not applicable.
3.4.2. Labeling of HSCs with Fluorophore

HSC labeling with fluorophore was reported in four of the selected studies [21,23–25], although in the study by Lopes [22], genetically modified animals were used that express green fluorescent protein (GFP), which, thus, required no labeling process, as described in Table 4. Most of the selected studies used fluorophore that emits radiation in the range of visible light wavelength, with the exception of the study by Ushiki [25], which used fluorophore that emits in the near-infrared fluorescence wavelength. The agent concentration and time of incubation during the labeling process varied among the studies, and two of the studies [22,25] reported a cellular toxicity evaluation after labeling, using mainly the flow cytometry technique.

3.4.3. Labeling of HSCs with Nanoparticles

HSC labeling with magnetic nanoparticle agents was reported in three of the studies (14%) [32–34] (Table 5). Of these studies, two [32,33] used commercial nanoparticles such as Feridex® [32] manufactured by Berlex Laboratories (Montville, New Jersey, USA), Resovist® [33] manufactured by Bayer Schering Pharma AG (Berlin, Germany), and Endorem® [33] manufactured by Guerbet S.A. (Roissy, France), with the exception the study by Sweeney [34], which synthetized nanoparticles of Gadolinium oxide. The nanoparticle characteristics of the selected studies were similar; the nanoparticles’ core sizes ranged from 3 to 5 nm [32,33], the coating was reported mainly with dextran [32,33], the hydrodynamic diameter ranged from 60 [33] to 180 [33,34], being monodisperse (polydispersity index-PDI lower than 0.5) in most of the studies [32,33]. The labeling process reported in the selected studies was performed with mainly two types of transfection agents (protamine sulfate [32,34] and lipofectamine [33]), and the nanoparticle concentration varied between 25 [33] and 125 [34] µg/mL, with mainly 4 h of incubation [33,34], with the exception of the study by Bengtsson [32], which used overnight incubation. Most of the selected studies [32,33] used labeling evaluation by Prussian blue.
Table 4. Labeling of hematopoietic stem cells with fluorophore.

| Ref. | Agent | Manufacture | Excitation/ Emission (nm) | Concentration | Incubation (min) | Evaluation |
|------|-------|-------------|---------------------------|---------------|-----------------|------------|
| Parada-Kusz et al. [23] | CellTrace™ Violet Cell Proliferation Kit | Thermo Fisher Scientific Inc. | 405/450 | 5 µM | 20–30 | NR |
| Hamilton et al. [21] | Fluorocine | NR | 495/519 | 5 µM | 20–30 | NR |
| Staal et al. [24] | PKH26 (#PKH26GL) | Sigma Aldrich | 551/567 | 2 µM | 2 | NR |
| Lopez et al. [22] | GFP™ | NA | 475/509 | NA | NA | FCT; PCR |
| Ushiki et al. [25] | Cy5.5 | GE Healthcare UK Ltd., Buckinghamshire, UK | 675/694 | 0.4 mg/mL | 15 | FCT |
| | AF750 | Invitrogen, Eugene, OR, USA | 749/775 | 0.1 mg/mL | 15 | FCT |

Abbreviations—Ref.: reference; CFSE: 5-(6)-carboxyfluorescein diacetate succinimidyl Ester; GFP: green fluorescent protein; Cy5.5: cyanine 5.5; AF750: Alexa Fluor 750 carboxylic acid, succinimidyl ester; PCR: Polymerase chain reaction; nm: nanometer; µM: micromolar; mg: milligrams; mL: milliliters; min: min; FCT: flow cytometry technique; NR: not reported; Note—* animals genetically modified that express GFP.

Table 5. Labeling of hematopoietic stem cells with nanoparticles.

| Ref. | Particle | Size core (nm) | Coating | Dₜ (nm) | Manufacture | PDI | Transfection agent | Concentration (µg/mL) | Incubation (h) | Evaluation |
|------|----------|----------------|---------|---------|-------------|-----|---------------------|-----------------------|-----------------|------------|
| Sweeney et al. [34] | Gd₂O₃-MSN with pores 24Å | NR | Pore functionalized with: (a) TRITC; (b) FITC; (c) TRITC and PEG | 177 | Synthetized | 0.535 | Polybrene or protamine sulfate | 125 | 2–4 | Fluorescent microscopy |
| Bengtsson et al. [32] | Magnetite | 4.8 | Dextran | 80–150 | Feridex (Berlex Laboratories, Montville, NJ, USA) | 0.29 | Protamine sulfate | NR | Overnight | Prussian blue |
| Niemeyer et al. [33] | Magnetite/maghemite | 3-5 | Carboxydextran | 60 | Schering Pharma AG, Berlin, Germany | 0.207 | Lipofectamine | 25 | 4 | Prussian blue |
| | Magnetite | 4.8 | Dextran | 120–180 | Endorem (Guerbet S.A.-Roissy, France) | 0.266 | Lipofectamine | 25 | 4 | Prussian blue |

Abbreviations—Ref.: reference; Dₜ: Hydrodynamic diameter; MSN: Mesoporous silica nanoparticle; Gd₂O₃: Gadolinium oxide; TRITC: tetramethyl rhodamine isothiocyanate; FITC: fluorescein isothiocyanate; PEG: Poly(ethyleneglycol); PDI: polydispersity index; NR: no reported; nm: nanometer; Å: ångström; µg: micrograms; mL: milliliters; h: hours.
3.5. Administration of HSCs Labeled/Transfected in the Bone Marrow Transplant Model

