Intermittent calorie restriction enhances epithelial-mesenchymal transition through the alteration of energy metabolism in a mouse tumor model

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Abstract. The effect of intermittent calorie restriction (ICR) on cancer is controversial. In this study, we examined the effects of ICR and food content in syngeneic BALB/c mice injected with CT26 mouse colon cancer cells. Mice were subjected to 24-h fasting once a week for 4 weeks, and then provided with a control, high-calorie, or trans fatty acid-rich diet. While ICR resulted in increases in tumor weights, metastasis and in the number of cancer stem cells (CSCs) in the tumors or blood of mice fed the control and high-fat diets, it had no effect on body weight after 4 weeks. In particular, we detected increases in the numbers of CSCs in the tumor or blood on the day after starvation, when food overconsumption was detected. Conversely, continuous calorie restriction had no effect on tumor weight, metastasis, or the number of CSCs in tumors or blood. In the post-starvation period, energy metabolism in the tumor was altered from oxidative phosphorylation to glycolysis/lactate fermentation, with the acquisition of the epithelial-mesenchymal transition (EMT) phenotype. Hyperglycemia at the post-starvation period induced the expression of insulin-like growth factor-1, hypoxia-induced factor-1α and Nanog, as well as the phosphorylation of Stat3. Taken together, these findings suggest that ICR induces an increase in the number of CSCs and enhances EMT by promoting the Warburg/Crabtree effect following post-fasting food overconsumption.

Introduction

Calorie restriction (CR) represents an effective means of regulating aging by inducing a healthy lifespan, as observed in various rodent and primate models (1). By contrast, obesity increases the risk, incidence and mortality rate of patients with various diseases, including breast and prostate cancer (2). Indeed, studies have examined the effects of reducing food consumption by 20-40% of normal intake (1). Intermittent CR (ICR) is usually performed by restricting energy intake for 1-3 days/week, followed by ‘free’ eating on non-restriction days. Given that ICR is easier to carry out than traditional CR or continuous CR (CCR), for which energy intake is limited every day, ICR has increased in popularity (3). Moreover, ICR is as effective as traditional CR in achieving weight loss, reducing fat mass and in maintaining the acquired status (4,5).

Weight loss has been shown to decrease the risk of cancer, and to inhibit disease progression and metastasis (6). Whereas traditional CR has been effective in this sense, the anticancer effect of ICR has yet to be proven (6,7). However, it has been proposed that weight cycling, exemplified by CR-induced weight loss followed by weight gain, may decrease the effectiveness of CR, and particularly ICR, on cancer inhibition (7).

Fatty acids, which are found in various food ingredients, influence a wide range of diseases, including cardiovascular disorders, metabolic diseases, such as type-2 diabetes, inflammatory diseases and cancer (8). Evidence indicates that a high-fat diet is associated with aggressive prostate cancer and that n-6 fatty acids enhance breast cancer invasion and metastasis (9). Trans fatty acids (TFAs) represent an established risk factor for cardiovascular diseases, and have been reported to increase the risk of certain types of cancers and to promote cancer cell growth, invasion, anti-apoptotic survival and metastasis (10). In an aim to reduce the incidence of cardiovascular diseases, the United States Food and Drug Administration has ruled that TFAs are not generally recognized as safe for human consumption and must be removed from prepared foods by June 2018 (11).

The overconsumption of saturated fatty acids or TFAs has also been shown to increase the risk of hyperglycemia induced by the enhancement of gluconeogenesis and insulin resistance (12,13). The energy metabolism of cancer cells exhibits a striking feature, referred to the Warburg effect, in which energy production depends predominantly on glycolysis and lactate fermentation, even under aerobic conditions (14,15).
The Warburg effect is enhanced by the glucose load and also by hyperglycemia induced by high-fat diets or the overconsumption of TFAs (16). As such, alterations in blood sugar levels and energy metabolism in cancer cells associated with ICR is an important subject of study (17). In this study, we therefore sought to characterize the effects of ICR on cancer cells, particularly in combination with a high-fat or high-TFA diet.

