Tumor Specific Hypoxia-Activated Expression of Cardiac Peptides Using an Engineered Subtype of *Escherichia Coli* Suppresses Tumor Growth, Angiogenesis and Metastasis in Mouse Model of Breast Cancer

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Research

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Abstract

Background

Application of genetically modified non-pathogenic bacteria expressing specific anti-tumor proteins under certain conditions specific to tumors is an effective approach for selective targeting of tumors. We developed here, for the first time, a novel spatiotemporal cancer targeted therapy applying engineered *E. coli* bacteria with capability of expressing cardiac peptides under hypoxic conditions of tumor.

Method

E. coli BW25133 was transformed with construction of co-expressing cardiac hormones and GFP. Bacteria bearing constructs were then IV administered in mice bearing tumors and then tumor localization, as well as tumor proliferation, invasion and angiogenesis biomarkers (Ki-67, VEGFR, CD31 and MMP9), changes in cytokine profile, suppression of tumor growth and survival were analyzed.

Results

IV Administered bacteria bearing constructs could specifically localize at tumor site and express cardiac peptides under hypoxic conditions. Administration of bacteria significantly enhanced survival rate, suppressed tumor progression and lowered expression levels of MMP-9, VEGFR2, CD31 and Ki67 as potent markers for angiogenesis, tumor proliferation and metastasis. Furthermore, applied bacteria resulted in significant reduction in the expression of IL-1 β , IL-6, GC-SF, IL-12 and TNF- α proinflammatory cytokines, whereas increasing IL-10, IL-17A and INF- γ cytokines.

Conclusion

Overall, administration of *E. coli* bearing cardiac hormone expression construct could effectively suppress tumor growth, angiogenesis, invasion and metastasis while enhancing survival rate in mice model of breast cancer.

Introduction

An excellent approach for spatiotemporal targeting of tumors, regardless of their origin or location, is application of genetically modified non-pathogenic anaerobic bacteria (e.g. *Salmonella*, *E. coli*, *Clostridium*, *Bifidobacterium*) which can be preferentially localized and proliferative in tumor site (Pawelek et al., 2003). Contrary to the conventional cancer therapies, bacteria can also penetrate easily in tumor site and be engineered in a way to respond in conditions specific to the tumors (Forbes, 2010). Despite this, rendering these genera of bacteria in to non-toxic forms for human results in attenuation of their toxicity against cancer cells (Nemunaitis et al., 2003). Therefore, numerous strategies have been applied for further improving their anti-tumoral activities. One such a strategy is to engineer bacteria to secrete specific group of cytolytic proteins such as toxins (e.g. *Staphylococcus aureus* alpha hemolysin) or

enzymes with capability of converting non-toxic prodrugs in to cytotoxic ones (e.g. *E. coli* cytosine deaminase, converting 5-Fluorocytosine in to 5-Fluorouracil) (Jean et al., 2014).

An even more specific cancer targeted therapy could be achieved through the application of specific promoters capable of controlling gene expression in response to specific inducers such as hypoxia. As a common feature of most aggressive tumors, hypoxia is clinically associated with poor clinical outcomes and development of resistance to most chemotherapeutics and radiotherapy (Vaupel and Mayer, 2007). Recently, Nasr and Eidgahi have successfully developed an engineered nirB promoter without any responding region for nitrite and nitrate, making it preferable and selective for expression only under hypoxic tumor condition but not in the presence of chemical inducers (Nasr and Eidgahi, 2014).

Long acting natriuretic peptide (LANP) together with kaliuretic peptide (KP) and vessel dilator (VDL) comprise three of the four cardiac peptides which have shown to suppress the growth of up to 97% of cancer cells in vitro. 80% of human pancreatic adenocarcinomas together with 86% of human small-cell lung carcinomas (SCLC) have shown to be successfully eradicated in athymic mice through the application of these peptides. Furthermore, administration of cardiac peptides significantly reduced the number of human breast adenocarcinoma cells within 24 h of application in vitro. Among these peptides, Vessel dilator appears to hold the strongest anticancer property, reducing up to 97% of human prostate cancer cell throughout the first 24 hours (Nguyen et al., 2012, Vesely et al., 2004, Vesely, 2006).

