

Bioengineered Let-7c Inhibits Orthotopic Hepatocellular Carcinoma and Improves Overall Survival with Minimal Immunogenicity

Joseph L. Jilek,^{1,3} Qian-Yu Zhang,^{1,3} Mei-Juan Tu,¹ Pui Yan Ho,¹ Zhijian Duan,¹ Jing-Xin Qiu,² and Ai-Ming Yu¹

¹Department of Biochemistry & Molecular Medicine, UC Davis School of Medicine, Sacramento, CA 95817, USA; ²Department of Pathology, Roswell Park Cancer Institute, Buffalo, NY 14263, USA

Hepatocellular carcinoma (HCC) remains a leading cause of cancer-related deaths, warranting better therapies. Restoration of tumor-suppressive microRNAs depleted in hepatocellular carcinoma represents a new therapeutic strategy. Herein, we sought to identify a potent microRNA (miRNA) agent that could alleviate HCC tumor burden and improve survival. Among a collection of bioengineered noncoding RNA molecules produced through bacterial fermentation, we identified let-7c agent as the most potent inhibitor of HCC cell viability. Bioengineered let-7c selectively modulated target gene expression (Lin-28 homolog B [LIN28B], AT-rich interactive domaincontaining protein 3B [ARID3B], B cell lymphoma-extra large [Bcl-xl], and c-Myc) in HCC cells, and consequently induced apoptosis and inhibited tumorsphere growth. When formulated with liposomal-branched polyethylenimine polyplex, bioengineered let-7c exhibited serum stability up to 24 h. Furthermore, liposomal polyplex-formulated let-7c could effectively reduce tumor burden and progression in orthotopic HCC mouse models, while linear polyethyleneimine-formulated let-7c to a lower degree, as revealed by live animal and ex vivo tissue imaging studies. This was also supported by reduced serum *a*-fetoprotein and bilirubin levels in let-7ctreated mice. In addition, lipopolyplex-formulated let-7c extended overall survival of HCC tumor-bearing mice and elicited no or minimal immune responses in healthy immunocompetent mice and human peripheral blood mononuclear cells. These results demonstrate that bioengineered let-7c is a promising molecule for advanced HCC therapy, and liposomal polyplex is a superior modality for in vivo RNA delivery.

INTRODUCTION

Hepatocellular carcinoma (HCC) is the most common malignancy of liver with an increasing incidence, a 5-year survival rate of 18%,¹ and an overall median survival of 29.8 months without improvement over time.² Given the molecular heterogeneity and complex etiology of HCC, development of pharmacological interventions has been slow.

Although nivolumab, and anti-PD1 monoclonal antibody, has recently been suggested as a promising therapy for HCC,³ the only molecularly targeted drugs indicated for HCC are the multi-kinase inhibitors, sorafenib and regorafenib, and these show only modest (approximately 3 months) improvements in overall survival.^{4,5} Over the past two decades, the discovery of aberrant microRNA (miRNA or miR) expression in cancer cells has evoked new targeted strategies for HCC therapy.⁶⁻¹² For example, tumor-suppressive miRNAs (e.g., miR-124, -199a, and -206) depleted in HCC tissues may be reintroduced into cancer cells to manage HCC progression.¹³⁻¹⁸ Because miRNA replacement presents a promising approach for cancer therapy,⁷ efforts to translate this strategy into clinical pharmacological interventions is warranted to improve upon current standards of care.

To investigate miRNA-based cancer therapies, we have developed a novel noncoding RNA (ncRNA) bioengineering platform to produce recombinant miRNA agents via bacterial fermentation,¹⁹ based upon a stable hybrid tRNA/pre-miRNA molecule (e.g., tRNA^{Met}/pre-miR-34a) identified in our lab.^{20,21} By replacing the miRNA (e.g., miR-34a) duplexes with target miRNA (e.g., let-7c) or small interfering RNA (siRNA; e.g., Nrf2-siRNA), we have produced a set of biologic miRNA or siRNA agents containing a common tRNA/pre-miR-34a molecular scaffold, which were all expressed at high levels in a common strain of Escherichia coli HST08 and purified to a high degree of homogeneity by anion exchange fast protein liquid chromatography (FPLC).²¹ These bioengineered miRNA agents are produced and folded in living cells, without or with minimal posttranscriptional modifications, and thus distinguished from conventional miRNA mimics or molecules that are produced by chemical synthesis and typically consisted of extensive degrees and various types of ribose and phosphate backbone modifications.^{19,22} Further studies have demonstrated that bioengineered noncoding RNA (ncRNA) agents are selectively processed to target miRNA or siRNA molecules in human cells, modulate target gene expression, and control cellular processes.^{20,21,23-26}

E-mail: aimyu@ucdavis.edu

Received 8 July 2018; accepted 15 January 2019; https://doi.org/10.1016/j.omtn.2019.01.007.

³These authors contributed equally to this work.

Correspondence: Ai-Ming Yu, PhD, Department of Biochemistry & Molecular Medicine, UC Davis School of Medicine, 2700 Stockton Blvd., Suite 2132, Sacramento, CA 95817, USA.



Figure 1. Bioengineered let-7c Molecule Is Identified as the Most Potent Inhibitor against HCC Cell Proliferation among a Small Collection of ncRNAs (A) Antiproliferative activities of a collection of bioengineered ncRNA agents (5 nM) against luciferase/GFP-expressing Huh7 cells were examined by luminometric ATP assay. Values were normalized to Lipofectamine 3000 transfection reagent/vehicle control (0% inhibition). (B) Dose-response curves of the top ranked ncRNA agents were further determined, and (C) their pharmacodynamic parameters were estimated, which indicate that let-7c is the most potent inhibitor of HCC cell viability in this collection of ncRNA agents. Values are mean \pm SD (n = 3 per group). *p < 0.05, compared with MSA control.

Systemic RNA therapy is hampered by the susceptibility of RNA molecules to serum RNases and the ability to cross the membrane barrier, warranting proper delivery systems. Polyethylenimine (PEI)-RNA complexes offer high delivery efficiency; however, PEI is cytotoxic with the increase of doses.²⁷ Lipidation of PEI-RNA polyplexes can reduce the toxicity of polyplexes²⁸ as the resultant lipopolyplexes (LPPs) exhibit more favorable biocompatibilities.^{29–31} Using a bioengineered GFP siRNA agent as model molecule, we have found that PEI-based cationic LPP nanocomplex offers efficient delivery of bioengineered RNA molecules in orthotopic HCC xenograft mouse models, leading to more consistent knockdown of target gene expression than polyplex in tumor tissue.³²

Herein, we present our findings on the identification of bioengineered let-7c as the most potent inhibitor against HCC cell viability among a small collection of recombinant miRNA or siRNA agents that are known to exhibit anti-proliferative activities. Following the validation and delineation of mechanistic actions of let-7c on target gene expression, as well as HCC cell stemness and apoptosis, our results demonstrate the utility of intravenously (i.v.) administered LPP/let-7c nanotherapeutics to reduce tumor progression and improve overall survival in orthotopic HCC xenograft mouse models. In addition, LPP-formulated let-7c treatment is well tolerated in mice, showing no or minimal immunogenicity in human peripheral blood mononuclear cells (PBMCs) and immunocompetent mice.

