

1 **High-fat diet-induced obesity accelerates the progression of spontaneous**  
2 **osteoarthritis in Senescence-Accelerated Mouse Prone 8 (SAMP8)**

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22 **Abstract**

23 Objectives: Aging and obesity are major risk factors for osteoarthritis (OA), a  
24 widespread disease currently lacking efficient treatments. Senescence-accelerated  
25 mouse prone 8 (SAMP8) display early-onset aging phenotypes, including OA. This  
26 study investigates the impacts of high-fat diet (HFD)-induced obesity on OA  
27 development in SAMP8.

28 Methods: SAMP8 at five weeks were fed either a normal chow diet or an HFD for ten  
29 weeks to induce obesity. Parameters related to obesity, liver function, and lipid and  
30 glucose metabolism were analyzed. At 14 weeks of age, knee joint pathology, bone  
31 mineral density, and muscle strength were assessed. Immunohistochemistry and  
32 TUNEL staining were performed to evaluate markers for cartilage degeneration and  
33 chondrocyte apoptosis.

34 Results: At 14 weeks of age, HFD-induced obesity increased liver and adipose tissue  
35 inflammation in SAMP8 without further exacerbating diabetes. Histological scoring  
36 revealed aggravated cartilage, menisci deterioration, and synovitis, while no further  
37 loss of bone mineral density or muscle strength was observed. Increased chondrocyte  
38 apoptosis was detected in knee joints following HFD feeding.

39 Conclusions: Ten weeks of HFD feeding promotes spontaneous OA progression in 14-  
40 week-old SAMP8, potentially via liver damage subsequent chondrocyte apoptosis.  
41 This aging-obese mouse model may prove valuable for further exploration of  
42 spontaneous OA pathophysiology.

43 Keywords: Apoptosis, High-fat diet, Obesity, Osteoarthritis, SAMP8

44

## 45 **Introduction**

46 Osteoarthritis (OA), the most prevalent joint disease that affects millions of people  
47 worldwide, is characterized by the loss of cartilage and bone remodeling, leading to  
48 pain, stiffness, and loss of mobility[1]. While OA can affect nearly any joint in the body,  
49 it predominantly occurs in weight-bearing joints, such as the hips and knees[2]. As a  
50 multifactorial disorder, spontaneous OA is initiated and perpetuated by a complex  
51 interplay of various contributing factors. Aging and obesity have been identified as  
52 significant risk factors for OA, because the risk of developing OA increases with age  
53 and overload, as the joints are subjected to years of wear and tear. The increased  
54 pressure on weight-bearing joints from carrying excess weight can cause inflammation  
55 and cartilage damage. Moreover, obesity is linked to chronic low-grade inflammation,  
56 which may further contribute to OA development[3]. In human OA cases, aging and  
57 obesity often cooperatively contribute to the progression of OA, imposing a substantial  
58 burden on healthcare systems[4], especially with the background of expanding elderly  
59 population in many countries.

60 Senescence-accelerated mouse (SAM), is a mouse strain derived from the AKR/J  
61 strain[5]. SAM consists of senescence-prone inbred strains (SAMP) and senescence-  
62 resistant inbred strains (SAMR) line which shows normal aging characteristic[5]. Thus,  
63 in SAM studies, SAMR has been used as control of SAMP. Among SAMP, SAMP8 has  
64 been used as a mouse model to investigate aging-related neurological and cognitive

65 deficits, such as Alzheimer's disease[6]. In C57BL/6 mice, moderate OA-like changes  
66 become detectable at 12-24 months, at least[7, 8]. However, we recently reported that  
67 SAMP8 exhibited significant meniscus and cartilage degeneration in knee joints with  
68 subchondral bone sclerosis and osteopenia from an early stage of about 14 weeks of  
69 age[9]. Furthermore, although various studies have examined the relationship between  
70 obesity and OA in ob/ob mouse, diet-induced obese C57BL/6 mice and/or post-  
71 traumatic OA models[10, 11, 12, 13, 14], the potential influence of obesity on the  
72 accelerated aging-related primary OA and the underlying mechanisms have yet to be  
73 determined. The purpose of the present study was to determine the effect of high-fat  
74 diet (HFD)-induced obesity to the development of OA in spontaneous OA model mouse,  
75 SAMP8.

76 In the present study, we demonstrated that in SAMP8, high-fat diet-induced obesity  
77 induces liver damage and further aggravates the severity of primary OA at an early  
78 age of 14 weeks through increased chondrocyte apoptosis, without further  
79 exacerbating remarkable metabolic dysregulation such as hyperlipidemia and diabetes.

80

## 81 **Materials and Methods**

### 82 **Animal Model and Experimental Design**

83 SAMP8 at 4 weeks of age were obtained from Japan SLC (Shizuoka, Japan), and only  
84 male mice were used in this study. After one week of acclimatization, forty SAMP8  
85 were randomly divided into two groups: the high-fat diet (HFD) group (n=20) and the

86 normal chow diet (ND) group (n=20). The HFD group was fed a diet of HFD-60  
87 (Protein:Fat:Carbohydrate = 18.2:62.2:19.6, Oriental Yeast Co., Ltd., Tokyo, Japan) for  
88 10 weeks, while the ND group was fed a normal chow diet (AIN93M;  
89 Protein:Fat:Carbohydrate = 14.1:10.0:75.9, Oriental Yeast) (Fig. 1A). All mice were  
90 housed in groups of five per cage (S143 mm × 293 mm × H148mm) with a sterilized  
91 beta-chip bedding and maintained at 23±1 °C with a 12-h light/dark cycle and acidified  
92 water and complete commercial pelleted food ad libitum. Body weight was measured  
93 weekly. Mice were sacrificed for histological and biological analyses at 14 weeks of  
94 age. All animal experiments were performed according to protocols approved by the  
95 Hiroshima University Animal Care and Use Committee.

96

### 97 **Histopathological Assessments of Knee Joints**

98 Knee joints from each mouse were embedded intact in paraffin after fixation in 4%  
99 paraformaldehyde phosphate buffer solution (PFA-PBS) for 48h at 4°C and  
100 decalcification in K-CX (FALMA, Japan) for 6 h at room temperature. Paraffin-  
101 embedded knee joints were sectioned at 4.5 µm in the coronal plane through the  
102 central weight-bearing region of the anterior and posterior femorotibial joint. Three  
103 different sections per joint were stained with Safranin O (MUTO PURE CHEMICALS,  
104 Tokyo, Japan) and Fast Green (Sigma-Aldrich, USA). Histological assessments were  
105 performed on each section, and the average scores from three different sections were  
106 used for statistical analysis. Three different researchers were blinded while performing

107 all scorings. Scoring systems for articular cartilage, meniscus, subchondral bone, and  
108 synovium were applied in this study. Damage of articular cartilage (maximum 24 points  
109 per knee joint section; 6 points for each quadrant of the medial and lateral femoral/tibial  
110 cartilage) was evaluated using the OARSI scoring system [15]. Subchondral bone  
111 changes, meniscus degradation, and the severity of synovitis were evaluated  
112 according to previously described histopathological scoring systems[16, 17, 18] using  
113 the left knee joints.

114

#### 115 **Hematoxylin and Eosin (H&E) Staining**

116 The liver, epididymal and subcutaneous fat tissue were resected from mice and  
117 immediately fixed in 4% PFA-PBS. The tissues were embedded in paraffin and sliced  
118 into 4.5 $\mu$ m thick sections for H&E staining. Slides were imaged using a Keyence BZ-  
119 X800 microscope (Keyence Cooperation, Japan). The H&E-stained sections of the  
120 epididymal adipose tissue were analyzed for adipocyte size determination and crown-  
121 like structures (CLS) quantification using Image J software (NIH, USA).

