

# Engineering of Human Lactoferrin for Improved Anticancer Activity

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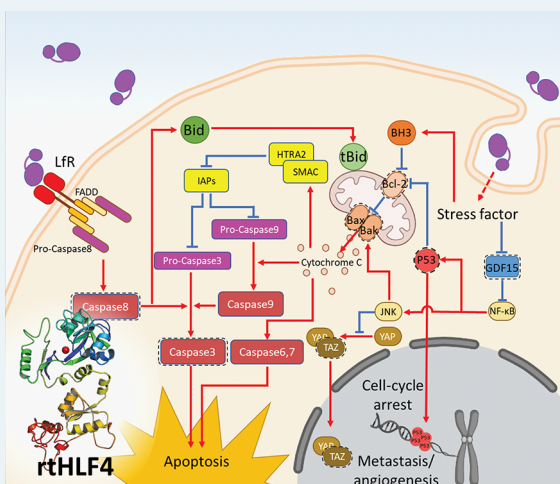
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**ABSTRACT:** Protease-digested lactoferrin fragments often exhibit improved therapeutic properties. However, there are limited studies investigating the anticancer properties of these fragments. The fragment with improved anticancer activities is an attractive alternative to chemotherapeutic drugs—presenting severe side effects. Herein, we report the isolation and characterization of recombinant engineered-lactoferrin (rtHLF4), exhibiting up to 100-fold improved anticancer activity compared to the full-length lactoferrin (flHLF). Further, rtHLF4 exerts its anticancer effect in a shorter duration. Through transcriptomic analysis of various cancer biomarkers, rtHLF4 was found to upregulate various pro-apoptotic markers and downregulate signaling proteins involved in angiogenesis and metastasis. We further determined that rtHLF4 showed no hemolytic activity at high concentrations. We believe that this anticancer protein can be further developed as a cancer treatment.



**KEYWORDS:** Lactoferrin, Anticancer, Tryptic-digestion

Cancer kills approximately 10 million patients each year, accounting for over 15% of deaths worldwide.<sup>1</sup> Thus, there is an urgency to search for new anticancer treatments. Current anticancer treatments often have undesirable side effects, whereas natural anticancer remedies have shown to be less efficacious, requiring a high local concentration to elicit the anticancer properties. The lactoferrin protein exerts multiple therapeutic properties, including anticancer, antimicrobial, antiparasitic, antiviral, and antifungal activity.<sup>2</sup> It was recently found that lactoferrin can hasten the recovery of patients infected with COVID-19.<sup>3,4</sup> This transferrin-class protein exhibits these therapeutic activities by recruiting immune-related cytokines and depleting nutrients in the diseased area through metal ion sequestration.<sup>5</sup>

Protease-digested lactoferrin often results in fragments that exhibit improved therapeutic activities. Lactoferricin (peptic-digested lactoferrin) has improved antimicrobial and antifungal properties compared to the full-length lactoferrin (flHLF).<sup>6</sup> These therapeutic activities are mainly attributed to the lactoferricin's amphipathic and cationic properties that are similar to bacteriocins and pyocins.<sup>6</sup> Other products such as lactoferrampin showed increased killing efficacy against certain fungal strains.<sup>7</sup> However, there are limited studies that suggest that protease-digested lactoferrin fragments show improved anticancer activities.

This study adopts a rational approach to investigate protease-digested lactoferrin fragments with improved anticancer properties. Previous studies found that lactoferrin is