In presented study, we developed a novel spatiotemporal cancer targeted therapy applying engineered *E. coli* bacteria with capability of expressing cardiac peptides under hypoxic condition of tumor. Creating a hypoxia-inducible system, the cardiac peptide genes were coupled to the synthetic nirB promoter sequence reported by Nasr and Eghbali. The stimulation of synthetic nirB promoter was demonstrated using fluorometry and the expression and function of cardiac peptides were demonstrated by immunoblotting and cell culturing. A murine breast cancer model was used to investigate the potency of hypoxia-inducible cardiac peptide secretion using *E. coli* in suppressing tumor growth.

Material And Method

Ethics statement concerning animal work

All animal experiments were complied with the ARRIVE guidelines and carried out in accordance to the National Institutes of Health guide for the care and use of Laboratory animals (NIH Publications No. 8023, revised 1978). Mice were euthanized by cervical dislocation following completion of tests. the whole study and included procedures were approved by the ethics committee of the Iranian academic center for education, culture and research (ACECR).

Development of plasmid constructs

A polycistronic cassette, expressing KP, VD and LANP genes in tandem under the control of synthetic nirB promoter reported by Nasr and Eidgahi was designed based on the ANP preprohormone sequence

(Genbank accession number: NP_006163). For visualization of bacteria at tumor sites, another polycistronic system expressing green fluorescent protein (GFP) was linked to the previous cassette through a Lambda transcription termination sequence. A ribosomal binding site (RBS) sequence was included upstream of GFP and each of the three gene sequences. For facilitating excretion of expressed cardiac peptides, an immunoglobulin κ chain signal peptide (IgK) leader sequence was embedded between RBS and each of the gene's sequence. A terminating codon was also located at the end of each gene sequence. Lambda transcription termination sequence (a rho independent transcription termination sequence) was allocated between the two polycistronic expressing systems to separate expression of corresponding mRNAs from each other. The DNA sequence of GFP was obtained from the GFP commercial plasmid pLOX-EWgfp and fused to the rest of the construct synthesized by Cinnagen Inc. (Iran) performing SOEing-PCR. At the end, the flanking sequences of pET-32 Ek/LIC vector was added to the endings of the construct performing another round of PCR and the construct was cloned into the vector based on previously described method (Farahmand et al., 2012).

Transformation of the cardiac peptide/GFP expressing plasmid DNA (pET-CP/GFP)

The *rpoS(Am) rph-1 λ^- rrnB3 Δ lacZ4787 hsdR514 Δ (araBAD)567 Δ (rhaBAD)568 rph-1 Escherichia coli* strain, BW25133, was grown in Luria-Bertani (LB) media at 37 °C until mid-log phase, and then harvested at 4 °C. After confirmation of correct cloning, pET-CP/GFP constructs were transformed in to *E. coli* BW25133 strain using calcium chloride method. Bacteria were then maintained and selected in LB media applying 50 mg/ml Ampicillin and 10 mg/ml tetracycline. Confirming function of synthetic nirB promoter, bacteria transformed with pET-CP/GFP were grown overnight on LB medium containing 50 mg/ml Ampicillin and 10 mg/ml tetracycline at 37 °C. The anaerobic condition was induced by bubbling filtered helium gas through the medium for 10 min. plates were then completely sealed in a way that no air could pass to the medium and incubated for 24h at 37 °C. Fluorescent microscope imaging was used to detect GFP proteins expressed by bacteria.

Toxicity of bacterial secreted cardiac protein on breast cancer cell line

MCF-7 breast cancer cell line was used for evaluating the anti-proliferative effects of expressed cardiac peptides. Briefly, 96 well microtiter plates were inoculated with 200

μ L growth media containing a density of 2×10^5 cells and then incubated for 24h under 5% CO₂ in humidified air. Bacteria bearing cardiac peptide construct and bacteria without construct were grow on LB agar media under hypoxic condition for 48h and then, the bacteria growth media were added to the cells. following 48h of exposure, 10 μ L of 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide (MTT) (Sigma Aldrich) with the concentration of 0.05 mg/mL was added to the media. After 4 h incubation at 37°C, whole medium was removed and replaced with 200 μ L DMSO to dissolve water-insoluble formazan salts. The Absorbance was then read at 570 nm using a 96-well plate spectrophotometer.