122

#### 123 **Blood Biochemical Analysis**

124 Blood samples were collected by cardiac puncture after sacrifice. After coagulating at  
125 room temperature for one hour, the blood samples were centrifuged under 1,500g for  
126 30 min. Supernatant serum were collected for subsequent analyses. Serum  
127 biomarkers of aspartate aminotransferase (AST), alanine aminotransferase (ALT), total

128 cholesterol (T-CHO), triglycerides (TG), glucose (GLU) and non-esterified fatty acid  
129 (NEFA) in serum were analyzed by Nagahama Life Science Laboratory (Nagahama,  
130 Japan) using routine laboratory methods.

131

### 132 **Glucose Tolerance Test (GTT)**

133 See Supplemental Materials.

134

### 135 **Dual-energy X-ray Absorptiometry (DEXA) Analysis**

136 See Supplemental Materials.

137

### 138 **Grip Strength Test**

139 See Supplemental Materials.

140

### 141 **TUNEL Staining**

142 TUNEL staining was completed using an in-situ detection kit for programmed cell death  
143 detection (MEBSTAIN apoptosis TUNEL Kit direct: MBL, United States) according to  
144 the manufacturer's instructions. Nuclei were stained by 4',6-diamidino-2-phenylindole  
145 (DAPI).

146

### 147 **Immunohistochemical Analysis**

148 Slides were pretreated with antigen-retrieval reagent (Immunoactive; Matsunami Glass

149 Ind, Osaka, Japan) at 60°C for 16 hours, followed by blocking serum for 30 minutes.  
150 Then, sections were immunostained with anti-C/EBP homologous protein (CHOP)  
151 antibody (proteintech, 15204-1-AP, 3.5µg/mL), anti-p16<sup>INK4a</sup> antibody (abcam, ab54210;  
152 0.1µg/mL), anti-ADAMTS5 antibody (GeneTex, GTX100332, 10µg/mL) and anti-  
153 MMP13 antibody (ThermoFisher Scientific, MA5-14328, 20µg/ml) diluted in Can Get  
154 Signal immunostaining solution (TOYOBO, Tokyo, Japan) using Vectastain Elite ABC-  
155 HRP kit and DAB substrate kit (Vector Laboratories, Burlingame, CA, USA) according  
156 to the manufacturers' instructions.

157

## 158 **Statistical Analyses**

159 Data are expressed as mean ± standard deviation or median ± 95% confidence interval  
160 (knee histopathological scores). Statistical analysis of data was performed using Prism  
161 v9.5.1 (GraphPad Software, San Diego) as follows: two-tailed unpaired Welch's t tests  
162 for tissue weights, blood biochemical markers, muscle strength and quantifications of  
163 histological data; 2-way ANOVA with Sidak's multiple correction for body weight;  
164 nonparametric Mann-Whitney tests for knee histopathological scores; multiple  
165 unpaired t tests with Holm-Sidak's multiple correction for bone mineral density. The  
166 results were considered statistically significant at a value of  $p < 0.05$ .

167

## 168 **Results**

### 169 **High-fat diet feeding induced obesity in SAMP8**

170 To assess the impact of a high-fat diet (HFD) on the body weight of SAMP8 in  
171 comparison to a normal chow diet (ND), we monitored the body weight of mice from  
172 both groups on a weekly basis. The body weight of the HFD group was significantly  
173 increased compared to the ND group (Fig. 1B, C). The average increase in body weight  
174 was 5.228 grams at 14 weeks of age. Furthermore, we observed an increase in the  
175 tissue weight of the liver (Fig. 1D), as well as epididymal and subcutaneous fat (Fig.  
176 1E). These results demonstrate that 10 weeks of HFD feeding, from 5 to 14 weeks of  
177 age, induced obesity and increased fat accumulation in SAMP8.

178

#### 179 **Lipid and glucose metabolism in SAMP8 by HFD**

180 Given the observed increase in liver and fat tissue weights in the HFD-fed mice, we  
181 investigated the effects of HFD on function of liver and adipose tissue next, as well as  
182 lipid and glucose metabolism, as obesity is a significant predisposing factor for  
183 metabolic dysregulation, such as non-alcoholic fatty liver disease (NAFLD),  
184 hyperlipidemia and type 2 diabetes. We observed enlarged hepatocytes and an  
185 increased number of lipid droplets in the livers of HFD-fed mice compared with those  
186 fed with a chow diet, as seen in the H&E-stained sections (Fig. 2A). Liver damage  
187 markers, aspartate aminotransferase (AST) and alanine aminotransferase (ALT), were  
188 significantly upregulated in the HFD group (Fig. 2B). Circulating lipids, including TG  
189 and T-CHO significantly reduced in serum of HFD group, indicating liver inflammation  
190 and damage induced by the HFD (Fig. 2C). Moreover, the HFD group exhibited

191 reduced serum albumin concentration and albumin/globulin (A/G) ratio, and  
192 unchanged total protein levels (Supplemental Fig. 1A), while the renal function markers  
193 were unaltered (Supplemental Fig 1B), suggesting an impaired hepatic function in  
194 these mice. Additionally, serum ammonia (NH<sub>3</sub>) levels were found to be elevated in the  
195 HFD group (Supplemental Fig. 1C), further supporting the notion of impaired liver  
196 function and ammonia clearance due to the HFD. Furthermore, to assess the influence  
197 of HFD on adipose tissue morphology and inflammation, we analyzed H&E-stained  
198 microscopic images of epididymal adipose tissue. Adipocyte size (Fig. 2D and E) and  
199 crown-like structure (CLS) ratio (Fig. 2F) were increased in the HFD group, indicating  
200 increased macrophage infiltration and systemic predisposition to inflammation[19, 20].  
201 Although 8 out of 20 SAMP8 in the ND group already displayed an impaired clearance  
202 of blood glucose and hyperglycemia (>150mg/dL) (Fig. 2G and H) at 14 weeks, HFD  
203 feeding did not further exacerbate the signs of diabetes mellitus, as demonstrated by  
204 the intraperitoneal glucose tolerance test (iGTT) (Fig. 2G) and fasting glucose levels,  
205 as well as non-esterified fatty acid (NEFA) levels (Fig. 2H). These results indicated that  
206 HFD led to liver and adipose tissue inflammation, as well as liver damage, in SAMP8.  
207 However, within the study timeframe, HFD did not cause considerable exacerbation of  
208 hyperlipidemia or diabetes mellitus.

209

#### 210 **Evaluation of knee pathological changes in HFD- and ND-fed SAMP8**

211 To determine whether HFD promotes the OA-like knee joint changes in SAMP8, we

212 assessed the histopathological changes in the knee joints of these mice using  
213 Safranin-O and Fast Green staining. At 14 weeks of age, the HFD group exhibited  
214 reduced Safranin-O staining, indicating proteoglycan loss, roughened surface, or  
215 cartilage erosion, which in turn indicate more severe cartilage loss compared to the  
216 ND group (Fig 3A a and e). The OARSI score revealed a significant elevation in the  
217 HFD group (Fig. 3B). OA severity was markedly increased in the medial compartment  
218 compared with the lateral compartment (Fig. 3C). Moreover, structural destruction of  
219 the medial menisci is a characteristic appearance during OA progression, which  
220 displayed even prior to the cartilage degeneration, according to our previous findings[9].  
221 The deterioration of medial menisci was more severe in the HFD group (Fig 3A b and  
222 f), as determined by the meniscus score (Fig. 3D), while the scores of lateral menisci  
223 in both groups remained at the same low level. Synovitis, which is commonly  
224 associated with OA, was significantly aggravated in the HFD group according to the  
225 synovitis scores (Fig. 3A c and g, Fig. 3E). Furthermore, both groups exhibited  
226 complete sclerotic changes, i.e., bone-bone marrow replacement at medial tibial  
227 subchondral bone (Fig. 3A d and h). The subchondral bone scores plateaued at 14  
228 weeks of age (Fig. 3F), which are consistent with our previous findings[9].