Materials and methods

Cells and reagents. The CT26 mouse colon cancer cell line was a kind gift from Professor I.J. Fidler (MD Anderson Cancer Center, Houston, TX, USA). The cells were cultured in Dulbecco’s modified Eagle’s medium (DMEM; Wako Pure Chemical Industries, Ltd., Osaka, Japan) supplemented with 10% fetal bovine serum (Sigma-Aldrich Chemical Co., St. Louis, MO, USA).

DMEM containing 75 mg/dl of glucose or 150 mg/dl glucose was generated by mixing glucose-free DMEM with high-glucose DMEM (glucose, 450 mg/dl) (both from Wako Pure Chemical Industries, Ltd.). The insulin-like growth factor-1 receptor (IGF-1R) inhibitor, pircropodophyllin (working concentration, 500 nM), and the hypoxia-inducible factor-1α (HIF-1α) inhibitor, chrysin (working concentration, 50 μM), were obtained from Santa Cruz Biotechnology, Inc. (Dallas, TX, USA), while MitoXpress and CellROX green were purchased from Luxcel Biosciences Ltd. (Cork, Ireland) and Thermo Fisher Scientific (Waltham, MA, USA), respectively.

For the measurement of oxygen consumption and reactive oxygen species (ROS) levels, the cells were seeded into 96-well plates (2,000 cells/well) and cultured with regular medium to the manufacturer’s instructions. The number of replicates was 30 cycles (94°C/1 min-66°C/1 min-72°C/1 min).

Reverse transcription-quantitative PCR (RT-qPCR). The extraction of total RNA was carried out using an RNaseasy mini kit (Qiagen), and total RNA (1 μg) was synthesized using the ReverTra Ace-α-RT kit (Toyobo, Osaka, Japan). qPCR was performed using a StepOne Real-Time PCR system with Fast SYBR®-Green Master Mix (Applied Biosystems, Life Technologies, Carlsbad, CA, USA); a relative standard curve quantification method was used for analysis (18,19). PCR products were visualized by agarose gel electrophoresis with ethidium bromide staining.

Animals. A total of 70 male BALB/c mice (4 weeks old, mean body weight, 17.6±2.4 g) were purchased from SLC Japan (Shizuoka, Japan). The animals were maintained in a pathogen-free animal facility in a 12/12 h light/dark cycle in a temperature (22°C)- and humidity-controlled environment, in accordance with the institutional guidelines approved by the Committee for Animal Experimentation of Nara Medical University, Kashihara, Japan following current regulations and standards of the Japanese Ministry of Health, Labor and Welfare (approval no. 9559).

Animal models. To establish a subcutaneous tumor model, CT26 cancer cells (1x10⁵) were inoculated into the scapular subcutaneous tissue of BALB/c mice. Mice were observed for 4 weeks following inoculation. For ICR, the mice were starved for 24 h (from 8 a.m. on the first day of the week to 8 a.m. on the second day of the week); 4 mice were examined at each time-point. For CCR, the mice were provided access to a low calorie (30% calorie reduction) diet ad libitum.

The animals were then subjected to the following diets: A control diet [CE-2 (containing 5% crude fat derived mainly from soy bean oil; CLEA Japan, Inc., Tokyo, Japan)], a high-fat diet [Quickfat (containing 14% crude fat derived mainly from beef tallow; CLEA Japan, Inc.)], an elaidic acid (EA)-rich diet [CE-2 + 10% EA (Sigma-Aldrich Chemical Co.), or a low calorie diet + 10% EA (70% w/w CE-2 + 30% w/w cellulose (Wako Pure Chemical Industries, Ltd.)]. In groups C and CI, mice were fed the CE-2 diet. In groups F and FI, mice were fed a high-fat diet. In groups E, EI, E-CNT and E-ICR, mice were fed an EA-rich diet. In group E-CRR, mice were fed a low calorie diet. In each group, 5 mice were used. The subcutaneous tumors were excised from the euthanized mice on day 28 to measure tumor weight and examine histologically.

