An Overview of RNA-Based Scaffolds for Osteogenesis

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Tissue engineering provides new hope for the combination of cells, scaffolds, and bifactors for bone osteogenesis. This is achieved by mimicking the bone’s natural behavior in recruiting the cell’s molecular machinery for our use. Many researchers have focused on developing an ideal scaffold with specific features, such as good cellular adhesion, cell proliferation, differentiation, host integration, and load bearing. Various types of coating materials (organic and non-organic) have been used to enhance bone osteogenesis. In the last few years, RNA-mediated gene therapy has captured attention as a new tool for bone regeneration. In this review, we discuss the use of RNA molecules in coating and delivery, including messenger RNA (mRNA), RNA interference (RNAi), and long non-coding RNA (lncRNA) on different types of scaffolds (such as polymers, ceramics, and metals) in osteogenesis research. In addition, the effect of using gene-editing tools—particularly CRISPR systems—to guide RNA scaffolds in bone regeneration is also discussed. Given existing knowledge about various RNAs coating/expression may help to understand the process of bone formation on the scaffolds during osseointegration.

Keywords: bone osteogenesis, RNA, tissue engineering, gene therapy, CRISPR

INTRODUCTION

Successful bone implants depend on well-established osteointegration which is highly relevant in implant design and/or coatings. Due to the bone complexity and dynamic structure, any large and unstable fractures may cause unsuccessful healing and require additional treatments before the bone regeneration occurs (Roseti et al., 2017). In tissue engineering, various scaffolding materials with different coatings have generated an enormous interest in developing an implants to match bone features (Leng et al., 2020). Implant characteristics, including the surface topography, chemistry, and mechanical properties, have a significant effect on osteogenesis and bacterial inhibition. For instance, nano-topographical surfaces, including nanorods, nanofibers, nanotubes, and nanowires, have demonstrated the ability to perform molecular-scale medical interventions for repairing damaged tissue (Bonilla-Represa et al., 2020). The possibility to functionalize the materials can be applied through different ways either physically such as surface wettability modification, or chemically, as with acid/alkaline treatment (Damiati et al., 2018).

Human bone mesenchymal stem cells (hBMSCs) are derived from the mesoderm during early embryonic development and are considered one of the most important seed cells for bone regeneration. The repair and regeneration of bone tissue is a complex procedure, and thus designing different biomaterials with load-related growth factors is one of the essential strategies in the bone regeneration field. For instance, bone morphogenic proteins (BMPs), including BMP2 (Damiati et al., 2018; Cheng et al., 2019), BMP3 (Daluisi et al., 2001), and BMP7 (Al-Jarsha et al., 2018), can induce stem cells differentiation into osteoblasts and chondrocytes. However, this
Approach also has some limitation in practical applications, such as the difficulty in transporting these growth factors to damaged areas and maintaining long-term high concentrations (Zhang et al., 2018). Due to that, the use of nucleic acids, including RNAs, as a bioactive coating for implants, has emerged recently and has been applied in bone implants (Miyamoto et al., 2018; Zhang et al., 2018).

RNA is a single strand molecule that forms secondary structures. RNA includes various types, such as messenger RNA (mRNA), which carries genetic information and form a protein as an end-product. Other non-protein coding RNA includes microRNA (miRNA), small interfering RNA (siRNA), and long noncoding RNA (IncRNA), which plays a more regulatory role in various cell functions (Mattick and Makunin, 2006).

In this review, we provide an overview of the effect of using different types of scaffolds based on RNAs family molecules as an organic coating, including mRNA, miRNA, siRNA, and IncRNA for bone formation applications. Further, the importance of using CRISPR based genome editing to guide the RNA for bone formation is also highlighted.

**Orthopedic Tissue Engineering**

Tissue engineering is an emerging multidisciplinary science that combines molecular biology, engineering, and chemistry that aids in cellular ex vivo and in vivo tissue regeneration. Orthopedic tissue engineering in particular aims to fabricate new functional bone tissue by using combinations of cells and bioactive molecules (e.g., RNA coating) that are seeded onto biomaterials scaffolds to create an implantable “osteogenic” implant (Awad et al., 2014). However, these biomaterials can be used as implants in bone plates, dental implants, and joint replacement. Bone is considered the second most transplanted tissue after blood transfusion which increase the importance of finding the optimal biomaterial to be used clinically (Campana et al., 2014). Biomaterial scaffolds can generally be divided into natural (e.g., collagen and chitosan), synthetic (e.g., polymers), or metals (e.g., Ti, gold, and stainless steel), each with its own benefits and limitations.