229 These results indicate that HFD-induced obesity accelerates cartilage and medial  
230 meniscus deterioration, as well as synovitis in 14-week-old SAMP8, suggesting a  
231 worsening of OA status due to the HFD.

232

233 **Bone density and muscle strength in HFD-fed SAMP8**

234 Osteoporosis and sarcopenia, not only OA, are common and significant age-related  
235 diseases of the musculoskeletal system in human, and appear earlier in SAMP8[9, 21].  
236 Thus, we assessed their bone density and muscle strength. Although SAMP8 already  
237 exhibited significantly low BMD in femur compared with SAMR1 at 9 weeks of age[9],  
238 HFD-fed SAMP8 had no further decline in bone density compared with ND-fed SAMP8  
239 at 14 weeks of age (Fig. 4A). Additionally, grip strength test did not show any early-  
240 onset of muscle strength decrease in HFD-fed SAMP8 (Fig. 4B). Therefore, these  
241 results suggested that obesity by HFD feeding did not exacerbate the phenotypes of  
242 osteoporosis or sarcopenia in SAMP8 at 14 weeks.

243

244 **Molecular mechanisms of HFD-induced OA progression in SAMP8**

245 To further elucidate the underlying mechanisms through which HFD-feeding  
246 exacerbates OA-like changes in SAMP8, we conducted a series of in situ detection  
247 focusing on potential molecular markers. Chondrocyte apoptosis has been reported to  
248 be induced by HFD through induced endoplasmic reticulum (ER) stress[22, 23], which  
249 strongly associates with cartilage degradation and OA[24], so we performed TUNEL  
250 staining to detect apoptotic chondrocytes. Our results revealed that SAMP8 subjected  
251 to 10 weeks of HFD feeding exhibited an increased number of apoptotic chondrocytes  
252 in their knee articular cartilage (Fig. 5A) , which was accompanied by a significant  
253 upregulation of C/EBP homologous protein (CHOP) (Fig. 5B), a mediator of ER-stress-

254 induced apoptosis[22]. We also assessed the expression of p16<sup>INK4a</sup>, a cell cycle  
255 regulator and a marker of chondrocyte senescence[25] that we previously found to be  
256 upregulated in SAMP8 compared to SAMR1. Immunohistochemistry analysis  
257 demonstrated that the rates of p16<sup>INK4a</sup>-positive chondrocyte remained unchanged in  
258 SAMP8 cartilage after HFD feeding (Fig. 5C). Additionally, we examined the  
259 expression of MMP13 and ADAMTS5, two hallmark metalloproteinase genes that are  
260 involved in cartilage degradation. However, immunohistochemical staining revealed no  
261 significant differences in the expression of these genes between the HFD and ND  
262 groups (Fig. 5D and E).

263 These results suggested that the increased severity of OA-like changes in HFD-fed  
264 SAMP8 may be primarily attributed to the elevated rate of chondrocyte apoptosis, while  
265 the expression levels of cellular senescence marker p16<sup>INK4a</sup> and cartilage degradation  
266 markers, MMP13 and ADAMTS5, appear to be unaffected by HFD feeding.

267

## 268 **Discussion**

269 OA is a heterogenous disorder, while aging is recognized as the predominant risk  
270 factor contributing to its development and progression. Nonetheless, at present, many  
271 studies utilize post-traumatic mouse OA models, such as destabilization of the medial  
272 meniscus (DMM), or rely on naturally aging mice, which require more than about 12  
273 months to induce OA-like changes[7, 8]. In our recent study, we demonstrated that the  
274 SAMP8 displays knee OA-like changes, and bone density loss in femur, at an early

275 age (14 weeks of age)[9]. Obesity represents another significant risk factor for OA and  
276 frequently coexists with aging. The combined effects of obesity and aging not only  
277 exacerbate OA through cumulative wear and tear on joints but also contribute to  
278 increased levels of inflammation[3]. In the present study, to unveil the combined effect  
279 of HFD-induced obesity and aging, we subjected SAMP8 to HFD feeding for 10 weeks,  
280 and found that HFD-induced obesity further exacerbates the OA in SAMP8 through  
281 elevated apoptosis. Thus, we suggest that SAMP8 serves as a valuable model for  
282 studying molecular mechanisms and developing preventive or therapeutic approaches  
283 for aging-related musculoskeletal diseases, particularly OA, because SAMP8  
284 spontaneously exhibits OA manifestations as early as 14 weeks of age which can be  
285 useful for evaluating the pathogenesis of OA influenced by various factors, such as  
286 HFD.

287 In the body parameters of ND- and HFD-fed SAMP8, after 10 weeks of feeding from  
288 5-week-old, the body, liver, and fat tissue weight of mice in the HFD group were  
289 significantly higher than those in the ND group. The difference in body weight (HFD:  
290 34.52g vs ND: 29.29g, +17.8%) meets the criteria for obesity (15%) according to  
291 previous research[26], and which is comparable with previous studies[27, 28]. The  
292 weight gains induced by HFD exacerbated the severity of OA in 14-week-old SAMP8.  
293 According to our previous study, compared to SAMR1, SAMP8 exhibited significant  
294 meniscal damage at 11 weeks and cartilage damage at 14 weeks, with early sclerosis  
295 of subchondral bone already present at 6 weeks[9]. In the present study, we compared

296 the cartilage damage in the knee joints by OARSI score, and SAMP8 fed with HFD  
297 displayed a more severe cartilage loss. OA severity was markedly increased in the  
298 medial compartment. Moreover, structural destruction of the menisci was observed,  
299 which, based on previous studies, could be due to mechanical damage resulting from  
300 increased weight load[29]. We also showed that increased levels of synovitis in the  
301 joints were consistent with the elevated inflammation levels observed in adipose tissue;  
302 however, this could also be a consequence of cartilage damage and repair[30],  
303 because synovitis was exhibited after cartilage degradation. In addition, the sclerosis  
304 of the subchondral bone in both groups already reached a plateau at 14 weeks of age,  
305 in line with our earlier observations[9]. Thus, HFD-induced obesity may promote the  
306 severity of OA in SAMP8 through locally overloading in medial compartment.  
307 Furthermore, we examined other age-related musculoskeletal disorders. Osteoporosis  
308 is an important aging-related phenotype in human and mouse, but its relationship with  
309 obesity is complex. Obesity increases bone density through elevated mechanical  
310 load[29], while on the other hand, enhances bone resorption through systemic  
311 inflammation by cytokines such as TNF- $\alpha$  and IL-6[31, 32]. In comparison to SAMR1,  
312 SAMP8 have a significantly more severe bone loss in femur at an early age of 9  
313 weeks[9]. However, at 14 weeks, our DEXA data revealed that HFD did not further  
314 decrease bone density, because bone loss already reached a plateau at 9 weeks of  
315 age[9]. Thus, the interplay between mechanical and inflammatory factors, along with  
316 the intrinsic properties of SAMP8 strain, results in the unaltered bone density loss in

317 HFD-SAMP8.

318 Sarcopenia is another prevalent aging-related condition, characterized by  
319 progressive deterioration in skeletal muscle functions and loss in mass[33]. The  
320 biomechanical effect resulting from cross-talk of joints and periarticular muscle could  
321 be associated with the development of OA. The prevalence of sarcopenia in people  
322 with knee OA was 45.2%, the prevalence of sarcopenia in knee OA was more than two  
323 times higher than in the control group[34]. SAMP8 exhibits sarcopenia around 35  
324 weeks of age[21]. At 14 weeks of age, however, HFD-SAMP8 did not exhibit  
325 acceleration of the early onset of sarcopenia. Thus, in SAMP8, muscle  
326 strength/sarcopenia might not be an initiator of OA.