Table 6 describes that most of the selected studies (57%) [9,11,16,18–20,25,27,28,32–34] used mice as the animal of choice as the cell receptor in the bone marrow transplant model. Among the different mice species (Nod scid gamma mouse [9,11,16,17]): C57Black 6 [19,20,28,32], Balb/c [18,33], and 129/SvJ [34]), the majority were female [18,27,28,32], mainly 8-weeks-old (ranging from 6 to 12 weeks). Following the popularity of the use of mice, four further studies (19%) [17,21,23,24] used zebrafish in early development [23] (3-5 h post-fetal) and adult (48 h [24] and 52 h [21]) stages, and two more studies (10%) [29,30] used male Lewis rats (7-weeks-old). Of the other studies, one used White Axolt Salamanders [22] that were more than 12 weeks of age, one used male Beagle Dogs [31] that were 1–3-years-old, and one used domestic swine [26] of both genders weighing between 38 and 70 kg. Some animals of the selected studies were irradiated by Cesium-137 in six studies [11,16,18,25,27,34] and by X-rays in four studies [16,17,29,30], exposing them to total body irradiation (TBI) in most of the studies (71%) [9,11,16–20,22,25,27–32,34] with varied doses of irradiation. The most used dose was 9.5 Gy [19,22,28–30,32], ranging from 2.25 Gy [27] to 20 Gy [17], although for the immunosuppressed animals (NSG mice), the dose used was lower than 3 Gy [9,11,16,27]. The study by Lin used radiation for spinal cord ablation and also used the associated O6-benzylguanine (BG) and 1,3-bis(2-chloroethyl) (BCNU) for non-myeloablative conditioning. A few studies reported a dose rate of irradiation that ranges between 0.9 [29,30] and 2.7 Gy/min.

After irradiation or not, the animals received the cell transplantation. The allo geneic type was most used (57%) [16–20,22,25,28–30,32,34] between the selected studies, followed by the xenogeneic type (38%) [9,11,21,23,24,26,27,33], and the study by Lange [31] used autologous transplantation. The time of cell implantation varied among studies; five studies [27–30,34] performed the implantation after 24 h of animal irradiation, three studies [9,18,25] used the acute stage between 1 and 8 h, and two studies [17,18] performed the implantation after 48 h of animal irradiation. The dose of cells implanted varied among studies; a dose 10^6 cells was more common in the selected studies [16,18,27,29–34], ranging from 5 \times 10^5 (in zebrafish embryos) [22] to 10^6 cells (in porcine) [26]. These cells were administered systemically in most of the selected studies (81%) [9,11,16,17,19,20,22,24–30,32–34], with the exception of the study by Lange [31], which used intrabone administration. Another four studies [16,25,26,29] compared systemic to intrabone administration, and in the studies that used zebrafish as a model, the cells were administered by yolk sac of Duct of Cuvier [21,24], intracardially [17], or retro-orbitally [21,34]. In vitro graft analysis was performed in 18 studies (86%) [9,11,16–20,22–25,27–30,32–34], and the method most used for this analysis was the flow cytometry technique (FCT), as reported in 14 of the studies (67%) [9,11,16–19,22–25,27,28,33,34]. The efficiency of the graft was evaluated in 13 of the selected studies (61%) [9,11,16,17,19,20,23,25,27,28,30,33,34], and the results reported that before one month of implantation, cell renewal was very low (less than 20%) [23,27,30,33,34], but after three months, more than 50% efficiency of the graft was reported [11,16,19,20,28], with the exception of the study by Wang [9] that reported 1.3% graft efficiency at 3.5 months. In the studies that used allo geneic cell transplant [16,19,20,25,28], the efficiency of the graft was greater than xenogeneic cell transplant [9,11,23,27,33], achieving high efficiency in a short time.
Table 6. Administration of hematopoietic stem cells (HSCs) labeled/transfected in bone marrow transplant model.