These scaffolds should include few key elements to achieve regenerative bone, including bioactivity, which induces the formation of a direct chemical bond between the implant and host tissue; biocompatibility, which indicates an ability to perform with an appropriate host response in a specific application; and biodegradability, which indicates the ability to dissolve fully or partially when in contact with the living organism without causing any toxicity (Damiati et al., 2018). There are various material approaches that can be used to add bioactivity to bulk materials. Broadly, these are changes in the chemistry (Trino et al., 2018), stiffness (Behaviors et al., 2020) and topography (Hasan et al., 2017; Damiati et al., 2018; Behaviors et al., 2020). Different scaffolds have been utilized to facilitate the delivery of RNA, such as polymers-based scaffolds, ceramic-based scaffolds, and metal-based scaffolds. In the next sections, we will describe the pros and cons of these scaffolds in the bone regeneration field, then we will introduce the different types of RNAs as a novel organic coating material.

**Polymer-Based Scaffolds**

Polymers have been broadly used for fabricated medical devices and tissue-engineering scaffolds due to their unique properties such as high porosity, biodegradability, and their mechanical properties (Ji et al., 2006). There are two types of polymers, natural polymers and synthetic polymers. Natural polymers can be considered as the first biodegradable materials that were used in medical applications. They can be classified as: i. proteins, such as collagen, gelatin, keratin, actin, myosin, fibrinogen, and elastin; ii. polysaccharides, such as cellulose and chitin; and iii. polynucleotides such DNA and RNA (Dhandayuthapani et al., 2011; Chocholata et al., 2019). Natural polymers are commonly used due to their high biocompatibility and biodegradability as well as low antigenicity and inflammation. However, they have certain limitations, such as the low structural and mechanical properties, which requires combination with other materials for use in biomedical applications (Perez-Puyana et al., 2020).

Collagen is one of the natural scaffolds that has been extensively used for bone osteogenesis applications. Collage is a natural, biodegradable material that enhances cell attachment and migration, and does not cause any negative host immune responses. In bones, collagen is up to 89% of the organic matrix and 32% of the volumetric composition (O’Brien, 2011). However, collagen scaffolds have a poor compressive strength compared to native bone. Due to that, collagen is typically combined with another material to provide more structural rigidity (Ryan et al., 2019). Previous studies have shown that the compressive and tensile mechanical properties of collagen and glycosaminoglycan (a polysaccharide) can produce a highly porous collagen-GAG (CG) scaffold through physical and chemical cross-linking methods (Haugh et al., 2009; Tierney et al., 2009; Cunniffe and O’Brien, 2011). Additionally, another study by Ryan et al. showed that collagen scaffolds functionalized with copper-eluting glass were able to reduce the implant infections by Staphylococcus aureus (S. aureus) and to improve the osteogenesis and angiogenesis in vitro and in vivo (Ryan et al., 2019).

A combination between collagen and hydroxyapatite was able to activate the adipose-derived multipotent stromal cell (ASC) osteogenesis signaling pathway (Duan et al., 2017). In addition, in nature, cellulose is found as a mixture of crystalline and amorphous strictures that organized in a fringed fiber arrangement (Hearle, 1958). However, cellulose has been used in bone tissue engineering applications as the cellulosic fibers to reassemble the collagen fibers of bone tissue. Shi et al. used the bacterial cellulose as delivery system to enhance the local concentration of cytokines, as the biocompatible scaffolds increased osteogenesis in the presence of BMP2 (Shi et al., 2012). Another study by Rescignano et al. used cellulose nanocrystals based on hydrogel composites and showed the ability to transport the biopolymeric nanoparticles to the bone marrow (Rescignano et al., 2014).

Synthetic polymers are very useful materials in biomedical applications due to their physical and mechanical properties that are similar to the natural polymers. In addition, synthetic polymers are much cheaper, and can be largely produced with a long-shelf time compared to the natural polymer's scaffolds.
Different Types of RNAs in Mammalian Cells

Mammalian cells naturally contain a tremendous amount of various RNAs, which are involved in numerous complex tasks vital to the cells. The mRNA journey starts in the nucleus with DNA transcription followed by the processing of immature RNAs and ending with the export of mature RNAs to the cytoplasm to be translated into proteins (Lodish et al., 2000). RNAs that do not

Gene Therapy in Bone Repair

A promising advantage of gene therapy is the local delivery of gene sequence coding that has an ability to promote bone reparative processes. Recent studies have begun to provide potential evidence of gene therapies to deliver lasting therapeutic benefits for the bone and cartilage defects, with treatments focused mainly on the delivery of genes encoding for morphogenetic proteins (Evans and Huard, 2015). For instance, a direct injection of adenovirus carrying BMP2 presented significant repair of femoral defects in rodents (Betz et al., 2006).