327 SAMP8 exhibits severe liver damage compared with SAMR1[35]. Furthermore, HFD-  
328 induced obesity in C57BL/6 or SAMP8 leads to metabolic disorders such as NAFLD,  
329 hyperlipidemia and hyperglycemia[36, 37, 38], which are strongly associated with OA  
330 and may predispose the body to an OA-prone environment through shared  
331 mechanisms of inflammation, oxidative stress, common metabolites and endothelial  
332 dysfunction[39, 40]. In the present study, although HFD-fed SAMP8 did not exhibit  
333 further exacerbation of diabetic changes, adipose tissue inflammation and NAFLD-like  
334 liver damage were identified. The phenotype could be attributed to the relatively short  
335 timeframe of the study (10 weeks), as HFD typically induces hyperglycemia in at least  
336 12 weeks[41]. However, NAFLD resulted in hepatic and systemic  
337 hyperammonemia[42], concordant with our findings in the present study that HFD-

338 SAMP8 with NAFLD and impaired hepatic function, exhibit elevated blood ammonia  
339 levels. Reduced TG and T-CHO in HFD group may indicate more severe liver damage  
340 by the HFD in SAMP8. Thus, NAFLD might be a potential mechanism of accelerated  
341 severity of OA in HFD-fed SAMP8. We should further examine the relation between  
342 liver damage and OA pathogenesis because these results might open a new insight in  
343 mechanisms of OA through NAFLD.

344 A marker of dysfunctional senescent cell including chondrocytes, p16<sup>INK4a</sup>, is a cell  
345 cycle inhibitor[25]. In our recent study, p16<sup>INK4a</sup> expression increased in SAMP8 at 6,  
346 9, and 11 weeks, indicating accelerated chondrocyte senescence in these[9]. In other  
347 OA-related studies, suppressing p16<sup>INK4a</sup> or clearance of p16<sup>INK4a</sup>-positive  
348 chondrocytes suppress OA[43, 44]. On the other hand, there are reports suggesting  
349 that the expression of p16<sup>INK4a</sup> does not contribute to the development of OA[25]. In  
350 our results, no differences were observed between the two SAMP8 groups, indicating  
351 that modulation of p16<sup>INK4a</sup> expression is not the way that HFD participate in OA  
352 pathogenesis. Moreover, MMP13 and ADAMTS5 are two of the most prominent  
353 metalloproteinases directly involved in the degradation of cartilage extracellular matrix  
354 components, regulated by upstream factors such as inflammation-related NF-κB  
355 pathway[45, 46]. However, in our study, we did not observe any significant changes in  
356 these two enzymes, suggesting that inflammation is not the primary factor for HFD-  
357 induced OA. Thus, HFD-induced obesity did not upregulate the local inflammatory  
358 condition in various tissues/cells of knee joint such as articular chondrocytes and may

359 not be strongly associated with the upregulation of ADAMTS5 and MMP13. The  
360 present study indicated the potential role of chondrocyte apoptosis in the HFD-induced  
361 exacerbation of OA-like changes in SAMP8. Although it has been reported that  
362 apoptosis often plays an important role in OA pathogenesis, apoptosis can be both a  
363 cause and a consequence of OA[24]. Our recent study showed that chondrocyte  
364 apoptosis in SAMP8 did not increase up to 11 weeks of age[9]. Moreover, both  
365 mechanical stress and HFD independently provoke chondrocyte apoptosis by inducing  
366 ER stress[22, 47]. The accumulation of mechanical stress or free fatty acids (FFAs),  
367 which are consequences of HFD-induced obesity, triggers ER stress in  
368 chondrocytes[22, 23, 47, 48]. Under excessive ER stress, apoptotic pathways will be  
369 activated primarily via CHOP[49]. The deletion of CHOP in mice has been shown to  
370 inhibit chondrocyte apoptosis and reduce the severity of OA in a surgical model[50].  
371 The ER stress-induced rise of CHOP activates a series of pro-apoptotic molecules,  
372 leading to the upregulation of apoptosis and, eventually, exacerbates OA  
373 progression[22, 47]. Indeed, excessive ER stress might be involved in apoptotic  
374 pathways because CHOP was increased in articular chondrocytes of HFD-fed SAMP8.  
375 As a limitation of the present study, although we did not clarify the main factors that  
376 induce chondrocyte apoptosis in HFD-fed SAMP8, it suggests that apoptosis via  
377 increased mechanical stress, accumulation of excessive FFAs, and waste products  
378 from liver, such as ammonia, might be a potential mechanism that accelerated severity  
379 of OA in HFD-fed SAMP8.

380 Together, our study demonstrated that HFD feeding further promotes the  
381 spontaneous OA progression via obesity with liver damage in SAMP8 at 14 weeks of  
382 age. Chondrocyte apoptosis is a potential mechanism underlying the promotion of OA  
383 development by HFD in SAMP8. These findings can be useful to better understand the  
384 mechanisms between aging, obesity and OA.

385

### 386 **Abbreviations**

387 OA: osteoarthritis; SAMP8: Senescence-Accelerated Mouse-Prone 8; BMD: bone  
388 mineral density; DAPI: 4',6-diamidino-2-phenylindole;

389

### 390 **Declarations**

#### 391 **Ethics approval and consent to participate**

392 All animal experiments were performed according to protocols approved by the  
393 institutional Animal Care and Use Committees at Hiroshima University.

394

#### 395 **Availability of data and materials**

396 The datasets used and/or analyzed during the current study are available from the  
397 corresponding author on request.

398

#### 399 **Competing interests**

400 The authors declare no competing interests.

401

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407 manuscript for publication.

408

409 **Author contributions**

410 C.D and S.M contributed to the conception and design of the study. C.D, D.Y, Y.M and  
411 Y.S performed the experiments. C.D, T.N, K.M, N.A and S.M contribute to the analysis  
412 and interpretation of data. C.D and S.M contributed to draft manuscript. All authors  
413 approved the submitted manuscript.

414

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543

544

545 **Figure Legends**

546 **Figure 1. Alterations in body parameters of SAMP8 subjected to HFD feeding.** A)

547 Schematic diagram of the experimental design. B) Monitoring of body weight changes  
548 during the experimental timeframe from 5 to 14 weeks of age (n=20 per group). C)  
549 Representative images of ND- and HFD-fed SAMP8 at 14 weeks of age prior to  
550 sacrifice. D) Liver tissue weights obtained post-sacrifice (n=14 per group). E)  
551 Combined weight of epididymal and subcutaneous adipose tissues and their proportion  
552 of total body mass (n=14 per group). Data represented as mean  $\pm$  S.D. Comparison of  
553 mean values was performed using 2-way ANOVA with Sidak's multiple correction (body  
554 weight) or Welch's t test. \* $p$ <0.05, \*\* $p$ <0.01, \*\*\* $p$ <0.001. iGTT: intraperitoneal glucose  
555 tolerance test.

556

557 **Figure 2. Metabolic alterations in SAMP8 following HFD feeding.** A)

558 Representative images of liver H&E staining. B) Serum levels of liver damage markers  
559 aspartate transaminase (AST) and alanine transaminase (ALT) (n=16 per group). C)  
560 Serum levels of circulating lipids triglyceride (TG) and total cholesterol (T-CHO) (n=16  
561 per group). D) Representative images of H&E-stained epididymal adipose tissue. E)  
562 Adipocyte areas in epididymal adipose tissue (n=3 per group). F) Ratio of crown-like  
563 structures (CLSs) to adipocytes in epididymal adipose tissue (n=3 per group). G)  
564 Changes in blood glucose levels during intraperitoneal glucose tolerance test (iGTT)  
565 (left) and area-under-curve of blood glucose (right) (n=12 per group). H) Fasting

566 glucose and non-esterified fatty acid (NEFA) concentrations in serum (n=16 per group).