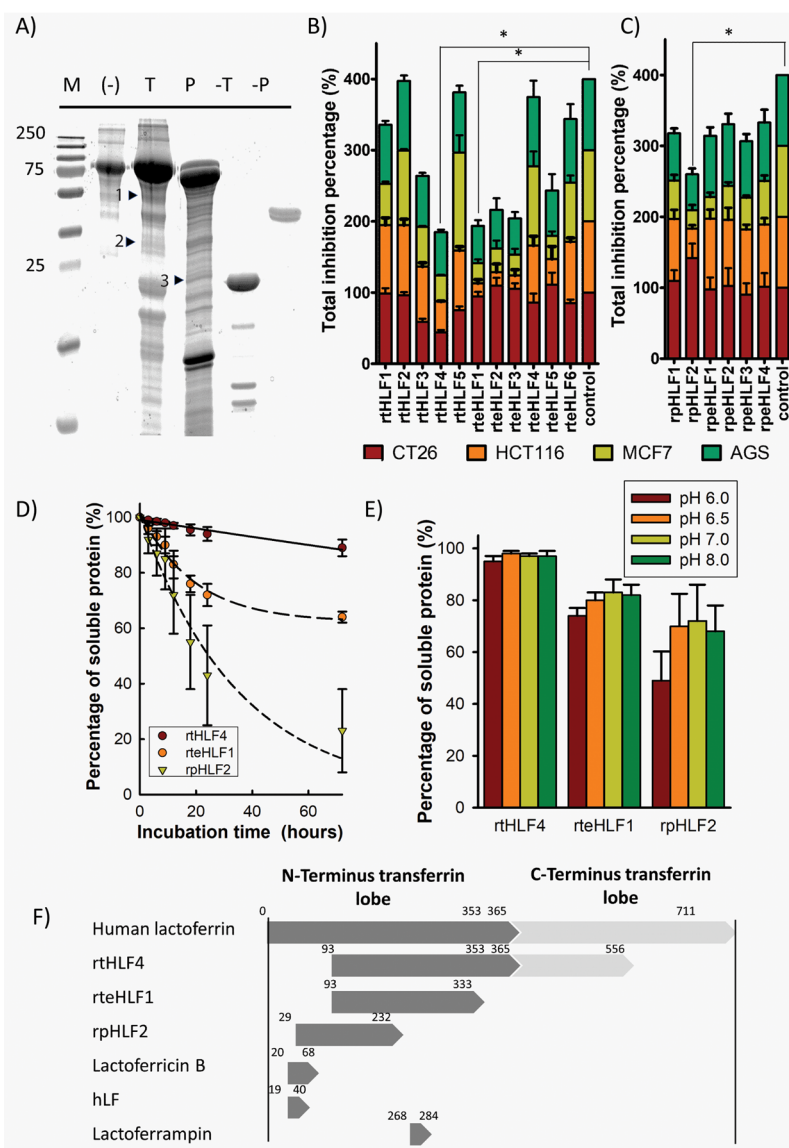
highly resistant to proteolytic degradation, resulting in incomplete digested fragments that exhibit different bioactivities.<sup>8</sup> Protease-digested recombinantly expressed human lactoferrin, sorted by fragment size (Figure 1A), were screened for anticancer activity against gastric (AGS), breast (MCF7), and colorectal cancer (HCT116 and CT26) cell lines (Figure 1B,C). Three promising candidates (rtHLF4, rteHLF1, and rpHLF1) were found to have improved inhibition against the four cancer cell lines, inhibiting over 50% of the cultured cancer cells. Out of the three fragments, only fragment rtHLF4 showed improved stability under human physiological temperature (Figure 1D) and pH (Figure 1E). Protein identification using MALDI-TOF of these fragments revealed that all three peptides are truncated at the N- and C-terminus (Figure 1F).

We focused our studies on rtHLF4 due to its stability exhibited under human-physiological conditions and its comparable anticancer properties to rteHLF1 and rpHLF2 (Figure 1B,C). We hypothesize that the rtHLF4 truncated C-terminus lobe functions to stabilize the lactoferrin protein, where rteHLF1 and rpHLF2 lacking the C-terminus domain