Additionally, the direct delivery of recombinant adeno-associated viral vector (rAAV) with insulin-like growth factor 1 (IGF-1) (Cucchiarini and Madry, 2014), fibroblast growth factor 2 (FGF-2) (Cucchiarini et al., 2005), or SRY-related high mobility group-box gene9 (SOX9) (Cucchiarini et al., 2013), has shown an improvements in bone repair in rabbits. Various scaffolds have been used in gene combinations and gene recombinants through gene transfer using viral or non-viral vectors to target the relevant cells of osteochondral tissue engineering in vivo and in vitro (Madry et al., 2020). However, Table 1 summarizes some of the RNA-scaffolds matrix strength, weakness, opportunities, and threats (SWOT analysis) that should be taken into account before clinical use.
encode proteins but have functions are collectively known as non-coding RNAs (ncRNAs). There are two classes of ncRNAs: housekeeping and regulatory ncRNAs. Housekeeping ncRNAs are expressed constitutively, including transfer ribonucleic acid (tRNA), ribosomal ribonucleic acid (rRNA), and small nuclear (snRNA). Many regulatory ncRNAs have been identified and have become a significant focus of research due to their role in gene regulation such as micro-RNA (miRNA), small interfering RNA (siRNA), small nucleolar RNA (snoRNA), Piwi-interacting RNA (piRNA), and long non-coding RNA (lncRNA) (Mattick

**TABLE 1 | The SWOT analysis of using scaffolds based on RNA-gene therapy.**

| Strengths | Weakness |
|-----------|----------|
| - Easily to introduce into cells with high efficiency. | - Cells might not be transfectable. |
| - Can be rapidly produced in the laboratory. | - Non-renewable resource. |
| - Cost efficient. | - Virus-mediated toxic effects. |
| - Chemical modification can be used to reduce the off-target effect. | - The uncertainty of the scaffold degradation rate may affect the efficacy of the RNAs. |
| - May have a long-time effect. | - RNAs release limitation due to the strong interaction between scaffolds and the vectors. |
| - Scaffolds can protect RNA complexes from endogenous RNases. | - Regulation policies may cause a delay to get clinical trials approvals. |
| - The local RNA delivery into the site of interest may use to avoid unwanted release in other sites. | |

| Opportunities | Threats |
|---------------|---------|
| - A new sector in the market to access that provides long-term revenue. | - Long-time follow-up is required to ensure the safety and efficacy of therapy. |
| - A collaboration between the digital market based on artificial intelligence (AI) and the currently available data may accelerate RNA treatment development. | - Pre- or post-immune reactivity may limit the clinical trials. |
| - Merge the field of personalized medicine and the gene therapy which targets the oligonucleotide of an individual’s genotype may become applicable for gene silencing and directing the gene-editing case. | - More studies are necessary to find the optimal RNA sequence to use for treatment. |
| | - Biosimilar competition will need to demonstrate the efficacy of new therapy comparing to the traditional therapies. |
| | - Significant investments are required to cover all the expenses needed for RNA-based therapy manufacturing. |

**FIGURE 1 | Schematic illustration showing different types of RNA in mammalian cells mRNA, miRNA, siRNA, and lncRNA. (A) Premature mRNA gets exported to the cytoplasm then translated into protein by ribosomes. (B,C) pre miRNA and pre siRNA are produced in the nucleus and then gets exported to the cytoplasm then processed by Dicer followed by RISC complex formation, finally the miRNA or siRNA binds to the target sequence by complementation. This causes the degradation of the target RNA or translation block. (D) lncRNA are produced in the nucleus then exported into the cytoplasm in which they can regulate the gene expression (Created with BioRender.com).**
and Makunin, 2006; Mercer et al., 2009; Ponting et al., 2009; Cech and Steitz, 2014).

The most common type of these RNA-delivered molecules is mRNA, which has been studied intensively. This RNA molecule is naturally synthesized in the nucleus as a pre-mRNA and is then processed and exported into the nucleus to be translated into proteins via the ribosome’s machinery (Figure 1A). Via the addition of a specific mRNA molecules into the cellular cytoplasm, certain proteins can be synthesized and supplemented for better bone osteogenesis, as seen by the addition of a chemically modified mRNA encoding BMP2 gene to enhance bone regeneration (Elangovan et al., 2015).

Gene silencing pathways (RNA interference (RNAi)) is another type of mechanism in which short segments of RNA of around 22 nucleotides are introduced into the cells, similarly to siRNA, or produced naturally, as with certain siRNA and miRNA. These small nucleotide segments can alter the gene expression of a certain osteogenesis and bone differentiation related genes through the inhibition of gene expression. miRNAs are naturally synthesized in the nucleus as a single stranded RNA than can form a hairpin structure. They are exported into the cytoplasm and processed by DICER. They form the RNA-induced silencing complex (RISC) and then binds to the Ago2 protein. The targeted mRNA sequence by complete base paring finally inhibits the target gene expression through mRNA cleavage or the inhibition of protein translation (Figure 1B).