567 Data represented as mean  $\pm$  S.D. Comparison of mean values was performed using

568 Welch's t test. \* $p$ <0.05, \*\* $p$ <0.01. ns: non-significant difference.

569

570 **Figure 3. Histopathological assessment of knee OA progression in HFD-fed**

571 **SAMP8.** A) Representative images of knee joint safranin O staining. a) and e): articular

572 cartilage; b) and f): medial meniscus; c) and g): synovium (arrowheads); d) and h):

573 subchondral bone of medial tibia. B) Histopathological OARSI scores for entire knee

574 joint cartilage at 14weeks of age (n=20 per group). C) OARSI scores of medial and

575 lateral tibial/femoral cartilage. D) Medial and lateral meniscus scores for left knees

576 (n=20 per group). E) Synovitis scores of left knees (n=20 per group). F) Scores of

577 subchondral bone sclerosis for medial tibias of left knees (n=20 per group). Scoring

578 data represented as median with 95% confidence interval and were compared

579 between ND and HFD groups by Mann–Whitney U test. \* $p$ <0.05, \*\* $p$ <0.01, \*\*\* $p$ <0.001,

580 \*\*\*\* $p$ <0.0001. ns: non-significant difference.

581

582 **Figure 4. Evaluation of bone density and muscle strength in HFD-fed SAMP8.** A)

583 Bone mineral density of right femurs in ND- and HFD-fed SAMP8 (n=8 per group). B)

584 muscle strength assessment in ND- and HFD-fed SAMP8 using grip strength test

585 (n=20 per group). Data represented as mean  $\pm$  S.D. Comparison of mean values was

586 performed using multiple unpaired t tests with Holm-Sidak's multiple correction (bone

587 mineral density) or Welch's t test (muscle strength). ns: non-significant difference.

588

589 **Figure 5. In situ staining of potential regulatory markers for OA.** A) TUNEL staining

590 was performed to assess knee joints. (n=8 per group). Green: TUNEL-positive cells;

591 Blue: DAPI nuclear stain. Immunohistochemistry of B) C/EBP homologous protein

592 (CHOP) (n=8 per group), C) p16<sup>INK4a</sup> (n=8 per group), D) MMP13 (n=8 per group) and

593 E) ADAMTS5 (n=8 per group) were performed and assessed on ND- or HFD-fed

594 SAMP8. Quantification by positive cell number/total cell number were represented as

595 mean  $\pm$  S.D. Comparison of mean values was performed using Welch's t test. \*\* $p < 0.01$ .

596 \*\*\*\* $p < 0.0001$ . ns: non-significant difference.

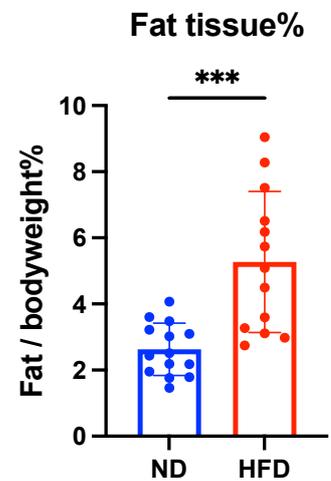
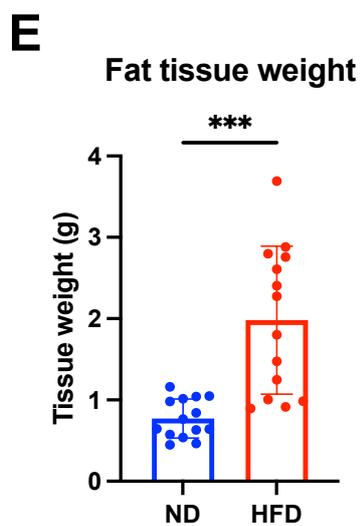
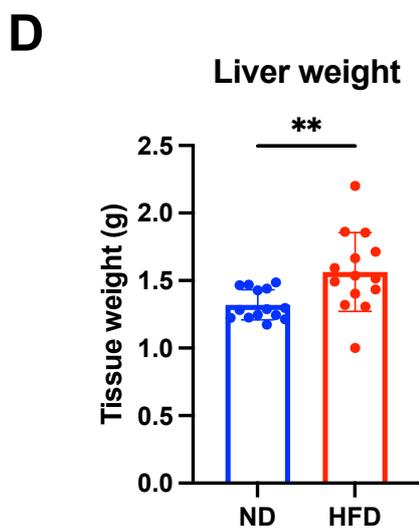
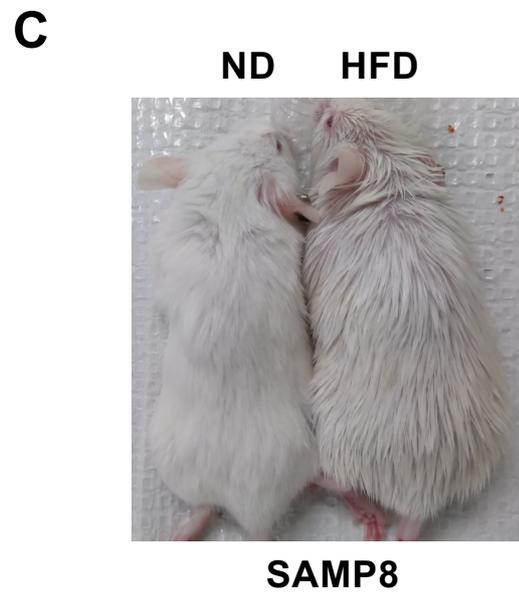
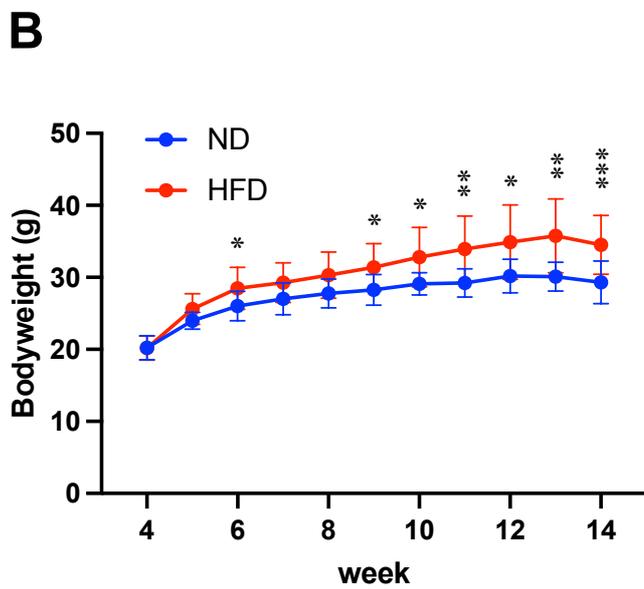
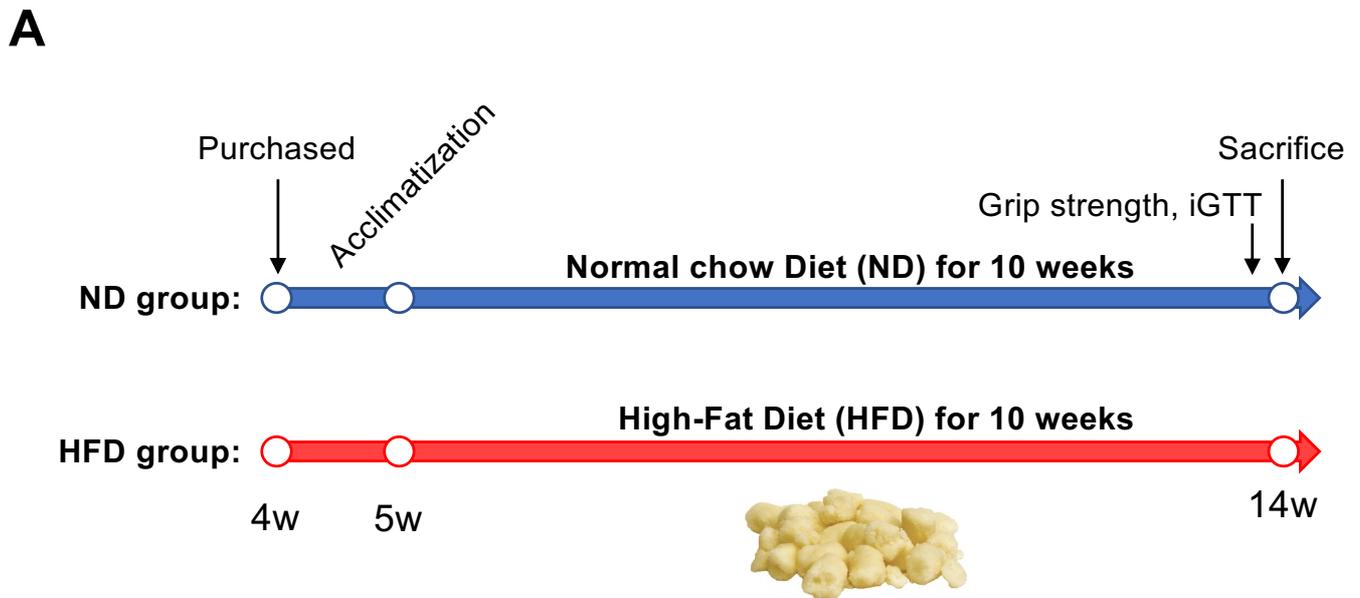