Annexin V-FITC/7-aminoactinomycin D (7-AAD) flow cytometry

Quantification of apoptosis was carried out by flowcytometry. Briefly MCF-7 cells were treated with 48h growth media of bacteria bearing construct and bacteria without construct as well as PBS for 48 h and then double stained with Annexin V-FITC/ 7-AAD. Fluorescence was then detected with a FACS Calibur flow cytometer (BD Biosciences, USA). Annexin V⁺/7-AAD⁻ cells were determined as early and Annexin V⁺/7-AAD⁺ cells as late apoptotic cells.

***In-vivo* Breast cancer tumor model**

In-vivo Breast cancer tumor model was developed based on a previously established method by Noori et al. (Noori and Hassan, 2011). A female Balb/c mouse bearing spontaneous mouse mammary tumor (SMMT) was obtained from Iranian institute of Pasteur. Previous data indicated that SMMT effectively resembles characteristics of Iranian patient's invasive ductal carcinoma (Langroudi et al., 2010). SMMT was then carefully separated, dissected into smaller pieces with sizes of less than 0.5 cm³ and subsequently, transplanted to 6-8 weak old healthy syngeneic Balb/c mice by surgery. 12 days after transplantation, when tumors reached to the size of 500 mm³, 10⁷ and 10⁸ cfu/g BW25133 bacteria bearing cardiac hormone expressing construct, as well as 10⁸ bacteria without construct and PBS as control were injected systemically to each mouse through the tail vein. A total of three Injections, each with a 4-day interval was performed. To evaluate the effect of treatment on tumor growth delay, tumor volume was measured on predetermined intervals according to the following formula:

$$V(\text{mm}^3) = a^2 \times b \times 0.5$$

In which a and b represents the shorter and longer diameters (mm) of the tumor respectively. The results were then compared to the PBS and bacteria without construct receiving groups.

Cytokine/chemokine profile measurement by multiplex [Enzyme-Linked ImmunoSorbent Assay \(ELISA\)](#) array

Alterations in expression of major pro- and anti-inflammatory cytokines and chemokines in response to treatment with injection of bacteria was evaluated by multiplex ELISA kits (Cat No. MEM-004A, Qiagen). Briefly, peripheral blood of tumor bearing mice receiving PBS and 10⁸ cfu/g bacteria bearing construct were collected in endotoxin-free EDTA containing collection tubes (BD Vacutainer Plus, US) immediately after completion of treatment period. Plasma was then collected by centrifugation of blood samples at 2500 × G for 10 min and kept at -80°C until further use. For analysis, 50 µL of thawed samples was added to each well of their respective row. The ELISA procedure was continued according to the manufacturer's instructions and at the end, absorbance of the 96-well plate was read at 450 nm and 570 nm utilizing a plate reader. For correction of autofluoresence, the 570 nm absorbance was subtracted from final absorbance readings. Expression of each cytokine and chemokine was reported as the fold increase above PBS receiving group levels.

Histological observations and Biodistribution of bacteria in normal and tumor tissues

Evaluation of Bacterial biodistribution in tumor and normal tissues was carried out 48 h after tail vein injection of 10^8 cfu/g BW25133 bacteria in mice. Livers as the highest perfused normal organ and tumors were first excised and fixed in 10% formalin solution. The specimens were then paraffin embedded, sectioned and stained with gram stain and visualized at 100 $\times$ using a light microscope. For counting the number of bacteria in normal and tumor tissues, liver and tumor samples were carefully weighed, homogenized in PBS and plated on LB agar. Bacterial concentration was determined by counting the number of CFUs after 24- to 48 h incubation of plates at 37 °C. Some of the sections obtained from specimens were stained with hematoxylin and eosin (H&E) staining solution to evaluate cytotoxic damages induced by cardiac peptide. Presence of *BW25133 E. coli* in sections was also confirmed by visualization of GFP under fluorescent microscope.

***In vivo* fluorescence reflectance imaging**

Fluorescence reflectance imaging was utilized to initially evaluate whether bacteria could specifically localize in tumor sites and then to monitor time-course of GFP expression as the indicator of successful expression of cardiac peptides in tumor site under hypoxic condition. Briefly, mice were IV injected with either PBS (control group) or *E. coli* BW25133 strain bearing construct with concentration of 10^8 cfu/g and images were taken on days 0, 1, 3 and 6, post injection using the FlouVison fluorescence planar imaging system *in vivo* (Tajhizafarinan Noori Parseh Co., Tehran, Iran). Images from whole body were taken by placing anesthetized mice in the supine position, in the center of the imaging cassette, inside the scanning field of the system. After proper positioning of animals, the imaging cassette was set to the appropriate depth to finely confine anesthetized mice. Finally, the animal body was subjected to the laser beam and emitted non-uniform fluorescent beams were recorded using a highly sensitive thermoelectrically cooled CCD camera located on the same side of the imaged animal.