RESULTS

Bioengineered let-7c Is the Most Potent Inhibitor against HCC Cell Proliferation among a Collection of ncRNA Agents

Screening of a small collection of bioengineered miRNA agents was predictive for their anti-proliferative activities (Figure 1A), in which the truncated RNA, namely MSA (methionine tRNA with Sephadex aptamer), just containing the tRNA portion that was proven as a good control to define the actions of tRNA-carried miRNAs^{20,21}

consistently yielded the least inhibition of cell viability. Several miRNA agents, including miR-298, miR-124, let-7c, miR-328, miR-144, and miR-126, showed greater antiproliferative activities in Huh7 cells, and thus were pursued for dose-response studies (Figure 1B). Let-7c was revealed as the most potent ncRNA, with the lowest EC₅₀ value (0.51 nM) in the inhibition of Huh7 cell proliferation (Figure 1C). Furthermore, let-7c was as pure (>97%, by high-performance liquid chromatography [HPLC]) as other tested ncRNAs purified by the same anion exchange FPLC method²¹ and had a low endotoxin level (Figure S1), suggesting minimal interference by impurities.

Bioengineered let-7c Reduces Protein Levels of Target Genes

To verify the actions of recombinant let-7c prodrug, we first examined protein levels of several known let-7c targets important in cancer. Lin-28 homolog B (LIN28B), a canonical target of let-7 family miRNAs,³³ was reduced by biologic let-7c over 60% and 85% in Huh7 and Hep3B cells, respectively (Figure 2A). This was associated with much higher levels of let-7c in cells treated with biologic let-7c- than control MSA (data not shown). Furthermore, AT-rich interactive domain-containing protein 3B (ARID3B) protein level³⁴ was suppressed consistently by bioengineered let-7c prodrug in both Huh7 and Hep3B cells, whereas downregulation of B cell lymphoma-extra large (Bcl-xl)35 and c-Myc³⁶ was obvious only in Huh7 cells at 72 h post-transfection (Figure 2A). Interestingly, significant downregulation of c-Myc in let-7c-transfected Hep3B cells was observed in Hep3B cells at 48 h after transfection (data not shown). In addition, the expression of HMGA2,^{37,38} as detected by immunofluorescence, was reduced significantly by biologic let-7c in both Huh7 and Hep3B cells (Figure 2B).

Induction of Apoptosis of HCC Cells by Bioengineered let-7c

Let-7 family miRNAs have been shown to induce apoptosis via targeting of Bcl-xl, among other mechanisms.³⁵ Likewise, our data showed



Figure 2. Suppression of Protein Levels of Target Genes by Bioengineered let-7c Agent in HCC Cell Lines (A) Immunoblot analyses of let-7c targeted ARID3B, c-MYC, LIN28B, and BcI-xI protein levels and (B) immunofluorescent analysis of HMGA2 in Huh7 and Hep3B cells treated with 15 nM let-7c or control MSA. Immunoblot intensity was normalized to total protein and Lipofectamine 3000/vehicle control for comparison between groups. ^ap < 0.05 compared with vehicle; ^bp < 0.05 compared with MSA (one-way ANOVA with Bonferroni's post hoc test). Immunofluorescent intensity of HMGA2 staining was normalized to total DAPI-positive cells. *p < 0.05 compared with MSA (Student's t test).

that a low dose (5 nM) of biologic let-7c prodrug induced a modest yet robust increase in early and late apoptotic cell populations in Huh7 cells (Figure 3), compared with either vehicle or MSA control treatments. Additionally, necrotic cell populations were not altered following the transfection of let-7c and MSA.

Biologic let-7c Suppresses HCC Cell Stemness

Negative feedback between let-7 and LIN28 influences the stemness of cancer cells,³³ which can have important impact on therapeutic efficacy. As such, we evaluated cancer stem cell (CSC) growth using a tumorsphere assay in Huh7 cells. Following transfection in adherent conditions and subsequent growth in ultra-low-attachment, serum-free conditions, we observed a significant half-diameter reduction in primary tumorsphere size, but not tumorsphere count, in let-7c-treated cells (Figure 4). Upon subsequent dissociation, transfection, and growth in ultra-low-attachment, serum-free conditions to form secondary tumorspheres, a similar 50% reduction in diameter was observed in biologic let-7c-treated cells. Although sphere formation efficiency from primary tumorspheres was not significantly reduced, sphere count and individual cell number were significantly reduced in secondary tumorspheres by let-7c treatment (Figure 4).

Preparation and Characterization of Biologic let-7c-Loaded LPP Nanocomplex

We thus employed LPP to load bioengineered let-7c molecule (Figure 5A) toward therapy study in animal models because our recent study demonstrated the effectiveness of LPP to deliver biologic RNA molecules to liver tissues.³² The size of biologic let-7c-loaded LPP was 98.35 ± 5.11 nm with a zeta potential value of $43.9 \pm$ 2.2 mV, which was complemented by transmission electron microscopy (TEM) examination (Figure 5B). Control MSA was formulated in the same manner, and LPP/MSA nanocomplex showed similar size (102.4 \pm 5.9 nm) and zeta potential (45.1 ± 1.2 mV) (Figure S2). In addition, LPP could effectively protect bioengineered let-7c agent from degradation in both FBS and human serum up to 6 h, whereas modest degradation had occurred by 24 h (Figures S3A and S3B).

LPP Efficiently Delivers Biologic let-7c Prodrug into HCC Cells to Elicit Inhibition of Cell Growth

We further assessed *in vitro* delivery efficiency by LPP in Huh7 cells, in parallel to Lipofectamine 3000 (LF3000). Our data demonstrated that biologic let-7c prodrug was efficiently delivered into Huh7 cells by LPP nanocomplex, as manifested by the increase in comparable levels of let-7c as those delivered by LF3000 (Figure 5C), which led



Figure 3. Apoptotic Cell Death Is Significantly Induced by Bioengineered let-7c Agent

Huh7 cells were transfected with 5 nM Lipofectamine 3000-carried MSA or let-7c for 48 h, stained with propidium iodide and Annexin V-FITC, and counted by a flow cytometer with a total cell gate of 10,000 events. A significant shift of the total population toward early and late apoptotic cells was observed, whereas total necrotic population showed no difference. Values are mean \pm SD (n = 3 per group). *p < 0.05; **p < 0.01; ***p < 0.001 (oneway ANOVA with Bonferroni's post hoc test).

to a sharp suppression of cell proliferation (Figure 5D). These data were also complemented by efficient delivery of another bioengineered ncRNA molecule, GFP-siRNA,^{20,21} by LPP nanoparticles, as indicated by the knockdown of target GFP levels in GFP/luciferaseexpressing Huh7 cells (Figure S4).