For detecting lung metastasis, indocyanine green-labeled (Dojindo Laboratories, Kumamoto, Japan) mouse anti-epidermal growth factor receptor antibody (EGFR, 1005, sc-03-G; Santa Cruz Biotechnology, Inc.) diluted in PBS (0.5 μg/200 μl) was injected into the caudal vein prior to euthanasia at day 28. Excised lungs were observed using a Clairvivo OPT in vivo imager (Shimadzu Corp., Kyoto, Japan) (18).

Reverse transcription-polymerase chain reaction (RT-PCR). In the weekly change studies, the blood was obtained from caudal vein of mice (n=5) at 2 p.m. each day, repetitively. In the energy metabolism experiment, the blood was obtained by cardiac centesis from euthanized mice (n=5) on day 2 of the ICR model, at 6 h after re-feeding. Total RNA (1 μg) was isolated from the blood samples using the PAX gene Blood RNA kit (Qiagen, Venlo, The Netherlands). PCR products were separated on a 2% agarose gel, visualized with ethidium bromide, and quantified using the QuantiTect Primer assay (Qiagen), according to the manufacturer’s instructions. The following primer sets (Sigma Genosys, Ishikari, Japan) were used for amplification: Mouse CDI33 (Prom1, accession no. BC028286.1) forward, 5'-gaa aag ttg ctc tgc gaa cc-3' and reverse, 5'-tct caa gct gaa cag aag ca-3'; mouse nucleostemin (NS) (Gltn3, accession no. AY181025.1) forward, 5'-cag gat gct gac gat caa cga ga-3' and reverse, 5'-ttg att gct gat gct gta cca gc-3; mouse β-actin (ActB, accession no. NM_007393.4) forward, 5'-agg cat gta cgt acg ca-3 and reverse, 5'-ctc tca gct gtt gtt gtt gtt aa-3'. PCR conditions were set in accordance with the manufacturer’s instructions. The number of replicates was 30 cycles (94°C/1 min-64°C/1 min-72°C/1 min).

Reverse transcription-quantitative PCR (RT-qPCR). The extraction of total RNA was carried out using an RNeasy mini kit (Qiagen), and total RNA (1 μg) was synthesized using the ReverTra Ace-α-RT kit (Toyobo, Osaka, Japan). qPCR was performed using a StepOne Real-Time PCR system with Fast SYBR®-Green Master Mix (Applied Biosystems, Life Technologies, Carlsbad, CA, USA); a relative standard curve quantification method was used for analysis (18,19). PCR was carried out under the conditions recommended by the manufacturer. Hif-1A RT-PCR primers were purchased from Santa Cruz Biotechnology, Inc. ActB mRNA was amplified as an internal control (GenBank Accession no. NM_001101). Each amplification reaction was evaluated by melting curve analysis. PCR products were visualized by agarose gel electrophoresis with ethidium bromide staining.
**Immunohistochemistry.** Consecutive 4-μm-thick sections were immunohistochemically stained via the immunoperoxidase technique, as previously described (20). Antibodies against mouse CD133 (18470-1-AP; Proteintech, Rosemont, IL, USA), NS (clone E-8, sc-166460; Santa Cruz Biotechnology, Inc.) and mindbomb E3 ubiquitin protein ligase 1 (M7240; MIB1; Ki-67 index) were used at 12.5% sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) and transferred onto nitrocellulose membranes. The membranes were then incubated with primary antibodies specific to Nanog (ab80892; Abcam, Cambridge, UK), Snail (clone C15D3, #3879; Cell Signaling Technology Japan, Tokyo, Japan), E-cadherin (clone HECD-1, ab1416), vimentin (ab24535) (both from Abcam), Stat3-phosphoTyr705 (clone D3A7, #9145), or Stat3 (#9132) (Cell Signaling Technology Japan), followed by peroxidase-conjugated IgG antibodies (#330 and #458; MBL, Nagoya, Japan). Anti-tubulin antibody was used to assess the levels of protein loaded per lane (clone DM1A, 05-829; Oncogene Research Products, Cambridge, MA, USA). Immune complexes were visualized using an ECL western blot detection system (Amersham, Aylesbury, UK).