IHC and histological examination

At the end of the 24-day treatment period, mice receiving PBS and *E. coli* BW25133 strain bearing construct with concentration of 10^8 cfu/g were sacrificed and tumors equivalent in size from both groups were selected for performing immunohistological analyses. Collected tumors were cut in to 7 μ m thick section on a microtome and stained using anti-Ki67 monoclonal antibody (ab15580, Abcam, USA), a marker for tumor proliferation; anti-MMP9 monoclonal antibody (ab38898, Abcam, USA), an enzyme involved in inducing angiogenesis and metastasis; anti-VEGF receptor 2 monoclonal antibody (ab2349, Abcam, USA), a proangiogenic growth factor; anti-CD31/PECAM-1 monoclonal antibody (ab24590, Abcam, USA), recognizing platelet–endothelial cell adhesion molecule-1 (PECAM-1) expressed on surface of endothelial cells; anti-CD8 monoclonal antibody (ab209775, Abcam, USA), an specific marker for cytotoxic T-cells; anti-CD4 monoclonal antibody (ab221775, Abcam, USA), an specific marker for helper T-cells; and their specific horseradish peroxidase-conjugated secondary antibody. Antigen recovery was performed in citrate buffer pH 6.0 and then blockade of endogenous peroxidase, as well as non-specific proteins was performed by 40 min incubation with 3% of H₂O₂ and another 40 min with 3% FBS. At the end, tumor sections were incubated with mentioned antibodies and extent of antigen expression was

evaluated using horseradish peroxidase (HRP)-conjugated streptavidin and 2-Solution DAB kit (Life Technologies) according to the companies' manual instructions. Finally, each antigen's expression level was quantified using ImageJ software (NIH, Bethesda, MD, USA). Results were expressed as the mean of at least 5 tumor sections for each antigen.

H&E staining was also performed for evaluating tumor infiltrating lymphocytes (TILs) according to the previously established protocols. In Brief, initially, percentage of stromal lymphocytes were determined by two separate observers and then stromal TILs were measured as the percentage of immune cells with mononuclear immunological infiltrate characteristics. Findings were classified based on three cut-off points for TIL proportions including 10%, 30% and 50%.

Statistical analysis

Data are represented as mean $\pm$ their standard deviation. Hypothesis testing was carried out applying Student's *t*-test and results considered significant in the case of $P < 0.05$. Survival analysis was done by drawing Kaplan–Meier curves and comparing groups with log-rank test. Hazard ratios were calculated by performing Cox's proportional hazards analysis.

Results

Design of construct and expression of cardiac peptides and GFP

The overall structure of the construct co-expressing the cardiac peptides and GFP under the control of synthetic *nirB* promoter has been represented in figure 1A. Presence of a single band at 550 bp in electrophoresis confirmed correct performance of SOEing PCR (figure 1B). results of colony PCR also confirmed the presence of constructs in recombinant bacteria (figure 1C). following induction of anaerobic condition, collected supernatant of bacteria was concentrated and subjected to SDS-PAGE analysis to evaluate expression of cardiac peptides. As demonstrated in figure 2A, presence of a single band at about 5 kDa confirmed expression of cardiac peptides. As all the expressed cardiac peptides possess a molecular weight very close to each other, three single bands could have been overlapped together and it is not surprising to observe only one band in SDS-PAGE analysis. Extracted proteins from cytoplasm of bacterial pellets in lysis buffer were also subjected to SDS-PAGE analysis for evaluation of expression of GFP. The band at 27 kDa relates to the molecular weight of GFP which is absent in the bacterial samples non-transformed with the cassette (figure 2B).