Bioengineered let-7c Prodrug Significantly Reduces HCC Tumor Progression in Orthotopic Xenograft Mouse Models and Is Well Tolerated

We thus established orthotopic HCC xenograft mouse models with luciferase/GFP-expressing Huh7 cells to investigate the efficacy of let-7c therapy (Figure 6A). As revealed by bioluminescent imaging in live animals (Figure 6B), HCC tumor burden was inhibited by approximately 50% by both i.v. administered LPP- and in vivo-jetPEI (IPEI)-formulated let-7c prodrug, compared with untreated mice, whereas control MSA had no impact. Ex vivo imaging of liver tumoral GFP signals (Figure 6C) further demonstrated the effectiveness of let-7c for the control of HCC, which was reduced over 70% by LPP/let-7c and around 33% by IPEI/let-7c as compared with untreated mice. Suppression of HCC was associated with higher levels of let-7c in both healthy livers and tumors isolated from let-7c-treated mice (Figure 6D). Moreover, the efficacy of biologic let-7c therapy in the inhibition of orthotopic HCC was supported by significantly lower serum α-fetoprotein (AFP) levels in let-7c-treated mice (Figure 6E), as well as less HCC tumors in let-7c-treated mouse liver slices with variable degrees of necrosis determined by histopathological analyses (Figure 6F). In addition, immunohistochemistry (IHC) studies on HCC xenograft tissues demonstrated that, although there is no significant difference in the level of cell proliferation (i.e., Ki-67) among various treatments, biologic let-7c prodrug therapy, especially the LPP formulation, led to an obvious higher degree of apoptosis (i.e., active-caspase-3), as compared with MSA control (Figure S5). It is also noteworthy that LPP/let-7c nanotherapeutics was efficient in the control of HCC, as indicated by a more ubiquitous and significantly greater degree of reduction of tumor burden determined by multiple measurements (Figure 6).

All treatments were well tolerated as body weights of all mice showed steady increases over time (Figure S6A). To further investigate the safety of bioengineered let-7c prodrug, blood biochemistry profiles were determined (Figures S6B–S6F). Biomarkers of hepatic and renal functions including alanine aminotransferase (ALT), aspartate aminotransferase (AST), blood urea nitrogen (BUN), and creatinine

were all within the normal ranges. To our surprise, blood total bilirubin levels in untreated and MSA-treated mice were highly variable and inclined to be above normal range, whereas they retained within normal range in biologic let-7c-treated mice, which may be another indication of the effectiveness of let-7c prodrug therapy in the control of HCC.

LPP/let-7c Nanotherapeutics Significantly Improves the Overall Survival of Orthotopic HCC Tumor-Bearing Mice

A separate cohort of orthotopic HCC Huh7 xenograft mice was established and employed to define the magnitude of benefit of i.v. administered LPP/let-7c nanotherapeutics on overall survival. After the development of HCC was confirmed by quantitative bioluminescence imaging of live mice, subjects showing the same degrees of tumor burden were randomized for LPP/let-7c and control LPP/MSA treatments (Figure 7A). Survival analysis showed that, compared with LPP/MSA, LPP/let-7c therapy significantly improved overall survival of HCC tumor-bearing mice (Figure 7B). This was also indicated by a longer median survival of LPP/let-7c-treated mice (26.0 days) than LPP/MSA controls (19.5 days). In agreement with the safety profiles of let-7c treatment in the other therapy study (Figure S6), LPP/let-7c treatment did not alter mouse body weights compared with LPP/MSA (Figure 7C).

LPP/let-7c Produces No or Minimal Immunogenicity in Human PBMCs and Immunocompetent Mice

Lastly, we assessed whether LPP/let-7c nanotherapeutics induces any immune response in human PBMCs and two different strains of healthy immunocompetent mice (BALB/c and CD-1). As expected, LPS treatment provoked a robust cytokine release in both human PBMCs (Figure 8A) and BALB/c (Figure 8B) and CD-1 (Figure 8C) mice, as indicated by a significantly sharp 1,000- to 30,000-fold elevation of interleukin (IL)-6 levels, as well as increase in tumor necrosis factor alpha (TNF- α) and IL-10 levels. By contrast, LPP/let-7c treatment did not alter the levels of IL-6, IL-4, or IL-10 in human PBMCs, whereas it slightly increased TNF- α levels, which is not statistically significantly different from untreated cells. Although LPP/vehicle, LPP/MSA, and LPP/let-7c all caused a mild increase in serum IL-6 levels in BALB/c and CD-1 mice, as compared with untreated mice, the elevated IL-6 levels were two to three orders of magnitude lower than those induced by LPS, indicating that LPP/let-7c shows minimal immunogenicity compared with LPS.



Figure 4. Bioengineered let-7c Prodrug Sharply Reduces Tumorsphere Growth

Following transfection with Lipofectamine 3000-formulated MSA or let-7c in adherent conditions, an equal number of Huh7 cells were grown in serum-free, ultra-lowattachment conditions for 7 days to yield primary tumorspheres. Primary tumorspheres were then digested to single cell, transfected again, and grown for another 7 days in serum-free, ultra-low attachment conditions to yield secondary tumorspheres. let-7c treatment resulted in smaller primary and secondary tumorspheres. Values are mean \pm SD (n = 3 per group). *p < 0.05; **p < 0.01; ***p < 0.001 (one-way ANOVA with Bonferroni's post hoc test).

DISCUSSION

miRNA replacement therapy represents a promising strategy for the control of tumor progression given the loss of expression of tumorsuppressive miRNAs in cancerous cells, including miR-122 and let-7 family miRNAs (e.g., let-7c and let-7a) in human HCC samples.^{35,39,40} However, due to the complexity in dysregulation of miRNAs, as well as other regulatory factors and pathways, reintroduction of functional miRNAs may not necessarily coincide with optimal efficacy. As such, although miR-122 is the most abundant hepatic miRNA and is frequently attenuated in HCC,⁴¹ we found that bioengineered let-7c showed the highest antiproliferative activity against HCC cells among a small collection of ncRNA agents including miR-122. Although the *in vivo* efficacy of other miRNAs is not compared with let-7c herein, this screening method is predictive of potential benefits of let-7c in relieving *in vivo* HCC tumor burden and improving overall survival revealed in this study.

Current miRNA research and drug development primarily uses synthetic miRNA mimics, which carry extensive and variable chemical modifications expected to improve systemic stability.¹⁹ Although these miRNA agents are assumed to retain the same "sequences" because of the presence of identical nucleobases, they are chemically different molecules and inevitably have distinct higher-order structures, as well as physiochemical and biological activities. Moreover, synthetic RNA agents pose risk for the induction of cytokine release syndrome.^{42,43} This is also in contrast to protein research and therapy, wherein recombinant or bioengineered proteins expressed in living cells have been preferably utilized instead of synthetic polypeptides and proteins. Bioengineered miRNA molecules presented herein represent a novel class of biologic miRNA agents, which are folded and tolerated in living cells, and thus may better capture the properties of cellular RNA macromolecules.^{19,22} With minimal natural modifications and exhibiting favorable stability in human cells,^{25,44} recombinant miRNA agents are selectively processed to target mature miRNAs that rewrite cellular miRNome profile and execute regulatory functions.²¹

The pleiotropic nature of miRNA-controlled gene regulation behind cancer cellular processes warrants extensive validation. The interplay between LIN28 and let-7 family miRNAs^{33,45} is a critical component in the regulation of pluripotency as well as HCC and other liver diseases.⁴⁶ LIN28, shown to be upregulated in stem-like cells, can reprogram cells into an undifferentiated state,⁴⁷ and thus LIN28 may be a druggable target for the suppression of CSCs and tumor initiation. By contrast, LIN28-regulatory *let-7* family miRNAs shown to inhibit pluripotency and favor differentiation may be employed to manage CSC maintenance and replication.^{48,49} This study demonstrated a consistent action of bioengineered let-7c prodrug in the inhibition of tumorsphere growth, which is likely attributable to the