| Ref. | Animal receptor | Specie | Age | Gender | Conditioning | Source | Dose (Gy) | Dose-rate (Gy/min) | Type of graft | Delay for cells infusion (hours) | Cell dose | Route | Vehicle | Graft assessment | Graft efficiency (% or number of cells) at time |
|------|----------------|--------|-----|--------|--------------|--------|-----------|-------------------|--------------|-------------------------------|-----------|-------|---------|----------------|---------------------------------|
| Parada-Kusz et al. [23] | Zebrafish (nacre<sup>−</sup>) | 3–5 h NR NA NA NA NA Xenogeneic NA 1-6 × 10<sup>4</sup> Blastoderm PBS FCT (CD45); TUNEL 0 at 4 d | | | | | | | | | | | | |
| Sweeney et al. [34] | Mice (129/SvJ) | NR NR TBI 137-Cs 2 × (7–8) NR Allogeneic 24 7.2 × 10<sup>6</sup> IV (RO) NR FCT (GFP) 0 at 20 d | | | | | | | | | | | | |
| Hamilton et al. [21] | Zebrafish (Tg-kdrl:EGFP) | 52 h NR NA NA NA NA Xenogeneic NA NR DC yolk sac NR NA NA | | | | | | | | | | | | |
| Asiedu et al. [28] | Zebrafish (C57BL/6 CD45.1) | 8–12 w F-M TBI NR 9.5 NR Allogeneic 24 2 × 10<sup>7</sup> IV NR FCT (CD45.1, CD45.2, lineage specify) 85 at 70 d | | | | | | | | | | | | |
| Lange et al. [31] | Dog (Beagle) | 1–3 y M NA NA NA NA Autologous NA 1.5 × 10<sup>9</sup>/kg IB (humerus) NR NA NA | | | | | | | | | | | | |
| Astuti et al. [17] | Zebrafish | NR NR TBI X-rays 20 2.7 Allogeneic 48 5–50 × 10<sup>4</sup> IV (IC) NR IF, FCT (GFP) 306.5 ± 136.6 cells at 14 d | | | | | | | | | | | | |
| Saia et al. [20] | Mice (B6 albino) | NR NR TBI NR 6.5 NR Allogeneic NA 5 × 10<sup>5</sup> IV (tail) NR qPCR (LUC2) 53 at 120 d | | | | | | | | | | | | |
| Faivre et al. [27] | Mice (NSG) | 8 w F TBI 137-Cs 2.25 NR Xenogeneic 24 2 × 10<sup>6</sup> IV (tail) Saline FCT (CD34, CD45) CD34: 1.5 ± 0.6; CD45: 20 ± 6 at 30 d | | | | | | | | | | | | |
| Staal et al. [24] | Zebrafish (caser Fli1-GFP) | 48 h NR NA NA NA NA Xenogeneic NA 0.5–5 × 10<sup>5</sup> DC yolk sac IV (RO) NR FCT (CD3, CD14, CD34, CD38, CD45) NR | | | | | | | | | | | | |
| Pantin et al. [26] | Porcine (domestic swine) | NR F-M NA NA NA NA Xenogeneic NA 2 × 10<sup>8</sup> IV (jugular) IB (iliac crest) IA (iliac) Saline NA NA | | | | | | | | | | | | |
| Lopez et al. [22] | Salamander (white mutant d/d) | >12 w NR TBI NR 9.5 NR Allogeneic NR 1-50 × 10<sup>4</sup> IV (IC) NR FCT (GFP); TUNEL; HC NA | | | | | | | | | | | | |
| Sambuceti et al. [30] | Rat (Lewis) | 7 w M TBI X-rays 9.5 0.9 Allogeneic 24 2 × 10<sup>6</sup> IV (NA) NR CA LV: 0.28 ± .018; SP: 0.19 ± 0.12; LG: 0.03 ± 0.42 | | | | | | | | | | | | |
| Lin et al. [18] | Mice (Balb/c) | NR F BG + BCNU; TBI 137-Cs 7.5 NR Allogeneic 48; 1–5 1–10 × 10<sup>5</sup> NA NR FCT; CFU NR | | | | | | | | | | | |
| Author(s) et al. | Species | Age | Sex | Treatment | Dose | Grade | Cell Type | NK Cell | IV Site | FCT Type | IHC Type | Notes |
|-----------------|---------|-----|-----|-----------|------|-------|-----------|---------|---------|-----------|----------|-------|
| Andrade et al. [16] | Mice (NSG) | 8–10 w | NR | TBI; RHLIR | 137-Cs; X-rays | 2.7 | NR | Allogeneic | NR | 1 × 10^6; 1 × 10^7 | IV (tail) | IB (femur) | NR | FCT (CD11b, Ly-6c/g, CD3e, B220, NK1.1) | 69 ± 5 at 98 d |
| Bengtsson et al. [32] | Mice (C57BL/6j) | 6–8 w | F | TBI | NR | 9.5 | NR | Allogeneic | NR | 1 × 10^3; 1 × 10^6 | IV (RO) | NR | HA | NA |
| Ohmori et al. [19] | Mice (C57BL/6Ly5.2) | 8–12 w | NR | TBI | NR | 9.5 | NR | Allogeneic | NR | 1 × 10^5 | IV (carotid) | NR | FCT; IF | 58 at 30 d |
| Ushiki et al. [25] | Mice (Balb/c) | 7–9 w | NR | TBI | 137-Cs | 8 | NR | Allogeneic | 6–8 | 1 × 10^6 | IV (tail) IB (tibia) | NR | FCT (eGFP); HA | 43.5 at 3 d |
| Massolo et al. [29] | Rat (Lewis) | 7 w | M | TBI | X-rays | 9.5 | 0.9 | Allogeneic | 24 | 2 × 10^6 | IV (tail) IB (tibia) | Saline | Blood count | NR |
| Niemeyer et al. [33] | Mice (Balb/c) | NR | NR | NA | NA | NA | NA | Xenogeneic | NA | 1–5 × 10^7; 5 × 10^7; 1 × 10^8 | IV (tail) | NR | HA; IHC; FCT (CD34, CD45, CD71) | 0.08 at 1d |
| Steiner et al. [11] | Mice (NSG) | 6–12 w | NR | TBI | 137-Cs | 2.7 | NR | Xenogeneic | NR | 8–35 × 10^4 | IV (tail) | NR | FCT (CD45, lineage specify); IHC | 39.6 at 365 d |
| Wang et al. [9] | Mice (NSG) | 8–10 w | NR | TBI | 137-Cs | 3 | NR | Xenogeneic | 2 | 4 × 10^4; 1 × 10^4 | IV (tail) | NR | FCT (CD45) | 1.3 at 105 d |