Measurement of blood sugar and lactate levels. In the energy metabolism experiment, blood, obtained by cardiac centesis from euthanized mice (n=5) on day 28 on 2 of the ICR model, at 6 h after re-feeding was subjected to the measurement of the sugar concentration. The sugar concentration was determined that only the mice in group FI exhibited a transient increase in CD133 expression levels on day 2, which then returned gradually to the control levels (Fig. 2B). Thus, increased CD133 mRNA blood levels might be affected by daily food intake, particularly by over-intake.

**Results**

Effect of a high-fat diet and intermittent starvation on tumor growth and cancer stem cells (CSCs). First, we assessed the effects of a high-fat (Quickfat) diet on the growth of CT26 colon cell subcutaneous tumors in BALB/c mice, with or without intermittent caloric restriction (ICR) (Fig. 1). The protocol is shown in Fig. 1A. No differences in body weight were observed between the groups (Fig. 1B); however, mice fed the control or high-fat diet had heavier tumors and more nodal metastases when subjected to ICR (groups C1 and F1, respectively), compared to the mice not subjected to ICR (groups C and F, respectively) (Fig. 1C and D). Likewise, the tumors of mice subjected to ICR (groups C1 and F1) contained significantly greater numbers of cells positive for CD133, a marker of CSCs, than those in the non-ICR groups (groups C and F) (Fig. 1E and F).

Effect of ICR on cancer cell stemness. To assess the temporal effects of ICR on the number of CD133+ cells in tumors, we compared CD133+ values on day 1 (fasting), day 2 (re-feeding) and day 4 (feeding) in each of the 4 mouse groups (Fig. 2A). Notably, while there was no difference in the number of CD133+ cells between the groups, by day 2, there was a significant increase in the number of CD133+ cells in the mice in groups C1 and F1, compared to those in groups C and F. However, the number of CD133+ cells in the mice in groups C1 and F1 decreased to baseline levels by day 4.

To examine the effect of ICR, we focused on the differences between group F and FI, which were markedly affected by ICR. Via daily comparisons of CD133 mRNA levels, we determined that only the mice in group FI exhibited a transient increase in CD133 expression levels on day 2, which then returned gradually to the control levels (Fig. 2B). Thus, increased CD133 mRNA blood levels might be affected by daily food intake, particularly by over-intake.

Effect of an EA-enriched diet with ICR on tumor growth and CSCs. We previously reported that EA (C18:2, trans) modulates CSCs and affects cancer metastability (22,23). Accordingly, in this study, the effects of EA on tumor growth were examined in the same mouse subcutaneous tumor model used above (Fig. 3A). Body weight was indistinguishable between the mice fed an EA-enriched diet with (group EI) or without ICR (group E) or the control diet with (group CI) or without ICR (group C) (Fig. 3B). Tumor weight and cell growth (MIB1 index) were most pronounced in the mice in group EI, followed by those in group E (Fig. 3C and D). Whereas the tumor weight did not differ between the two control groups, the MIB1 index was higher in the mice in CI than in those in group C. NS+ cells were also most abundant in mice in group EI, followed by those in groups C and CI (Fig. 3E). By contrast, the CD133+ cells were most frequent in mice in group EI, followed by those in groups CI and E (Fig. 3F).