Figure 5. Biologic let-7c Is Efficiently Delivered into HCC Cells by LPP Nanocomplex to Control Cell Proliferation

(A) Schematic illustration of let-7c-loaded lipopolyplex (LPP). (B) TEM image of let-7c-loaded LPP (indicated by arrows) nanocomplex, as well as the size and zeta potential measured by dynamic light scattering. Scale bar indicates 500 nm. (C) Efficient delivery of let-7c (15 nM) into Huh7 cells led to (D) sharp suppression of cell growth. Lipofectamine 3000 (LF3000) treatments were used for comparison. Values are mean \pm SD of triplicate treatments (n = 3 per group). **p < 0.01; ***p < 0.001 (one- or two-way ANOVA with Bonferroni's post hoc test).

strong suppression of LIN28B expression in Huh7 cells that is consistently shown in Hep3B cells as well. Moreover, induction of apoptosis is a common mechanism of antineoplastic agents, and resistance to apoptosis is a common feature of CSCs. let-7 family miRNAs have also been shown to either induce or sensitize cells to apoptosis via attenuation of anti-apoptotic proteins, including Bcl-xl.³⁵ In this study, we found the suppression of Bcl-xl expression by biologic let-7c in Huh7 cells, which is consistent with the induction of apoptotic, but not necrotic, cell populations by a low dose of let-7c prodrug. It is also interesting to observe variable effects of let-7c on different targets (e.g., c-MYC versus LIN28B) in the same cell line and/or the same target (e.g., Bcl-xl) in different cell lines (e.g., Huh7 versus Hep3B). This is understandable because a specific gene is often regulated by multiple regulatory mechanisms; among them, miRNA-controlled posttranscriptional regulation may have certain degree of contribution to that gene. In addition, genotypes and regulatory signaling pathways, as well as miRNA machinery, may be variable in different HCC cell lines, highlighting the challenges in molecular therapy and demands for precision medication.

RNA drugs for systemic administration currently under clinical investigation are mainly delivered by lipid-based systems (e.g., liposomes), given their excellent biocompatibility and favorable lipid composition.^{50–52} For example, a phase I trial is under way to evaluate a small activating, double-stranded RNA targeting the transcription factor C/EBP- α formulated in SMARTICLES liposomal nanoparticle for advanced HCC (https://clinicaltrials.gov/ct2/show/NCT02716012). Among lipid-based delivery systems, LPPs convey the favorable properties of both liposomes and polyplexes.^{28,29,53} Our recent studies have demonstrated that IPEI is able to deliver biologic RNAs to livers to achieve target gene knockdown,²⁰ as well as tumor tissues to control disease.^{21,23,26,54} In the present study, we identified a more favorable serum stability of let-7c formulated in LPP nanocomplex, which was improved to even greater degrees as compared with IPEI,³² owing to the outer PEGylated lipid coating of polyplex. As a result, LPP showed high in vitro delivery efficiency in HCC cell lines. Most notably, LPP/let-7c provided a significantly greater extent of suppression of orthotopic HCC tumor burden in vivo, consistently supported

by multiple independent endpoints including *ex vivo* GFP reporter intensity, serum AFP level, and histological tumor area. In addition, we revealed that LPP/let-7c nanotherapeutics significantly improved the median survival of orthotopic HCC mice by 6.5 days. Although modest, the HCC xenograft used in this study is incredibly aggressive, occupying more than 50% of total liver tissue after 2 weeks.

Consistent with previous findings,^{25,26} this study demonstrated that bioengineered RNAs were well tolerated in HCC tumor-bearing immunodeficient mice and caused no or minimal degree of cytokine release in immunocompetent mice. Interestingly, serum bilirubin level, an indicator of liver damage, fell within normal range in let-7c-treated mice only. This is likely attributable to the effectiveness of let-7c therapy in the control of HCC tumor growth, suppressing further liver damage, and highlighting the aggressive nature of this HCC model.⁵⁵ Moreover, the present study showed for the first time that bioengineered RNAs show minimal immunogenicity in human PBMCs, an addition to the safety profile of recombinant miRNA molecules that are produced and folded in living cells.

In conclusion, we have demonstrated the efficacy of LPP/let-7c nanotherapeutics in an aggressive orthotopic HCC tumor mouse model, showing marginal immunogenicity in mice and human PBMCs. The first-in-class biologic let-7c prodrug was identified as the most potent among a small set of miRNAs in inhibiting HCC cell viability via interference of specific targets and critical cellular functions. Our findings suggest that LPP-formulated biologic let-7c may serve as an effective and safe treatment for HCC, warranting clinical verification.

MATERIALS AND METHODS

Materials

In vivo-jetPEI (linear 22-kDa PEI; *in vivo*-jetPEI polyethylenimine [lPEI]) was purchased from Polyplus Transfection (Illkirch, France). Branched PEI with molecular weight 10,000 Da (bPEI10k) was bought from Alfa Aesar (Wardhill, MA, USA). 1,2-Di-O-octade-cenyl-3-trimethylammonium propane (DOTMA) and cholesterol were purchased from Avanti Polar Lipids (Alabaster, AL, USA).



Figure 6. Systemic Administration of LPP/let-7c Nanotherapeutics Largely Reduces Tumor Growth in Orthotopic HCC Huh7 Xenograft Mouse Models (A) Timeline of establishment of HCC xenograft mouse models and drug treatment. (B) Suppression of orthotopic HCC progression by LPP/let-7c was demonstrated by live animal imaging of luciferase bioluminescent signals. *In vivo*-jetPEI (IPEI)-formulated biologic let-7c and MSA were used for comparison. (C) *Ex vivo* GFP fluorescence images of HCC-bearing livers further demonstrated the effectiveness of bioengineered let-7c prodrug, which was associated with high levels of tumoral let-7c (D). (E) Serum AFP levels were significantly reduced in let-7c-treated mice. (F) Representative H&E staining images (×40) of tumor-bearing liver tissues and quantitative measurement of the percentage of tumors in corresponding liver slices. Areas circled in blue lines are necrotic areas, and the red lines are applied to distinguish tumors from healthy liver (L) tissues. Values are mean \pm SD (n = 4–6 in each group). *p < 0.05; ***p < 0.001 (one- or two-way ANOVA with Bonferroni's post hoc test).

1,2-Dimyristoyl-sn-glycerol, methoxypolyethylene glycol (DMG-PEG2000) was purchased from NOF America Corporation (White Plains, NY, USA). LF3000, TRIzol RNA isolation reagent, and bicinchoninic acid assay (BCA) protein assay kit were purchased from Thermo Fisher Scientific (Waltham, MA, USA). Direct-zol RNA MiniPrep Kit was from Zymo Research (Irvine, CA, USA). Celltiter-Glo assay was purchased from Promega (Madison, WI, USA). Matrigel was purchased from Corning (Corning, NY, USA). Human AFP ELISA kit was purchased from R&D Systems (San Diego, CA, USA). All other chemicals and organic solvents of analytical grade were purchased from Sigma-Aldrich or Thermo Fisher Scientific.