**Abbreviations** — Ref.: reference; h: hours; min: minute; w: week; y: year; mg: milligram; kg: kilogram; TBI: total body irradiation; RHLIR: right hind limb irradiation; BCNU: 1,3-bis(2-chloroethyl)-1-nitrosourea; BG: O6-benzylguanine; 137-Cs: Cesium-137; Gy: gray; IV: intravenous; IB: intrabone; RO: retro-orbital; IC: intracardiac; DC: duct of Cuvier; PBS: phosphate-buffered saline; qPCR: quantitative polymerase chain reaction; TUNEL: terminal deoxynucleotidyl transferase dUTP nick end labeling; CFU: colony forming units; FCT: flow cytometry technique; IHC: immunohistochemistry; IF: immunofluorescence; HC: histochemical; HA: histological analysis; CA: Compartm ental analysis; LV: liver; SP: spleen; LG: lungs; NR: not reported; NA: not applicable; **Note** — Mice (129/SvJ): mice have mutated CD23 and hyper IgE; Mice (NSG) — Nod scid gamma mouse; Mice—Balb/c: an albino mice; Mice (C57BL/6) wild type (expressing CD45.2) and congenic (expressing CD45.1).
3.6. Imaging Techniques Used in HSC Migration Homing and Tracking Analysis

HSC tracking was evaluated by different noninvasive imaging techniques after their transplant in the animals. The technique most used in the selected studies was bioluminescence in seven of the studies (33%) [9,11,16–20], followed by PET/SPECT used in six studies (29%) [26–31], fluorescence imaging in five studies (24%) [21–25], and MRI in the other three studies (14%) [32–34].

3.6.1. Bioluminescence

Table 7 shows that the bioluminescence images were acquired after luciferin administration; this substrate was applied in most of the studies (86%) by intraperitoneal administration [11,16–20], using a dose that ranged from 0.75 to 150 mg/kg with different time intervals (2-15 min) to begin the image acquisition. The Xenogen IVIS system equipment was used in all studies, and although the model varied, the same software was used to process the images; that is, the Living Image software. The selected studies that used mice [9,11,16,18,20] acquired images mainly in the dorsal and ventral animal positions; the study by Steiner [11] also used the lateral position. Among the few BLI acquisition parameters reported by the selected studies, the exposure time used in the image acquisition varied among studies in accordance with the signal finding, but three studies [9,11,16] reported a decrease in the exposure time over the weeks (3 to 1 min/1 s) due to rise of signal intensity. Binning acquisition parameters was reported only in the study by Astuti [17], with the 4 or 8 values. The HSC tracking was performed from 1 day after bone marrow transplant in three studies (43%) [16,17,20], and in another four studies [9,11,18,19], bioluminescence imaging only was detectable at the end of the first week. Bioluminescence imaging was used for follow-up evaluation of longitudinal HSC tracking in all studies [9,11,16–20].

3.6.2. PET/SPECT

Of the six studies that tracked cells with radioisotope agents (Table 8), three of them (50%) [29–31] used the single-photon emission computed tomography (SPECT) technique with 99mTc-HMPAO by gamma camera equipment, two [29,30] used a window energy of 5% centered over 140 KeV and dynamic acquisition, and the final study [31] used whole-body acquisition. The time of the cell homing evaluation of these studies was 30 min [29,30] and 24 h [31], detecting uptake distribution commonly to the heart, liver, lung, spleen, and other regions.

Another three studies (50%) [26–28] used hybrid positron emission tomography–computed tomography (PET-CT) scanning by a micro-PET-CT scanner [27,28] or a Philips clinical PET-CT scanner [26]. The studies by Asiedu [28] and Pantin [26] used 89Zr radioisotopes, a whole-body image with a 400 to 700 KeV energy window, and the homing evaluation was reported in intervals of 2-4 h or 1–7 days in the former study and 5-15 h in the latter study. The study by Faivre [27] used 18F-FDG with dynamic image acquisition and a 511 KeV energy window for 3 h of homing evaluation. Regarding the uptake distribution of cells after transplantation, the selected studies [26–30] that implanted cells by systemic administration (intravenous) reported greater concentrations of HSCs labeled in the lung. In two studies [26,29] that implanted the cells intraosseously, the cell distribution to the lung was lower, and posterior cell migration for other organs was reported.

3.6.3. Fluorescence Imaging

Fluorescence imaging was performed in five of the studies [21–25] for HSC tracking analysis after the bone marrow transplant (Table 9). In the studies that used zebrafish or axolotl models, microscopy was used for imaging acquisition [21–24], and in the study by Ushiki [25], which used mice as receptors, the HSCs were tracked using the IVIS Spectrum system (Xenogen, Alameda, CA, USA). FLI acquisition parameters were reported only in the study by Ushiki [25]. Two studies reported the acute evaluation of HSCs after transplantation (until 24 h), but the study by Lopes reported longer homing evaluation (until 6 days) with positive detection until three days. Only the study by Ushiki [25], which used mice as cell receptors, was the uptake distribution reported in the
bone marrow, lung, spleen, liver, and kidney, corroborating the ex vivo analysis [25]; in the studies that used zebrafish, the uptake distribution was reported mainly for the tail [21,23,24].