Effect of an EA-enriched diet with ICR on metastasis. To examine the effects of an EA-enriched diet with ICR on metastasis, mice in groups E and EI were starved on day 1 (Fig. 4A and B, respectively) and then provided the appropriate diet for the remainder of the week and for 4 weeks. Whereas no significant changes in metastasis were observed in the mice in group E, those in group EI exhibited increased numbers of NS+ cells and an increased MIB1 index on day 2. By contrast, there was no change in the numbers of CD133+ cells in either group at this time-point. As
shown in Fig. 4C, the mice subjected to ICR (groups CI and EI) exhibited a surge in NS mRNA levels in the blood on the day after starvation. At 4 weeks after inoculation, micrometastatic foci were detected in the lungs by immunofluorescence using anti-EGFR antibody (18). The intensity of fluorescence increased in the following order: C < CI < E < EI (Fig. 4D). Specifically, the intensities in CI and EI groups increased to levels 1.8- and 6-fold greater than those observed in groups C and E, respectively.

Comparison of EA-associated pro-tumoral effect between ICR and CCR. Subsequently, we examined the effects of ICR on the pro-tumoral effect of EA in comparison with CCR (Fig. 5). We prepared the low calorie diet by mixing the standard diet (CE-2) with edible cellulose at a 7:1 ratio. This food therefore contained 70% of the calories of the CE-2 diet. As shown in the protocol (Fig. 5A), the mice were divided into 3 groups as follows: the no calorie restriction with 10% EA (E-CNT), ICR with 10% EA (E-ICR) and CCR with 10% EA (E-CCR) groups. While there were no differences in body weight between the 3 groups on day 29, the mice in the E-ICR group exhibited increased tumor weights and MIB1 index scores, compared to those in the E-CNT or E-CCR groups (Fig. 5C and D).

The examination of daily caloric intakes indicated that mice in the E-CCR group ate a smaller volume of food to reach 80% of the calorie intake of mice in the E-ICR group on non-starvation day (the expected calorie intake was 70% in E-CCR group) (Fig. 5E). Specifically, the mice in the E-CCR group consumed 83% of the calories as those in the E-ICR group weekly. However, this over-intake of food resulted in an enhanced EA intake (117%) in mice provided with E-CCR (Fig. 5F), compared to those provided the E-ICR diet.

To confirm the observed surge in stem cell marker expression in the blood (Figs. 2 and 4), we measured the mRNA expression levels of NS in the blood samples from each mouse on day 2 (Fig. 5G). Notably, the mice in the E-ICR group, but not those in the E-CCR group, exhibited an enhanced NS mRNA expression. Likewise, lung micrometastasis was increased in mice provided the E-ICR diet, but not in those given the E-CCR diet (Fig. 5H).

Effect of post-starvation overconsumption on energy metabolism, stemness and the EMT phenotype in CT26 cells. To
examine the energy metabolism and stemness phenotypes of CT26 tumors in the post-starvation phase, tumors harvested between the starvation and post-starvation phase were compared in mice fed the control or 10% EA diet (Table I). In both groups, blood sugar concentrations were elevated 2-fold in the post-starvation (over-intake) phase, compared to the starvation phase. Likewise, the levels of cell proliferation (MIB1) and stemness (NS) markers were also increased in the post-starvation phase. Notably, while energy was primarily produced via oxidative phosphorylation (OXPHOS) during starvation, cells exhibited a switch to glycolysis/lactate fermentation (GL/LF)-mediated energy production during the post-starvation phase, which was accompanied by a decrease in ROS production.

Table I. Comparison of tumors in starvation phase with those in the post-starvation (over-intake) phase.

|                     | Control diet |                          | 10% EA diet |                          |
|---------------------|--------------|---------------------------|-------------|---------------------------|
|                     | Starvation   | Over-intake               | P-value     | Starvation               | Over-intake               | P-value     |
| Blood sugar (mg/dl) | 78±8.2       | 148±16                   | <0.0001     | 84±9.1                   | 167±18.4                 | <0.0001     |
| MIB1 positivity (%) | 48±3.1       | 65±5.8                   | <0.0001     | 62±6.3                   | 88±7.4                   | <0.0001     |
| NS positivity (%)   | 44±3.2       | 62±7.1                   | <0.0001     | 55±6.4                   | 76±6.2                   | <0.0001     |
| O₂ consumption (kpiu) | 7.5±0.51 | 4.5±0.25                 | <0.00001    | 8.9±0.59                 | 4.2±0.18                 | <0.0001     |
| Lactate (pM)       | 2.1±0.31     | 7.2±0.60                 | <0.00001    | 2.4±0.15                 | 8.7±0.66                 | <0.00001    |
| ROS (%)            | 100±12       | 57±7                     | <0.00001    | 121±11                   | 63±5                    | <0.00001    |