Cell Culture

Huh7 cells were purchased from the Japanese Collection of Research Bioresources and grown in DMEM (GIBCO, Grand Island, NY, USA), and Hep3B and Sk-Hep-1 cells were obtained from American Type Culture Collection and grown in Eagle's minimal essential medium (Cellgro, Manassas, VA, USA). All of the cell lines were supplemented with 10% fetal bovine serum (GIBCO, Grand Island, NY, USA) and 1% antibiotic/antimycotic (Cellgro, Manassas, VA, USA). GFP/luciferase-expressing cell lines were established after transduction of parental cells with pCCLc-Luc-EGFP lentiviral constructs (Vector Core; UC Davis Medical Center, Sacramento, CA, USA). All cells were transfected by using the LF3000 (Invitrogen, Carlsbad, CA, USA) per the manufacturer's instructions for *in vitro* RNA delivery, unless otherwise indicated.

Production of Recombinant miRNA Agents

Bioengineering of miRNA molecules was conducted as described recently.²¹ In brief, inserts encoding target miRNA- or siRNA-containing ncRNAs were cloned into a pBSMrna vector using In-Fusion cloning technology and transformed into HST08 *Escherichia coli*. Recombinant ncRNAs were purified by anion exchange FPLC^{21,24} to >96% purity as determined by a HPLC assay.²⁵ Bioengineered let-7c less than 95% pure was re-purified by the same FPLC method to reach >96% homogeneity (Figure S1). Endotoxin activity was examined with Pyrogent-5000 kinetic LAL assay (Lonza, Walkers-ville, MD, USA).



Figure 7. LPP/let-7c Nanotherapeutics Significantly Improves Overall Survival of Orthotopic HCC Xenograft Tumor-Bearing Mice

(A) Bioluminescence images of HCC tumor-bearing animals before the i.v. treatment with LPP/let-7c and control LPP/MSA, and quantitative measurement of bioluminescent intensities. (B) Survival analysis showed that LPP/let-7c-treated mice lived much longer than the control (**p < 0.01; n = 10 per group; log rank [Mantel-Cox] test). The median survival was 26.0 days for LPP/let-7c-treated mice and 19.5 days for LPP/MSA-treated animals. (C) Mouse body weights during the treatment. \checkmark indicates days on which mice received treatments.

c-Myc rabbit mAb [CST 13987], and LIN28B rabbit mAb [CST 11965] from Cell Signaling; ARID3B rabbit polyclonal antibody [pAb] [AB 92328] from Abcam) overnight at 4°C at 1:1,000 dilution in 5% BSA in TBS-T, followed by horseradish peroxidase-conjugated goatanti-rabbit secondary antibody (1:10,000 dilution [111-035-003] from Jackson Immunoresearch) for 2 h at room temperature prior to

Cell Viability Assay

GFP/luciferase-expressing Huh7 cells (5,000 cells/well) were seeded in 96-well plates and grown overnight. Biologic let-7c or MSA was administered in triplicate with either LF3000 or LPP. Cell viability was measured using Cell Titer-Glo kit. Inhibition of cell viability was determined as relative to vehicle control (0% inhibition), and pharmacodynamics parameters were estimated by fitting the data to a normalized dose-response equation with variable slope:

$$Y = \frac{100}{1 + 10^{(LogEC_{50} - X) * Hill Slope}}$$

Given low efficacy, MSA (E_{max} , $E_{min} = 100$, 20.64%) and miR-144 (E_{max} , $E_{min} = 44.42$, 17.27%) in Huh7 cells were best fit to the full dose-response equation:

$$Y = E_{min} + \frac{E_{max} - E_{min}}{1 + 10^{(LogEC_{50} - X) * Hill Slope}}$$

Immunoblot and Immunofluorescence Analyses

Huh7 and Hep3B cells were seeded in a six-well plate at 300,000 cells/well and transfected with 15 nM RNA. After 72 h, cells were harvested and lysed in radio immunoprecipitation assay (RIPA) buffer with protease inhibitor (Pierce, Rockford, IL, USA). Protein levels were determined by BCA assay (Pierce, Rockford, IL, USA). After separation on a 12% SDS-PAGE gel (Bio-Rad, Hercules, CA, USA), proteins were transferred onto a polyvinylidene difluoride membrane and blocked in 5% milk/0.1% Tween 20 in Tris-buffered saline. Total immobilized protein was imaged per the manufacturer's instructions. Membranes were incubated with primary antibodies (Bcl-xl rabbit monoclonal antibody [mAb] [CST 2764],

chemiluminescent imaging with Clarity ECL (Bio-Rad, Hercules, CA, USA). Relative band intensity was normalized to total immobilized protein.

To assay HMGA2 expression, Huh7 and Hep3B cells were grown on glass chamber slides and transfected with MSA or let-7c. After 72 h, cells were fixed with 10% formalin, permeabilized with 0.3% Triton X-100, and incubated with HMGA2 rabbit mAb (CST 8179) (1:400 dilution in 5% BSA) overnight at 4°C. Antigen was detected with Alexa 488-conjugated anti-rabbit IgG Fab fragment (CST 4412), and nuclei were counterstained with DAPI (CST 4803).

Flow Cytometry

Huh7 cells were plated in six-well plates at a density of 150,000 cells/ well and transfected with 5 nM RNA. After 48 h, cells were stained with propidium iodide (PI) and Annexin V-FITC (fluorescein isothiocyanate) per manufacturer's instructions (Trevigen, Gaithersburg, MD, USA). Cell count and fluorophore intensity were measured using a BD Biosciences Fortessa 20 color cytometer. Total event count was gated at 10,000 events, and quadrant gating was set relative to vehicle control. Healthy cells were defined by low PI/low Annexin V staining, early apoptotic cells were defined by low PI/high Annexin V, late apoptotic cells were defined by high PI/high Annexin V, and necrotic cells were defined by high PI/low Annexin V intensity.

Tumorsphere Assay

Huh7 cells were seeded under adherent conditions at a density of 300,000 cells/well in six-well plates and transfected with 15 nM



Figure 8. Safety of LPP/let-7c Nanotherapeutics

LPP/let-7c has no or limited impact on cytokine release in human PBMCs (A) and two stains of immunocompetent mice (B and C). LPS was used as positive control to induce cytokine release storm, whereas untreated and LPP vehicle-treated mice or cells were considered as negative controls. Values are mean \pm SD. For BALB/c and CD-1 mice, three females and three males were included in each group (n = 6). Blood was harvested 1 h after i.v. administration of formulated RNA molecules, and serum samples were prepared for cytokine measurement. For human PBMCs, each treatment was conducted in triplicate (n = 3), and cell culture medium was collected 24 h post-treatment. **p < 0.01; **p < 0.001 (two-way ANOVA with Bonferroni's post hoc test). n.d., non-detectable.

RNA. After 48 h, live cells were transferred to 24-well ultra-lowattachment plates (Corning, Kennebunk, ME, USA) at a density of 2,500 cells/well and grown in DMEM/F12+B27 with penicillin/streptomycin, GlutaMAX (GIBCO, Grand Island, NY, USA), 20 ng/mL human epidermal growth factor, and 10 ng/mL human basic fibroblast growth factor (PeproTech, Rocky Hill, NJ, USA). After 7 days, primary tumorspheres (>10 μ m diameter) were counted, and sphere diameter was measured in ImageJ and dissociated with trypsin to a single cell. After all cells were transferred to new wells in ultra-lowattachment, serum-free conditions, cells were transfected again with 15 nM RNA. After 7 days, secondary tumorspheres were again counted, and diameter was measured and dissociated to count individual cells. Sphere formation efficiency (%) was calculated relative to total single cells seeded from the previous generation.