3.6.4. Magnetic Resonance Imaging

Of the selected studies, MRI was performed in three studies [32–34] to evaluate HSC tracking after the bone marrow transplant (Table 10). The study by Niemeyer [33] only evaluated the acute stage from 2–24 h, using low magnetic field equipment (1.5 Teslas), but the other two studies [32,34] reported a long evaluation, from 3 h [34] to 14 days [32], using equipment with a higher magnetic field (4.7 [34] and 17.6 [32] Teslas). These three studies [32–34] reported uptake distribution for the bone marrow, liver, and spleen. MRI acquisition parameters used in the selected studies showed that the T2 weighted image was widely used with variations in the field of view, matrix, slice thickness, and others. In addition, this technique was the most detailed in terms of image acquisition parameters among the selected studies.

All aspects of the analysis of the noninvasive tracking of hematopoietic stem cells in a bone marrow transplant model are shown in Figure 2.

Figure 2. Schematic illustration of the bone marrow transplantation process, from the isolation of HSCs to their implantation, and tracking evaluation by noninvasive imaging techniques. (A) Main cell donors; (B,C) main cell sources for HSC extraction; (D) HSC characterization after isolation; (E) contrast agent used in the HSC labeling; (F) animal pre-condition (irradiation) before HSC transplantation; (G) cell route used in the transplantation; (H) type of graft; (I) noninvasive imaging technique used in HSC tracking; (J) imaging modality features; and (K) the graphic of the percentual distribution of the main element analyzed in the systematic review.
Table 7. HSCs migration homing and tracking by bioluminescence.

| Ref.           | Substrate | Dose (mg/kg) | Substrate administration route | Time before image (min) | Equipment | Software | Animal position | Exposure time (min) | Binning | Homing evaluation time | Outcome (cells migration) |
|----------------|-----------|--------------|---------------------------------|--------------------------|-----------|----------|----------------|---------------------|---------|------------------------|--------------------------|
| Astuti et al.  | Luciferin | 75§          | IP                              | 10-15                    | Xenogen IVIS50 system (Caliper Life Sciences) | Living Image | Dorsal and lateral | 1                   | 4; 8                | 98 d                   | +                        |
| Saia et al.    | Luciferin | 80           | IP                              | 15                       | Xenogen IVIS Lumina system (Caliper, PerkinElmer) | Living Image | Ventral and dorsal | 1                   | NR (20, 30, 40-120 d) | +                      |
| Lin et al.     | Luciferin | 125          | IP                              | 7                        | Xenogen IVIS 200 Imaging system (Caliper Life Sciences) | Living Image | Ventral and dorsal | 5                   | NR                  | 146 d                  | +                        |
| Andrade et al. | Luciferin | 125          | IP                              | 8                        | Xenogen IVIS 100 imaging system (Caliper Life Sciences) | Living Image | Ventral and dorsal | 3-1                 | NR                 | 1-65 d                 | +                        |
| Ohmori et al.  | Luciferin | 1.5§         | IP                              | NR                       | IVIS Imaging system (Xenogen, Alameda, CA) | Living Image | Ventral and dorsal | NR                  | NR (3, 7, 14, 21, ... | 256 d                 | +                        |
| Steiner et al. | Luciferin | 150          | IP                              | NR                       | IVIS-IVIS Imaging system (Caliper Life Sciences Hopkinton, MA) | Living Image | Ventral, dorsal and lateral | 3-1                 | NR                 | 14 d (3, 6, 12 months) | +                        |
| Wang et al.    | Luciferin | 125          | IV                              | 2                        | IVIS 3-D optical imaging system (Xenogen, Alameda, CA) | Living Image | Ventral and dorsal | NR                  | 1, 8 d               | 7-15 weeks             | +                        |

Abbreviations — Ref.: reference; IP: intraperitoneal; IV: intravenous; NR: not reported; mg: milligrams; kg: kilograms; µg: microgram; d: days; min: minutes. Note — Units: §: mg/body. #: µg/body.

Table 8. HSCs migration homing and tracking by PET/SPECT.

| Ref.                 | Image modality | RP          | Equipment                                                                 | Reconstruction parameters     | Type of image | Energy window (KeV) | Image acquisition time | Homing evaluation time | Uptake distribution                  | Outcome (cells migration) |
|----------------------|----------------|-------------|---------------------------------------------------------------------------|-------------------------------|---------------|---------------------|------------------------|------------------------|---------------------------------------|--------------------------|
| Asiedu et al. [28]   | PET-CT         | 89Zr-oxime  | MicroPET-CT (BioPET, Bioscan)                                              | 3D; Iterative reconstruction  | Wholebody     | 400-700             | 5 min per bed for 4 h; 6.5, 7.5, 12.5, 15.5 min per bed for 1, 2, 5.7 d | 0, 2, 4, 24, 48 h (5, 7 d) | Bone marrow, spleen and liver          | +                        |
| Lange et al. [31]    | SPECT          | 99mTc-HMPAO | PRISM 2000 XP gamma camera (Phillips, Hamburg, Germany)                    | NR                            | Wholebody     | 140±5%              | NR                     | 1, 6, 24 h               | Bone marrow, lungs, ribs and spines     | +                        |
| Faivre et al. [27]   | PET-CT         | 18F-FDG     | Micro-PET-CT (Inveon, Siemens)                                             | OSEM 3D                       | Dynamic       | 511±5%              | Dynamic study for 2.5 h | 0, 1, 2, 3 h             | Lung, kidney, spleen, liver, femur and vertebrae | +                        |
| Pantin et al. [26]   | PET-CT         | 89Zr        | Gemini TF clinical PET/CT (Philips)                                       | 3D; Iterative reconstruction; Scatter and attenuation | Wholebody     | 400-700             | NR                     | 5, 10, 15 h              | Lungs, bone marrow                     | +                        |
Table 9. HSC migration homing and tracking by fluorescence imaging.