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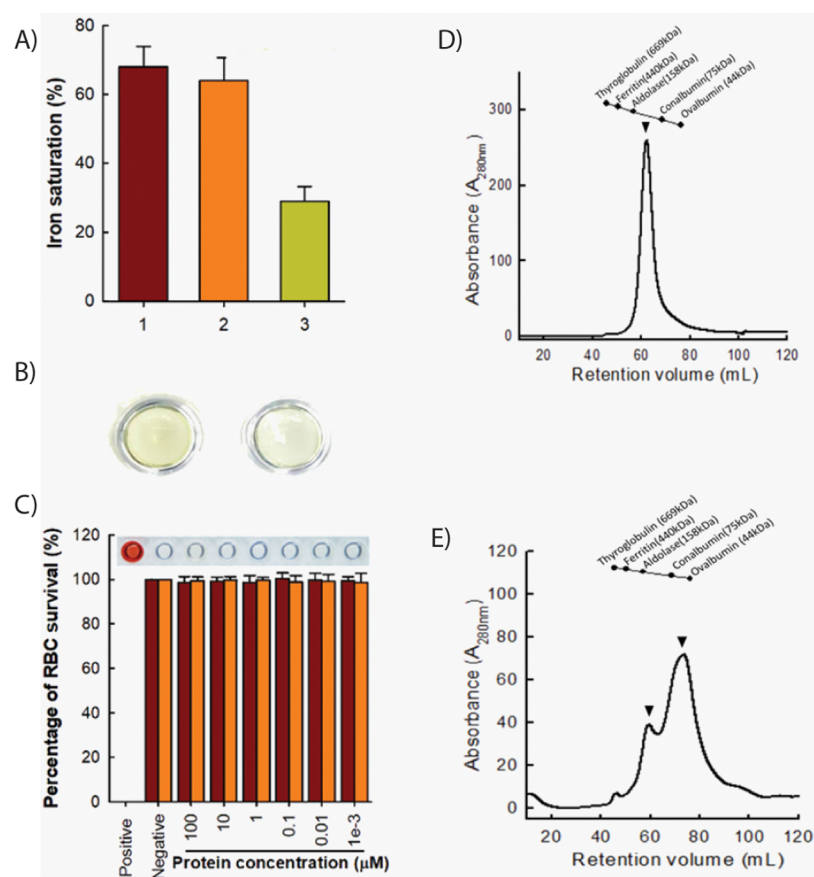
**Figure 1.** Isolation of anticancer lactoferrin fragments. (A) SDS-PAGE of trypsin (T) and pepsin (P) digested lactoferrin. Fragments with positive anticancer activity are indicated with arrows (M, marker; T, Trypsin; P, Pepsin; 1, rtHFLF4; 2, rteHFLF1; 3, rpHFLF2). (B) Anticancer activity of trypsin-digested lactoferrin fragments. (C) Anticancer activity of pepsin-digested lactoferrin fragments. (D) Isolated lactoferrin fragment stability at 37 °C. (E) Purified lactoferrin fragment stability under different pH after 12 h. (F) The sequence length of isolated fragments compared to lactoferrin and other lactoferrin-derived fragments (\* $P \leq 0.0083$  after Bonferroni correction; each group performed in triplicate).

are less stable under different pH and temperature changes. Recombinant expression of rtHFLF4 showed that the level of ferric occupancy in the binding pockets is 3-fold lower than that of flHLF (Figure 2A), leading to the color of rtHFLF4 being much lighter than that of flHLF under the same concentration (Figure 2B). The flHLF comprises two globular lobes, each with one iron-binding site.<sup>9</sup> The 3-fold lower ferric ion occupancy in rtHFLF4 suggests that the truncated N-terminus and disrupted C-terminus lobe cannot retain the ferric-binding activity due to the loss of its conserved ferric-binding residues. The modeled rtHFLF4 structure showed perturbation of its C-terminus lobe affecting the ferric binding pocket (Figure S1b)<sup>10</sup> as compared to the flHLF structure (Figure S1c). Additionally, rtHFLF4 can exist in both the monomeric and dimeric conformation (Figure 2E), whereas flHLF has the preference of being in the dimeric state (Figure 2D). These findings indicate that the truncated C-terminus

lobe helps in protein stability, but incurs the loss of ferric-binding and interferes with the dimerization interface (Figure S1a).

We conducted a hemolytic toxicity assay to evaluate the safety of using recombinant rtHFLF4 on healthy cells. The hemolytic toxicity assay using blood acquired from 10 donors (aged between 25–40 years old) found no hemolytic activities in the erythrocytes treated with 100  $\mu$ M of recombinantly expressed flHLF or rtHFLF4 (Figure 2C). These results suggest that the observed anticancer activities result from direct rtHFLF4 interactions with the cancer cells and are not the result of cell toxicity.

We compared the anticancer activities of both rtHFLF4 and flHLF against human cancer cell lines including breast (MCF7), stomach (HGC27 and AGS), and colorectal (Caco2, LoVo, and HCT116) cancer cell lines. We evaluated the anticancer activity by observing the treatment time (3, 24,