Anti-proliferative and pro-apoptotic effects of bacterial secreted cardiac protein

Incubation of MCF-7 human mammary carcinoma cells with growth media of bacteria bearing construct under hypoxic condition for a 48-h period, promoted both cell death and apoptosis. The results of MTT assay demonstrated that administration of the growth media of bacteria bearing construct, resulted in inhibition of growth in about 40% of MCF-7 cells which was significantly higher compared to growth media of bacteria without construct and PBS (25% and 5% inhibition respectively). (fig. 3A). The Annexin-

V/7-AAD flow cytometry analysis also demonstrated that treating MCF-7 cells with the growth media of bacteria bearing construct significantly increases rate of apoptosis compare to bacteria without construct and PBS treated group (Fig. 3B).

Preferential colonization of *E. coli* BW25113 within tumor

Systemic administration of *E. coli* BW25113 in syngeneic mice bearing spontaneous mammary tumors resulted in elicit and preferential colonization of bacteria at tumor site. As depicted in figure 4A, bacteria accumulated in tumor site with an approximately 15-fold greater density compare to the liver. Furthermore, microscopic examination of tissue slides also confirmed that *E. coli* was present in all analyzed tumors. Accumulation of *E. coli* BW25113 was more significant at necrotic sites of tumor.

Confirming hypoxia-activated expression of cardiac peptides following tumor specific localization of *E. coli* BW25113 using *In vivo* fluorescence reflectance imaging

E. coli BW25113 bearing cardiac hormone expressing constructs and PBS were injected to the mice and images were collected immediately and 24 h, 3 days and 6 days post injection. As depicted in figure 4B, no significant differences in emitted fluorescent intensity was observed between two group during the first 24 h post injection. On day 3 however, emitted fluorescent from bacteria receiving group slightly grew and become a bit higher than those observed in PBS receiving group. Continuing similar trend, on day 6 post injection, emitted fluorescent from bacteria receiving group reached to its maximum and was completely restricted to the tumor site. Regarding other organelles, no significant changes in fluorescent intensity was observed during image acquisition period neither in control nor in bacteria receiving groups. These observations further confirm site specific colonization of bacteria and expression of GFP as an indicator of successful expression of construct in hypoxic condition.

Tumor growth retardation potential of bacteria expressing cardiac protein

The delay in tumor growth was evaluated over a 24-day period after the administration of bacteria bearing construct with concentrations equal to 10^7 and 10^8 cfu/g, bacteria without construct in concentration equal to 10^8 cfu/g and PBS as the control group. Based on results, administration of bacteria without construct didn't result in any significant tumor growth retardation compared to PBS receiving group. Therefore, bacteria per se were not capable of suppressing tumor growth speed. Contrarily, administration of bacteria bearing constructs, even at concentration equal to 10^7 cfu/g resulted in a significantly greater delay in tumor growth compare to PBS group. Furthermore, increasing the concentration of bacteria from 10^7 cfu/g to 10^8 cfu/g resulted in more enhanced tumor suppression. As depicted in figure 5A, differences in delaying tumor growth between two concentrations of bacteria bearing construct become significant on day 15 from beginning of treatment. Furthermore, the differences between delaying in tumor growth in group receiving concentrations of bacteria equal to 10^8 cfu/g and PBS group become significant on day 12. However, similar significance for the group receiving bacteria bearing construct at 10^7 cfu/g took place on day 15. Based on these observations we concluded

that first, the delay in tumor growth observed with administration of bacteria bearing construct is mainly due to the secretion of cardiac proteins. Second, increasing concentration of administered bacteria may result in enhancement of production and secretion of cardiac peptides. Administration of bacteria bearing constructs at both concentrations also significantly enhanced survival rate as observed in Kaplan-Meier analysis (figure 5B).

Alteration in pro-inflammatory cytokine profile

We also examined the effect of bacterial treatment on expression of pro-inflammatory cytokines. As depicted in figure 6, administration of bacteria bearing construct at concentrations equal to 10^8 cfu/g to the mice bearing spontaneous breast tumors, resulted in a significant decline in expression of pro-inflammatory cytokines, including IL-1 α , IL-1 β , IL-6, IL-12, TNF- α , G-CSF and GM-CSF while increasing expression of IL-17A, IL-10 and INF- γ compared to the PBS receiving group. No significant changes were observed in expression profile of IL-2 pro-inflammatory cytokine. Therefore, treatment with bacteria bearing cardiac hormone expressing construct could increase expression of some of the most important cytokines of both T_H1 and T_H2 cells (INF- γ and IL-10 respectively).