Formulation and Characterization of LPP Nanocomplex

Details of formulation and characterization of LPP nanocomplex, as well as serum stability and *in vitro* delivery efficiency, are provided in the Supplemental Methods and as previously described.²⁹

Therapy Studies in Orthotopic HCC Xenograft Mouse Models

All animal procedures were approved by the Institutional Animal Care and Use Committee of the University of California, Davis.

Establishment of Orthotopic HCC Xenograft Mouse Model

Luciferase/GFP-expressing Huh7 cells were mixed with Matrigel to a final concentration of 1×10^8 cells/mL. Four-week-old male athymic nude mice (Jackson Laboratory, Bar Harbor, ME, USA) were anesthetized, and an incision (~1 cm) along the linea alba in the midline of the abdominal muscle layer was made. 20 µL of Huh7 cells in Matrigel suspension (2×10^6 cells) was injected into the left lobe of liver. Successful engraftment of Huh7 cells was confirmed by bioluminescent imaging using ChemiDoc MP Imaging System (Bio-Rad, Hercules, CA, USA), following the intraperitoneal injection of D-luciferin (150 mg/kg) (BioVision, Milpitas, CA, USA).

Tumor Progression Study

One week post-inoculation, mice were assigned into five groups (untreated, IPEI/MSA, LPP/MSA, IPEI/let-7c, and LPP/let-7c) according to tumor sizes determined by *in vivo* bioluminescence imaging, and treated i.v. (40 μ g RNA) three times per week through tail vein injection. Mice were imaged once per week to monitor tumor growth. Mice were sacrificed 2 h after the last dose on day 15. Livers with engrafted tumors were harvested and imaged for GFP fluorescence using ChemiDoc MP Imaging System. RNA was extracted from healthy livers and HCC tumors, and the levels of let-7c were quantified using stem loop qRT-PCR using gene-selective primers (Table S1). Blood was collected for blood chemistry profiling (Comparative Pathology Laboratory, UC Davis), and serum AFP was examined by ELISA (R&D Systems, Minneapolis, MN, USA). H&E histopathological study was performed by the Clinical Immunohistochemistry Laboratory at Roswell Park Cancer Institute (Buffalo, NY, USA).

Survival Study

One week post-inoculation, a separate cohort of tumor-bearing mice was assigned into two groups (LPP/MSA and LPP/let-7c) according to tumor sizes determined by bioluminescence imaging, and treated i.v. with 40 μ g RNA three times per week for 3 weeks. Body weights were recorded twice a week to assess animal health. Survival was analyzed by Kaplan-Meier method and compared by log rank (Mantel-Cox) test.

Induction of Cytokine Release

Human PBMCs were purchased from Lonza (Walkerville, MD, USA) and maintained in RPMI 1640 supplemented with 10% human AB serum (Sigma, St. Louis, MO, USA). PBMCs were seeded onto a 96-well plate at a density of 2×10^5 cells/well and allowed to grow overnight. Cells were treated with 10 ng/mL or 100 ng/mL LPS (positive control), LPP/MSA (5 nM), LPP/let-7c (5 nM), or LPP vehicle. Twenty-four hours post-treatment, medium was harvested and cell debris was removed by centrifugation. IL-6, TNF- α , IL-4, and IL-10 levels were quantified using corresponding human cytokine ELISA assay kit (Invitrogen, Carlsbad, CA, USA).

Healthy BALB/c and CD-1 mice (5–6 weeks old) were randomly assigned into different groups (three females and three males per group) and injected i.v. with 40 µg of either MSA or let-7c-loaded LPP nanocomplex, 20 µg of LPS (positive control), or LPP vehicle (negative control). Blood was collected 1 h post-treatment, and serum IL-6 and IL-4 levels were quantified using a mouse IL-6 and IL-4 ELISA assay kit (Invitrogen, Carlsbad, CA, USA).

Statistical Analysis

All values are mean \pm SD. Statistical analysis was performed using one- or two-way ANOVA with Bonferroni's post hoc test or Student's t test where appropriate (GraphPad Prism, San Diego, CA, USA).

SUPPLEMENTAL INFORMATION

Supplemental Information includes Supplemental Methods, one table, and six figures and can be found with this article online at https://doi.org/10.1016/j.omtn.2019.01.007.

AUTHOR CONTRIBUTIONS

Conceptualization, A.-M.Y., J.L.J., Q.-Y.Z., and M.-J.T.; Methodology, J.L.J., Q.-Y.Z., M.-J.T., P.Y.H., Z.D., and A.-M.Y.; Resources, A.-M.Y., J.L.J., Q.-Y.Z., P.Y.H., and M.-J.T.; Investigation, J.L.J., Q.-Y.Z., P.Y.H., M.-J.T., and J.-X.Q.; Writing, J.L.J., Q.-Y.Z., M.-J.T., A.-M.Y., and all other authors; Visualization, J.L.J., Q.-Y.Z., J.-X.Q., and A.-M.Y.; Funding Acquisition, A.-M.Y.; Supervision, A.-M.Y.

ACKNOWLEDGMENTS

This study was supported by the National Institute of General Medical Sciences (grant R01GM113888 to A.-M.Y.) and in part by the National Cancer Institute (grants U01CA175315 and R01CA225958 to A.-M.Y.) of the NIH. J.L.J. was supported by an NIGMS-funded Pharmacology Training Program grant (T32GM099608). The authors appreciate the access to the Flow Cytometry and Molecular Pharmacology Shared Resources funded by the UC Davis Comprehensive Cancer Center Support Grant (CCSG) awarded by the National Cancer Institute (P30CA093373).