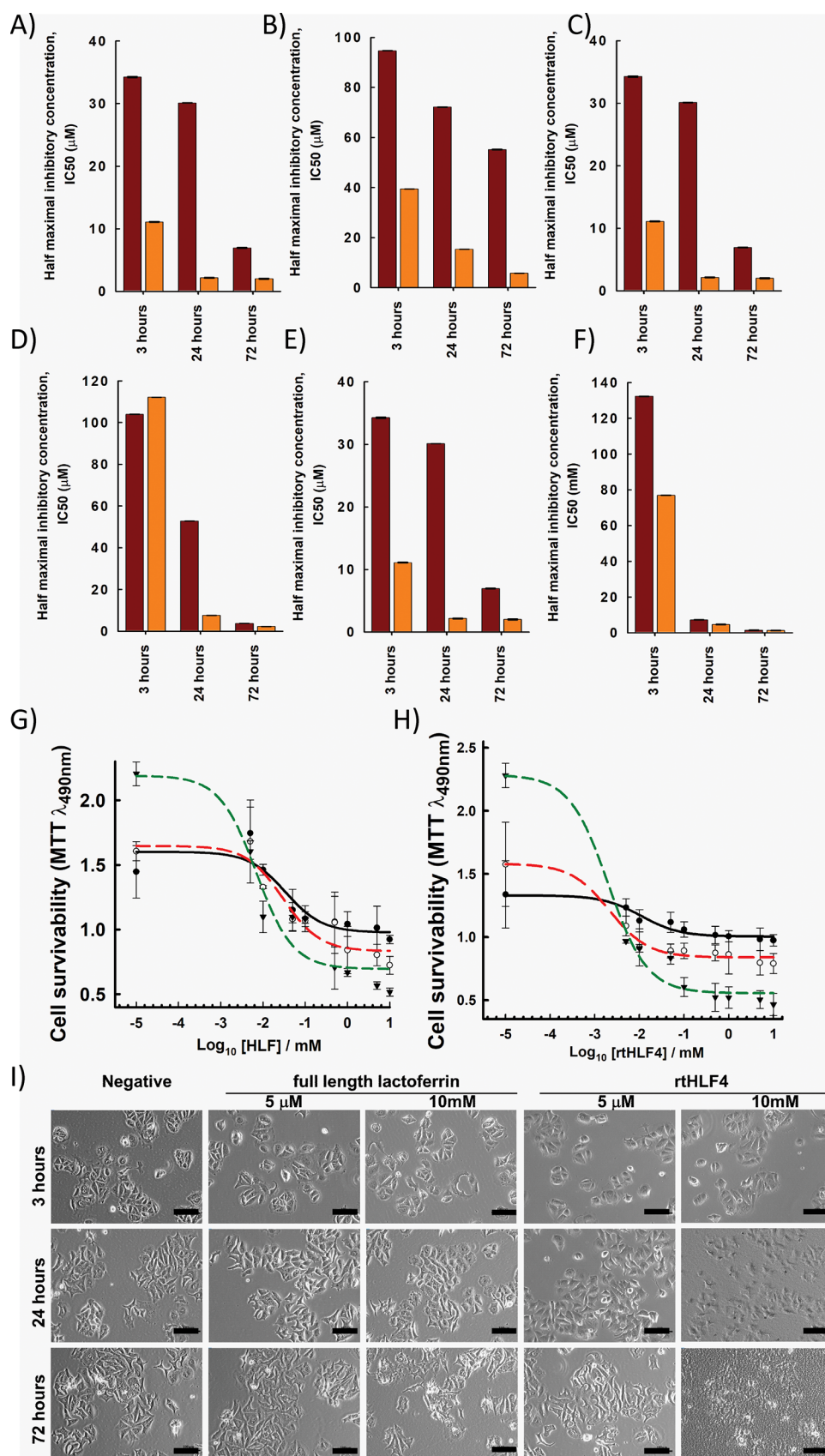
**Figure 2.** Structural and biosafety analysis of rHHLF4. (A) Percentage of ferric ion saturation of lactoferrin and rHHLF4 (1, commercial lactoferrin; 2, fHHLF; 3, rtHHLF4; each group performed in triplicate). (B) Color of recombinant lactoferrin (left) and rHHLF4 (right; samples prepared at 0.1 mM). (C) Hemolytic assay of purified lactoferrin (maroon) and rHHLF4 (orange) on human red blood cells (inset: hemolytic results using the corresponding rHHLF4 concentrations;  $n = 10$ , positive control, 0.1 mg/mL indomethacin). (D) Size exclusion chromatographic separation of fHHLF (protein weight, 80 kDa) (arrows indicate dimeric conformation of the isolated protein). (E) Size exclusion separation of rHHLF4 (protein weight, 51 kDa; arrows indicate dimeric and monomeric conformation of the isolated protein).