| Ref. | Tracking agent | Equipment | Parameters | Homing evaluation time | Uptake distribution | Outcome (Cells migration) |
|------|----------------|-----------|------------|------------------------|---------------------|----------------------------|
| Parada-Kusz et al. [23] | CellTrace™ Violet Cell; CellTrace™ CSFE | Epifluorescence microscope (Zeiss Axio Observer), an AIR; C2 (Nikon) confocal microscope; Eclipse Ti (Nikon) spinning disk confocal microscope | NR | [1, 2, 3 d]*, 4, 5, 6 d | Yolk sac, tail, ICM, PBI and AGM | + |
| Hamilton et al. [21] | FITC | Spinning Disk confocal microscope | NR | 1, 4, 7, 10, 13 h | Tail | + |
| Staal et al. [24] | PKH26 | Leica fluorescent microscope | NR | NR | Yolk sac, tail | + |
| Lopez et al. [22] | GFP | Leica MZ16FA microscope, using a Hamamatsu digital camera model C7780 and Volocity Imaging software (Perkin Elmer) | NR | NR | NR | + |
| Ushiki et al. [25] | Cy5.5; AF750 | IVIS Spectrum system (Xenogen, Alameda,CA, USA) | Excitation/emission (nm): 640/700 for Cy5.5, 710/780 for AF750. Exposure time: 5 s, lamp level: high, binning: medium; FOV: 12.96x12.9 cm, and f/stop: 1. | 0.5, 1, 3, 6, 12, 18, 24 h | Bone marrow, lung, spleen, liver, kidney | + |

**Abbreviations**—Ref.: reference; CFSE: 5-(6)-carboxyfluorescein diacetate succinimidyl Ester; FITC: Fluorescein isothiocyanate; GFP: green fluorescent protein; Cy5.5: cyanine 5.5; AF750: Alexa Fluor 750; ICM: intermediate cell mass; PBI: posterior blood island; AGM: aorta-gonad-mesonephros; NR: not reported; s: seconds; cm: centimeter; d: day; h: hours; min: minutes; nm: nanometer, FOV: field of view. **Note**—* means the FLI intensity signal detectable.
### Table 10. HSC migration homing and tracking by magnetic resonance imaging.

| Ref.         | Equipment                                      | Software          | MF (T) | Sequence                  | Weighted images (TR/TE; ms) | FOV;MT;ST | Homing evaluation time | Uptake distribution | Outcome (cells migration) |
|--------------|-----------------------------------------------|-------------------|--------|---------------------------|----------------------------|-----------|-----------------------|----------------------|-------------------------|
| Sweeney et al. [34] | Varian® Unity/INOVA 4.7 T small animal scanner | MIPAV             | 4.7    | Fast Spin Echo; Gradient Echo | T2: 2100/15 T2*: NR         | NR; 256x512; NR | 3, 24 h (6, 9 d)       | Bone marrow, spleen, and liver | +                      |
| Bengtsson et al. [32] | Bruker BioSpin, Madison, WI, USA             | Paravision 4.0 (Bruker, Madison, WI, USA) and OsiriX v.3.5 | 17.6   | 3D Gradient Echo          | T2*: 80/2.5                | 1.1x0.6x0.5cm³; 393x214x83; NR | 7, 14 d              | Bone marrow             | +                      |
| Niemeyer et al. [33] | 1.5-T imaging MR scanner (ACS NT; Philips, Best, the Netherlands) | NR                | 1.5    | 3D Fast Field Echo        | T2*: 32/14                 | 100x80mm; 512x512; 0.4mm | 2, 24 h              | Bone marrow and liver   | +                      |

**Abbreviations**—Ref.: reference; 3D: three dimensional; MF: magnetic field; TR: repetition time; TE: echo time; T2: transversal relaxation time; T2*: transversal relaxation time star; FOV: field of view; MT: matrix; ST: Slice thickness; NR: not reported; T: tesla; ms: millisecond; mm: millimeter; d: day; h: hours.
4. Discussion

The results of this review show an increase in using noninvasive imaging techniques in HSC tracking in the bone marrow transplant model. The most selected studies used bioluminescence, followed by fluorescence, PET/SPECT, and MRI. The use of noninvasive imaging techniques can reveal that hematopoietic activities in both steady-state and pathological conditions are dynamic, and that their sequence is regulated spatiotemporally by interaction with the niche [55]. Successful clinical trials show that the engraftment depends on adequate HSC numbers, their homing, and the subsequent short-term and long-term engraftment of these cells in the niche (bone marrow). Enhancing the homing capability of HSCs could have a great impact on improving transplantation procedures and patient survival [56]. The main challenge of leukemia or the dysfunctional or depleted bone marrow preclinical model is the mimicking of the bone marrow niche or the environment. Experimental findings in mice often correlate with human biology and, as such, they serve as a research stand-in for human patients [57]. In this review, most of the selected studies used mice as the animal model, and regarding the type of graft, allogeneic cells were the most used among the studies. In exploring this aspect, it was reported that when comparing allogeneic and xenogeneic grafts, using the cell route by the tail vein, the allogeneic cells arrive at the niche faster (3 h after transplantation) than in the xenogeneic cells (15 h after transplantation), and in a greater amount, being detectable in the endosteal region of the femur (13% of cell quantity) and in the central marrow region (58% of cell quantity), associated with efficiency and marrow repopulating ability [58].