REFERENCES

- Siegel, R.L., Miller, K.D., and Jemal, A. (2018). Cancer statistics, 2018. CA Cancer J. Clin. 68, 7–30.
- Kim, N.G., Nguyen, P.P., Dang, H., Kumari, R., Garcia, G., Esquivel, C.O., and Nguyen, M.H. (2018). Temporal trends in disease presentation and survival of patients with hepatocellular carcinoma: a real-world experience from 1998 to 2015. Cancer 124, 2588–2598.
- Iňarrairaegui, M., Melero, I., and Sangro, B. (2018). Immunotherapy of hepatocellular carcinoma: facts and hopes. Clin. Cancer Res. 24, 1518–1524.
- 4. Bruix, J., Qin, S., Merle, P., Granito, A., Huang, Y.H., Bodoky, G., Pracht, M., Yokosuka, O., Rosmorduc, O., Breder, V., et al.; RESORCE Investigators (2017). Regorafenib for patients with hepatocellular carcinoma who progressed on sorafenib treatment (RESORCE): a randomised, double-blind, placebo-controlled, phase 3 trial. Lancet 389, 56–66.
- Llovet, J.M., Ricci, S., Mazzaferro, V., Hilgard, P., Gane, E., Blanc, J.F., de Oliveira, A.C., Santoro, A., Raoul, J.L., Forner, A., et al.; SHARP Investigators Study Group (2008). Sorafenib in advanced hepatocellular carcinoma. N. Engl. J. Med. 359, 378–390.
- 6. Ambros, V. (2004). The functions of animal microRNAs. Nature 431, 350-355.
- Bader, A.G., Brown, D., and Winkler, M. (2010). The promise of microRNA replacement therapy. Cancer Res. 70, 7027–7030.
- Klingenberg, M., Matsuda, A., Diederichs, S., and Patel, T. (2017). Non-coding RNA in hepatocellular carcinoma: mechanisms, biomarkers and therapeutic targets. J. Hepatol. 67, 603–618.
- Kota, J., Chivukula, R.R., O'Donnell, K.A., Wentzel, E.A., Montgomery, C.L., Hwang, H.W., Chang, T.C., Vivekanandan, P., Torbenson, M., Clark, K.R., et al. (2009). Therapeutic microRNA delivery suppresses tumorigenesis in a murine liver cancer model. Cell 137, 1005–1017.
- Rupaimoole, R., and Slack, F.J. (2017). MicroRNA therapeutics: towards a new era for the management of cancer and other diseases. Nat. Rev. Drug Discov. 16, 203–222.
- Yu, A.M., Tian, Y., Tu, M.J., Ho, P.Y., and Jilek, J.L. (2016). MicroRNA pharmacoepigenetics: posttranscriptional regulation mechanisms behind variable drug disposition and strategy to develop more effective therapy. Drug Metab. Dispos. 44, 308–319.
- 12. Xie, M., Ma, L., Xu, T., Pan, Y., Wang, Q., Wei, Y., and Shu, Y. (2018). Potential regulatory roles of microRNAs and long noncoding RNAs in anticancer therapies. Mol. Ther. Nucleic Acids 13, 233–243.
- 13. Callegari, E., D'Abundo, L., Guerriero, P., Simioni, C., Elamin, B.K., Russo, M., Cani, A., Bassi, C., Zagatti, B., Giacomelli, L., et al. (2018). miR-199a-3p modulates MTOR and PAK4 pathways and inhibits tumor growth in a hepatocellular carcinoma transgenic mouse model. Mol. Ther. Nucleic Acids 11, 485–493.
- Fu, X., Rivera, A., Tao, L., De Geest, B., and Zhang, X. (2012). Construction of an oncolytic herpes simplex virus that precisely targets hepatocellular carcinoma cells. Mol. Ther. 20, 339–346.
- 15. Sandbothe, M., Buurman, R., Reich, N., Greiwe, L., Vajen, B., Gürlevik, E., Schäffer, V., Eilers, M., Kühnel, F., Vaquero, A., et al. (2017). The microRNA-449 family inhibits TGF-β-mediated liver cancer cell migration by targeting SOX4. J. Hepatol. 66, 1012–1021.
- Wu, H., Tao, J., Li, X., Zhang, T., Zhao, L., Wang, Y., Zhang, L., Xiong, J., Zeng, Z., Zhan, N., et al. (2017). MicroRNA-206 prevents the pathogenesis of hepatocellular

carcinoma by modulating expression of met proto-oncogene and cyclin-dependent kinase 6 in mice. Hepatology 66, 1952–1967.

- 17. Zhang, J., Yang, Y., Yang, T., Yuan, S., Wang, R., Pan, Z., Yang, Y., Huang, G., Gu, F., Jiang, B., et al. (2015). Double-negative feedback loop between microRNA-422a and forkhead box (FOX)G1/Q1/E1 regulates hepatocellular carcinoma tumor growth and metastasis. Hepatology 61, 561–573.
- Lu, Y., Yue, X., Cui, Y., Zhang, J., and Wang, K. (2013). MicroRNA-124 suppresses growth of human hepatocellular carcinoma by targeting STAT3. Biochem. Biophys. Res. Commun. 441, 873–879.
- Yu, A.M., Jian, C., Yu, A.H., and Tu, M.J. RNA therapy: are we using the right molecules? Pharmacol. Ther. Published online December 4, 2018. https://doi.org/10. 1016/j.pharmthera.2018.11.011.
- 20. Chen, Q.X., Wang, W.P., Zeng, S., Urayama, S., and Yu, A.M. (2015). A general approach to high-yield biosynthesis of chimeric RNAs bearing various types of functional small RNAs for broad applications. Nucleic Acids Res. 43, 3857–3869.
- Ho, P.Y., Duan, Z., Batra, N., Jilek, J.L., Tu, M.J., Qiu, J.X., Hu, Z., Wun, T., Lara, P.N., DeVere White, R.W., et al. (2018). Bioengineered noncoding RNAs selectively change cellular miRNome profiles for cancer therapy. J. Pharmacol. Exp. Ther. 365, 494–506.
- 22. Ho, P.Y., and Yu, A.M. (2016). Bioengineering of noncoding RNAs for research agents and therapeutics. Wiley Interdiscip. Rev. RNA 7, 186–197.
- 23. Jian, C., Tu, M.J., Ho, P.Y., Duan, Z., Zhang, Q., Qiu, J.X., DeVere White, R.W., Wun, T., Lara, P.N., Lam, K.S., et al. (2017). Co-targeting of DNA, RNA, and protein molecules provides optimal outcomes for treating osteosarcoma and pulmonary metastasis in spontaneous and experimental metastasis mouse models. Oncotarget 8, 30742–30755.
- 24. Li, P.C., Tu, M.J., Ho, P.Y., Jilek, J.L., Duan, Z., Zhang, Q.Y., Yu, A.X., and Yu, A.M. (2018). Bioengineered NRF2-siRNA is effective to interfere with NRF2 pathways and improve chemosensitivity of human cancer cells. Drug Metab. Dispos. 46, 2–10.
- 25. Wang, W.P., Ho, P.Y., Chen, Q.X., Addepalli, B., Limbach, P.A., Li, M.M., Wu, W.J., Jilek, J.L., Qiu, J.X., Zhang, H.J., et al. (2015). Bioengineering novel chimeric microRNA-34a for prodrug cancer therapy: high-yield expression and purification, and structural and functional characterization. J. Pharmacol. Exp. Ther. 354, 131–141.
- 26. Tu, M.J., Ho, P.Y., Zhang, Q.Y., Jian, C., Qiu, J.X., Kim, E.J., Bold, R.J., Gonzalez, F.J., Bi, H., and Yu, A.M. (2019). Bioengineered miRNA-1291 prodrug therapy in pancreatic cancer cells and patient-derived xenograft mouse models. Cancer Lett. 442, 82–90.
- Lv, H., Zhang, S., Wang, B., Cui, S., and Yan, J. (2006). Toxicity of cationic lipids and cationic polymers in gene delivery. J. Control. Release 114, 100–109.
- Rezaee, M., Oskuee, R.K., Nassirli, H., and Malaekeh-Nikouei, B. (2016). Progress in the development of lipopolyplexes as efficient non-viral gene delivery systems. J. Control. Release 236, 1–14.
- 29. Ewe, A., Panchal, O., Pinnapireddy, S.R., Bakowsky, U., Przybylski, S., Temme, A., and Aigner, A. (2017). Liposome-polyethylenimine complexes (DPPC-PEI lipopolyplexes) for therapeutic siRNA delivery in vivo. Nanomedicine (Lond.) 13, 209–218.
- 30. Ewe, A., Schaper, A., Barnert, S., Schubert, R., Temme, A., Bakowsky, U., and Aigner, A. (2014). Storage stability of optimal liposome-polyethylenimine complexes (lipopolyplexes) for DNA or siRNA delivery. Acta Biomater. 10, 2663–2673.
- Schäfer, J., Höbel, S., Bakowsky, U., and Aigner, A. (2010). Liposome-polyethylenimine complexes for enhanced DNA and siRNA delivery. Biomaterials 31, 6892–6900.
- 32. Zhang, Q.Y., Ho, P.Y., Tu, M.J., Jilek, J.L., Chen, Q.X., Zeng, S., and Yu, A.M. (2018). Lipidation of polyethylenimine-based polyplex increases serum stability of bioengineered RNAi agents and offers more consistent tumoral gene knockdown in vivo. Int. J. Pharm. 547, 537–544.
- Nam, Y., Chen, C., Gregory, R.I., Chou, J.J., and Sliz, P. (2011). Molecular basis for interaction of let-7 microRNAs with Lin28. Cell 147, 1080–1091.
- 34. Liao, T.T., Hsu, W.H., Ho, C.H., Hwang, W.L., Lan, H.Y., Lo, T., Chang, C.C., Tai, S.K., and Yang, M.H. (2016). let-7 modulates chromatin configuration and target gene repression through regulation of the ARID3B complex. Cell Rep. 14, 520–533.
- 35. Shimizu, S., Takehara, T., Hikita, H., Kodama, T., Miyagi, T., Hosui, A., Tatsumi, T., Ishida, H., Noda, T., Nagano, H., et al. (2010). The let-7 family of microRNAs inhibits