siRNA is a double stranded segment of RNA that works through partial binding of the mRNA targets, followed by mRNA cleavage via the RISC complex (Figure 1C) (Wilson and Doudna, 2013). RNAi has been widely introduced to cells as a therapeutic agent or for the inhibition of gene expression of a specific gene aiding in bone regeneration. IncRNAs are a group of RNAs transcribed in the nucleus with a length longer than 200 nucleotides. Some of these IncRNAs remains in the nucleus, while other are exported into the cytoplasm to play vital regulatory roles (Figure 1D). These RNA molecules play various roles, such as the regulation of gene expression and epigenetic regulation (Mercer et al., 2009). IncRNAs are also delivered into tissues to alter the gene expression of osteogenesis-related genes.

mRNA-Based Therapy

Advancements in the field of synthetic biology have enabled researchers to implement novel applications of artificial nucleic acid and its analogs as biomaterials. Synthesized mRNA can be delivered into cells for in vitro transcription (IVT mRNA) to repair and enhance bone regeneration using chemical or physical methods of delivery. They can be used to induce and modulate the expression of specific osteogenesis-related genes (Zhang et al., 2018). The host immune system can recognize the foreign mRNA, subsequently causing its degradation, and henceforth a chemical modification of its nucleic acids is required.

Elangovan and colleagues in 2015 successfully delivered the first chemically modified mRNA encoding BMP2 gene with a polyethylenimine polymer into BMSCs. They found a significant enhancement in bone regeneration in vivo with the chemically modified mRNA-polymer complex in a rat model with calvarial bone deficiency (Elangovan et al., 2015). Another study showed that the chemically modified mRNA encoding BMP2 and vascular endothelial growth factor (VEGF-A) genes in collagen-based scaffolds enhanced bone regeneration by driving bone osteogenesis in BMSCs (Geng et al., 2021). Geng et al. found that a chemically modified mRNA encoding BMP9 in a collagen scaffold enhanced osteogenesis at a calvarial bone deficient site in rats (Geng et al., 2021). Serval other studies investigated the role of chemically modified mRNA BMP2 in osteogenesis in vivo and in vitro, showing that mRNA can be considered a very useful tool to enhance bone osteogenesis in the collagen or hydrogel-based scaffolds (Badieyan et al., 2016; Balmayor et al., 2017; Elangovan et al., 2015; Khorsand et al., 2017; Zhang et al., 2019).

RNAi-Based Therapy

Tissue engineering implements the organism’s own gene expression to aid in bone osteogenesis with the use of bone scaffolds. Therefore, using miRNA and siRNA can play vital roles in regulation of gene expression via the gene silencing pathway. They can be used as a biomolecule in bone tissue engineering by entering cells using a viral or a non-viral vectors such as lentivirus and Lipofectamine (Arriaga et al., 2019).

miRNA-based therapy uses two main methods. The first is in silencing the cellular miRNA that binds to the target mRNA. In this method, the delivered miRNA binds by complementation to the cellular miRNA causing a loss of function. This subsequently causes the expression of the target gene (anti-miR) (Figure 2A). The second method is by direct down regulation of the gene via inducing the gene silencing pathway mediated by miRNA. In this case, the miRNA is designed to inhibit the target gene expression via complementary binding (Zhang et al., 2018) (Figure 2B).

Serval miRNAs (miR-100, miR-125b, miR-13, miR-196a, miR-218, and miR-22) were shown to promote osteogenesis through their action upon osteogenic target genes, while miR-126 was found to suppress osteogenesis. let-7b, let-7g, miR-133a and miR-29a were found to aid in collagen-fiber formation as summarized by (Sartori et al., 2019). miRNA can be added into the bone scaffolds in order to maintain stable long-lasting effects of these miRNAs upon the expression of the target RNA. The commonly used elements in miRNA scaffold-based tissue engineering are listed in Table 2, and Table 3 summarizing the effect of miRNA addition into scaffolds upon osteogenesis differentiation in MSCs (Leng et al., 2020).

Synthesized siRNA can also be used to silence specific osteogenesis-related genes. These double stranded siRNA could be introduced into the cells through lipid-based vectors, such as Lipofectamine. Other polymer-based delivery methods are available, such as the use of poly(lactic-co-glycolic) acid (PLGA), 3D polymeric hydrogels, and Atelocollagen scaffolds (Ghadakzadeh et al., 2016). These siRNA have shown to be a very useful tool in better understanding of osteogenesis genes, as seen in Table 4. The efficacy and lasting effects of the introduced siRNA were shown to increase in combination with scaffolds, such as lyophilized chitosan sponge (Ghadakzadeh et al., 2016).
The use of miRNA and siRNA in gene therapy has certain drawbacks: the small size of these RNA molecules leaves them unprotected from endogenous RNAase and prone to degradation; also, they also have an unstable structure and a short half-life. Therefore, chemical modifications are needed to protect them in the cells and to increase the stability, such as the use of a locked amino acid or the addition of 2-O-methoxyethyl phosphonothioate (2′-MOE) or cholesterol to modify the RNA (Zhang et al., 2018).