and 72 h) needed to elicit complete inhibition of the cancer cells. rHHLF4 is more potent against cancer cells by 1 to 2 orders of magnitude when compared to fHHLF. Additionally, rHHLF4 elicits complete cancer cell inhibition within 24 h of treatment (Figure 3A–F) compared to the fHHLF that needs a longer incubation time. The dose-dependent-cell viability assay of rHHLF4 against MCF-7 (Figure 3G,H) and other cancer cells (Figures S2–S6) revealed improved efficacy and sensitivity against these cancer cell lines. Similarly, the cell morphology of MCF7 (Figure 3I) and other cancer cells (Figures S2c–S6c) showed an indication of observable cell death annotated by detachment and cellular apoptosis within 24 h of treatment with rHHLF4, while fHHLF requires a higher concentration and longer incubation time to elicit a similar level of cellular death. Studies have shown that fHHLF's anticancer activity results from the ferric ion sequestration, causing gradual killing activity arising from nutrient starvation.<sup>8</sup> As rHHLF4 has lesser ferric ion binding ability, we hypothesize that the mechanism of action attributing to rHHLF4's improved anticancer properties differs from that of fHHLF.

In order to understand the mechanism of action, we investigated the gene expression of anticancer-associated biomarkers in the cancer cell cultures treated with rHHLF4 and fHHLF. The gene expression of pro-apoptotic, angiogenesis, and metastatic proteins was monitored using quantitative PCR and Western blot (Figure 4A). In comparison to fHHLF,

millimolar and micromolar rHHLF4 concentration administered to the cancer cells revealed changes in protein expression level involved in the death receptor pathway (caspase3 and caspase8), mitochondrial outer membrane permeabilization (MOMP; Bak/Bax, and Bcl2), and P53-related pathway (P53, GDF15, and TAZ).

rHHLF-treated HGC27, HCT116, and MCF7 showed elevated caspase3 and caspase8 expression, suggesting that rHHLF4 triggers activation of caspase8, leading to the proteolytic cleavage of caspase3 (Figures 4a, S7, S8, S12). Studies have shown that lactoferrin-binding receptors with Fas-associated death domain (FADD) trigger the activation of caspase8, triggering the death receptor pathway (Figure 4b).<sup>11,12</sup> Caspase8 then activates caspase3 that subsequently cleaves critical cellular protein, leading to cancer cell apoptosis. These results are consistent with previous studies that indicated that the presence of FADD-associated lactoferrin-binding receptors such as CD91, TLR2, TLR4, and intelectin-1 activates the death receptor pathway upon lactoferrin binding.<sup>13</sup> It was found that these receptors are present in various cancer cell lines including colorectal epithelial cancer, breast cancer, and gastric cancer.<sup>14</sup> We hypothesize that the improved upregulation of the caspase8 can be attributed to the predominant monomeric state of rHHLF, allowing better receptor–ligand interactions.



**Figure 3.** Anticancer activity of lactoferrin and rHLF4. (A) The half maximal inhibitory concentration, IC<sub>50</sub>, of fHLF (maroon) and rtHLF4 (orange) tested against breast cancer (A) MCF7; gastric cancer (B) HGC27 and (C) AGS; and colorectal cancer (D) Caco2, (E) LoVo, and (F) HCT116. Dose–response of MCF7 against varying concentrations of (G) fHLF and (H) rHLF4. (I) Cell morphology of MCF7 cell lines treated with varying concentrations of fHLF and rHLF4 (bar = 100 μm).



the upregulation of P53 and JNK proteins that repress Bcl-2 and upregulate Bak/Bax activities, respectively.<sup>16,17</sup> Evidence shows that the cytochrome C released from the mitochondria following MOMP triggers the upregulation of HTRA2/SMAC activation that represses Survivin, a major inhibitor of apoptosis protein (IAP).<sup>18</sup> IAP blocks the effect of caspases 3, 7, and 9, preventing cellular apoptosis. Previous studies revealed that bovine lactoferrin significantly decreases IAP expression in colorectal cancer cell lines while observing the upregulation of IAP inhibitors SMAC and HTRA2.<sup>18</sup> Our results reveal similar traits of caspase3 activation in both fHHLF and rtHHLF4, where the latter showed an approximately 2-fold increase of caspase3 in four cancer cell lines (MCF7, HGC27, AGS, and LoVo). The rtHHLF4-treated LoVo cells showed upregulated caspase3 and Bax/Bak proteins, while showing downregulation of GDF15, suggesting that the repression of GDF15 relieves the NF- $\kappa$ B activating JNK, leading to Bax/Bak activation. Through the Bak/Bax activation, the caspase3 proteolytic cleavage is triggered by cytochrome C–caspase9 activation.<sup>19</sup>