*aStatistical significance was calculated by a two-tailed Student’s t-test; * ′measured by MitoXpress dye staining; kpiu, kilo phosphorescence intensity; ′reactive oxygen species levels were measured by CellROX Green dye staining. Fluorescence intensity is standardized by that of control diet-starvation group, which is set to 100%. Anti-Ki-67 protein antibody clone MIB1; NS, nucleostemin; ROS, reactive oxygen species.

Figure 2. Temporal changes in CD133 expression. (A) Percentage of CD133+ cells on day 1 (fasting), day 2 (re-feeding) and day 4 (feeding). (B) Time-course analysis of CD133 mRNA levels in the blood and food intake in groups F (high-fat diet) and FI (high-fat diet with ICR). mRNA levels and food calorie intake were standardized to those on day 1 for group F, which were set as 100%. Error bars indicate the means ± standard deviations. Group C, controls not subjected to intermittent calorie restriction (ICR); group CI, controls subjected to ICR.
To confirm the effects of sequential increases in blood glucose concentrations on energy metabolism and stemness, the CT26 cells were cultivated in medium containing 75 mg/dl glucose, followed by medium containing 150 mg/dl glucose (Table II). These glucose concentrations were selected to simulate those observed in the blood of mice, as shown in Table I. The CT26 cells cultivated in the presence of 150 mg/dl glucose exhibited increased cell proliferation, stemness and GL/LF, and decreased OXPHOS and ROS production, compared to those cultivated in the presence of 75 mg/dl glucose.

Subsequently, CT26 tumors in the animal model and CT26 cells in in vitro treatment were subjected to immunoblot analysis to compare the expression levels of EMT- and stemness-specific proteins between the starvation phase and post-starvation phase and between cultivation with 75 and 150 mg/dl glucose, respectively (Fig. 6A and B). The tumors harvested during the post-starvation phase or the cells after cultivation in medium containing 150 mg/dl glucose exhibited increased cell proliferation, stemness and GL/LF, and decreased OXPHOS and ROS production, compared to those cultivated in the presence of 75 mg/dl glucose.

To examine the signaling pathway associated with the alterations observed during the post-starvation phase, the CT26 cells were implanted into the scapular subcutaneous tissues of BALB/c mice. Mice were subjected to a 24-h fasting period once each week for 4 weeks. Tumors were excised and examined on day 29.

Table II. Comparison of CT26 cells cultured under hypoglycemic and normoglycemic conditions.

| Glucose concentration (mg/dl) | 75 | 150 | P-valuea |
|-------------------------------|----|-----|----------|
| MIB1 positivity (%)           | 45±4.9 | 70±6.2 | <0.01 |
| NS positivity (%)             | 40±4.1 | 58±6.1 | <0.05 |
| O2 consumption (kpiu)b         | 8.75±0.626 | 4.78±0.302 | <0.001 |
| Lactate (pM)                  | 1.9±0.22 | 8.1±0.71 | <0.001 |
| ROS (%)c                      | 100±9 | 62±7 | <0.01 |

aStatistical significance was calculated by a two-tailed Student’s t-test; bmeasured by MitoXpress dye staining; kpiu, kilo phosphorescence intensity; creactive oxygen species levels were measured by CellROX Green dye staining. Fluorescence intensity is standardized by that of control diet-starvation group, which is set to 100%. Anti-Ki-67 protein antibody clone MIB1; NS, nucleostemin; ROS, reactive oxygen species.