**lncRNA-Based Therapy**

lncRNA-based research has increased in the last few years as more functional roles of them have emerged. lncRNAs can either promote or inhibit the gene expression of serval genes or miRNAs (Ju et al., 2019). Studies have shown that lncRNA such as (MALAT1 (metastasis-associated lung adenocarcinoma transcript 1), HOTAIR (HOX transcript antisense RNA), H19, MODR, MIAT and MEG3) play essential roles in osteogenic differentiation. DANCE—another lncRNA—was found to regulate osteoclast differentiation in MSCs (Peng et al., 2018). Generally, lncRNAs are essential regulators for many biological processes; however, the exact roles of MSCs osteogenic differentiation remain unclear (Li et al., 2021).

The use of lncRNA combined with scaffolding has only been investigated in certain recent publications. Mingyue Wang et al. and Zheng et al. and revealed that the lncRNAs HIF1A-AS1 and PWRN1-209 promoted the bone formation of MSCs on Ti implants (Wang et al., 2020; Zheng et al., 2020). The lncRNA LOC103691336 was found to be upregulated in magnesium-based biodegradable implants, and competed with the BMP2 for miR-138-5p-binding in MSCs to change the inhibitory effect of miR-138-5p on BMP2 expression (Li et al., 2019).

In general, various RNAs molecules, such as mRNA, miRNA, siRNA, and lncRNA, can be implanted as biomolecules in...
different types of scaffolds to enhance the bone osteogenesis, and some examples are summarized in Table 5.

RNA Delivery
RNA delivery is a challenging task due to the following reasons: i) RNA molecules are negatively charged with a complex structure to pass across the cell membrane, and ii) the single stranded RNA is highly susceptible to degradation via endogenous cellular enzymes (Sahay et al., 2010). However, the use of RNA-based therapies has increased in the last few decades to repair bone defects. Due to the advancements in nanotechnology and molecular biology these RNA particles can be easily synthesized and delivered through various vectors into the targeted bone. The addition of these RNAs in the implant relays to the different indispensable roles in gene expression and regulation, including molecular triggers, signaling pathways, cellular processes, and the transcriptional regulators in bone osteogenesis (Zhang et al., 2018; Leng et al., 2020) (Table 6).

Two RNA delivery methods that are commonly used are systematic and local delivery. In systematic delivery, different vectors are used to deliver therapeutic RNA into scaffolds, such as viruses, dependent factors, or independent factors, like lipids, and polymers. In local delivery, the defect site primarily utilizes a non-viral biocompatible scaffold (Figure 3). For nanoparticles-specifically polymers-non-viral delivery is the most common method of RNA delivery due to the high ability to protect the RNA from degradation and to support the in-cellular uptake and endosomal escape (Anderson et al., 2003). Lipids and lipid-like

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**TABLE 3** | List of miRNA and role in osteogenesis differentiation in MSCs cells modulated by miRNA scaffold therapy.

| miR | Target gene | Study | miR | Target gene | Study |
|-----|-------------|-------|-----|-------------|-------|
| miR-26a | Smad 1/5/8 (drosophila mothers against decapentaplegic) | Trompeter et al. (2013) | miR-26a | Osx through Gsk-β | Luzi et al. (2008) |
| miR-3960 | BMP | Hu et al. (2011) | miR-93 | 3 (glycogen synthase kinase) suppression | Yang et al. (2012) |
| miR-148B | NOG (noggin) | Mykhaylyk et al. (2008); Vosen et al. (2016) | miR-31 | Osx (osterix) | Baglio et al. (2013) |
| miR-135 | Smad 1/5/8 | Vosen et al. (2016) | miR-214 | | Shi K. et al. (2013) |
| miR-31 | Satt2 (special AT-rich sequence-binding protein 1) | Deng et al. (2013) | miR-637 | | Zhang et al. (2011) |
| miR-135 | Hoxa2 (homeobox 2) | Xie et al. (2016) | miR-145 | | Jia et al. (2013) |
| miR-2281 | | Diomede et al. (2016) | miR-143 | Runx2 (run-related transcription factor 2) through Hoxa 10 (homeobox a10) suppression | Godfrey et al. (2018) |
| | | | miR-27a | | Hassan et al. (2010) |
| | | | miR-23a | Runx2 through Sabt2 suppression | Qi et al. (2014) |
| | | | miR-27a | | Chen et al. (2014) |
| | | | miR-24 | Runx2 through FAK (focal adhesion kinase) suppression | Huang et al. (2012) |
| | | | miR-138 | Runx2 through TAG1 (transient axonal glycoprotein 1) suppression | |
| | | | miR-34a | Runx2 through HDAC6 (histone deacetylase 6) suppression | Huang et al. (2012) |