In addition, in the experiment of rtHHLF4's anticancer activities, we found that rtHHLF4 slows cancer cell growth better than fHHLF (Figure 3A–F) by 1 to 2 orders of magnitude. We hypothesize that rtHHLF4 additionally inhibits the PDZ-binding motif (TAZ) and growth differentiation factor 15 (GDF15). rtHHLF4 showed better inhibition of TAZ in HGC27, AGS, and LoVo cells and improved inhibition of GDF15 in AGS and LoVo (Figures 4a, S8, S9, S11). TAZ conventionally regulates mesenchymal stem cell differentiation, but it is also a known coactivator yes-associated protein (YAP) of the Hippo-pathway genes that upregulate multiple tumorigenic genes including cell migration, invasion, and anchorage-independent growth.<sup>20,21</sup> The TAZ/YAP activation plays an essential role in cancer initiation and solid tumor formation through the activation of Survivin expression (Figure 4b).<sup>22</sup> Thus, rtHHLF4's suppression of TAZ prevents both the cancer metastatic potential and angiogenesis. GDF15, a protein from the transforming growth factor beta (TGF- $\beta$ ) superfamily, is upregulated and secreted by various metastatic cancers including glioblastoma, ovarian, prostate, breast, and colorectal cancers. While the corresponding role in cancer metastasis is not fully understood, upregulated GDF15 is associated with poor patient survival after cancer remission.<sup>23</sup> The ability of rtHHLF4 to inhibit GDF15 indicates the potential use of the lactoferrin fragment for the prevention of cancer metastasis and upregulation of P53 mediated cell apoptosis (Figure 4b).

In summary, we identified three protease-digested human lactoferrin fragments that present improved anticancer properties as compared to fHHLF. Among the three, rtHHLF4 showed the best stability and tolerance to human physiological conditions. Upon closer inspection on rtHHLF4's mode of mechanism, we found that rtHHLF4 elicits a wide range of anticancer effects on various cancer cell types via lactoferrin receptor interactions, internalization of rtHHLF4, and inhibition of metastatic genes, thus indicating the potential use of this protein in future cancer therapy. Further investigation would be needed to better understand the anticancer mechanism prior to development of rtHHLF4 as a viable treatment. These include a closer investigation of receptor proteins and binding proteins involved in the internalization of rtHHLF4 to the cytoplasm. We intend to further investigate such interactions using thermal proximity coaggregation to profile the protein

complex dynamics within these cancer cells.<sup>24</sup> The rtHHLF4 potentially can also function as an output mechanism for synthetic biology-driven cellular engineering for anticancer therapy.<sup>25,26</sup>

## ■ ASSOCIATED CONTENT

### Supporting Information

The Supporting Information is available free of charge at <https://pubs.acs.org/doi/10.1021/acspsci.1c00134>.

Experimental details and supporting figures and tables; details of the materials and methods involved in the experiments, including the list of RT-PCR primer sequences; the sequence and structure of fHHLF and rtHHLF4, including all anticancer activities of the protein against the different cancer cells (PDF)

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### Author Contributions

<sup>§</sup>These authors contributed equally. C.L.H. designed the study. Y.P., N.C. and K.L. performed the experiments. Y.P., N.C., and C.L.H. analyzed the data. C.L.H. wrote the manuscript. C.L.H. supervised the project. All authors discussed the results and commented on the manuscript.

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### Notes

The authors declare no competing financial interest.

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## ■ ABBREVIATIONS

rtHLF4, recombinant engineered-lactoferrin; fHLF, full-length lactoferrin; MALDI-TOF, matrix assisted laser desorption ionization-time-of-flight; FADD, Fas-associated death domain; TLR, toll-like receptor; BH3, Bcl-2 homology 3; MOMP, mitochondrial outer membrane permeabilization; TAZ, PDZ-binding motif; GDF15, growth differentiation factor 15; YAP, yes-associated protein; TGF- $\beta$ , transforming growth factor beta; SDS-PAGE, sodium dodecyl sulfate-polyacrylamide gel electrophoresis.

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