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were treated with inhibitors of IGF-1R or HIF-1α (Table III). Both treatments, respectively, abrogated the effects observed in CT26 cells cultured in medium containing 150 mg/dl glucose, following medium containing 75 mg/dl glucose.

Table III. Effects of the inhibitors of IGF-1R and HIF-1α on energy metabolism in CT26 cells.

| Glucose concentration (mg/dl) | 150  | 150  | P-value* | 150  | P-value* |
|------------------------------|------|------|----------|------|----------|
| Inhibitor                    |      |      |          |      |          |
| MIB1 positivity (%)          |      |      |          |      |          |
| None                         | 71±6.5 | 45±4 | <0.01    | 42±3.1 | <0.01    |
| IGF-1R                       |      |      |          |      |          |
| HIF-1α                       |      |      |          |      |          |
| MIB1 positivity (%)          |      |      |          |      |          |
| None                         | 61±6.3 | 42±4.4 | <0.05 | 40±3.5 | <0.01    |
| IGF-1R                       |      |      |          |      |          |
| HIF-1α                       |      |      |          |      |          |
| O₂ consumption (kpiu)*       | 4.67±0.298 | 7.44±0.752 | <0.001 | 7.58±0.788 | <0.001 |
| Lactate (pM)                 | 7.9±0.84 | 2.2±0.24 | <0.001 | 2.0±0.25 | <0.001 |
| ROS (%)³                     | 64±7 | 99±10 | <0.01    | 103±11 | <0.01    |

Statistical significance was calculated by a two-tailed Student's t-test; *measured by MitoXpress dye staining; kpiu, kilophorescence intensity; reactive oxygen species levels were measured by CellROX Green dye staining. Fluorescence intensity is standardized by that of control diet-starvation group, which is set to 100%. Anti-Ki-67 protein antibody clone MIB1; NS, nucleostemin; ROS, reactive oxygen species; IGF-1R, insulin-like growth factor-1 receptor; HIF-1α, hypoxia-inducible factor-1α.

Figure 4. Effect of an EA-enriched diet with intermittent calorie restriction (ICR) on cancer metastasis. (A) Temporal changes in food intake and the number of nucleostamin-positive (NS⁺) and CD133⁺ cells in group E (EA-rich diet without ICR) tumors. (B) Temporal changes in food intake and the number of NS⁺ and CD133⁺ cells in group EI (EA-rich diet with ICR) tumors. Food calorie intake was standardized to that on day 1 for group E, which was set as 100%. Positivity of NS or CD133 was calculated as the percentage of positively stained cells among 200 cells observed. (C) Time-course analysis of NS mRNA levels in the blood. (D) Micrometastasis in the lung on day 29. Cancer cells were detected by ICG-labeled anti-EGFR antibody and measured using an in vivo imager. Error bars indicate the means ± standard deviations. Group C, controls not subjected to ICR; group CI, controls subjected to ICR; group E, mice given EA-rich diet not subjected to ICR; group EI, mice given EA-rich diet subjected to ICR.
Discussion

In this study, we examined the effects of ICR on tumor growth using a mouse syngeneic subcutaneous tumor model. Our findings demonstrated that instead of having a tumor suppressive effect, ICR promoted tumor growth and increased the number of CSCs. Such a pro-tumoral effect was suggested to depend on the induction of a starvation-overeating cycle.

In designing the ICR experiments, starvation for 1 day/week was expected to decrease dietary calorie intake by 14%. However, overeating after starvation abrogated CR, and dietary calorie loss was only 1.5%. This was clearly confirmed by the lack of any difference in body weight between the mice subjected to ICR and those not subjected to ICR. Strictly speaking, our experimental design therefore achieved dietary restriction, but not CR. Dietary restriction is defined as intermittent fasting without CR (1) and has been shown to effectively prolong lifespan with minimal adverse effects (1), as well as to ameliorate obesity and type-2 diabetes (24,25).