Figure 1

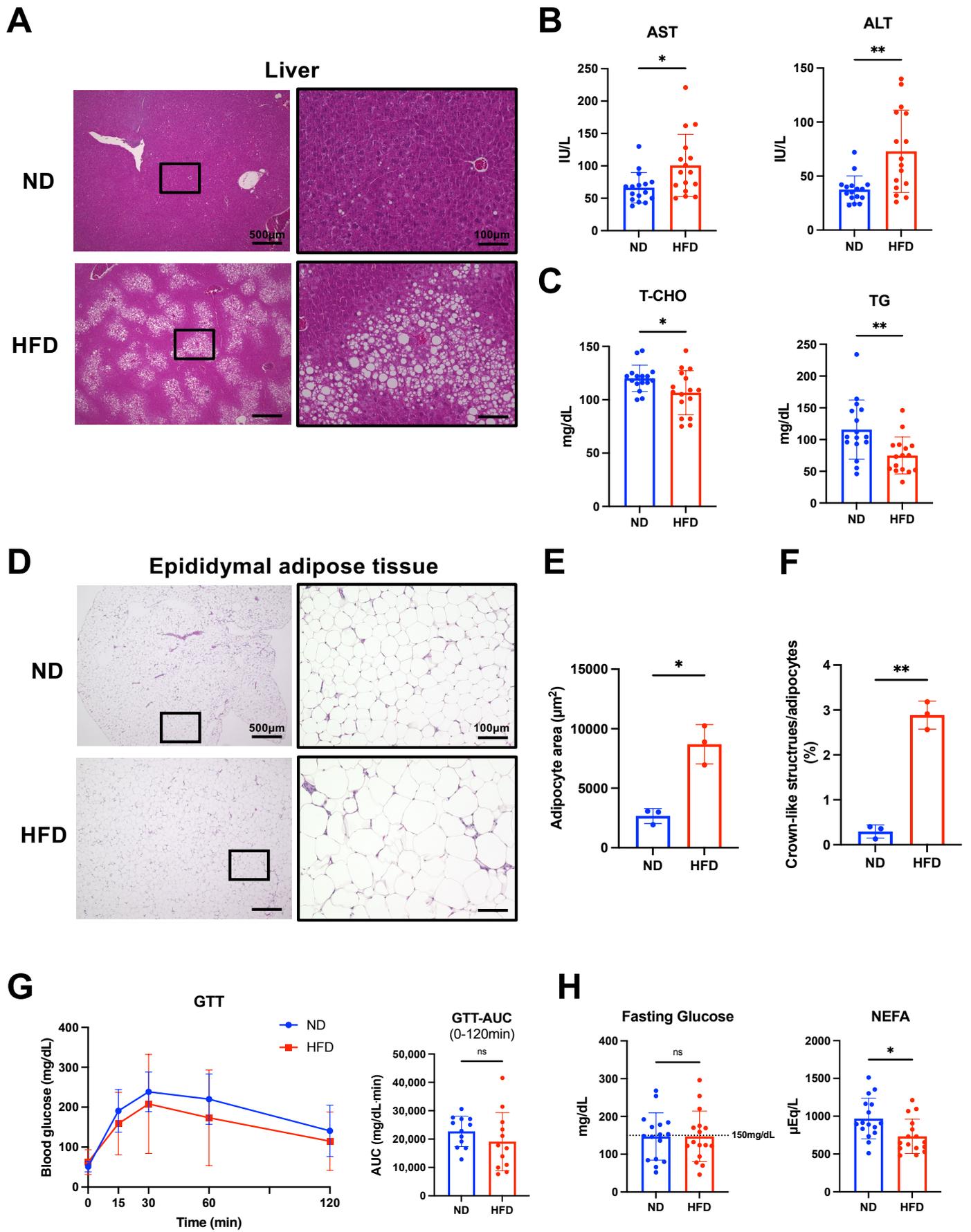


Figure 2

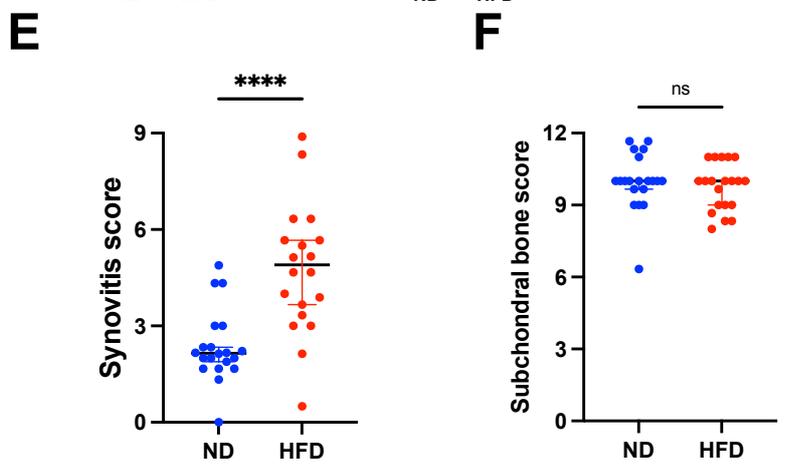
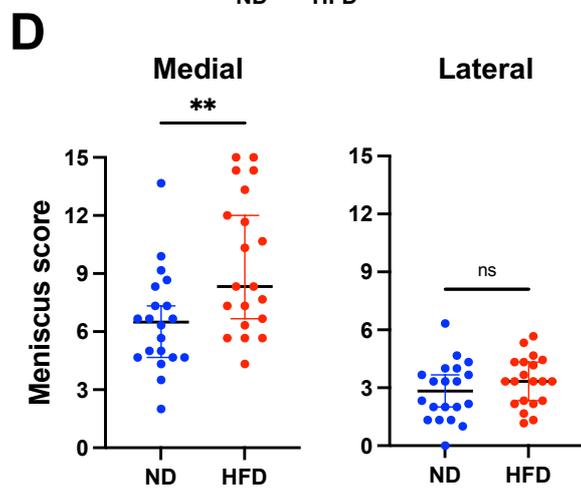
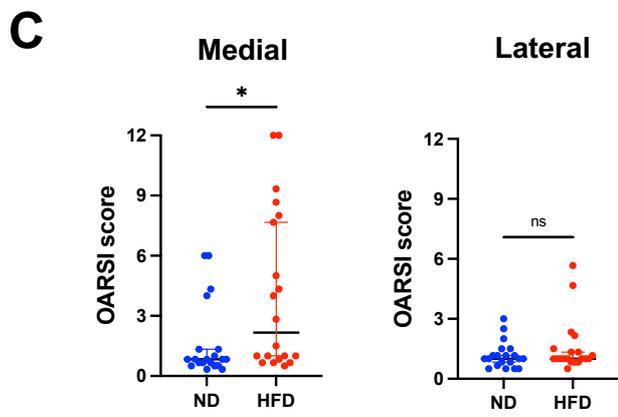