Immunohistochemistry, quantification of micro-vessel density and H&E staining results

As illustrated in Figure 7, MMP9 expression was significantly decreased in mice receiving bacteria bearing construct at concentrations equal to 10^8 cfu/g compared to PBS receiving group. Furthermore, expression of VEGF receptor 2 (VEGFR2) and Ki-67 were also significantly decreased in response to therapy. Also, comparing state of CD31 expression as a marker of micro-vessels density demonstrated a significant decline in mice receiving bacteria bearing construct at concentrations equal to 10^8 cfu/g compare to control group. Finally, the number of TILs and CD8+ TILs were significantly increased in mice treated with bacteria bearing construct at concentration equal to 10^8 cfu/g.

Discussion

Based on observed results, it is elicited that *E. coli* BW25113 is capable of expressing and secreting cardiac peptides including LANP, VD and KP as well as GFP as an indicator protein under the control of hypoxic or anaerobic condition. Secreted cardiac peptides by interacting with their receptors on carcinoma cells result in induction of apoptosis and cell death. The main advantage of presented approach is selectivity in both targeting tumor site and inducing cytotoxicity only in hypoxic areas of tumor. As demonstrated in sections collected from tumor tissue and liver, bacteria preferentially localized in tumor site with a significantly greater density. Furthermore, activation of protein expression only at hypoxic condition, prohibits expression of cardiac proteins and observation of undesired side effects at normally perfused body organs including liver, lung, heart, etc. Other studies, including the one performed by Stritzker et al., have also demonstrated that different strains of *E. coli* are capable of preferentially colonizing in tumor site (Stritzker et al., 2007).

Based on the study performed by Kim et al, contrary to the *salmonella typhimurium* which is frequently applied in bacterial tumor targeting, *E. coli* strain K-12 (MG1655) is not capable of suppressing tumor growth per se (Kim et al., 2015). Our results also clearly demonstrated that administration of *E. coli* BW25113 even at high doses didn't significantly suppressed tumor growth. As the only difference between bacteria bearing construct and untransformed ones is the capability of expressing cardiac proteins, differences in tumor growth delay may be attributed to the expressed cardiac peptides under hypoxic condition of tumor.

Several in vitro studies have clearly indicated the cytotoxic effects of cardiac peptides against malignant cells. Based on studies performed by Vesely et al, four cardiac peptides including LANP, KP, VD and ANP demonstrate significant anti-proliferative effects against breast, prostate, lung, brain and renal carcinomas (Vesely, 2013). Reported by Sun et al., VD and KP demonstrate their anti-cancer activities in prostate cancer through inhibiting RAS activity (Sun et al., 2010). Furthermore, LANP is capable of inhibiting 80–90% of MEK1/2 and ERK1/2 activity in prostate cancer cells (Sun et al., 2006, Sun et al., 2007). As ERK plays an important role in controlling apoptosis, inhibition of its phosphorylation and activation by LANP may partly explain its pro-apoptotic mechanism of action (Vesely et al., 2005).

Currently, multiple studies have addressed the role of cytokines in cancer proliferation, invasion and metastasis. For instance, it has been shown that high circulating levels of IL-1 β and IL-6 are unfavorable prognostic indicators and directly interrelate with higher rate of recurrence and clinical stage in patients with breast cancer (Kovacs, 2001, Mettler et al., 2004, Kozłowski et al., 2003). Contrarily, overexpression of INF- γ has shown to promote cell death through up-regulating caspases and demonstrate significant antiangiogenic activity (Connett et al., 2005, Bouker et al., 2005, Sidky and Borden, 1987). Administration of bacteria bearing construct resulted in a complex effect on expression of pro-inflammatory cytokines. Many studies have shown that cardiac peptides demonstrate an anti-inflammatory behavior. For instance, it has been shown that ANP could down regulate expression of TNF- α which has an important role in alloreactivity of T-cells (Kierner et al., 2000, Ladetzki-Baehs et al., 2007). Furthermore, ANP can also enhance expression of IL-10 which in turn, may result in downregulation of IL-2 and proliferation of T-cells (Leitman et al., 1988, Moro et al., 2007). IL-6 concentration has also been shown to decline in response to administration of ANP (Moro et al., 2007). Finally, administration of ANP together with LPS decreases production of IL-12 by dendritic cells and generates fewer INF- γ positive T cells and much more IL-4 positive ones (Mohapatra et al., 2004). Consistent with these reports, here we also observed a significant decline in peripheral blood concentrations of pro-inflammatory cytokines including IL-1a, IL-1b, IL-6, IL-12 and TNF-a. contrarily the concentrations of IL-10 and INF- γ were significantly increased in response to therapy.