**TABLE 4** | Examples of genes targeted by siRNA used to understand osteogenesis.

| siRNA targeted gene | Finding | Study |
|-------------------|---------|-------|
| S100A4 | Silencing it induce osteogenic differentiation in periodontal ligament cells, via increase expression of osteoblastic markers (osteopontin and osteocalcin). | Kato et al. (2004) |
| Guanine nucleotide-binding protein (G protein) alpha subunit 1 (GNAS1) | Osteogenesis suppressor in MSCs; expression induction was detected by qRT-PCR and western blots of osteogenesis markers such as bone-specific sialoprotein (BSP), Cbfα1 and Osx. | Zhao and Ding (2007) |
| Nogging (NOG) | BMP2 expression increases causing induced osteoblastic differentiation in C2C12 cells, and enhance calvarial bone defects in rats. | Takayama et al. (2009); Nguyen et al. (2018) |
| NOG and GNAS | A high dose of BMP2, NOG, and GNAS delivery increased the cell death of human fetal osteoblast cell line (hFOB1.19) to more than 90% and the 50% less of cell proliferation comparing to the control. | Ramasubramanian et al. (2015) |
| Scaffolds                                                                 | Cell type                          | Gene            | Findings                                                                                                                                  | Study                                      |
|--------------------------------------------------------------------------|------------------------------------|-----------------|------------------------------------------------------------------------------------------------------------------------------------------|--------------------------------------------|
| SMAT-Ti (surface mechanical attrition treatment)                          | hBMSCs                             | miRNA, miRNA,   | The genes expression was upregulated (has-circ-0032599, has-circ-0032600, and has-circ-0032601) in SMAT-Ti scaffolds comparing to the annealed Ti. | Zhu et al. (2020)                          |
| Poly (ethylene glycol) (PEG)                                             | hMSCs                              | miRNA, siRNA    | Bone formation was improved in the rat calvarias bone defect after PEG gel implantation containing hMSCs and miRNA-20a compared to the hydrogels without siRNA or with negative control siRNA. | Nguyen et al. (2018)                       |
| 3D hybrid scaffolds (Composite ink made of polycaprolactone (PCL)/ poly(D,L-lactide-co-glycolide) (PLGA)/ hydroxyapatite nano-particles| Rat bone marrow stem cells (rBMSCs)| miR-148b        | In vitro: a significant upregulation of Runx2 levels for the miR-14b group comparing to the control, which indicates an early stage of bone differentiation during the bone remodeling, but not with osteocalcin (OCN) and alkaline phosphatase (ALP) expression. | Moncal et al. (2019)                       |
| 8-tricalcium phosphate (8-TCP)                                           | Mice bone marrow stem cells (mBMSCs)| miRNA-26a       | The micro-computed tomography, eosin, and toluidine blue staining showed an improvement in the bone repair after 8-TCP scaffolds co-cultured with the MSCs. High expression for ALP, Runx2, and osteocalcin was also observed on the transfected implant. | Liu et al. (2018)                          |
| Chitosan (Cs)/ hyaluronic acid (HA) nanoparticles (NPS) cross linked onto gel culture plate | hBMSCs                             | miR-21          | The combination of Cs/HA/miR-21 NPs delivery on the hBMSCs sheets showed an improvement on the osterogenic differentiation markers (OCN and OPN) and enhanced the ALP activity, collagen secretion, and bone nodule formation. | Wang et al. (2016)                         |
| CS/nano HA/ nano-zirconium dioxide (nZrO2)                              | Mouse MSCs                         | miR-590-5p      | The combination of CS/nHA/nZrO2/mBMSCs/ miR-590-5p suggested the potential of osteoconductive properties, by activating various signaling pathways, such as Runx2, Collagen type 1, and ALP. | Balagangadharan et al. (2018)               |
| Collagen-nHA                                                             | hMSCs                              | miR-16          | miR-16 may play an inhibitory role in osteogenesis due to its ability to directly target Smad5 and AcvR2a, which also could be used as a potential of a scaffold with the known potential for bone repair applications. | Mencia Castaño et al. (2019)               |
| CS sponge                                                                | MSCs                               | siRNA           | The CS sponge with siRNA significantly upregulated the OCN, ALP, and the vascular endothelia growth factor in vitro. In vivo: the critical size defect in the rat skull showed a marked bone regeneration using the CS sponge and siRNA treatment. | Jia et al. (2014)                          |
| Collagen sponge                                                          | C2C12 cells (osteoblast)            | siRNA           | BMP2 enhanced the osteoblast differentiation by noggin-targeted siRNA in vitro. In vivo, the collagen-retaining BMP2 discs was implanted (after noggin-silencing siRNA) and the bone mineral contents were improved after 2 weeks of surgery. | Takayama et al. (2009)                     |
| PEG/ poly (lactic acid)-dimethacrylate (PEG/PLA-GM) hydrogel            | In vivo (mice)                     | siRNA           | For the siRNA/NP that embedded within the gel, the diffusion could be controlled via encapsulation with tunable kinetics degradation and modeled for a delivery depot. | Wang et al., 2018)                         |
| Sand blasted, large-grit, acid-etched Ti (SLA-Ti)                        | hBMSCs                             | IncRNA          | IncRNA PWIRN1-209 enhanced ALP activity and osteogenic markers (e.g., Runx2, Col1, and Bsp) of MSCs cultured on microtopographic Ti comparing to the cells cultured on the flat Ti in vitro. | Wang et al. (2020)                         |
| SLA-Ti                                                                  | hBMSCs                             | IncRNA          | MSCs cultured on the SLA-Ti scaffolds showed high levels of HIF1A-AS1 and VEGFA expression, while the knockdown of HIF1A-AS1 inhibited the osteogenic differentiation by regulating the p38 MPK cascade proteins. | Zheng et al. (2020)                        |
materials are the second major approach of nanoparticle-based RNA delivery (Kaczmarek et al., 2017). Lipids are positively charged at acidic pH, which enhances the efficacy of endosomal escape (Schroeder et al., 2010), reducing the toxicity (Kanasty et al., 2013), and they have the capability to self-assemble into well-ordered nanoparticle structures called lipoplexes (Desigaux et al., 2007). In addition to the nanoparticles, for the direct conjugate a bioactive ligand such as N-acetylgalactosamine (Yu et al., 2016), antibodies (Xia et al., 2009), vitamins (Nishina et al., 2008), or cholesterol (Lorenz et al., 2004), can be used as an alternative method of RNA delivery. Additionally, another effective method of nucleic acid delivery are the chemical modifications made to the RNA itself that can impart degradation resistance to the RNAase, making them unrecognizable by the immune system (Soutschek et al., 2004; Morrissey et al., 2005). RNA chemical alterations to the ribose sugar, phosphate linkage, and individual bases can be used to deliver nucleic acids to the target receptors (Prakash et al., 2005; Wittrup and Lieberman, 2015; Li et al., 2016).