Irradiation is an important aspect of HSC homing analysis. The study by Xie [59] showed that only 7% of cells after 6 h of transplantation arrive at the bone marrow niche compared to the irradiated hosts, which obtained better efficiency with 35% of cells detectable in the niche. In addition, immunoassays identified the endosteal zone of irradiated recipients as a site of increased HSC proliferation after transplantation. In this review, it was observed that most rodent models of the selected studies applied bone marrow irradiation with cesium-137 or X-ray, and cell homing analysis was possible and detectable by noninvasive image and other techniques, particularly the flow cytometry technique.

Optical imaging certainly holds utility in assessing hematopoietic stem cell tracking of xenografts, allografts, autologous grafts applied in the bone marrow transplant model [35]. The same study reported that bioluminescence imaging is certainly more sensitive than fluorescence imaging. However, one strength of fluorescent fluorophores is the ability to perform high-resolution microscopy of GFP-expressing cells. As a result, if tracking of small numbers of cells at high magnification and perhaps in real-time is desired, fluorescence-based methods can be considered. Besides these techniques, this review reported the used of other techniques, such as PET/SPECT and MRI in cell homing, where PET/SPECT has a small temporal window in signal detection and MRI requires a magnetic contrast agent that may interfere with cellular growth biology and the interaction of the environment in a niche (bone marrow), decreasing the sensitivity of this technique.

Most of the selected studies of this review used bioluminescence due to its reported high sensitivity, with a wide temporal detection window (0 days to 1 year), but a limitation of this technique is that the light propagation through tissue currently restricts the application of bioluminescence imaging to small animals, where the signal can easily penetrate at all depths, and it is a technique that has not yet been well studied in vivo [9]. Another prevalent technique used by six of the selected studies was PET/SPECT imaging, which provides three-dimensional information with high sensitivity, excellent penetration depth, and a capability for whole-body imaging, but which requires radioisotopes. An advantage of this technique is that it can also be applied in small animals as well as in humans, and the disadvantages include the high cost of the cyclotron and limited temporal follow-up due to the radioisotopes' half-life—a factor that was reported in the selected studies, where cell homing was commonly evaluated at 15 h. In addition, in the selected studies, the radioisotope most used was the 99mTc-HMPAO, which has high translation capacity, but it is important to bear in mind the potential effects of radiation on therapeutic/biology cell function [60].
The fluorescence imaging technique has a high sensitivity (lower than bioluminescence), a high temporal resolution, a low cost, and it is activatable, but the disadvantages in relation to the other image techniques described before include the attenuation of sensitivity by overlying tissues, low spatial resolution, and poor penetration depth. Therefore, this technique is the most important imaging modalities for live-cell imaging at single-cell resolution using light sources to excite the fluorescent molecules [61]. In this review, the fluorescence technique observed HSC homing into the niche up until 24 h.

MRI has a high spatial and temporal resolution, no tissue penetrating limit, and no radiation, but the disadvantages include relatively low sensitivity and low contrast, requiring a high load of cells with a magnetic label and comparatively long imaging times, with the possibility of affecting cell viability or biological interaction with the contrast agent [43,62–64]. In this review, three studies used HSC tracking analysis by MRI with different magnetic fields, in which the highest magnetic field (17.6 Teslas) provided longer follow-ups (14 days after transplantation) when using low nanoparticle concentrations and incubation times in the cell labeling process. Our group provided evidence, in previous studies, that a low nanoparticle concentration in the labeling process was enough to detect stem cell migration homing and tracking. Additionally, nanoparticle sizes of between 35 and 200 nm used in our previous studies showed magnetic characteristics adequate for detection by MRI [65–67]. This review corroborates the nanoparticle size findings among the selected studies (60 to 180 nm).

Regarding the molecular imaging techniques approaches in this review, their physical principles, applicability, advantages, and limitations (Figure 3)[43,46,47,68–73], show a wide potentiality not only for in vitro studies and pre-clinical applications but also in the translation of some techniques in clinical studies, such as nuclear images (PET and SPECT) and MRI [47,73]. Comparing the molecular imaging technique characteristics and the HSC tracking evaluation shown in Figure 3, the optical techniques allowed a live cell tracking for a long period of time by BLI (at 1 day to 12 months) [9,11,16–20] and short time by FLI (at 30 min to 6 days) [21,23–25], but in the latter, the signal is not necessarily from living cells. In nuclear techniques, the time window varies with the radioisotopes used due to their half-life, in SPECT the radioisotopes have a shorter half-life, and therefore, the time allowed for tracking the cells was also shorter (at 30 min to 1 day) [29–31], compared with cell tracking made by PET (at 1 h to 7 days) [26–28], in which the radioisotopes used have an average half-life of 78.4 h, as is the case with Zr. MRI has high spatial resolution and a wide temporal window for cell tracking; however, for all this, a high nanoparticle load is required for its detection, as its sensitivity is low (10^3 - 10^5 mol/L) [72] compared to the other techniques, in addition to decreasing the signal temporally due to cell division, making it difficult to keep track of the cells [74].