Angiogenesis is a firmly controlled process, crucial for tumor expansion beyond 2 to 3 mm in diameter and a critical step of tumor invasion and metastasis (Carmeliet and Jain, 2000). Different studies have now suggested a critical role for MMP9 in formation of a proper microenvironment for promotion of tumor growth and angiogenesis through enhancing the association between VEGF and VEGFR2 (Rodríguez-Manzanque et al., 2001, Mira et al., 2004). This mainly takes place through the proteolytic

cleavage and subsequent release of extracellular matrix (ECM) bound VEGF by MMP9 activity which is an important step of angiogenic switch. Based on Bergers et al. inhibitors of MMP9 effectively suppressed tumor growth and reduced angiogenic switching (Bergers et al., 2000). It has also been shown that MMP-cleaved VEGF plays an important role in tumors vascular patterning (Lee et al., 2005, Nguyen et al., 2012). Based on previous reports, cardiac peptides effectively reduce expression of VEGF in vitro (Nguyen et al., 2012). In present study, we demonstrated that administration of bacteria expressing cardiac peptides under hypoxic condition could reduce MMP-9 expression in vivo. In parallel, VEGFR2-triggered angiogenesis is considered as a hallmark of cancer progression and metastasis (Olsson et al., 2006, Herbert and Stainier, 2011). Consistent with previous studies, we also found that expression of VEGFR2 was significantly reduced in tumors from mice receiving bacteria bearing cardiac peptides expressing construct (Nguyen et al., 2012).

Results from different reports show that high tumor micro-vessel density measured by immunohistochemical analysis of CD31 expression, is correlated with higher rates of lymph node metastasis and poor prognosis (Weidner et al., 1991, Hansen et al., 2000, Jacquemier et al., 1998). In addition, MMP-9 has widely been mentioned as an important driver of cancer's malignant progression, invasion and metastasis (Huang, 2018). Also, determination of proportion of cancer cells undergoing proliferation by immunohistochemical analysis of Ki-67, has shown that high Ki-67 expression is associated with increased risk of metastasis development as well as poor overall survival (Green et al., 2016). As administration of bacteria expressing cardiac peptides could effectively reduce expression of Ki-67, CD31 and MMP-9, it may be an effective therapeutic for preventing from invasion and future development of metastasis. Finally, presence of TILs (especially CD8 + TILs), which demonstrate effective anti-tumoral activity in tumor sites, has shown to be together with a favorable prognosis value in different neoplasms (Mahmoud et al., 2011, Wang et al., 2016, Laghi et al., 2009, Galon et al., 2006). Therefore, significant increase in survival rate as observed with Kaplan-Meyer analysis may be partly attributed to the higher number of CD8 + TILs in tumor site, as well as lower expression values of above-mentioned markers.

Overall, in present study, we demonstrated that IV administration of *E. coli* BW25113 strain bearing cardiac hormone expression construct could result in tumor specific localization and subsequent hypoxia derived expression of cardiac peptides. This resulted in suppression of tumor growth, significant increase in survival rate and decreased expression of MMP-9, VEGFR2, Ki-67 and CD31, all of which propose an antiangiogenic and anti-metastatic activity for the proposed therapy.

Conclusion

Overall, administration of *E. coli* bearing cardiac hormone expression construct could effectively suppress tumor growth, angiogenesis, invasion and metastasis while enhancing survival rate in mice model of breast cancer.