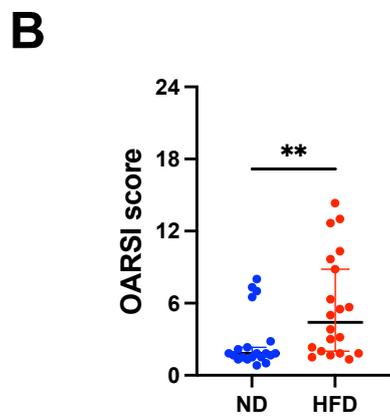
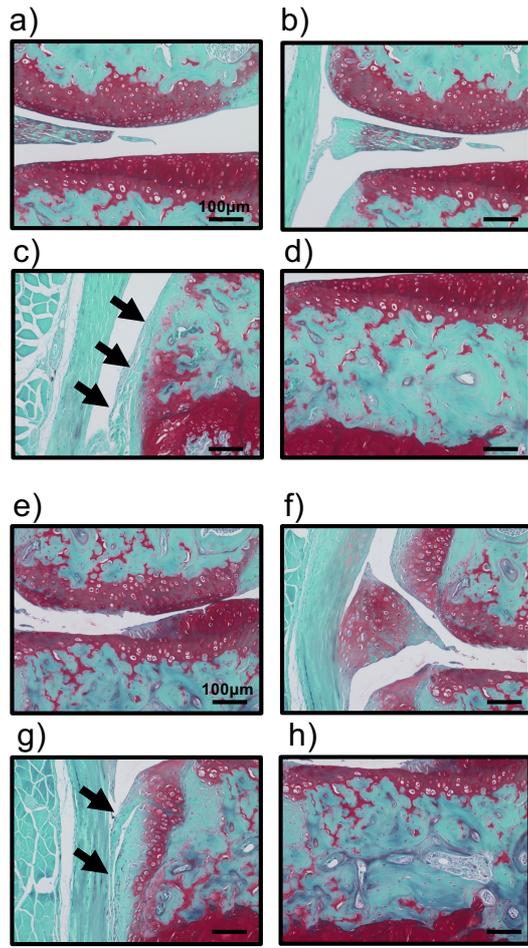
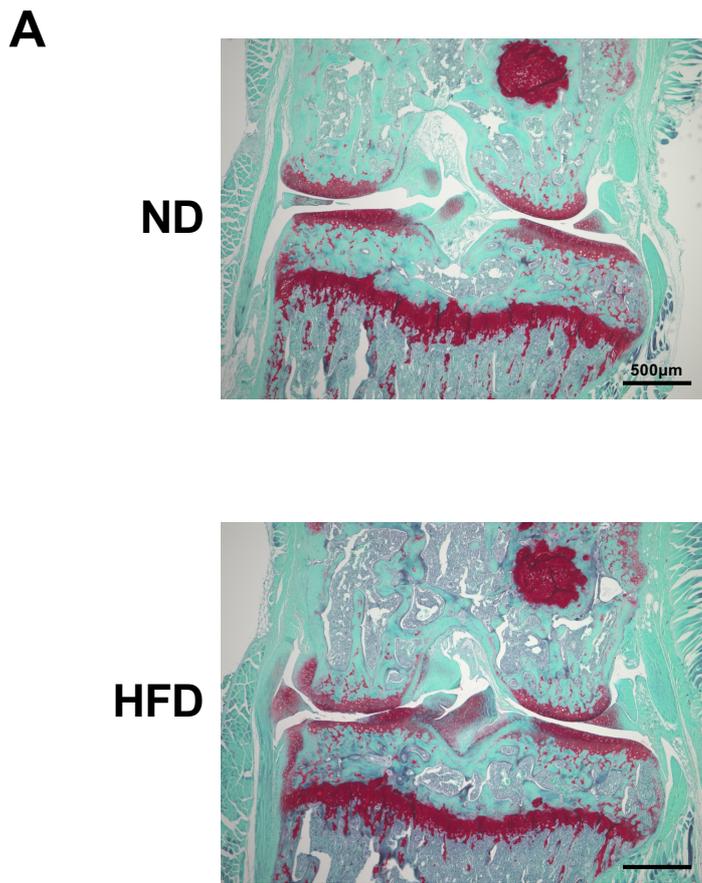
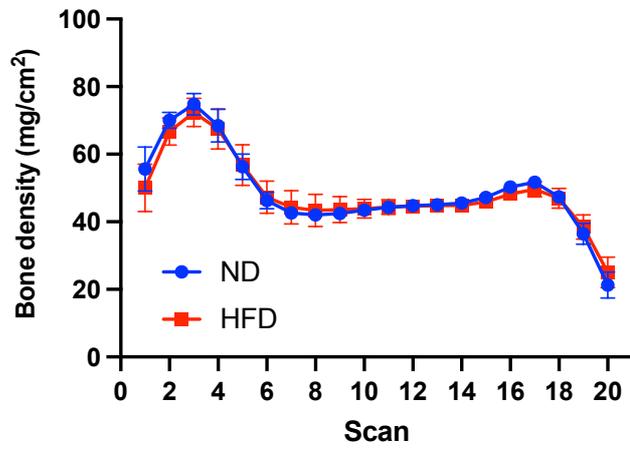