Therefore, for preclinical studies of the cell transplantation model, all molecular imaging techniques covered in this review have good applicability in cell tracking for early assessment, but the BLI optical technique stands out in the prolonged tracking assessment [11]. In the clinical application of oncological diseases, the PET nuclear technique and MRI would be the most suitable for early assessment of cell tracking [26–28,33,34]. However, cell graft was not evidenced by any of the molecular techniques mentioned in this review, and for cell graft to occur, we must have good control of cell tracking. The studies selected in this review used other techniques to quantify chimerism upon transplantation. The technique most used to access chimerism in the studies selected in this review was the FCT to determine the percentage of grafted donor cells grafted using specific cell line markers such as CD45 expressed by all lymphocytes [9,11,19,23,24,27,28,33], or GFP expressed by cells after genetic modification [17,22,25,34]. The CD45, in addition to making it possible to analyze the occurrence of engraftment, also allows checking the percentage of chimerism through discrimination of its two different alleles (CD45.1 and CD45.2), which are functionally identical. CD45.1+ donor cells can be readily detected when transplanted into CD45.2 mice, as reported in two studies in this review [19,28]. In these studies, the efficiency assessment was performed at early (2 to 4 days after implantation) [23,25,33] and late time points after transplantation (after 20 days to 365 days after implantation) [9,11,16,19,20,27,28]. The evaluation of cell grafting is one of the most
relevant aspects of clinical application in cancer patients, as the biodistribution and efficiency of implantation are the targets for the success of the treatment, aspects that still have many gaps in clinical research.

Figure 3. Molecular imaging modalities comparing their techniques features, tracers, applications, highlighting the follow time of each technique in the HSC tracking and the analysis of cell graft by other techniques after transplantation. Abbreviations: sec—seconds, min—minutes, cm—centimeters, mm—millimeters, mol—mole, L—liter, NIRF: near-infrared fluorescence; GFP: green fluorescent protein; qPCR: quantitative polymerase chain reaction; TUNEL: terminal deoxynucleotidyl transferase dUTP nick end labeling; CFU: colony forming units; FCT: flow cytometry technique; IHC: immunohistochemistry; IF: immunofluorescence; HC: histochemical; HA: histological analysis; CA: compartmental analysis; MSN: mesoporous silica nanoparticle; Gd2O3: Gadolinium oxide; Fe3O4: magnetite, γ-Fe2O3: maghemite. Note: if the cell graft was analyzed by other techniques, without being the molecular imaging mentioned in the Figure.

Successful transplantation depends on the formation of engraftment, in which donor cells are integrated into the recipient's cell population, as well as a supportive hematopoietic stromal microenvironment [75]. Evaluation of this was performed in most of the selected studies, which showed that the studies that used allogeneic cell transplantation had greater graft efficiency in a shorter time period than xenogeneic transplantation in the experimental model. Another relevant aspect of the success of transplantation is the functionality of cells after the graft, because graft failure and poor graft function are important issues in the current comprehensions of the interaction between the immune and hematopoietic compartments in these conditions [76]. Therefore, besides the quantitative analysis of the graft, the graft functionality should be adequate in the evaluation of graft efficiency, but this aspect was poorly approached in most of the selected studies.

Unfortunately, there are not enough clinical HSCs available to set up phase I/II clinical trials to test the tracking of these new cells by noninvasive imaging techniques. Most investigators conducting such trials are “wed” to their personal favorite procedure. If, in the future, we can deal with this problem and find means for additional clinical efforts, it is possible that several new procedures could be used together [34], such as hybrid equipment or imaging systems. This, however, adds additional
logistical problems versus the use of one procedure alone, such as requiring multifunctional probes, smart processes, and improvement of the technical limitations of imaging equipment [27]. Therefore, there is viability to translate these experimental findings into bedside application. The cell tracking improvement of niches provides a reduction in therapeutic time with high efficiency of hematopoietic cellular renewal.

**Author Contributions:** Conceptualization, F.A.O., M.P.N., and L.F.G.; methodology, F.A.O., M.P.N., I.S.F., J.M.F., L.P.N., J.B.M., F.A., and L.F.G.; validation, F.A.O., M.P.N., J.M.F., L.P.N., J.B.M., F.A., G.N.A.R., and L.F.G.; formal analysis, F.A.O., M.P.N., I.S.F., and L.F.G.; investigation, F.A.O., M.P.N., J.M.F., and L.F.G.; resources, L.F.G.; data curation, M.P.N., and L.F.G.; writing—original draft preparation, F.A.O., M.P.N., L.P.N., L.E.B.S., and L.F.G.; writing—review and editing, F.A.O., M.P.N., L.P.N., L.E.B.S., and L.F.G.; visualization, F.A.O., M.P.N., I.S.F., J.M.F., J.B.M., F.A., A.T.K., N.H., and L.F.G.; supervision, L.F.G.; project administration, L.F.G.; funding acquisition, L.F.G. All authors have read and agreed to the published version of the manuscript.

**Funding:** This research was funded by “Amigos da Oncologia e Hematologia Einstein” AMIGOH, CNPq (465259/2014-6), FAPESP (2014/50983-3;2016/21470-3) and SisNANO 2.0/ MCTIC.

**Conflicts of Interest:** The authors declare no conflict of interest.

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