Abbreviations

TNBC: triple negative breast cancer

LANP: Long acting natriuretic peptide

KP: kaliuretic peptide

VD: vessel dilator

SCLC: human small-cell lung carcinomas

E. coli: Escherichia coli

GFP: green fluorescent protein

RBS: ribosomal binding site

pET-CP/GFP: cardiac peptide/GFP expressing plasmid DNA

LB: Luria-Bertani

MTT: 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide

FITC: Fluorescein isothiocyanate

7-AAD: 7-aminoactinomycin D

SMMT; spontaneous mouse mammary tumor

ELISA: Enzyme-Linked ImmunoSorbent Assay

CFU: colony forming units

H&E: hematoxylin and eosin

IHC: Immunohistochemistry

TILs: tumor infiltrating lymphocytes

SDS-PAGE: sodium dodecyl sulfate polyacrylamide gel electrophoresis

IL: Interleukin

TNF- α : Tumor necrosis factor α

INF- γ : Interferon γ

MMP9: matrix metalloproteinase 9

VEGFR2: vascular endothelial growth factor receptor 2

ECM: extracellular matrix

CD31: cluster of differentiation 31

Declarations

Disclosure statement

The authors declare that there exist no conflict of interests.

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Description of the contribution

Keivan Majidzadeh: conceived and designed research. **Leila Farahmand;** conducted new reagents or analytical tools. **MAlihe S, Mitra S, Neda J, Zeinab N, Marjan M, Ahad M, Azadeh SH and SHima Moradi:** contributed conducted experiments. **Behrad darvishi:** analyzed data, wrote the manuscript. All authors read and approved the manuscript.

Ethical approval

"All applicable international, national, and/ or institutional guidelines for the care and use of animals were followed."

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Figures

Figure 1

(A) The overall structure of the construct co-expressing the cardiac peptides and GFP under the control of synthetic nirB promoter. (B) results of GFP and Cardiac peptide PCR. Lane 1: the DNA size marker, Lane 2: result of GFP PCR, Lane 3: result of cardiac peptides PCR, Lane 4: Negative control (C) Result of SOEing PCR. Presence of a single band at 550 bp in electrophoresis confirmed correct performance of SOEing PCR. Lane 1: the DNA size marker, Lane 2: result of SOEing PCR Lane 3: Negative control (D) Results of colony PCR, confirming the presence of constructs in recombinant bacteria. Lane 1: The negative control (prepared PCR mixture applying water instead of DNA template), Lane 2: the DNA size marker, Lane 3: the positive colonies comprising the cardiac hormone expressing construct.

Figure 2

Results of SDS-PAGE analysis. (A) Presence of a single band at about 5 kDa confirmed expression of cardiac peptides. Lane 1: protein marker, Lane 2: before induction of hypoxic condition, Lane 4: 24 h after induction of hypoxic condition. (B) The single band at 27 kDa relates to the molecular weight of GFP which is absent in the bacterial samples non-transformed with the cassette. Lane 1: before induction of hypoxic condition, Lane 2: protein marker, Lane 3: 24 h after induction of hypoxic condition.

(A)

(B)

Figure 3

(A) In vitro cytotoxicity of purified cardiac peptides from 48h growing media of bacteria bearing construct under hypoxic condition. (B) The results of Annexin-V/7-AAD flow cytometry analysis. Treating MCF-7 cells with cardiac peptides mixture significantly increased rate of apoptosis compare to PBS receiving group (Left: PBS, Middle: media of bacteria without construct, Right: cardiac peptides purified from media of bacteria bearing construct).

Figure 4

(A) Bacterial concentration in tumor tissue and liver of mice 48 h after IV administration of 108 cfu/g bacteria. Accumulation in tumor site was about 15-fold greater compare to the liver. (B) Results of In vivo fluorescence reflectance imaging 0, 1, 3 and 6 days post injection. on day 6 post injection, Maximum New fluorescent emissions were completely restricted to the tumor site and reached to its maximum on day 6 post injection.

Figure 5

(A) suppression of tumor growth by administration of different concentrations of bacteria bearing construct. Increase in concentration of administered bacteria significantly enhanced tumor suppression potency. Ns: not significant, ****: $P < 0.0001$ compared to PBS, ####: $P < 0.0001$ compared to bacteria bearing construct at concentrations equal to 10^7 cfu/g (B) Results of Kaplan-Meier analysis demonstrating a higher rate of survival in mice receiving bacteria bearing construct compared to PBS and bacteria without construct.

Figure 6

Alteration in expression of pro-inflammatory cytokines in response to bacterial therapy. Results are expressed as fold changes compared to control group receiving PBS. *: $P < 0.05$, **: $P < 0.01$, ***: $P < 0.001$ compared to control group.

Figure 7

Results of immunohistochemical analysis, quantification of micro-vessel density and H&E staining.