Several promising results have been found in various experimental studies implementing gene manipulated of MSC for treating bone defects, however these studies are still limited due to experimental caveats, and the safety and efficacy of the experiments need to be illustrated in the near future (Oryan et al., 2017). Also, developing a clinical-grade vector is a complicated, expensive process. No scaffold is currently in routine clinical use to deliver gene vector to the defect site. All the clinical trial results were not entirely satisfying, or were very limited to a few case studies, which require more investigations with longer follow-up (Kon et al., 2014; Madry et al., 2020).

**CRISPR to Guide RNA-Based Scaffolds**

To obtain a successful bone implants in tissue engineering, all osteogenesis parameters are ought to be controlled and
understood at molecular level. Traditional molecular methods can aid in this process; however, they have some limitations and require much experienced molecular biologist to obtain a genetically modified cell. MSCs are considered the primary used cell type used in studying bone regeneration and osteogenesis either to study the involved gene or to be included with scaffolds. However, some limitations were found in using it due to their ability to differentiate and the transplantation efficiency (Oryan et al., 2017; Arriaga et al., 2019). Henceforth, a novel and relatively easy genome editing approach has been implanted recently in the field of tissue engineering to control and understand osteogenesis at the molecular level. The bacteria adaptive immune system known as clustered regulatory interspaced short palindromic repeat (CRISPR) CRISPR-associated protein 9 (Cas9) (CRISPR/Cas9) has been mimicked recently to apply specific genome cuts in human cell lines (Yang et al., 2013).

This can occur by introducing into cells the Cas9 nuclease and a chimeric single guide RNA (sgRNA) complementary to the targeted genome segment, directed by the presence of the protospacer adjacent motif (PAM) sequence. The Cas9 nuclease guided by the sgRNA and the PAM sequence produces double strand breaks in the target genome sequence. The cells then repair this break via the non-homology end-joining (NHEJ) pathway, which may result in a frame shift mutation (insertion/deletion) that can affect the gene expression of the targeted gene (Figure 4A). The high success rate, low-relative cost and low off-target effects made this system widely used by researchers to introduce specific cuts to the genome and to change the gene expression.

Several other types of gene editing methods have emerged adapting the CRISPR/Cas9 system, such as CRISPR interference (CRISPRi) and CRISPR activation (CRISPRa) relying on the use of a modified Cas9 enzyme to alter the gene expression. CRISPRi works using a modified inactive Cas9 nuclease (dCas9) that blocks the targeted DNA transcription via sgRNA mediated binding. This results in silencing the targeted gene. On the other hand, CRISPRa stimulates gene expression of the target gene by fusion of the dCas9 with transcription activators, such as VP64, and this results in the gain of function of the targeted gene (Figures 4B,C) (Kampmann, 2018; Truong et al., 2019).