Bcl-xL expression and potentiates sorafenib-induced apoptosis in human hepatocellular carcinoma. J. Hepatol. 52, 698–704.

- 36. Sampson, V.B., Rong, N.H., Han, J., Yang, Q., Aris, V., Soteropoulos, P., Petrelli, N.J., Dunn, S.P., and Krueger, L.J. (2007). MicroRNA let-7a down-regulates MYC and reverts MYC-induced growth in Burkitt lymphoma cells. Cancer Res. 67, 9762–9770.
- Lee, Y.S., and Dutta, A. (2007). The tumor suppressor microRNA let-7 represses the HMGA2 oncogene. Genes Dev. 21, 1025–1030.
- Mayr, C., Hemann, M.T., and Bartel, D.P. (2007). Disrupting the pairing between let-7 and Hmga2 enhances oncogenic transformation. Science 315, 1576–1579.
- 39. Gramantieri, L., Ferracin, M., Fornari, F., Veronese, A., Sabbioni, S., Liu, C.G., Calin, G.A., Giovannini, C., Ferrazzi, E., Grazi, G.L., et al. (2007). Cyclin G1 is a target of miR-122a, a microRNA frequently down-regulated in human hepatocellular carcinoma. Cancer Res. 67, 6092–6099.
- 40. Hou, J., Lin, L., Zhou, W., Wang, Z., Ding, G., Dong, Q., Qin, L., Wu, X., Zheng, Y., Yang, Y., et al. (2011). Identification of miRNomes in human liver and hepatocellular carcinoma reveals miR-199a/b-3p as therapeutic target for hepatocellular carcinoma. Cancer Cell 19, 232–243.
- Nakao, K., Miyaaki, H., and Ichikawa, T. (2014). Antitumor function of microRNA-122 against hepatocellular carcinoma. J. Gastroenterol. 49, 589–593.
- 42. Robbins, M., Judge, A., and MacLachlan, I. (2009). siRNA and innate immunity. Oligonucleotides 19, 89–102.
- 43. Yu, H., Wang, Z., Sun, G., and Yu, Y. (2012). Recognition of nucleic acid ligands by toll-like receptors 7/8: importance of chemical modification. Curr. Med. Chem. 19, 1365–1377.
- 44. Li, M.-M., Addepalli, B., Tu, M.-J., Chen, Q.-X., Wang, W.-P., Limbach, P.A., LaSalle, J.M., Zeng, S., Huang, M., and Yu, A.M. (2015). Chimeric microRNA-1291 bio-synthesized efficiently in Escherichia coli is effective to reduce target gene expression in human carcinoma cells and improve chemosensitivity. Drug Metab. Dispos. 43, 1129–1136.
- 45. Guo, Y., Chen, Y., Ito, H., Watanabe, A., Ge, X., Kodama, T., and Aburatani, H. (2006). Identification and characterization of lin-28 homolog B (LIN28B) in human hepatocellular carcinoma. Gene 384, 51–61.
- 46. McDaniel, K., Hall, C., Sato, K., Lairmore, T., Marzioni, M., Glaser, S., Meng, F., and Alpini, G. (2016). Lin28 and let-7: roles and regulation in liver diseases. Am. J. Physiol. Gastrointest. Liver Physiol. 310, G757–G765.
- 47. Yu, J., Vodyanik, M.A., Smuga-Otto, K., Antosiewicz-Bourget, J., Frane, J.L., Tian, S., Nie, J., Jonsdottir, G.A., Ruotti, V., Stewart, R., et al. (2007). Induced pluripotent stem cell lines derived from human somatic cells. Science 318, 1917–1920.
- Reinhart, B.J., Slack, F.J., Basson, M., Pasquinelli, A.E., Bettinger, J.C., Rougvie, A.E., Horvitz, H.R., and Ruvkun, G. (2000). The 21-nucleotide let-7 RNA regulates developmental timing in Caenorhabditis elegans. Nature 403, 901–906.
- Viswanathan, S.R., and Daley, G.Q. (2010). Lin28: a microRNA regulator with a macro role. Cell 140, 445–449.
- Kim, H.J., Kim, A., Miyata, K., and Kataoka, K. (2016). Recent progress in development of siRNA delivery vehicles for cancer therapy. Adv. Drug Deliv. Rev. 104, 61–77.
- Sullenger, B.A., and Nair, S. (2016). From the RNA world to the clinic. Science 352, 1417–1420.
- Xue, H.Y., Guo, P., Wen, W.C., and Wong, H.L. (2015). Lipid-based nanocarriers for RNA delivery. Curr. Pharm. Des. 21, 3140–3147.
- Xia, Y., Tian, J., and Chen, X. (2016). Effect of surface properties on liposomal siRNA delivery. Biomaterials 79, 56–68.
- 54. Zhao, Y., Tu, M.J., Yu, Y.F., Wang, W.P., Chen, Q.X., Qiu, J.X., Yu, A.X., and Yu, A.M. (2015). Combination therapy with bioengineered miR-34a prodrug and doxorubicin synergistically suppresses osteosarcoma growth. Biochem. Pharmacol. 98, 602–613.
- 55. Carr, B.I., Guerra, V., Giannini, E.G., Farinati, F., Ciccarese, F., Ludovico Rapaccini, G., Di Marco, M., Benvegnù, L., Zoli, M., Borzio, F., et al.; Italian Liver Cancer (ITA.LI.CA) Group (2014). Association of abnormal plasma bilirubin with aggressive hepatocellular carcinoma phenotype. Semin. Oncol. 41, 252–258.