We also compared our ICR mouse model with a CCR mouse model. While CCR model achieved an ~80% loss in dietary calorie intake each day, the mice in the CCR group...
exhibited no suppression of tumor growth, compared with those receiving a normal diet (group CNT). In the context of an EA-enriched diet, CCR may therefore not be effective for tumor suppression; however, CCR did not enhance tumor growth. The animals subjected to CCR also exhibited no increase in lung micrometastasis in comparison to the mice in the ICR group. These findings suggest that CCR may not affect cancer growth or metastasis.

Mice receiving either an EA-enriched diet or the control diet with ICR exhibited marked increases in lung micrometastasis, compared to the control animals. Likewise, both groups of mice exhibited surges in the mRNA expression of stemness-associated genes within the blood. These findings suggest that the release of CSCs into the blood may be associated with the starvation-overeating cycle.

To assess the mechanisms underlying the effect of post-starvation food overconsumption on metastasis, we compared energy production, stemness and EMT phenotype in CT26 cells both in vivo and in vitro. From the starvation phase to the post-starvation phase, blood sugar concentrations were increased 2-fold in mouse tumors in vivo. The transition from the low- to high-glucose conditions altered the primary energy production pathway from the OXPHOS pathway to GL/LF, which is compatible with the Crabtree effect. The Crabtree effect (26) is associated with an increase in stemness and acquisition of the EMT phenotype through the IGF-1/HIF-1α/Nanog/Stat3 signaling pathway, and HIF-1α expression is upregulated by hyperglycemia (27). IGF-1 and AKT are known as the non-hypoxic HIF-1α induction pathway (28). In particular, the hypoglycemic-hyperglycemic transition promotes IGF-1 expression (29,30), and AKT is located downstream of IGF-1 in this pathway (31). Moreover, HIF-1α is a key factor necessary for switching the energy production pathway from OXPHOS to GL/LF through the activation of pyruvate dehydrogenase (32,33). This switching induces higher levels of side population, and promotes the expression of Nanog and Oct3/4 (34), which in turn activates Stat3, resulting in the induction of Snail and a subsequent EMT phenotype (35,36). Thus, the Crabtree effect, a temporal increment of the Warburg effect, may be closely associated with the acquisition of metastatic potential in cancer cells.

EA is a main component of industrial dietary TFAs, which play an important role in the progression and metastasis of cancer, as well as in cardiovascular diseases (8,37). The consumption of TFAs is associated with overeating behaviors (38), which may result in a preference for fast food or snacks containing TFAs (39). In this study, we focused on the intake of EA under conditions of ICR. Such conditions can be generalized to the intake of cancer-promoting compounds accompanied by intermittent overeating.

Our data indicated that ICR led to a starvation-overeating cycle. Intermittent feeding increases the expression of orexigenic neurotransmitters, agouti-related peptide and neuropeptide Y. In addition, ICR inhibits the expression of
pro-opiomelanocortin in the hypothalamus of rodents, even on feeding days, thus explaining the overeating pattern observed in this study (40,41). Feeding uncertainty may induce ‘food addiction’, particularly to fat- and sugar-rich diets (42). This conclusion is supported by the weight loss exhibited by the control diet group subjected to ICR, but not by the high-fat and EA diet groups subjected to ICR in this study.

In conclusion, overeating associated with ICR abrogated CR and weight reduction in our models. Moreover, overeating after ICR induced the Crabtree effect in cancer cells, resulting in increased numbers of CSCs and cancer cells circulating within the blood. The Warburg effect or glycolytic energy production is linked closely with malignant phenotypes, such as metastasis (43,44). Our data on ICR are compatible with the mechanism. We also demonstrated that dietary EA was associated with an increment in the metastatic potential of tumors by the acquisition of the EMT phenotype. We have already previously reported the mechanism of EA-induced EMT with the enhancement of stenness (23). Restriction-associated overeating may be the result of an instinctive behavior. Our observations suggest that an irregular dietary intake evoking the starvation-overeating cycle may endow cancer cells with metastatic ability.

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