Other systems applied the CRISPR/Cas9 system for the live imaging of proteins, guided by the sgRNA to locate specific regions on the genome (Ma et al., 2018) as done by Narai et al. in which they used CRISPR technology to localize osteogenic differentiation in MSCs through the monitoring of bone gamma-carboxyglutamate protein (BGLAP) expression in vivo via an enhanced green fluorescent protein (EGFP) reporter (Narai et al., 2020).

**CRISPR/Cas9 in Bone Osteogenesis**

CRISPR/Cas9 gene silencing could be implemented to study the cellular control of osteogenesis genes, contributing to a better understanding of this vital cellular process.

A study by Lee el al. demonstrated that the CRISPR/Cas9-mediated gene silencing of PUMILO2 (PUM2, a conserved posttranscriptional regulator) inhibited lipid accumulation and induced excessive bone formation by blocking MSC adipogenesis and enhancing the osteogenesis. They also showed that PUM2 works as a negative regulator on the 3′-untranslated regions of...
**CRISPRi and CRISPRa in Bone Osteogenesis**

CRISPRi and CRISPRa have been used for different applications, such as genome-scale genetic screening (Bester et al., 2018), genetic interaction mapping (Du et al., 2017), cell signaling engineering (Liu et al., 2017), disease remodeling (Mandegar et al., 2016), and cell fate regulation (Black et al., 2016), also they can also be used to affect the gene expression of osteogenesis-related genes. Truong et al. developed a CRISPRai system that comprises active Cas9, activation/repression proteins complexes, and two single guide RNAs (sgRNAs) as a scaffold for recruiting activators (sgRNAa) or inhibitors (sgRNAi). They found that the CRISPR system delivered by the hybrid baculovirus stimulated chondrogenesis, and repressed the adipogenesis of rat BMSCs in 2D cultures, and stimulated the formation of engineered cartilage in 3D cultures, which may be of use to improve the calvarial bone healing (Truong et al., 2019). A more recent work by Hsu et al. showed that the hybrid baculovirus robustly activated endogenous Wnt10b and Fox2 for a long period of time and that the coactivation of Wnt10b and Fox2 successfully stimulated osteogenesis and repressed adipogenesis in vitro.

In vivo, the implantation of the CRISPRa-engineered BMSCs into the critical-sized calvarial defects in rat significantly improved bone healing (Hsu et al., 2020a). Another study from the same group reasoned that Noggin gene (Nog) inhibition, concurrent with BMP2 overexpression by using the CRISPRi system, could enhance the osteogenesis of adipose-derived stem cells and could improve calvarial bone healing (Hsu et al., 2020b).

There are some drawbacks to the use of CRISPR tools that can limit its in vivo applications, such as off-target effects if any of the sgRNAs were poorly designed. This could be avoided by the use of several sgRNAs for the same gene to increase the results validation or by using an enhanced version of Cas9 that has less off-target effects. Another tool is the use of a mutated Cas9 nuclease “Cas9 nickase (Cas9n)” that can induce a single strand break in two regions on the genome flanking the target gene sequence (Fu et al., 2013; Shen et al., 2014; Wu et al., 2019).

**CONCLUSION**

Despite the rapid evolution in bone tissue engineering, many challenges need to be solved to find the optimal bone implants in clinical applications. Numerous materials have been utilized in bone tissue engineering applications such as polymers and metals, and each has benefits and limitations. However, Ti materials were demonstrated to be the best implants in orthopedical and dental applications in vivo, due to their biocompatibility and mechanical properties that are close to the human bones.

In recent decades, the RNA-based scaffolds have shown promising bone osteogenesis findings as therapeutic molecules coated or delivered to the scaffolds. In this review, we summarized the effects of different types of RNAs on the bone formation of different types of scaffolds. RNAs are starting to have a significant role as biomarkers for bone osteogenesis. A better understanding of RNA upregulation, downregulation, and silencing will increase bone remolding, improve treatments, and enhance patient quality of life by finding a better solution for implant loss.

We also discussed using the CRISPR-based genome editing technology, which offers a new tool to understand osteogenesis in many possible ways in a cost and time-efficient manner. CRISPR/Cas9 had proven to be a successful tool in understanding osteogenesis and bone healing, as well as providing a novel method to control bone infection. The utilization of this cutting-edge technology in the future will not only be limited to understand osteogenesis by obtaining a genetically modified cells (e.g., MSCs), but it will also provide a new tool in in vivo therapeutics gene editing in defective bone cells. Generally, this technology provides insights at the molecular and cellular level and aids in directing the cells cultured on the scaffolds to enhance bone formation, which provides a new technology to be used clinically for bone implants. Future applications based on RNA-scaffolds-cell interactions may accelerate bone osteogenesis and control implant failure.

**AUTHOR CONTRIBUTIONS**

LD and SE-M contributed equally to the preparation of this manuscript.
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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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