Figure 3

**A**



**B**

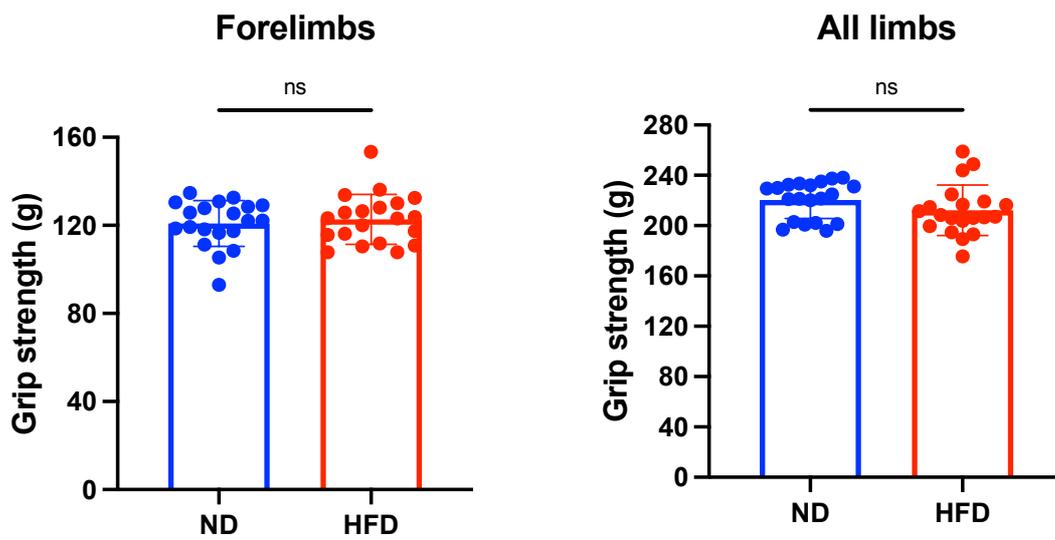


Figure 4

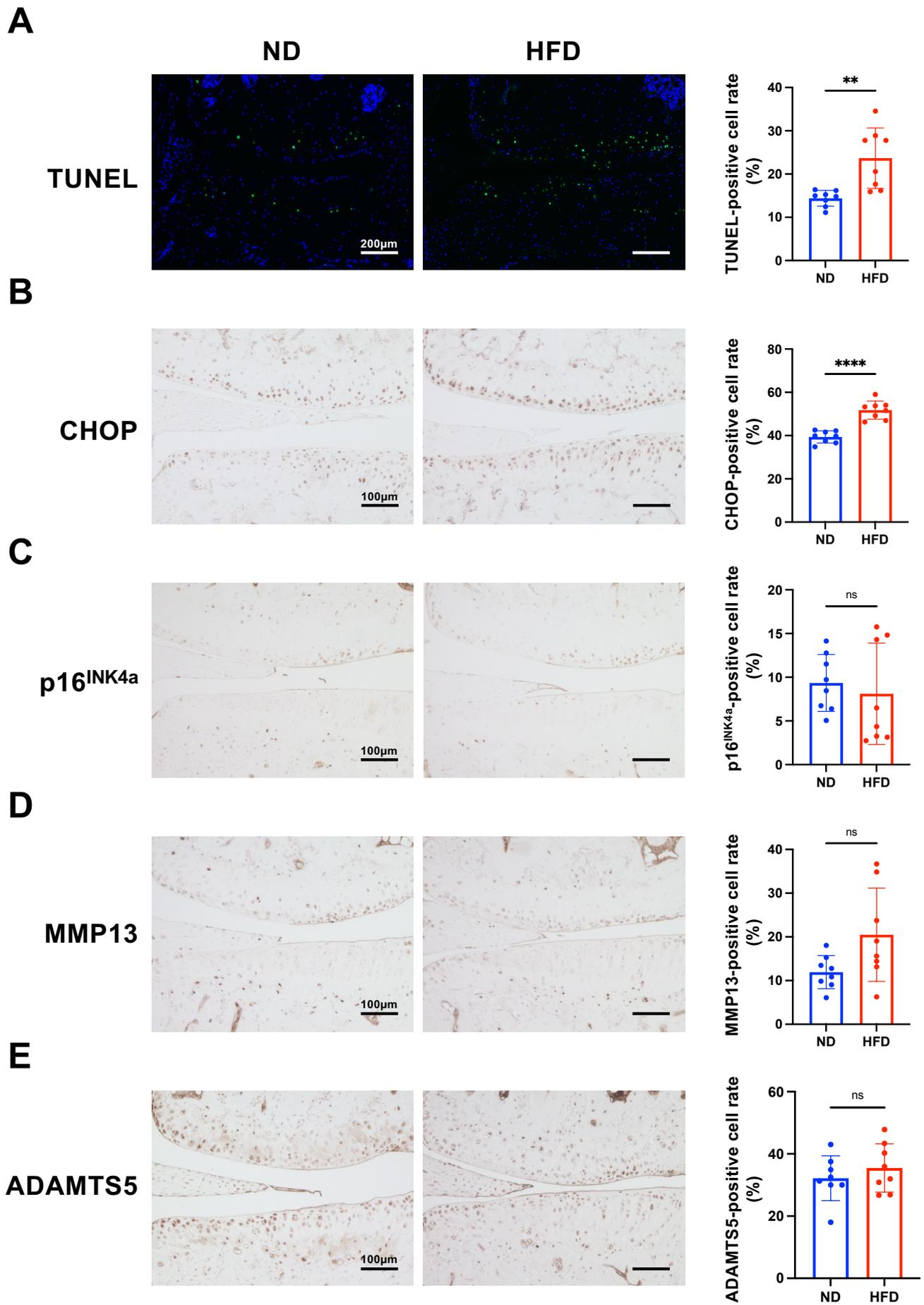


Figure 5

## **Supplemental Materials**

### **Materials and Methods**

#### **Glucose Tolerance Test (GTT)**

The intraperitoneal glucose tolerance test (iGTT) was performed one day before sacrificing at the end point of the animal study. The SAMP8 underwent fasting for 16 hours prior to the test. The mice were weighed and injected intraperitoneally with 10% D-glucose solution (1.5 g/kg body weight). The blood samples were obtained from the tail veins at 0, 15, 30, 60 and 120 min following the administration and glucose levels were measured using a glucose meter MEDISAFE (TERUMO, Japan).

#### **Dual-energy X-ray Absorptiometry (DEXA) Analysis**

The skin and muscles were removed from the right hind limbs which were fixed in 4% PFA-PBS for 48 h at 4°C. Bone mineral density (BMD) in femur bone was measured using dual-energy X-ray absorptiometry (DEXA) densitometry (Aloka DCS600EX, Aloka Co., Tokyo, Japan). Bone density measurements of right femur were taken from sixteen consecutive images with a scan pitch of 2 mm by DEXA scan from the distal femur.

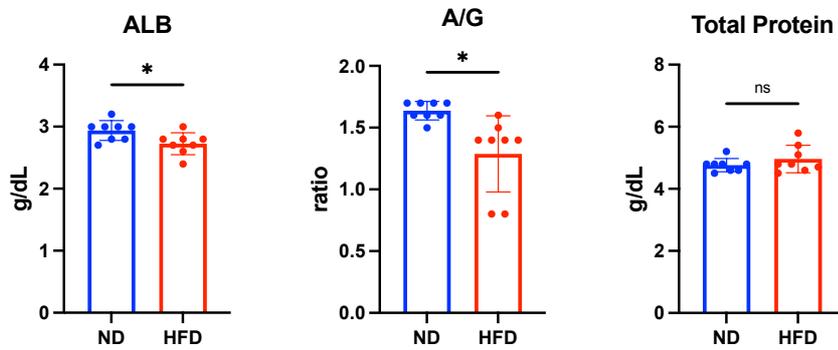
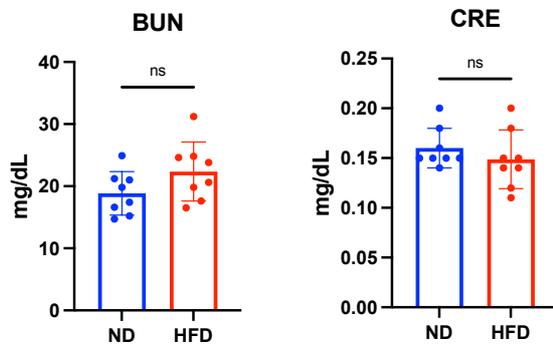
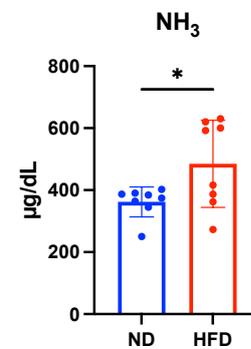
#### **Grip Strength Test**

Muscle strength was evaluated by measuring grip strength at 14 weeks of age (MK-380Si; MUROMACHI Kikai Co., Ltd. Japan). The wire mesh was grasped by the forelimb or all limbs of the mouse and pulled backward until it was released. The value

of the spring scale linked to the wire mesh was recorded. The measurement was performed 5 times per mouse, and the average value was used for statistical analysis.

### **Blood Biochemical Analysis**

Blood samples were collected by cardiac puncture after sacrifice. After coagulating at room temperature for one hour, the blood samples were centrifuged under 1,500g for 30 min. Supernatant serum were collected for subsequent analyses. Serum biomarkers of total protein, albumin (ALB), albumin/globulin (A/G) ratio, blood urea nitrogen (BUN), creatinine (CRE) and ammonia (NH<sub>3</sub>) in serum were analyzed by Nagahama Life Science Laboratory (Nagahama, Japan) using routine laboratory methods.

**A****B****C**

**Supplemental Figure 1. Alterations in markers for hepatic and renal function of HFD-fed SAMP8.** A) Serum levels of circulating albumin (ALB), albumin/globulin (A/G) ratio and total protein (n=8 per group). B) Serum levels of blood urea nitrogen (BUN) and creatinine (CRE) (n=8 per group). C) Serum levels of ammonia (NH<sub>3</sub>) (n=8 per group). Data represented as mean ± S.D. Comparison of mean values was performed using Welch's t test. \**p*<0.05. ns: non-significant difference.